

# JAX Colony Management System



**The Jackson  
Laboratory**

*Leading the search  
for tomorrow's cures*

Welcome To JAX JCMS Release: 6.1.0

Logged in as: mtsadmin

JCMS

[Visit JCMS home page](#)

[Click here for support](#)

**JAX  
JCMS™**

Start  
(Workstation)

Start  
(Handheld Device)



The development and distribution of this software is made possible by generous grants from  
The National Institute for General Medical Sciences (NIGMS) [Grant number: 2R01GM072863-05A1]  
The National Cancer Institute (NCI) [Grant number: P30 CA034196-20]  
and  
The Howard Hughes Medical Institute

# USER GUIDE

## Release 6.1



## JAX Colony Management System (JCMS)

### User Guide JCMS Release 6.1

Prepared by:

Beth Sundberg, Scientific Software Engineer  
Charles Donnelly, Scientific Software Engineer  
Michael McFarland, Scientific Software Engineer  
Abigail Ames, Scientific Software Engineer  
Dave Springer, Scientific Software Engineer  
Kavitha Rama, Scientific Software Engineer  
Peter Blauth, Software QA Engineer  
Rick Palazola, Scientific Software Engineer  
Michael Amato, Scientific Software Engineer  
Computational Sciences  
The Jackson Laboratory

The development and distribution of this software is made possible by generous grants from  
The National Institute for General Medical Sciences (NIGMS)  
[Grant number: 2R01GM072863-05A1]  
The National Cancer Institute (NCI) [Grant number: P30 CA034196-20]  
and  
The Howard Hughes Medical Institute

---

#### Document modification history

Revision number	Date	Who	Comments
1	10/14/05	bas	Document started
10	11/19/07	dsj	Add sample tracking
11	11/23/07	aames	Add sample tracking guide
13	9/18/08	mmm	Added colony report and MLV
14	9/19/08	BAS	Added experimental data import, updated experimental data section, cage cards
15	12/10/08	BAS	Added plug dates
16	10/23/09	aames	Release 3.4.0 updates
17	1/12/10	BAS	Release 4.0.0 updates
18	6/4/10	BAS	Release 4.1.0 updates
19	6/3/11	BAS	Release 4.2.0 updates
20	9/7 - 14/11	RPP	Release 4.3.0 Add Sample updates

<b>21</b>	10/3/11	BAS	Release 4.3.0 updates
<b>22</b>	11/16/11	BAS	Release 4.4.0 updates
<b>23</b>	6/13/12	BAS	Release 4.5.0 updates
<b>24</b>	9/10/12	BAS	Release 4.6.0 updates
<b>25</b>	2/20/13	BAS	Release 4.7.0 updates
<b>26</b>	3/19/13	BAS	Release 4.8.x updates
<b>27</b>	5/3/13	BAS	Release 5.0.0 updates
<b>28</b>	5/10/13	MKA	Release 5.0.0 installation information
<b>29</b>	7/5/13	BAS	Release 5.1.0 updates
<b>30</b>	9/23/13	BAS	Release 6.0.0 updates
<b>31</b>	10/9/13	BAS	Release 6.1.0 updates
<b>32</b>	2/26/14	BAS	Release 6.1.2 updates

## About this document

This User Guide is now available from the JCMS Colony Management System Web site ([colonymanagement.jax.org](http://colonymanagement.jax.org)) as a separate download from the installation files for those who only need a copy of the documentation.

### Document Purpose

This document provides a user manual for the JAX Colony Management System (JCMS) database, primarily focusing on the Microsoft Access user interface.

### Related Documents

JCMS\_ReleaseNotes.pdf  
ReadMe.pdf

### Audience

Users and Database Administrator.

### Table of Contents

1	Overview.....	12
1.1	What is JCMS? .....	12
1.2	Types of Information Collected by JCMS .....	14
2	Getting Started .....	15
2.1	Quick Start .....	15
2.1.1	Administrator .....	15
2.1.2	Regular Users: Owners and Secretaries.....	15
2.2	Installation .....	15
2.2.1	System Requirements .....	15
2.2.2	Download the JCMS Files .....	16
2.2.3	ReadMe Files .....	17
2.3	Configuration.....	18
2.3.1	Required Configuration Steps .....	18
2.3.2	MS Access 2007 Configuration .....	18
2.3.3	MS Access 2010 Configuration .....	21
2.3.4	Removing Security Alerts and Warnings.....	22
2.3.5	Configure some Database Options .....	23
2.3.6	Creating a Multi-User Environment .....	23
2.3.7	Moving JCMS to a New Location .....	24
2.3.8	Re-linking the Tables.....	24

2.3.9 System.mdw File .....	26
2.3.10 Installing a Multi-User Client .....	26
2.3.11 Backing up.....	27
2.3.12 Installing the Bar Code 128 Font.....	27
2.3.13 Configuration Issues and Answers to Common Problems.....	28
3     Administrator Setup.....	30
3.1 Owner and Secretary Accounts .....	30
3.1.1 Rule 1: Every User of JCMS Must Have a Logon to MS Access.....	30
3.1.2 Rule 2: Each logon name that JCMS uses must be defined as either an owner (of mice) or a secretary. ....	30
3.2 Changing the Forms Used by Secretaries and Owners .....	32
3.3 Initializing Controlled Value (CV) Tables .....	32
3.3.1 Simple Controlled Value Tables.....	32
3.3.2 Strain Table .....	35
3.3.3 Mouse Line Viability .....	37
3.3.4 Ear Tag Ranges .....	38
3.3.5 Approved Strains for Matings.....	39
3.3.6 Life Status .....	40
3.3.7 Gene, Gene Class, and Allele Tables .....	40
3.3.8 Rooms, Health Level, and Pens.....	43
3.3.9 Mouse Use Types .....	43
3.4 Setup Variables Table (DbSetup) for Customizing the Installation.....	43
3.5 Cage Card Setup .....	47
3.5.1 Printing Cage Cards .....	47
3.5.2 Blank Cage Cards .....	48
3.5.3 JCMS-Provided Cage Card Formats .....	48
3.5.4 Creating Custom Cage Cards .....	60
3.5.5 User-designed Cage Cards.....	60
3.6 Setting up an Experimental Plan.....	61
4     User Setup.....	61
4.1 Logging On and Passwords.....	61
5     Basics on Using JCMS.....	62
5.1 What are Owners and Secretaries?.....	62
5.2 Changing Passwords .....	62
5.3 Button Bars .....	63
5.4 Using the Forms.....	64
5.4.1 Special Features of Some Forms.....	65
5.5 Navigation Buttons .....	65
5.6 Session Reports.....	65
5.7 Printing .....	65
5.8 Statistics.....	66
6     Mice .....	66
6.1 Mouse IDs .....	66
6.1.1 Automatically Incrementing Mouse IDs.....	66
6.1.2 Base Mouse Numbers.....	67
6.2 Adding Mice .....	67
6.3 Editing Mice.....	69
6.3.1 Changing the ID of a Mouse .....	69
6.4 Importing or Bulk Adding Mice .....	70
6.5 Adding Pups.....	71
6.6 Changing Life Status or Diet of a Group of Mice .....	73
6.7 Changing Life Status using the Handheld Forms .....	73
6.8 Change Life Status of an Individual Mouse using Handheld.....	73
7     Pens and Cage Cards .....	75
7.1 Manage Pen Configuration Form.....	75
7.2 Rooms and Health Level.....	77

7.3 Adding Pens.....	77
7.3.1 Pen Info Form.....	78
7.3.2 Add Pen Form .....	78
7.4 Editing and Retiring Pens .....	79
7.5 Moving Mice between Pens .....	79
7.6 Moving Mice Using Handheld .....	80
7.7 Printing Blank Cage Cards.....	80
7.8 Retiring Pens Automatically .....	80
7.9 Bulk Retire Pens .....	81
7.10 Correcting Pen Status and Date Errors.....	81
7.11 Cage Use Reports .....	82
7.11.1 Cage Use Report.....	82
7.11.2 Cage Use Summary .....	83
8 Matings .....	85
8.1.1 Automatic Litter Number Generation.....	85
8.2 Technical Matings .....	85
8.3 Which Mating Forms to Use?.....	86
8.4 Approved Mating Strains.....	86
8.5 Designing a New Mating .....	87
8.6 Activating a Mating.....	88
8.7 Add a Natural Mating .....	89
8.8 Add a Technical Mating .....	90
8.9 Edit a Mating .....	92
8.10 Design Matings to be Retired.....	92
8.11 Retire a Mating.....	93
8.12 Working with Matings using a Handheld.....	94
8.13 Automatic Retiring of Matings .....	94
9 Litters.....	95
9.1 Automatic Litter Number Generation .....	95
9.1.1 Turning Automatic Litter Numbering On or Off .....	95
9.1.2 Setting the Number of Litter Numbers that are Generated for Each Mating.....	95
9.2 Which Litter Forms to Use? .....	95
9.3 Adding Litters .....	96
9.4 Wean Report.....	96
9.5 Editing Litters .....	97
9.6 Wean Litters .....	98
9.7 Adding Litters with Pups at Weaning .....	98
9.8 Weaning Mice using a Handheld .....	100
10 Embryo Litters.....	101
10.1 Add Embryo Litter w/Samples.....	101
10.2 Edit Embryo Litter .....	103
11 Plug Dates and Pregnancy Checking.....	104
11.1 Add Plug Date.....	104
11.2 Edit Plug Date .....	105
11.2.1 Automatically mark plug dates as obsolete.....	105
11.3 Plug Date and Pregnancy Check Work Reports.....	106
11.3.1 Plug Date / Pregnancy Stage Report.....	107
11.3.2 Plug Check Work Report.....	107
11.3.3 Pregnancy Check Work Report.....	108
11.3.4 Plug Date History .....	108
12 Breeding Performance Reports.....	109
12.1 Full Breeding Performance Report .....	110
12.2 Summary Breeding Performance Report.....	111
12.3 Breeding Performance Report Tabular Spreadsheet .....	112
12.4 Breeding Performance Report Litter Detail Pivot Table.....	112
12.5 Breeding Performance Report Litter Summary Pivot Chart.....	113

12.6 Breeding Performance Report Mating Count Pivot Chart.....	113
12.7 How to Use the Dynamic Features of a Pivot Table or Chart.....	114
12.7.1 Changing the Chart Type .....	114
12.7.2 Showing Only Some of the Criteria .....	114
12.7.3 Adjusting the Axis Scale.....	115
12.7.4 Switch from Pivot Chart to Pivot Table.....	116
12.7.5 Viewing Values for Portions of a Pivot Chart or Table .....	116
13     Genotyping .....	117
13.1 How Does Genotyping Work?.....	117
13.2 Adding a Genotype to a Mouse or Sample.....	117
13.2.1 Mouse-Genotype Documents .....	118
13.3 Editing a Genotype .....	119
13.4 Adding a Genotype to a Group of Mice .....	120
13.5 Genotype String Format.....	120
13.6 Genotype Work Report .....	121
14     Genotype Loader .....	122
14.1 Import genotypes from a comma separated value (CSV) file.....	122
14.1.1 Input File Format .....	123
14.1.2 Alleles.....	125
14.1.3 Import File Integrity Checks .....	125
14.1.4 Select the type of Import .....	126
14.1.5 Verification.....	127
14.1.6 Genotype Load Report.....	129
15     Scheduling Procedures (Uses).....	130
15.1 How do Mouse Uses Work? .....	130
15.2 Adding a Use to a Mouse.....	130
15.3 Mouse Use Work Report.....	131
15.4 Editing a Mouse Use .....	132
15.5 Adding or Editing a Mouse Use for a Group of Mice .....	133
16     Calendar .....	135
16.1 Edit a Mouse Use Shown on the Calendar .....	137
16.2 Mouse Use Details Form .....	138
16.3 Print a Calendar Report .....	139
16.3.1 Suggestions for speeding up the calendar form .....	139
17     Documents.....	140
17.1 Document Storage .....	140
17.2 Adding (Uploading) Documents.....	140
17.3 Selecting Documents to Associate .....	142
17.4 Edit Document .....	143
17.5 Bulk Add Document Associations.....	144
18     Queries .....	145
18.1 Colony Summary Report.....	145
18.2 What are Queries used for? – or How to Search the Database .....	145
18.2.1 Basics on using the Query Forms .....	146
18.2.2 How to Select Query Criteria.....	147
18.2.3 Like Criteria for Mouse ID .....	148
18.2.4 Mouse Age .....	148
18.2.5 Filtering Strains or Stock #'s .....	149
18.3 Mouse Query .....	149
18.3.1 Query by Genotype (QGT).....	149
18.3.2 Interpreting the Genotype Output.....	150
18.3.3 Restricting Genotypes in the Query Output .....	150
18.3.4 Listing Documents Associated with Genotypes .....	150
18.3.5 Listing Other Associated Documents .....	151
18.3.6 Query by Mouse Use.....	151
18.4 Mating Query .....	153

18.5 Experimental Data Query.....	154
18.5.1 Documents Associated with Experimental Data .....	155
18.6 Sample Query .....	155
18.7 Microsoft Query.....	155
19     Experimental Plans.....	156
19.1 How to use an Experimental Plan.....	157
19.2 How to Create Experimental Data without using an Experimental Plan.....	158
19.3 Setting up Test Types (Data Descriptions) .....	159
19.4 Editing a Test Type .....	160
19.5 Setting up Default Data for Experiments .....	160
19.6 Editing Default Data .....	161
19.7 Adding an Experimental Plan.....	161
19.8 Adding an Experimental Test.....	162
19.9 Editing an Experimental Test.....	163
19.10 Selecting Mice for an Experimental Plan .....	163
19.11 Adding Mice to a Plan using the Mouse Complex Query Form .....	164
19.11.1 Example of the report format:.....	165
19.11.2 Error message: "XX mice were selected by the query, more than can be held in the Query Results box." .....	165
19.12 Managing an Experimental Plan .....	166
19.12.1 Choosing Mice for Experimental Tests .....	166
19.12.2 Removing Mice from an Experimental Plan .....	167
19.12.3 Editing Experimental Plan Fields .....	167
19.12.4 Deleting an Experimental Plan .....	168
19.12.5 Deleting an Experimental Test .....	168
19.13 Adding or Editing Experimental Data for a Mouse.....	169
19.14 Editing Experimental Data .....	170
19.15 Adding Experimental Data to Several Mice at Once .....	171
19.16 Experiment Work Report for Scheduling Procedures .....	171
19.16.1 Connector/ODBC Window Pops-up When Using the Experiment Work Report....	173
19.16.2 Setup Variables Used by the Experiment Work Report .....	174
19.17 Experimental Plan Query .....	174
19.18 Importing Experimental Data .....	174
19.18.1 Input File Format .....	175
19.18.2 Create a Test Type.....	176
19.18.3 User Interface .....	177
19.18.4 Troubleshooting Notes .....	183
20     Samples.....	184
20.1 Set up Controlled Vocabulary for Sample Tracking.....	185
20.2 Pooled Samples.....	188
20.3 Add Samples.....	189
20.4 Edit Samples .....	191
20.5 Bulk Change Samples .....	192
20.6 Query Samples .....	193
20.6.1 Searching for sample genotypes.....	194
20.7 Browse Sample Storage Locations.....	195
20.8 Print Sample Labels .....	196
21     FAQ's (Frequently Asked Questions) .....	197
21.1.1 JCMS starts up and shows the database, but no welcome window appears.....	197
21.1.2 Every time I add, edit, or delete a record in JCMS I'm prompted with a dialog box.	197
21.1.3 What to do about an "end/debug" error message .....	197
21.1.4 Error messages when editing date fields .....	197
21.1.5 Error message: "user Admin does not have permission to use this form" .....	197
21.1.6 Error message about "could not find file" .....	198
21.1.7 A note about session boxes .....	198
21.1.8 List boxes of mice information are scrambled!.....	198

21.1.9 Warning message about not saving a record.....	198
21.1.10 The database keeps telling me it is "Read only".....	198
21.1.11 The Connector/ODBC Window is Popping Up.....	198
21.1.12 Other FAQs .....	199
<b>22 Technical Guide.....</b>	<b>199</b>
22.1 Security .....	199
22.2 Changing Security Access to Forms.....	200
22.3 Data Integrity.....	200
22.4 Back up the Database.....	200
22.5 The dblInfo Table.....	200
22.6 Access to the Primary Data Tables .....	200
22.6.1 Editing Records in Datasheet View.....	200
22.7 Temporary Tables.....	201
22.8 Temporary Queries .....	201
22.9 Screen Resolution.....	201
22.10 Printer Notes .....	201
<b>23 Appendix 1: Automated TGS Genotype Loader Submissions .....</b>	<b>203</b>
23.1 Adding a TGS Genotype Request .....	203
23.2 Editing a TGS Genotype Submission Request.....	207
23.3 Importing TGS Genotyping Results .....	208

## Table of Figures and Tables

Figure 1-1 Database Overview Diagram .....	14
Figure 2-1 File browser.....	17
Figure 2-2 Setting full control.....	17
Figure 2-3 Logon Password .....	18
Figure 2-4 Link to the database source .....	19
Figure 2-6 JCMS Welcome Window.....	20
Figure 2-5 Link Tables dialog box .....	20
Figure 2-7 MS Access 2010 Users and Permissions dialog boxes.....	21
Figure 2-8 MS Access 2010 Linked Table Manager .....	21
Figure 2-9 Access 2007 Trust Center.....	22
Figure 2-10 Access 2007 Confirm Options .....	23
Figure 2-11 Access 2007 configure record-level locking.....	23
Figure 2-12 JCMS Failed to start up correctly.....	24
Figure 2-13 Office 2007 Linked Table Manager .....	25
Figure 2-15 Sample of a bar code for the pen ID number.....	27
Figure 2-14 JCMS client shortcut .....	27
Figure 2-16 Error: release numbers don't match.....	28
Figure 3-1 Dialog Box: Personal ID .....	30
Figure 3-2 Form: Administrator Buttons .....	31
Figure 3-3 Form: Owner and Secretary Accounts .....	31
Figure 3-4 Form: Edit Form Privileges.....	32
Figure 3-5 Sample Simple CV Table .....	33
Figure 3-6 Forms: Default Strain Types and Generations .....	34
Figure 3-7 Form: Add or Edit Strains.....	35
Figure 3-8 Open Mouse Line Viability Report .....	37
Figure 3-9 Example mouse line viability report .....	38
Figure 3-10 Ear tag ranges.....	39
Figure 3-11 Add Approved Strain Form .....	40
Figure 3-12 Form: Add or Edit Gene .....	41
Figure 3-13 Form: Add or Delete an Allele.....	42
Figure 3-14 Add or delete mouse use types .....	43
Table 3-1 JCMS Setup Variables .....	47
Figure 3-15 Form: Print blank cage cards .....	48

Figure 3-16 OS_2PWeanCageCard.....	49
Figure 3-17 TS_1PWeanCageCard or TS_1PWeanCageCardWithBarCode .....	49
Figure 3-18 CC_JCMS_WeanCageCardUsingPenNames .....	52
Figure 3-19 TS_MatingCAgeCard or TS_MatingCageCardStyle1With BarCode .....	53
Figure 3-20 TS_MatingCageCardStyle2With BarCode.....	53
Figure 3-21 OS_MatingCageCard.....	53
Figure 3-22 MW_MatingCageCardLandscape .....	53
Figure 3-23 CC_JCMS_MatingCageCardUsingPenNames.....	53
Figure 3-24 TS_DetailCageCard .....	57
Figure 3-25 TS_DetailCageCardWithBarCode .....	57
Figure 3-27 MW_DetailCageCardLandscape .....	57
Figure 3-26 (left) CC_JCMS_DetailCageCardUsingPenNames .....	57
Figure 3-28 Office 2007 reports list shown using F11 key .....	60
Figure 4-1 User Logon Screen .....	61
Figure 5-1 Dialog Box: Change Password .....	63
Figure 5-2 Main Button Bar .....	63
Figure 5-4 Example of regular form buttons.....	64
Figure 5-3 Example of a text box.....	64
Figure 6-1 Add Mouse Form.....	67
Figure 6-2 Add mouse form after a successful submit.....	68
Figure 6-3 Change Mouse ID Form.....	69
Figure 6-4 Edit Mouse With Browser Functions .....	70
Figure 6-5 Bulk Add Mice Form.....	71
Figure 6-6 Add Pup from Litter Form.....	72
Figure 6-7 Form: Bulk Change Life Status or Diet .....	73
Figure 6-8 Individual Mouse Change Life Status Handheld Form .....	74
Figure 6-9 Bulk Exit Handheld Form .....	74
Figure 7-1 Manage Pen Configuration Form .....	75
Figure 7-2 Administrator Forms for Managing Room and Health Level.....	77
Figure 7-3 Pen Info Form .....	78
Figure 7-4 Add Pen Form .....	78
Figure 7-5 Edit Pen Form .....	79
Figure 7-6 Move Mouse Form .....	79
Figure 7-7 Handheld Move Mice Form .....	80
Figure 7-9 Edit Pen Status/Location Form .....	81
Figure 7-8 Pen Maintenance Form.....	81
Figure 7-10 Cage Use Report .....	82
Figure 7-11 Cage Use Summary Report .....	83
Figure 7-12 Cage Use Summary (bottom) vs. Cage Use Report (top) .....	83
Figure 8-1 Mating to Weaning Diagram .....	85
Figure 8-2 Add Approved Strains Form.....	86
Figure 8-3 Design Mating Form.....	87
Figure 8-4 Activate Mating Form .....	88
Figure 8-5 Add Mating Form.....	89
Figure 8-6 Add Ovary Transfer Mating .....	90
Figure 8-7 Add Embryo Transfer Mating .....	91
Figure 8-8 Add In vitro Fertilization Mating.....	91
Figure 8-9 Design Retire Matings Form .....	92
Figure 8-10 Retire Matings Form.....	93
Figure 8-11 Handheld Form: Pair Mating .....	94
Figure 9-1 Add Litter Form .....	96
Figure 9-2 Wean Work Report.....	96
Figure 9-3 Edit Litter Form.....	97
Figure 9-4 Add Litter w/Pups Form.....	98
Figure 9-6 Handheld Forms: Weaning and Add litter and mice .....	100
Figure 10-1 Add Embryo Litter with Sample records.....	101

Figure 10-2 Edit Embryo Litter form .....	103
Figure 11-1 Add Plug Date Form.....	104
Figure 11-2 Edit Plug Date Form.....	105
Figure 11-3 Confirm plug date expiration .....	106
Figure 11-4 Form: Request plug date or pregnancy reports .....	106
Figure 11-5 Sample plug date / pregnancy stage report.....	107
Figure 11-6 Sample tabular spreadsheet format plug date / pregnancy stage report.....	107
Figure 11-7 Plug Check Report.....	107
Figure 11-8 Sample pregnancy check work report.....	108
Figure 12-1 Select the Breeding Performance Report .....	109
Figure 12-2 Form: Request a breeding performance report .....	109
Figure 12-3 Detail section of the report .....	110
Figure 12-4 Strain total section of the report .....	111
Figure 12-5 Owner and grand total section of the report for all strains .....	111
Figure 12-6 Summary shown for one strain .....	111
Figure 12-7 Tabular spreadsheet format.....	112
Figure 12-8 Litter detail pivot table .....	112
Figure 12-9 Litter summary pivot chart.....	113
Figure 12-10 Mating counts by week .....	113
Figure 12-11 Chart changed to 3-D and only one owner .....	114
Figure 12-12 Changed "drill buttons".....	115
Figure 12-13 Properties dialog box .....	115
Figure 12-15 Hover over a column with the mouse to see the values .....	116
Figure 12-14 View button .....	116
Figure 13-1 Form: Add Genotype .....	117
Figure 13-2 Add genotype documents .....	118
Figure 13-3 Form: Edit Genotype .....	119
Figure 13-4 Form: Bulk Add Genotype.....	120
Table 13-1 Examples of genotype strings .....	121
Figure 13-5 Request a genotype work report.....	121
Figure 13-6 Genotype Work Report .....	121
Figure 14-1 Form: The main button bar with the "Import Genotype" button visible .....	122
Figure 14-2 Spreadsheet: the input file as seen in MS Excel .....	124
Table 14-1 The mapping of the input fields to the data tables .....	124
Table 14-2 Genotype loader allele representations .....	125
Table 14-3 Genotype loader error messages.....	125
Figure 14-3 Select import type dialog box .....	126
Figure 14-4 The file open dialog.....	127
Figure 14-5 The user notification screen during validation .....	127
Figure 14-6 The user notification after validation .....	128
Figure 14-7 The user notification after import .....	128
Figure 14-8 Genotype Load Report.....	129
Figure 15-1 Form: Add Use .....	130
Figure 15-2 Adding a use based on a plug date .....	131
Figure 15-3 Request mouse use work report.....	132
Figure 15-4 Form: Edit Use .....	132
Figure 15-5 Bulk Add or Edit Use Form .....	133
Figure 15-6 Warning about making a change to the "done" field.....	134
Figure 16-2 Form: Request mouse use calendar.....	135
Figure 16-1 Open the calendar.....	135
Figure 16-3 Example of a mouse use calendar.....	136
Figure 16-4 Single day format .....	137
Figure 16-5 Opening the edit mouse uses form .....	137
Figure 16-6 Mouse use details form .....	138
Figure 16-7 Example of a calendar printout .....	139
Figure 17-1 Possible data file setup for documents .....	140

Figure 17-2 Add document form.....	140
Figure 17-3 File browser.....	141
Figure 17-4 Upload New button.....	141
Figure 17-5 Associate Documents form .....	142
Figure 17-6 Associate Existing button .....	142
Figure 17-7 Edit Document form .....	143
Figure 17-8 Bulk Add Document Associations form.....	144
Figure 17-9 Experimental plan category .....	144
Figure 18-1 Colony Summary Report.....	145
Figure 18-2 Mouse Query Form .....	146
Figure 18-3 Datasheet: Mouse Query Results .....	146
Figure 18-4 Query: Do you want to save changes to Form? .....	147
Figure 18-5 Query Form: List Box Criteria .....	147
Figure 18-6 Query Form: Range Criteria.....	147
Figure 18-7 Query Form: Check Box Criteria.....	147
Figure 18-8 Query Form: Like Criteria.....	148
Figure 18-9 Mouse age result choice .....	148
Figure 18-11 Query Form: Genotype .....	149
Figure 18-10 Strain or Stock # Filter .....	149
Figure 18-12 Query Datasheet: Restricted Genotype Output .....	150
Figure 18-13 Requesting documents in the query output .....	150
Figure 18-14 Query Form: Mouse Use.....	151
Figure 18-15 Query Datasheet: Mouse Uses .....	151
Figure 18-16 Mouse query - use term error message.....	152
Figure 18-17 Query Mating Form .....	153
Figure 18-18 Mating type selection .....	153
Figure 18-19 Query Datasheet: Mating .....	153
Figure 18-20 Query Experiment Form.....	154
Figure 18-21 Experiment Query Datasheet.....	155
Figure 19-1 Diagram: Experimental Plan Tables .....	156
Figure 19-2 Diagram: Experimental Plan .....	157
Figure 19-3 Form: Add Test Type .....	159
Figure 19-4 Form: Add Test Type Defaults .....	160
Figure 19-5 Form: Add Experimental Plan .....	161
Figure 19-6 Form: Add Experimental Test .....	162
Figure 19-7 Form: Bulk Add Mice from Query.....	164
Figure 19-8 Report: Add Mice to Plan .....	165
Figure 19-9 Form: Manage Experimental Plan .....	166
Figure 19-10 Report: Edit/Mange Experimental Plan.....	167
Figure 19-11 Form: Add Experimental Data.....	169
Figure 19-12 Form: Bulk Add Experimental Data.....	171
Figure 19-13 Request Experiment Work Report .....	172
Figure 19-14 Experiment Work To Do Report Based on the Mouse Age Range .....	173
Figure 19-15 Experiment Work Report Spreadsheet .....	173
Figure 19-16 Connector/ODBC window .....	174
Table 19-1 MySQL setup variables .....	174
Figure 19-17 Process for importing experimental data from a user file .....	175
Figure 19-18 Create a test type to match the input fields .....	176
Figure 19-19 Import Exp Data button .....	177
Figure 19-20 Import Experimental Data with no experimental plan .....	177
Figure 19-21 Validate input .....	178
Figure 19-22 Sample validation report .....	179
Figure 19-23 Maximum number of validation errors.....	179
Figure 19-24 Import Experimental Data using an Experimental Test .....	180
Figure 19-25 Data validation failure.....	181

Figure 19-26 Allow experimental data to be imported to an experimental test without preselecting the mice .....	181
Figure 19-27 Enabled load data button .....	182
Figure 19-28 Final load report .....	182
Figure 19-29 Changing the experimental test status.....	183
Figure 20-1 Simple controlled vocabulary buttons for sample tracking.....	185
Figure 20-2 Example of editing simple controlled vocabulary .....	185
Figure 20-3 Complex controlled vocabulary buttons for sample tracking .....	186
Figure 20-4 Manage sample types screen .....	186
Figure 20-5 Manage preservation vocabularies screen .....	187
Figure 20-6 Manage sample locations screen .....	187
Figure 20-7 Dummy strain for a pooled sample .....	188
Figure 20-8 Add Sample form .....	189
Figure 20-9 Edit Sample Form .....	191
Figure 20-10 Bulk Sample Update Form .....	192
Figure 20-11 Query Samples Form .....	193
Figure 20-12 Sample genotype criteria .....	194
Figure 20-13 Sample genotype results in multiple columns.....	194
Figure 20-14 Genotype results when no gene criteria .....	195
Figure 20-15 Browse Sample Storage Locations Form .....	195
Figure 20-16 Print Sample Labels .....	196
Figure 21-1 Could Not Find File Error .....	198
Figure 21-2 Error: You can't save this record at this time .....	198
Figure 22-1 Administrator - Owner - Secretary Security Relationship .....	199
Figure 23-1 The Add Genotyping Request button.....	203
Figure 23-2 Select mice whose IDs begin with "A1" .....	204
Figure 23-3 Setting the plate ID.....	204
Figure 23-4 A well plate with samples from seven mice of two strains .....	205
Figure 23-5 The genotyping request ID field .....	205
Figure 23-6 To seal the plate select "Sealed" .....	206
Figure 23-7 The Submit button .....	206
Figure 23-8 Notification of a successful submission. ....	206
Figure 23-9 To edit an existing genotyping request .....	207
Figure 23-10 Select a request to be edited. ....	207
Figure 23-11 Import Genotype data .....	208
Figure 23-12 Select the format.....	208
Figure 23-13 To import the results from the selected file. ....	209

# 1 Overview

## 1.1 What is JCMS?

The Jackson Laboratory's Colony Management System (JCMS) is a multi-user relational database for managing mouse colonies in a research environment. It was developed in response to increased demand within The Jackson Laboratory for a system that would execute the core functionality of colony management from an intuitive, easy-to-use interface.

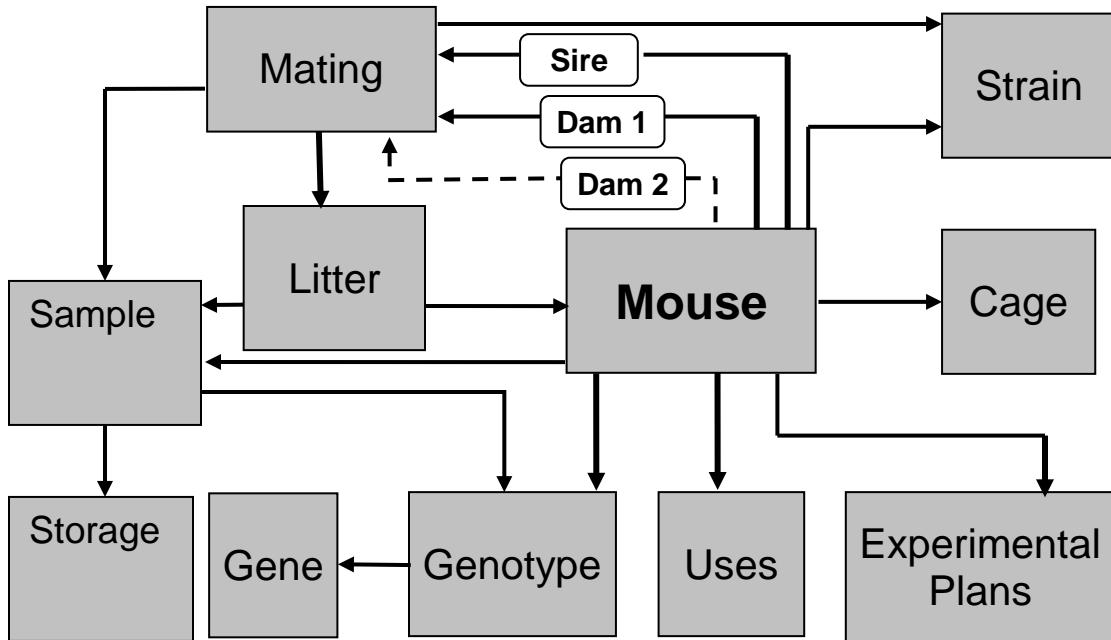
Since its initial release in 1998 it has been widely used within The Jackson Laboratory and adopted by a number of outside institutions. Some of the salient features that JCMS offers are:

- Tracking individual mice
  - Genotype
  - Uses and status
  - Pedigrees
  - Breeding performance

- Mating records
- Litter records
- Animal cage/pen management
  - Preformatted or user-designed cage cards with optional bar codes
- Experiment setup and tracking
  - Schedule mice
  - User designed metadata
- Track biological samples relating to mice, matings, or litters
- Bulk data entry
- Sophisticated query capabilities
- Data export to Excel
- Hard copy reports
- Handheld computer support
- User configurable

JCMS has an underlying data model that is independent of any specific database management system technology. The software engineers of Computational Sciences worked closely with the scientific research staff to derive a data model that was comprehensive yet flexible enough to fully satisfy the requirements of multiple research labs.

## 1.2 Types of Information Collected by JCMS



**Figure 1-1 Database Overview Diagram**

The above diagram shows the main types of information stored by the JCMS database. The mouse is the central entity. A mouse will have a specific strain and multiple genotypes associated with it. Each mouse is also associated with its current cage (pen). Two or three mice are set up in a mating (breeding pair or trio). Each mating can be tracked individually from pre-design to retirement based on a particular laboratory's workflow. Matings have their own strain, which may be different from the strains of the sire and dams. Each mating produces one or more litters. The litters of pups will become individual mice in the database at weaning/tagging. An embryo litter will produce biological samples. Technical matings involving samples (embryo, egg, sperm, or ovary), host dam, and sire may be recorded. These can produce samples or pups.

Two methods are provided for keeping track of experimental workflow (procedures, tests, routine activities, etc.) that may be scheduled. 1. The simple method is called **Mouse uses** and provides a method of assigning various “uses”, proposed dates, and simple data results to individual mice. 2. The more comprehensive method is called **Experimental plans**. This section of JCMS provides a method of defining experimental plans. Each plan contains various user-designed experimental tests, user-designed data fields, default data values, and scheduling information. Individual mice can have experimental data results associated with them and can be scheduled for various experimental tests and various experimental plans. See the section on [Experimental Plans](#) for a more comprehensive diagram and description.

Biological samples that are related to mice, matings, and litters may be entered. Users may also define their own types of samples. See the section on [Sample Tracking](#) for more information.

## 2 Getting Started

### 2.1 Quick Start

BEFORE BEGINNING, YOU MUST (1) Create a password for user ‘Admin’, (2) Create the user ‘mtsadmin’, and make it a member of the Admins group.

JCMS defines a security hierarchy with three levels of permission: Administrator, Owner, and Secretary. All users of JCMS are assigned into one of these groups. The Administrator (mtsadmin) has the overall responsibility for the database. Owners are usually the people who are maintaining the colony, including any breeding activities, and/or are performing experiments, which may include creating and storing biological samples. An owner has direct responsibility for groups of mice, matings, samples, and/or for experimental plans and data. An owner may have limited access to perform activities only with the mice, samples, experiments, that they “own”. Secretaries are users that have more limited security access to the database and normally are restricted to entering data, executing queries, and viewing reports.

#### 2.1.1 Administrator

One individual must be assigned to be the database Administrator (mtsadmin). The Administrator will be responsible for the installation and setup of the database and for creating other user accounts. The Administrator has full access to all data. The Administrator is not expected to be an IT professional, so they may need assistance from their IT department to do a networked installation.

The Administrator must complete the [Installation](#) and [Administrator Setup](#) sections of this manual before other users may begin. The other users are divided into two groups: Owners and Secretaries.

#### 2.1.2 Regular Users: Owners and Secretaries

Users have restrictions within the database depending on their security level. The Administrator (see above) has full access to the data and forms. Owners have access only to forms at the “owner” or “secretary” security level. These include the ability to enter and edit the mice, matings, samples, experimental plans, and experimental data that they “own”.

A secretary is a user who has only limited security access to the database and is restricted to forms at the “secretary” level. Secretaries usually do not have the authority to make changes to data. This type of user may request reports and run queries. A secretary is assigned to specific owners and may only enter data or make changes to mice, matings, etc. that belong to those owners.

It is possible for a user to be both an owner of their own mice and to act as the “secretary” for one or more other owners.

To get started, first the Administrator will provide a new user with either an owner or secretary logon. Then begin with the [User Setup](#) and [Basics on Using JCMS](#) sections.

### 2.2 Installation

#### 2.2.1 System Requirements

- Single user
  - PC computer with Microsoft Office 2007 or 2010; a version that includes an English language Microsoft Access.
  - MySQL 5.5

- Disk space requirements are minimal; 100 MB should be plenty to get started.
  - Memory requirements are minimal; however, large amounts of RAM will boost performance when JCMS has large amounts of data in it.
  - Bar Code 128 from Elfring Fonts Inc. [www.barcodingfonts.com](http://www.barcodingfonts.com) is required only if using the optional bar codes on cage cards or for sample labels.
- Multi-user
  - Either a client or file server computer with Microsoft Office 2007 or 2010; a version that includes an English language Microsoft Access.
  - A file server computer running MySQL 5.5
  - A network environment set up so that all client computers can access the file server.
  - PC computers, Macintosh computers, wireless handheld computers, and UNIX/LINUX computers may act as clients if supported by the network software.
  - It is recommended that network thin client software such as Citrix® Metaframe Client/Server be used if computers other than PCs are used.
  - Disk space requirements are minimal; 100 MB should be plenty to get started.
  - Memory requirements are minimal; however, large amounts of RAM will boost performance when JCMS has large amounts of data in it.
  - Bar Code 128 with a site license from Elfring Fonts Inc. [www.barcodingfonts.com](http://www.barcodingfonts.com) is required only if using the optional bar codes on cage cards or for sample labels.

## 2.2.2 Download the JCMS Files

Download from the Jackson Laboratory JCMS Web site ([colonymanagement.jax.org](http://colonymanagement.jax.org)) the Windows installer file **JCMSInstaller.msi** and place it on the desktop. Double click the file to run it. It will ask you to specify an installation folder. The default is C:\Program Files\The Jackson Laboratory, but a different folder may be used for the installation. At the same level as the installation folder, the subfolders \data and \docs are created (see diagram).



The following files are included in the folder named *docs*:

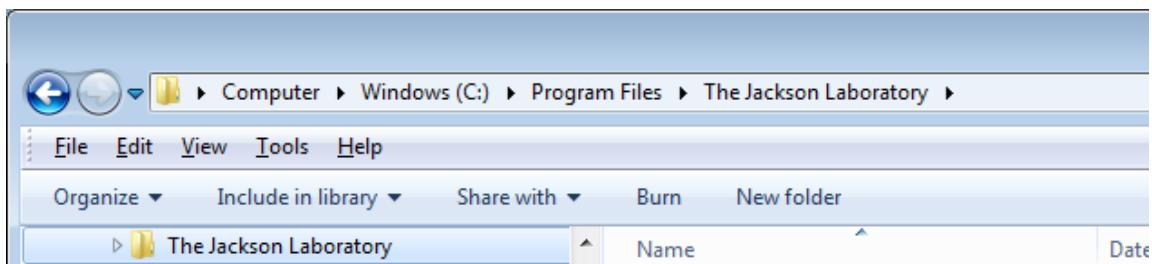
- ReadMe.pdf
- JCMS\_UserGuide.pdf
- JCMS\_UserGuide.docx
- JCMS\_ReleaseNotes.pdf
- JCMSInstall-Upgrade-guide.pdf
- License.txt

The data folder is used to hold the user files (referred to as documents) that are associated with the mice, mouse genotypes, samples, and/or experimental data in JCMS. JCMS does not incorporate the documents into its database. It moves a copy of the original file into this special "data" folder and maintains a link to the file in the JCMS database. You must change the JCMS\_DATA\_FILE\_DIRECTORY setup variable to match the path of the data directory.

### 2.2.2.1 Windows 7 and MS Access 2010

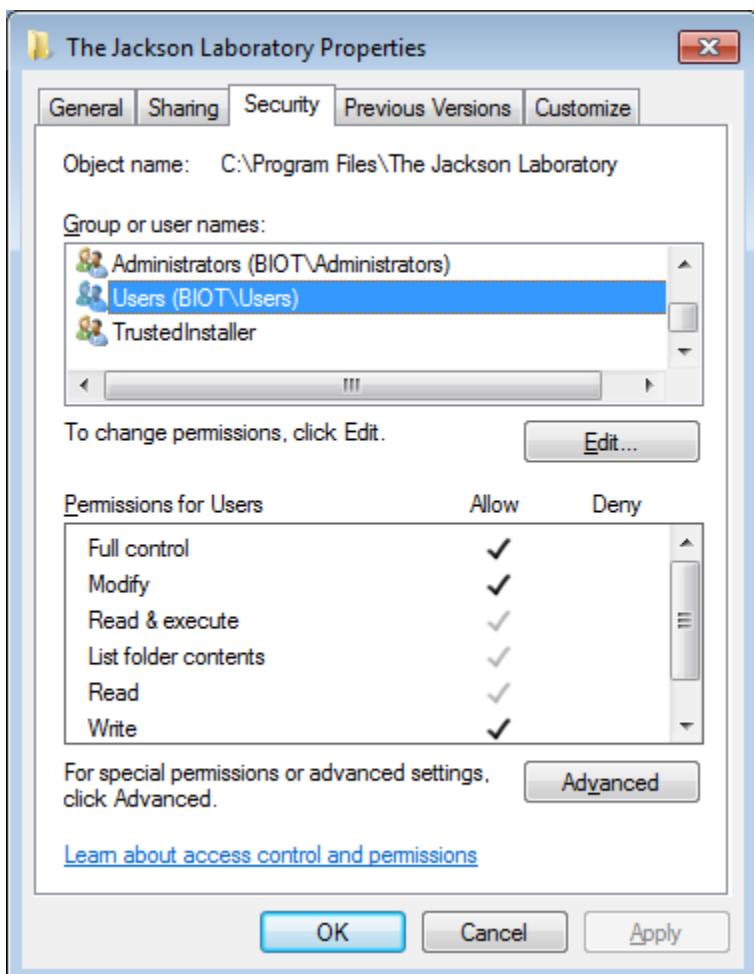
It is necessary to set the folder permissions in Windows 7 to allow the user "Full" access to the installation folder, otherwise everything opens in read-only mode and you can't complete the setup.

Using a file browser, right click the installation folder name and select "Properties".



**Figure 2-1 File browser**

If on the Security tab, the user does not have “full control” click the Edit button to change it.



**Figure 2-2 Setting full control**

Microsoft has moved the location of some of the features on the ribbon. The new locations are described in the MS Access 2010 configuration section below.

### 2.2.3 ReadMe Files

We suggest you first read the files **ReadMe.pdf**, **JCMS\_ReleaseNotes.pdf**, and online JCMS discussion forum for updated information. These should contain up-to-date information on changes not included in this documentation. We recommend you become a member of the JCMS discussion forums at: [http://community.jax.org/jcms\\_discussion\\_forum/default.aspx](http://community.jax.org/jcms_discussion_forum/default.aspx). This is a

place where JCMS users can post questions and answers plus review answers to previous questions. To receive notices of upgrades and other important user information we recommend also subscribing to the JCMS mailing list by sending an email to: [jaxcms-subscribe@lists.jax.org](mailto:jaxcms-subscribe@lists.jax.org). No subject or email body is required.

## 2.3 Configuration

### 2.3.1 Required Configuration Steps

BEFORE BEGINNING, YOU MUST (1) Create a password for user ‘Admin’, (2) Create the user ‘mtsadmin’ and make it a member of the Admins group.

Solutions to some minor problems that may occur are given at the end of this section (see the configuration issues section 2.3.13.) A new installation of JCMS will need special logon passwords created (system.mdw). The directions below are specific to different versions of MS Access. If the version of MS Access in use has password protection already set up, skip down to the *Setup the Administrator User* step.

### 2.3.2 MS Access 2007 Configuration

#### (1) Initialize Using Passwords for MS Access (Admin)

- Open the JCMS interface by double clicking on the desktop shortcut.
- Select the “Database Tools” tab.
- Select “Users and Permissions” from the ribbon.



- Select “User and Group Accounts”.
- Select “Change Logon Password” tab.
- Old password <blank>.
- New password “Admin” (or your choice).
- Verify (repeat your choice).
- Select “OK”.

#### (2) Setup the Administrator User (mtsadmin)

- Select the “Database Tools” tab.
- Select “Users and Permissions” from the ribbon.
- Select “User and Group Accounts”.
- Select “Users” tab.
- Select “New” button.
- Put “mtsadmin” in the Name field.
- Enter any random value in the Personal ID field.
- Select “OK”.
- Use the “Add>>” button to set user mtsadmin as

**Figure 2-3 Logon Password**

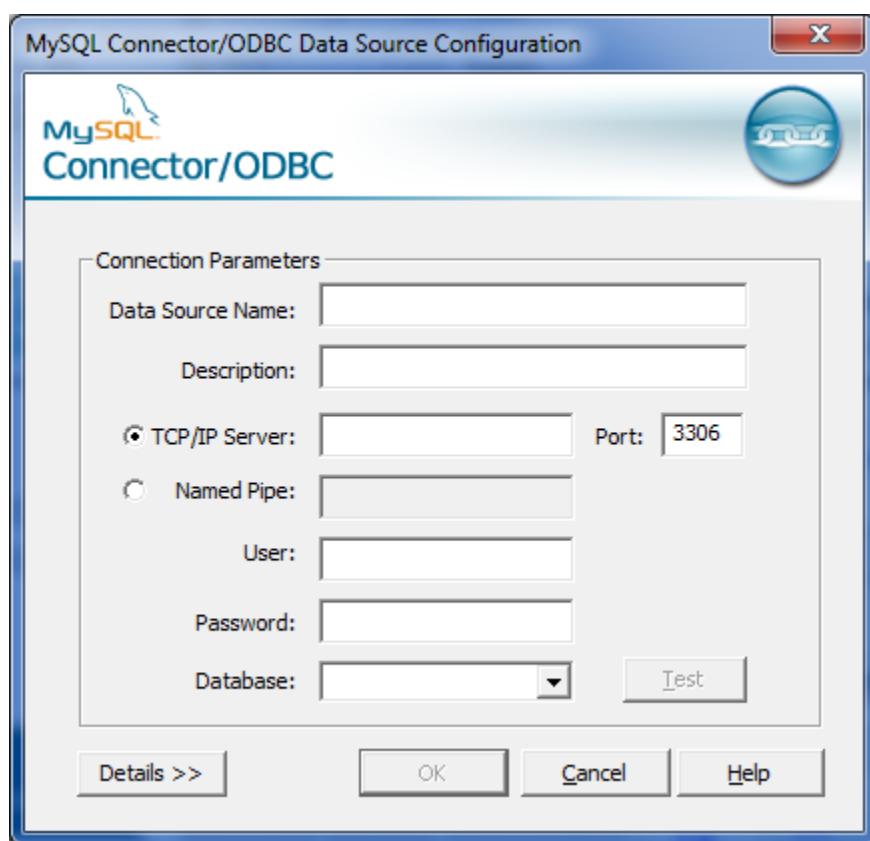
a member of the *Admins* and *Users* groups.

- Close MS Access.
- Restart JCMS and logon as mtsadmin. Leave the password blank.
- Now give mtsadmin a password using the same procedure as described above to give the Admin user a password. This is not required, but it is very strongly encouraged.

#### (3) Link the Database Tables (Only necessary if linking failed during install)

- On the toolbar, select the “External Data” tab.
- Select “ODBC Database” in the Import section of the ribbon.

- Select “Link to the data source by creating a linked table” radio button (DO NOT IMPORT the tables).
- Click on the “Machine Data Source” tab and click “New”
- Select either User Data Source or System Data Source and click “Next >”
- Select “MySQL ODBC 3.51” Driver from the driver list, click “Next” followed by “Finish”
- Fill out the information in the form. Data Source Name and Description are up to you, TCP/IP Server is the name of the machine with MySQL on it and Username/Password is your MySQL username/password (Figure 2-4).
- Select the DSN you created from the list of datasources and click OK.
- The “Link Tables” window opens (Figure 2-5).
- Select “Select all”.
- Select “OK”.
- Close JCMS.mdb and re-open it. The welcome window should appear as shown below with the proper release number (Figure 2-6). The installation is now ready to be used.



**Figure 2-4 Link to the database source**

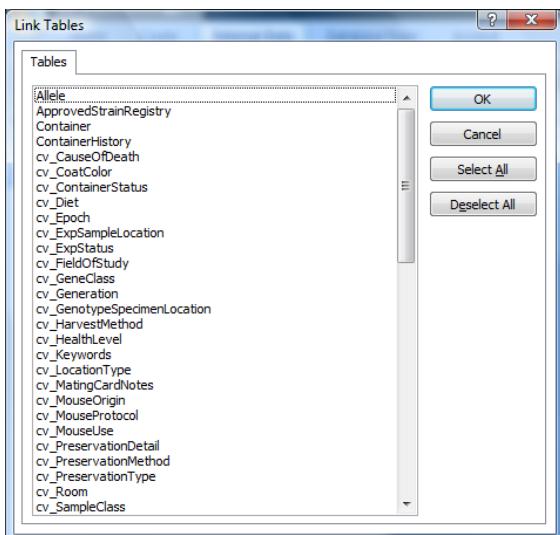


Figure 2-5 Link Tables dialog box

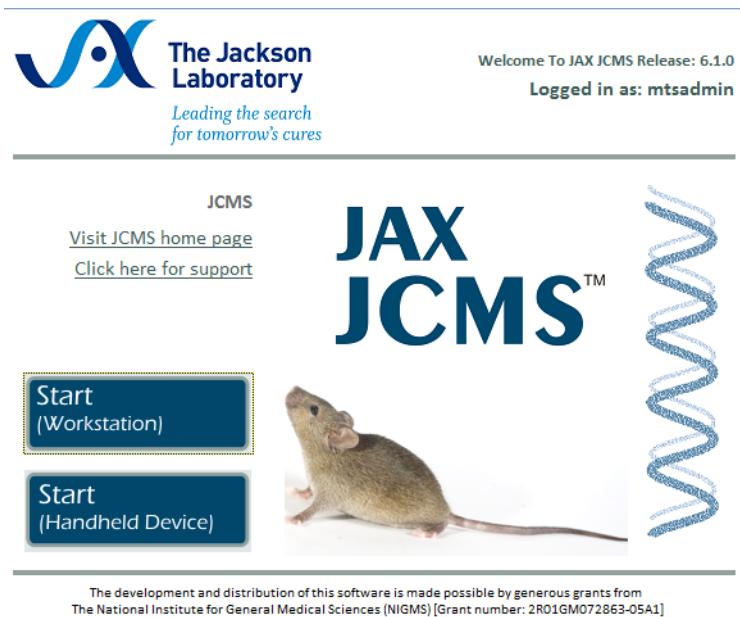


Figure 2-6 JCMS Welcome Window

### 2.3.3 MS Access 2010 Configuration

The steps are the same as indicated above for Access 2007. However, some of the features have been moved.

#### Users and Permissions

Use the red File tab to find the Users and Permissions tool.

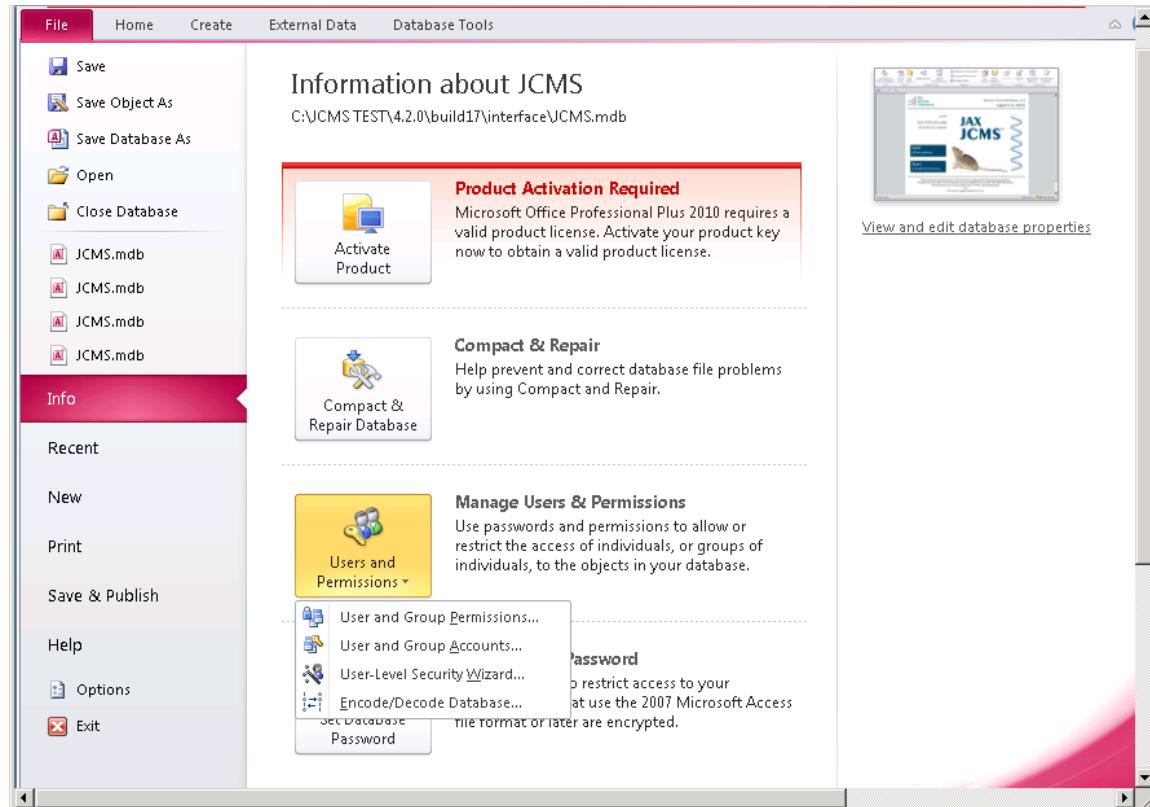


Figure 2-7 MS Access 2010 Users and Permissions dialog boxes

#### Linked Table Manager

Use the External Data tab to find the Linked Table Manager.



Figure 2-8 MS Access 2010 Linked Table Manager

#### Deleting Tables that are not linked

In MS Access 2010, certain configuration changes may require logging on as Admin, for example, if tables need to be deleted because they were imported instead of linked. Only the Admin logon can perform this function. However, remember that the Admin user logon is not able to open the regular JCMS forms.

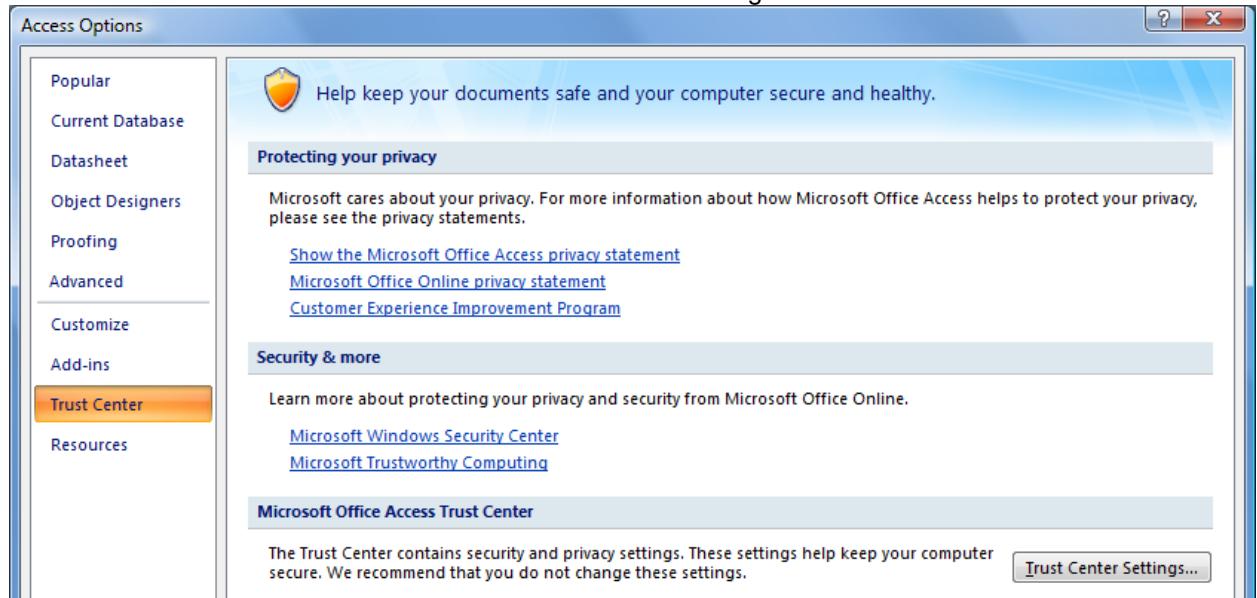
## 2.3.4 Removing Security Alerts and Warnings

In Access 2007, you may see a security alert.



The folder location of JCMS must be added to the “trusted locations” as follows.

- Select the Office button (top left of window).
- Select the Access Options button.
- Select “Trust Center” and then the “Trust Center Settings” button.

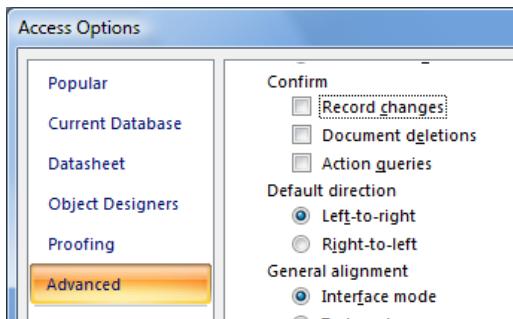


**Figure 2-9 Access 2007 Trust Center**

- Select “Trusted Locations”.
- Check “Allow Trusted Locations on my network”.
- Select “Add new location”.
- Browse to the folder containing JCMS.
- Check “Subfolders of this location are also trusted”.
- Select “OK” to save.

### 2.3.5 Configure some Database Options

- In Access 2007, click the Office button, select the Access Options button, and then the Advanced section on the left. Scroll down to the Confirm choices. Make sure the Confirm options are all un-checked as in Figure 2-10.



- Scroll further down the Advanced dialog to find the “open databases by using record level locking” and make sure it is not checked (Figure 2-11).

Figure 2-10 Access 2007 Confirm Options

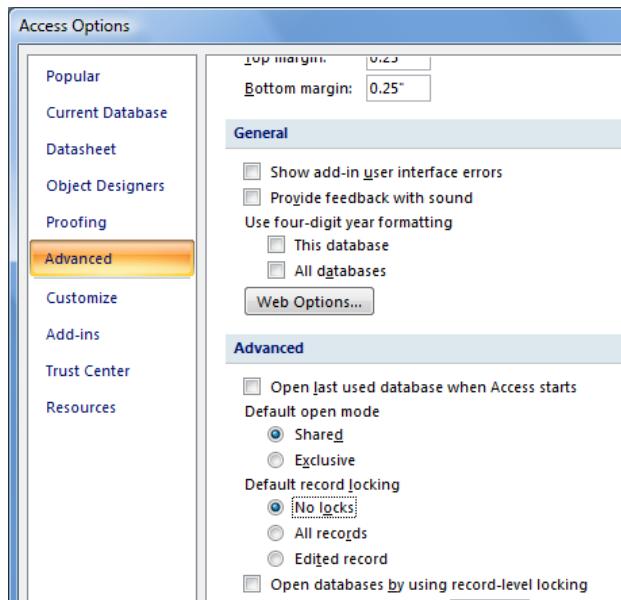


Figure 2-11 Access 2007 configure record-level locking

### 2.3.6 Creating a Multi-User Environment

JCMS is designed to be used by many people simultaneously (multi-user). To implement this, the database is placed on a computer (server) that all users may access. A copy of the interface may be placed on each user's computer (client) and the tables are then linked to the database.

However, we recommend that both the MySQL data source and JCMS.mdb (interface) files sit on one central machine (server). Each user's computer (client) will have a shortcut on the desktop that invokes the JCMS.mdb file through a share folder over the network. That way you don't have multiple copies of the JCMS.mdb file to maintain. The “best” way to access share folders from clients is by using Uniform Naming Convention (UNC) instead of direct paths with drive letters. Thus if your server machine is named *goofy* and the share folder is named *JCMS\_Share* then the UNC would be

\goofyJCMS\_Share

Often creating a multi-user environment involves moving an existing installation of JCMS onto a server. Therefore, the instructions for moving are included below. These may be skipped if the original installation is on the server and the goal is to create client shortcuts.

### 2.3.7 Moving JCMS to a New Location

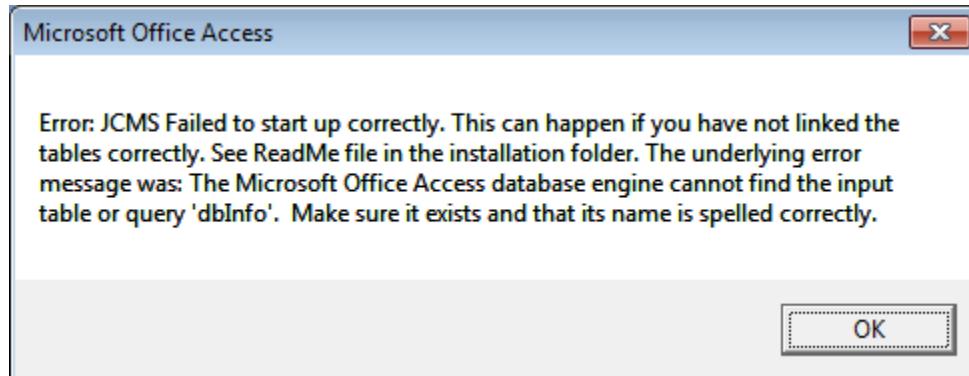
#### Access-MySQL

1. Install MySQL on the new computer.
2. Dump the MySQL JCMS database on the old machine and move the dump file onto the new machine.
3. Open a terminal and navigate to the directory containing the dump file.
4. In the terminal type in "mysql -uroot -p < dumpFileName.sql" where dumpFileName.sql is the name of the dump file you moved onto the new machine.
5. Relink the tables following the instructions in Configuration Section 2.3.
6. For support please visit our Forums at:  
[http://community.jax.org/jcms\\_discussion\\_forum/default.aspx](http://community.jax.org/jcms_discussion_forum/default.aspx)

The standard installation location for the JCMS.mdb file is  
C:\Program Files\The Jackson Laboratory\JAX-CMS\interface

If it is installed somewhere else, that's okay; just adjust the paths accordingly.

### 2.3.8 Re-linking the Tables



**Figure 2-12 JCMS Failed to start up correctly.**

The message in Figure 2-12 is displayed when the MySQL database is moved to a new location. Tell JCMS where to find the moved tables by re-linking them.

## Office 2007

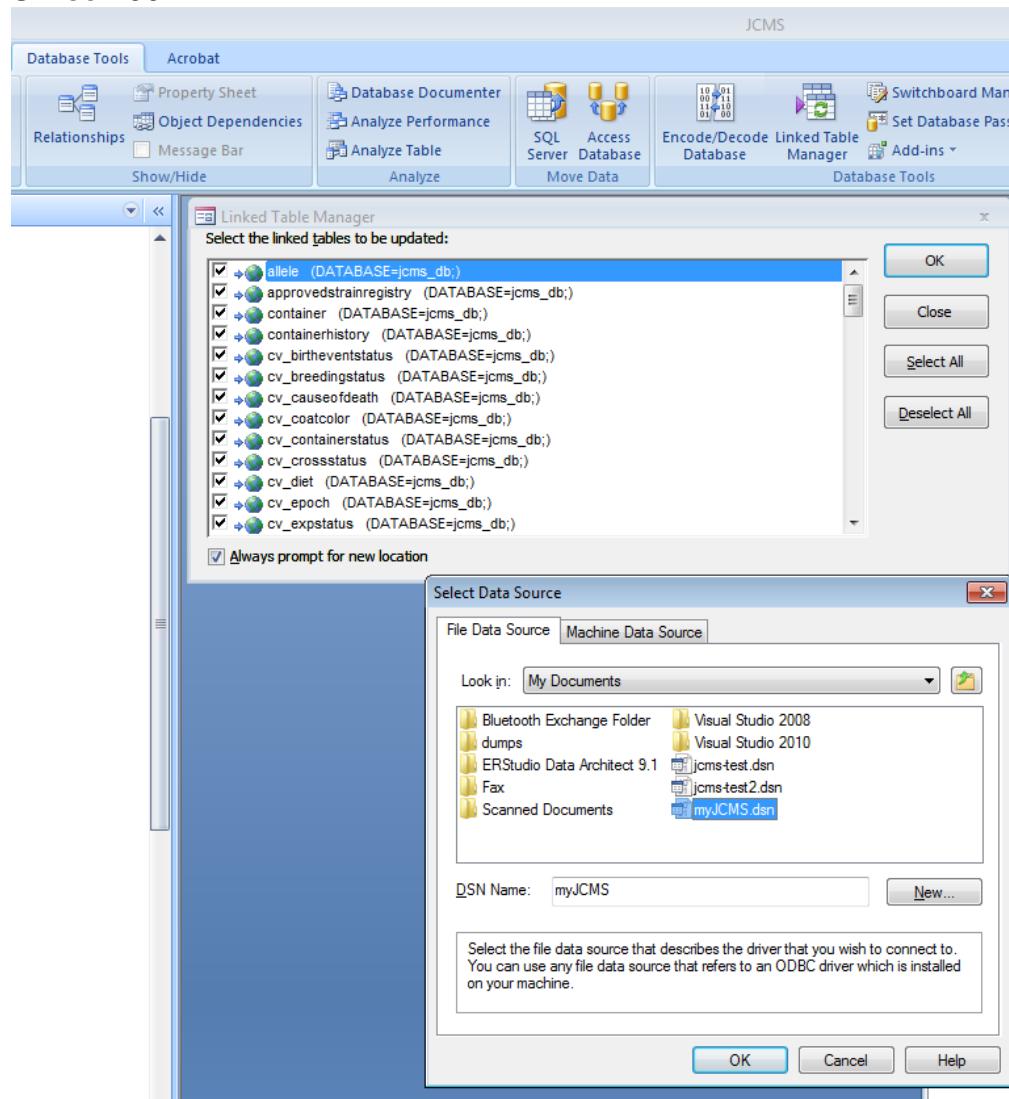


Figure 2-13 Office 2007 Linked Table Manager

- Select the “Database Tools” tab
- Select “Linked Table Manager” on the Database Tools ribbon.
- In the Linked Table Manager dialog box, click the “Select All” button.
- Check “Always prompt for new location”
- Select “OK”
- Browse to the new location of the data source; Select “Open”. If using a UNC such as \\goofy\JCMS\_Share it may be necessary to type the path.
- Select “OK” and the “Close”

## Office 2010

In Office 2010 use the External Data tab to find the Linked Table Manager and follow the steps above.

### **2.3.9 System.mdw File**

When the Administrator creates new users in MS Access, this information is stored in a “workgroup” file called **system.mdw** on the local computer (client). Every time a client computer is added, the Administrator has to set up mtsadmin and the users again in the system.mdw file on the new client machine. MS Access does provide a method of avoiding repeating this setup. The database must use a special workgroup file instead of the default system.mdw. We recommend naming this file JCMS.mdw. This file is set up once on the file server and contains all the users’ MS Access logons. The Administrator can set up a new user from any machine and only has to do it once.

Locate the system.mdw file currently in use, copy it into the JCMS installation folder, and rename it.

Hint: To locate the system.mdw file, use advanced search options to search system folders, hidden files, and subfolders. It can typically be found in:

Windows XP: C:\Documents and settings\<your user name>\Application Data\Microsoft\Access  
Vista and Windows 7: C:\Users\<your user name>\AppData\Roaming\Microsoft\Access

### **2.3.10 Installing a Multi-User Client**

Let’s assume that a copy of the system.mdw file has been put it into the JCMS installation folder and renamed JCMS.mdw. Of course, the mdw file can be put anywhere and named anything you like.

When MS Access is started, it must “know” to use the special workgroup file. There are several ways of implementing this depending on the version of MS Access and if other MS Access applications are in use.

Assuming everything is installed in the standard spot and JCMS\_Share is the shared folder, then the next thing to do is create a shortcut for JCMS.mdb that uses the JCMS.mdw workgroup file. This is easy, first create the shortcut (we’ll assume you can do that). Open the shortcut properties by right clicking the mouse on the shortcut and select properties. Now just edit the shortcut properties so that the target string in the shortcut will look like this

```
Target = "C:\Program Files\Microsoft Office\OFFICE12\MSACCESS.EXE"  
"\goofy\JCMS_Share\JCMS.mdb" /wrkgrp "\goofy\JCMS_Share\JCMS.mdw"
```

Now just put a copy of the shortcut on any client's desktop.

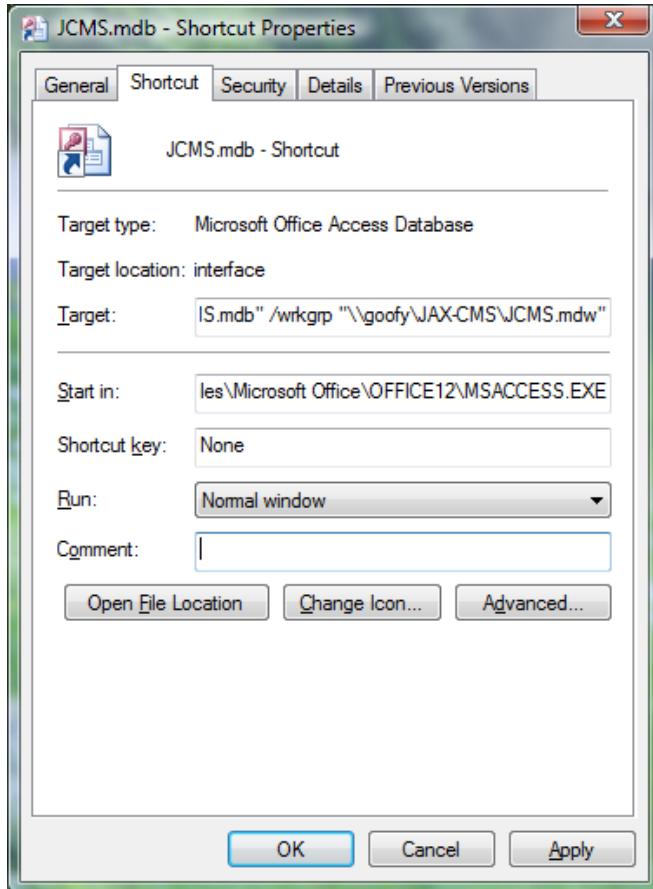


Figure 2-14 JCMS client shortcut

#### Notes:

- The client machine must have MS Access installed on it.

### 2.3.11 Backing up

Make sure to have a reliable and routine back up strategy in order to avoid loss of data. See section 22.4 for suggestions.

### 2.3.12 Installing the Bar Code 128 Font

JCMS offers optional bar codes on cage cards or for sample labels. To use this feature, Bar Code 128 with a site license from Elfring Fonts Inc. ([www.barcodingfonts.com](http://www.barcodingfonts.com)) will need to be installed on any clients that print cage cards, sample labels, or read bar codes. Follow the manufacturer's installation instructions.

P.I.	PI Name 555-1212 (office)	Owner OWN1	
Activation date: 12/20/2005	Sec:	Count / sex: 3 F	Pen# P57
Put card note here (dbsetup)			
Expt #:			

Figure 2-15 Sample of a bar code for the pen ID number

### **2.3.13 Configuration Issues and Answers to Common Problems**

#### **2.3.13.1 File MSCOMCT2.OCX is missing**

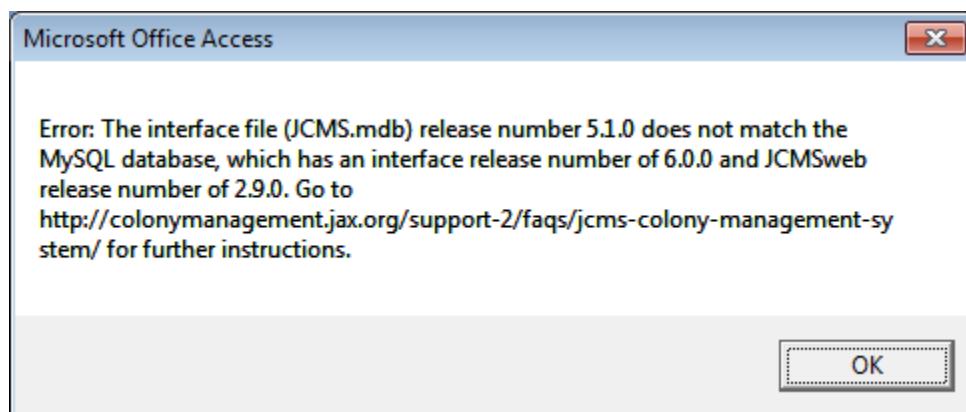
#### **2.3.13.2 File MSCOMCTL.OCX is missing**

#### **2.3.13.3 File MSFLXGRD.OCX is missing**

#### **2.3.13.4 Error messages about DTPicker**

Some versions of MS Access may not have a necessary Windows file installed. The JCMS installer should have installed these files for you. To add the controls to a client machine, look on the JCMS website (<http://colonymanagement.jax.org/support-2/jcms-user-tips/>.) User Tip #4 has a link to download a special installer called JCMS\_Controls\_v4.1.msi. Run this on all clients with this problem.

#### **2.3.13.5 When JCMS is started it states the release numbers do not match.**



**Figure 2-16 Error: release numbers don't match**

This message indicates that the linked database tables are for a different version of JCMS. There are several possible causes, which include opening an old version of the JCMS access interface, opening a new version of the access interface that needs to have the tables relinked, or that upgrading JCMSweb has upgraded the tables and now an upgraded version of the Access interface is needed. The link to the online [FAQ](#) page will provide the most up-to-date instructions.

To relink the tables in Access 2007 use the Database Tools tab. The Linked Table Manager is in the Database Tools section of the ribbon. In Access 2010 the Linked Table Manager is found on the External Data tab, Import & Link section.

#### **2.3.13.6 The MS Access menu bar has virtually no options on it.**

- MS Access has an option that allows menus (such as the tools menu) to be displayed or not. If the menu bar shows only the bare minimum of buttons, then enable full-menus as follows:
  - Close JCMS (if it isn't already)
  - Start JCMS and hold down the shift key as it is starting up. A full set of menu options should now appear.
  - Go to the Tools/Startup menu option and check the box that says *allow full menus*.

#### **2.3.13.7 The Welcome window is not displayed**

If all tables have been linked or re-linked and no error messages are displayed at startup it is possible that the startup form needs to be reset.

- Select the “Office button”
- Select “Access Options” button
- Select “Current Database”

In the application options “Display Form” field enter “MTS Welcome”. Select “OK” to save. Close MS Access and start the application again.

#### **2.3.13.8 I need to delete the tables and the delete button is grayed out**

In MS Access 2010, certain configuration changes may require logging on as Admin, for example, if tables need to be deleted because they were imported instead of linked, that only Admin can perform this function. However, remember that the Admin user is not able to open the regular JCMS forms.

#### **2.3.13.9 Didn't find your question here?**

Also check the FAQs in section 21, the online [JCMS Discussion Forum](#), and the online colony management website [FAQs](#).

## 3 Administrator Setup

JCMS uses the logon name **mtsadmin** as the database Administrator. Anyone accessing JCMS as mtsadmin can perform Administrator functions in JCMS.

### 3.1 Owner and Secretary Accounts

JCMS uses a very simple user account system. There are two rules to know.

- 1) Every user of JCMS must have a logon to MS Access.
- 2) Each logon name that JCMS uses must be defined as either an owner (of mice) or a secretary. The logon name and the JCMS owner/secretary names must be identical.  
To add owners and secretaries you must be logged on as mtsadmin.

#### 3.1.1 Rule 1: Every User of JCMS Must Have a Logon to MS Access.

An Administrator (user *Admin* or *mtsadmin*) can set up new MS Access logon accounts for starting MS Access. Setting up new MS Access accounts is not a JCMS function; it is an MS Access function. JCMS uses the logon name of the user to determine if the user is an owner, secretary, or Administrator (mtsadmin.)

##### 3.1.1.1 How to Add a New User Logon.



Figure 3-1 Dialog Box: Personal ID

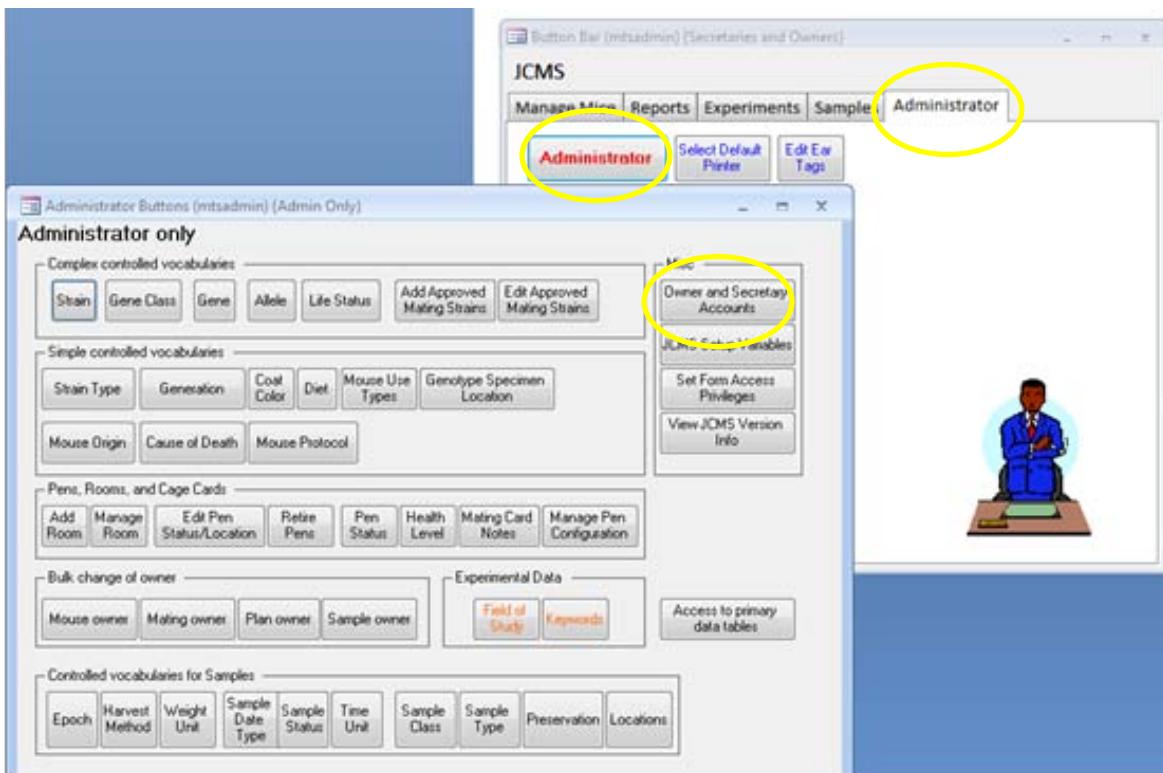
With MS Access running, select the Database Tools tab. In the Administer section of the ribbon select Users and Permissions, and then select User and Group Accounts.

Select the Users tab and then the “New” button. Note: the Personal ID (PID) is not used by JCMS at this point. It is recommended to use the logon name also for the personal ID and add numbers as necessary to make it long enough. Keep all logon names short with no spaces, commas, semicolons, or quotes in them; this is especially important for mouse owners since owner logon IDs are stored with each mouse. Do not make owners and secretaries part of the Admins group; they are part of the Users group by default. Users can later set their own passwords by logging on MS

Access and setting the password from this same menu. When a user is first added, his/her password is blank.

#### 3.1.2 Rule 2: Each logon name that JCMS uses must be defined as either an owner (of mice) or a secretary.

This rule applies to owners and secretaries. User mtsadmin is neither an owner nor a secretary. The Administrator needs to tell JCMS if the user is an owner or a secretary or both. Owners own mice, matings, experimental plans, experimental data, and samples. Secretaries are associated with one or more owners and have limited edit access to their associated owner's data. NOTE: Owners can be secretaries of other owners and thus have limited editing capability for other owners.



**Figure 3-2 Form: Administrator Buttons**

To set up a user (other than mtsadmin) as an owner or secretary, open JCMS. Open the Administrator button bar from the Administrator tab and select the button labeled *Owner and Secretary Accounts*. From this account maintenance form owners and secretaries can be added or deleted (see example below.)

**Figure 3-3 Form: Owner and Secretary Accounts**

NOTE: the names entered in this form must agree exactly with the names used for logging on to JCMS. NOTE also, do not delete owners that have mice in JCMS as their mice will no longer be accessible (unless you add the owner back using this form). Bulk changes of the owner of mice, matings, samples, and experimental plans may be done using the buttons provided on the Administrator button bar.

## 3.2 Changing the Forms Used by Secretaries and Owners

JCMS allows the Administrator to configure access to the forms in the interface by the type of user. For example, secretary users may be set up to only use certain add forms and none of the edit forms. Note that some forms are required by JCMS to be available to all users or only to the Administrator.

To change the access or “privilege” level for one or more forms, use the Set Form Access Privileges button on the Administrator button bar.

Form name	Privilege level	Description
Bulk Add Genotype	Owner	Add Genotype: Add new genotype information for a group of mice.
Add Genotype	Owner	Add Genotype: Add new genotype information for a particular mouse.
Add Litter w/Pups	Sec	Add Litter: Add a litter and optionally create mouse records for each pup at the same time.
Add Litter	Owner	Add Litter: Add a litter.
Do Matings	Admin	Add Mating: Add a mating record after it has been done in the mouse room.
Design Matings	Owner	Add Mating: Create a mating “on paper” before doing it for real.
Activate Matings	Owner	Add Mating: Finalize the creation of a mating that was designed previously by entering the mating date.
Import Mice	Owner	Add Mice: Add a group of mice at once. All must be similar.
Wean	Owner	Add Mice: Add a new mouse from a specific JCMS litter.
Add Mouse	Sec	Add Mice: Add new mice, one at a time.
Add Use	Owner	Add Mouse Use: Add a new use for a particular mouse. Uses can store small amounts of associated data.
Bulk Add or Edit Use	Owner	Add Mouse Use: Add or edit mouse uses for a group of mice.
Edit Genotype	Owner	Edit Genotype: Make changes to the genotype information for a mouse.
Edit Litter	Sec	Edit Litter: Make changes to the information about a litter of mice.
Edit Matings	Owner	Edit Mating: Make changes to the information about a mating.
Bulk Change Life Status or Diet	Owner	Edit Mice: Make a change to either the life status or diet of a group of mice.
Edit Mouse	Owner	Edit Mice: Make changes to the information about a mouse.
Edit Use	Owner	Edit Mouse Use: Make changes to the information about a mouse's use and add or update associated data.
Bulk Add Mice from Query	Owner	Experiment: Add a group of mice into an experimental plan or test - mice are selected with the mouse query form.
Add Exp Plan	Owner	Experiment: Add a new experimental plan.
Add Exp Test	Owner	Experiment: Add a new experimental test.
Add Test Type Defaults	Owner	Experiment: Add a new set of default values for the data corresponding to a particular test type.
Add Test Type - Data Description	Owner	Experiment: Add a new test type and describe the data to be collected.
Bulk Add Data to Specific Test	Sec	Experiment: Add data values that are the same to a group of mice all at once.
Add Exp Data	Sec	Experiment: Add experimental data for a mouse.
Edit Test Type	Owner	Experiment: Make changes to a test type.
Manage/ Edit Exp Plan	Owner	Experiment: Make changes to an experimental plan or manage the mice and tests that are part of the plan.
Edit Exp Test	Owner	Experiment: Make changes to an experimental test.
Edit Test Type Defaults	Admin	Experiment: Make changes to the default data values for a particular test type.
Edit Exp Data	Sec	Experiment: Make changes to the experimental data associated with a mouse.
Add Litter w/Mice	Sec	Handheld: Add a litter and add mouse records for the pups at the same time.

Figure 3-4 Form: Edit Form Privileges

All forms listed may have the access privilege level changed to secretary (Sec), owner (Owner), or Administrator (Admin). The Administrator may use any form; owner level forms are restricted to only owners or the Administrator; secretary level forms may be used by secretaries, owners, and the Administrator. Any change made to the access privilege level for a form will NOT take effect immediately. You must quit and restart JCMS for the change to be implemented.

## 3.3 Initializing Controlled Value (CV) Tables

Before any mice can be added into the database, values for certain required fields must be established. These values are kept in a set of tables referred to as the controlled value or CV tables. A few of the very simplest controlled vocabularies are stored in the user interface forms instead of in tables and cannot be changed (for example, sex can only be “m”, “f”, or “-”).

Controlled vocabularies are changed from the **Administrator** button bar (see Figure 3-2) and may only be changed by the mtsadmin user.

### 3.3.1 Simple Controlled Value Tables

The **Strain Type**, **Generation**, **Coat Color**, **Diet**, **Genotype Specimen Location**, **Mating Card Notes**, **Mouse Origin**, **Cause of Death**, **Mouse Protocol**, **Field of Study**, and **Keywords** buttons each open a table for adding or deleting choices from the lists. These lists of choices appear in pull-down menus on many forms. These simple CV terms should be kept minimal in length since they are copied into the database records.

The following list shows all JCMS simple controlled vocabulary tables.

- Generation: valid mouse generation terms (e.g. F1)
- Diet: list of diets (e.g. 4%)
- Coat Color: list of valid coat color names
- Genotype Specimen Location: specifies where genotype samples are located (e.g. a freezer number)
- Cause Of Death: terms that a user can select to specify why a mouse died
- Mating Card Notes: some mating cards allow notes to be printed on them
- Mouse Origin: list of terms that specify where mice can originate, including the local colony. Every mouse has an origin. When mice are brought in from other institutions, they too should be listed in this table.
- Strain Type: Used in the strain table to specify the type of strain (e.g. congenic)
- Mouse Protocol: Some institutions need to link mice to protocol numbers
- Field of Study: Used to associate experimental plans with each other
- Keywords: Used to describe an experimental plan and may include keywords used for publication of the results

Add or Delete Coat Color (mtsadmin)	
Coat Color	Description
Only the description field is editable, to change a coat color delete it and add a new one.	
Agouti	Agouti
Albino	Albino
Black	Black (non-agouti)
BrAgouti	Brown agouti
Dominant	Dominant

**Figure 3-5 Sample Simple CV Table**

A value in the simple controlled vocabulary tables cannot be changed, instead add or delete a value. Thus, to change a value in one of the tables, delete it, and then add the value back with the change. Deleting a choice will not remove it from any data records where it has already been used within the database; the choice is simply removed from the list of current possible choices.

**IMPORTANT NOTE:** think carefully before deleting (or changing) a controlled vocabulary value if it has already been used. Controlled vocabularies may be used in queries, and thus the ability to find records based on a CV term may be lost if the term is deleted. This could result in a loss of what is known in database terminology as referential integrity, a mortal sin in relational database systems. This means that some data items will reference no longer existing data objects (the deleted CV term) resulting in undefined consequences.

**To add a choice** to a table, scroll to the bottom of the list. Type the new choice into the blank space at the bottom.

**To delete a choice** from the table, click on the box at the left side of the row in the table. The whole row will be highlighted. Press the delete key. A dialog box will ask for confirmation that the record should be deleted.

Some of the CV tables have two fields, the choice and a description. This description field may be edited.

DATA CONSTRAINTS on CV terms: CV terms **cannot** contain single or double quote characters, commas or semicolons. Thus, “Fred’s chair” would be illegal because of the “ and ’ symbols.

Some CV tables also come with pre-set values that may be deleted or added to.

Add or delete strain type terms	
	strainType
	coisogenic
	congenic
	conplastic
	consomic
	F1 hybrid
	F2 hybrid
	inbred
	outbred
	recombinant congenic
	recombinant inbred
	wild-derived inbred
▶	

Add or Delete Generation Name	
	generation
▶	F01
	F02
	F03
	N01
	N02
	N03
	N04
	N05
	N06
	N07
	N08
	N09
	N10

**Figure 3-6 Forms: Default Strain Types and Generations**

The **Strain, Gene Class, Gene, Allele, and Life Status** buttons open forms that are more complicated than those used for maintaining the simple CV tables described above. These are described in more detail below. **Owner and Secretary Accounts** are described above. Other sections are dedicated to the Pen, Room, and Cage Card terms and the Samples terms.

### 3.3.2 Strain Table

Strain names often use unusual characters to delimit fields in the strain name. Some characters may confuse some of the JCMS displays. It is best to avoid using the following characters in the strain names: double quote ("), single quote ('), and comma (,).

The screenshot shows the 'Add or Edit Strain' window. Key fields include:

- \*Strain: B6.129P2-Apoe<tmUnc>/J
- \*Active: Yes
- \*Strain Status: A
- JR # / Stock #: 2052
- Frozen Embryos: 0
- FE Max Generation: 0
- \*Frozen Sperm: 0
- FS Max Generation: 0
- \*Frozen Ovaries: 0
- FO Max Generation: 0
- Section: (empty)
- Card color: (empty)
- Strain type: congenic
- Comments: (empty)

**Line Viability Constraints**

YELLOW	RED
Minimum Number of Males:	Minimum Number of Males:
Minimum Number of Females:	Minimum Number of Females:
Maximum Age in days for Males:	Maximum Age in days for Males:
Maximum Age in days for Females:	Maximum Age in days for Females:

Record: < 1 of 17 > No Filter Search

**Figure 3-7 Form: Add or Edit Strains**

There are three “name” fields in each strain record: Strain, Abbreviation, and Formal Name. Currently, JCMS only uses the “Strain” field. The other two fields can have information added to them for reference purposes, but it will not be used anywhere in the JCMS system.

By default, a strain is active. Set *active* to “No” if a strain is no longer in use. Some of the add forms have an option to display only active strains as choices. A strain may be changed back to active when it is in use again. Note that the *strain status* of A (active), F (frozen), D (discarded), and C (see comments) is not used to limit the strains displayed as choices.

Use the navigation buttons to move from one strain to another. The strains are listed in alphabetical order. Or click in the strain field and use the find icon (it looks like binoculars) to search for a particular strain. If the exact strain is not known, use the “Match start of field” option.

To **add a strain**, click on ▶ \* or move to the last record. Only the strain name and status are required. The strain will not be added unless values are entered into those fields. Adding will occur when you navigate to another existing record or press ▶ \* to move to a new record.

To **edit a strain**, type in the field and change it. If a mistake is made, undo using the standard undo icon. Note that pressing the ESC key acts the same as the undo icon.

You cannot **delete a strain** from the database unless it is not associated with any matings or mice. If the strain name is incorrect, edit the name. Everywhere in the database where this strain is used, the name will change.

The strain name field contains the name that will appear for all the mice or litters. The strain abbreviation is the shorthand that may be used to refer to this strain within a user group. The formal name is the one assigned by the nomenclature committee. JR # stands for Jackson Laboratory Resource number. Stock # and JR # is the same field. This field may also be called JRnum. Users may enter their own stock # when not using Jackson Laboratory strains or leave the number as 0 if not using a numbering system. This number is always associated with a particular mouse strain. The strain nomenclature may change over time, but the Stock # will remain the same. The Stock # may be useful in the future to look up nomenclature changes to the strain name. Additional data is stored about whether or not frozen embryos, sperm, and ovaries are available.

### 3.3.2.1 Strain table fields

- Strain: name of strain as it will appear on JCMS forms and reports
- Formal Name: formal name of the strain assigned by the nomenclature committee
- Strain Abbr.: a short string abbreviation for the strain name for use in lab only
- Active: indicates if this strain is available to use (yes/no)
- Strain status: a controlled vocabulary that specifies how the strain is currently maintained (e.g., frozen, on shelf, etc.)
- Min tag, Max tag, Last tag: specifies ear tag numbers as discussed below
- JR # / Stock # / JRnum: the Jackson Laboratory Resource Number or user's stock number
- Frozen Embryo: number of frozen embryos
- FE Max Gen: maximum generation stored as frozen embryo
- Frozen Sperm: quantity of frozen sperm
- FS Max Gen: maximum generation stored as frozen sperm
- Frozen ovaries: number of frozen ovaries
- FO Max Gen: maximum generation stored as frozen ovaries
- Section: section number in the mouse room where this strain is kept
- Card color: color of cage cards used for this strain (displayed on several forms from which cage cards are printed)
- Strain type: from Strain Type controlled vocabulary table, a term that specifies this strain type such as "congenic"
- Comments: free form text
- For *Line Viability Constraints* values, see the next section: **Mouse Line Viability**.

### 3.3.3 Mouse Line Viability

The mouse line viability function provides the colony manager the ability to configure warning levels when the breeders in her colony are:

- approaching a critical age threshold or
- when the numbers of potential breeders have fallen below certain thresholds.

Thresholds are set at the strain level. Therefore the configuration is done by the super user, *mtsadmin*, within the Strain controlled vocabulary function.

The **Strain** configuration function is accessible from the **Administrator** button bar.

There are two levels of thresholds: yellow (warning) and red (dire).

To bring up the report, click on the **Line Viability Report** button on the reports tab.

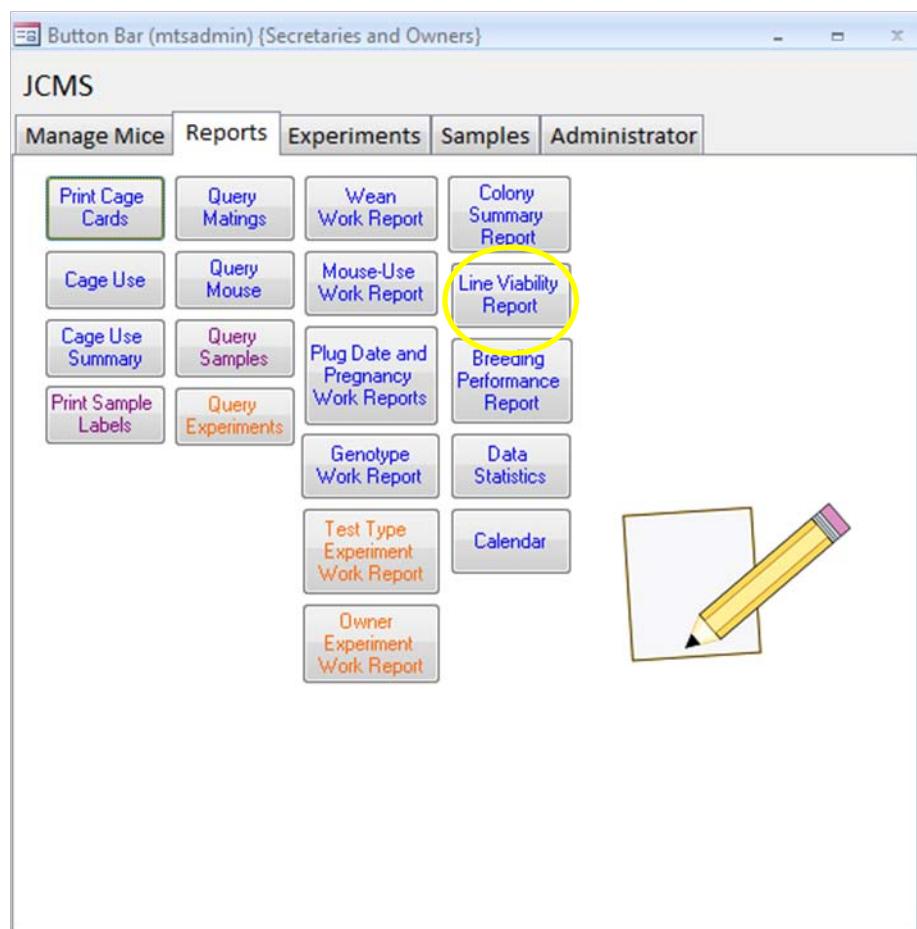


Figure 3-8 Open Mouse Line Viability Report

## Line Viability Report

Strain: B6D2F1/J		Warning Level	Not At Risk
Age Limits	Red	Yellow	
Female	100	70	
Male	100	70	
<b>Minimum Number Mice Required</b>			
	Red	Yellow	
Female	25	10	
Male	25	10	
<b>Number of Mice in Inventory</b>			
	# Yellow Mice (between yellow and red age)	# Green Mice (<= yellow age)	
Female	25	20	
Male	25	20	
Strain: BALB/cJ		Warning Level	YELLOW
Age Limits	Red	Yellow	
Female	65	35	
Male	65	35	
<b>Minimum Number Mice Required</b>			
	Red	Yellow	
Female	4	9	
Male	4	9	
<b>Number of Mice in Inventory</b>			
	# Yellow Mice (between yellow and red age)	# Green Mice (<= yellow age)	
Female	0	10	
Male	30	0	

**Figure 3-9 Example mouse line viability report**

### 3.3.4 Ear Tag Ranges

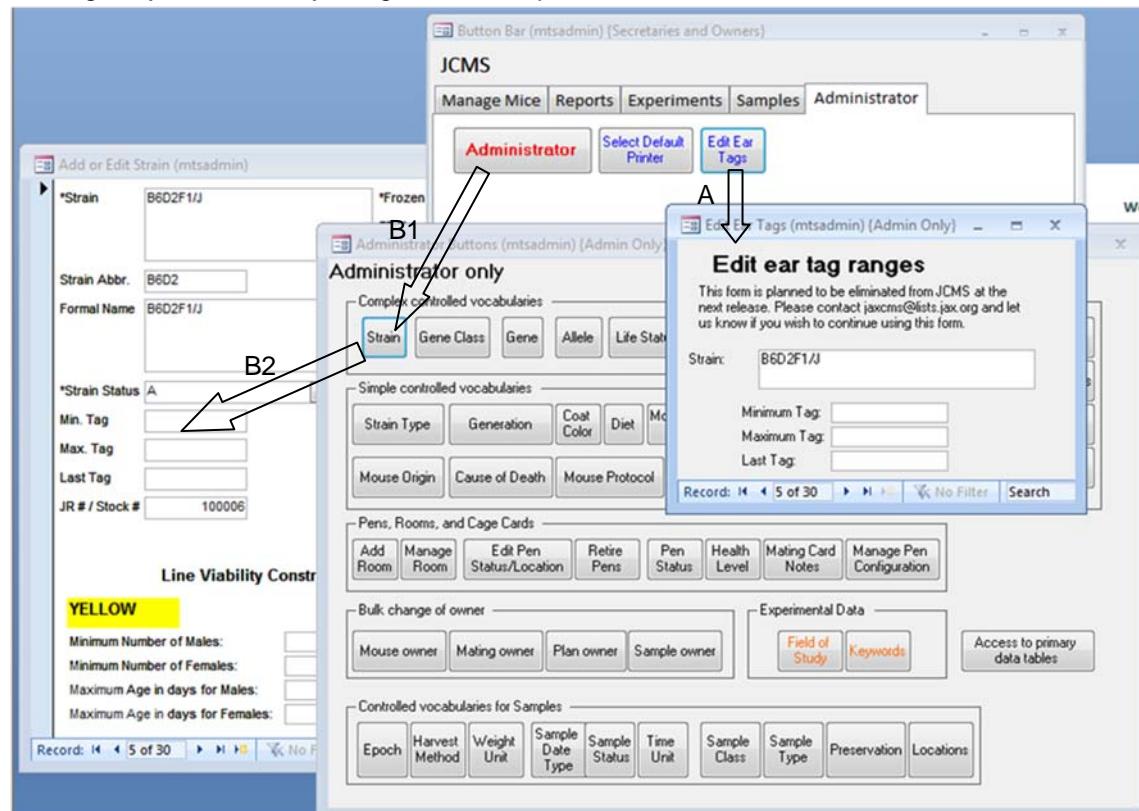
If the pups from a particular strain are all given ear tags from a pre-set group of tags, then the ear tag ranges can be used to keep track of which ear tags are used as a function of strain. This range is called the **minimum tag** and **maximum tag**. The **last tag** field is used to record the highest tag number from that range that is currently in use.

To update the ear tag range, use the **Edit Ear Tags button** on the Administrator tab. To find the correct strain, use the navigation buttons or click in the strain field and use the find icon (it looks like binoculars). If you do not know the exact strain, use the “Match start of field” option.

If a range has been completely used, either blank it out (if no new range has been assigned) or enter the new range, replacing the old one.

All fields in the strain table, with the exception of the strain, stock #, section, and color fields are simply for user tracking purposes. The four fields that JCMS uses may be printed on cage cards, or will show up in pull down lists.

There are two places where ear tag ranges may be entered. To eliminate this redundancy, it is proposed to eliminate the Edit Ear Tag ranges form. If this form is important to you, please send a message to [jaccms@lists.jax.org](mailto:jaccms@lists.jax.org) as soon as possible.



**Figure 3-10 Ear tag ranges**

Method A: use the edit ear tags button (A) on the Administrator tab.

Method B: use the Administrator button (B1) on the Administrator tab and then the Strain button (B2) to open the Edit strain form.

### 3.3.5 Approved Strains for Matings

The JCMS Administrator has the ability to create a table of approved litter strains to limit the mating forms to pre-approved strains. This will establish better breeding control and prevent inadvertent errors. If necessary, the user is able to override the restriction. To implement using approved strains, the Administrator must change JCMS\_ENFORCE\_APPROVED\_MATINGS in the setup variables to "True".

An approved litter strain field is a function of the dam strain and the sire strain. A sire strain and dam strain combination is **not** unique. That is, a sire and dam strain combination may result in multiple litter strains. The special case of when the sire and dam strains are the same will result in that strain always being 'approved' and need not be entered in the database. Click the **Add combination** button to create a new approved strain.

The **New combination w/ reversed parent strains** button tells the software to reverse the dam and sire strains and create a new record in the database, that is, switch the dam and sire strains but produce the same litter strain. This must be done to make that combination also valid.

The **Swap dam/sire** button is a convenience that will reverse the strains listed in the Dam Strain and Sire Strain boxes.

Dam Strain	Sire Strain	Litter Strain
BALB/cJ	DBA/2J	New mutant
C57BL/6J	C3H/HeJ	New mutant
DBA/2J	BALB/cByJ	New Mutant 2
BALB/cByJ	DBA/2J	New Mutant 2

**Figure 3-11 Add Approved Strain Form**

Approved strain records can be active or inactive. Inactive records are ignored as candidates for litter strains. They are kept in the database for documentation purposes and possible future use.

Use the **Edit Approved Strains** button to make combinations inactive or to delete combinations.

### 3.3.6 Life Status

The life status controlled vocabulary has three fields. The first field is the lifeStatus field. Use one or two letter entries to specify a life status. The second field gives a description of the life status so users will know which one to select from pull-down menus. The third field, “exitStatus” is a Boolean (true or false) field that specifies if the associated life status implies that the mouse has exited the colony. JCMS uses this Boolean value to enforce rules about whether or not a cause of death or an exit date may be entered. JCMS will prevent entering an exit date if the mouse life status is not an exit status.

The Life Status controlled vocabulary table must include the following pre-set terms in order for JCMS to work correctly.

VOCABULARY TERM	DESCRIPTION	EXIT STATUS?
A	Alive	no
K	Killed	yes
D	Dead	yes
M	Missing	yes
S	Shipped	yes

### 3.3.7 Gene, Gene Class, and Allele Tables

In the database, alleles are associated with specific genes or gene classes. The JCMS forms that display alleles that may be associated with a gene use both criteria (association to the gene and

association to the gene class) to come up with a list of alleles that may be associated with a given gene.

Any mouse or sample may have several genotypes. Each genotype consists of a gene and the two alleles that were found for that gene.

### 3.3.7.1 Genes

In order to set up a genotype for a mouse or sample, **first the gene must be entered** into the database.

Formal Symbol	*Lab Symbol	Gene Class	Chromosome	cM	megabase	Comments
lsl/lsl	lsl	A				
cpdm/cpdm	cpdm	A				
*						

**Figure 3-12 Form: Add or Edit Gene**

The gene table will be shown as a form. Genes may only be added or edited, not deleted. The field called "Lab symbol" is the gene name that will be used when setting up alleles and genotypes. Gene class is optional. Certain gene classes have been preset. Any new gene classes should be added before the gene is added.

To **add a gene**, scroll to the bottom of the list on the form and enter the new gene into the empty bottom row or use the navigation buttons to move to a new record. The formal symbol and lab symbol are required and the gene will not be added unless values are entered in them. Adding will occur after clicking in a field that is not part of this row.

To **edit a gene**, type in the field that should be changed. If a mistake is made, pressing the ESC key once will revert that field back to the original value. Pressing the ESC key twice in a row will revert the whole record back to the original values.

**A gene cannot be deleted** from the database. If the gene name is incorrect, edit the name. Everywhere in the database where this gene is used, the name will change.

### 3.3.7.2 Gene classes

Gene classes are stored as a controlled vocabulary in a controlled vocabulary table. As with all controlled vocabularies, do not delete or change a gene class term unless no records in the database use that term.

The following Gene Class values are preset. These terms may be used to further describe a specific gene. Note that if alleles are assigned to a gene class, all genes in that class are able to have that allele.

Class name	Comments
E	endogenous
MKO	multi allele knock out
MTG	multi allele transgene
TG	transgene
KO	knock out
KI	knock in
Floxed	tissue specific knock out

Class name	Comments
CTK	Combination transgene + KO

### 3.3.7.3 Alleles

Once a gene is added, the list of possible alleles for this gene must be entered into the allele table. Click on the **allele button** on the Administrator button bar to open the add or delete allele form. Alleles may be associated with either a specific gene or a particular *gene class*. If an allele is associated with a gene class it will also be associated with all genes of that class. The user can select which association to use by clicking the appropriate radio button in the “Associate allele with” group box.

To associate an allele with a gene, enter the lab gene symbol and the list of alleles already available for this gene will appear. To add another allele to the list, type it into the new allele field and click the submit button. To remove an allele from the list, check the delete box, type the name of the allele to delete into the “new allele” box, and press the submit button (or double click the allele name in the list box and the name of the allele will appear in the “new allele” box).

Allele	Class
A	none
het	none

Associate allele with ..

Gene    01-167008   

Gene class    geneclass1   

New Allele:

Delete

Figure 3-13 Form: Add or Delete an Allele

### 3.3.7.4 Generic alleles

To save time entering alleles, it is possible to create generic alleles that are available for a whole gene class. These alleles will be automatically available as a choice for any gene that is associated with this gene class.

To associate an allele with a gene class, click on the “Gene class” radio button and perform the same operations as described above.

NOTE: Allele names cannot be edited and an allele that is associated with a gene class cannot be deleted.

NOTE: Since each genotype record links to a record in the gene table, changes to a gene name will be reflected in the genotype records that “point” to the gene. Allele names, on the other hand, are stored directly in each genotype record (not pointed to by a genotype record). If there ever is a need to search for mice based on specific alleles, then those alleles must still exist in the Allele table. Otherwise they will not be found. Therefore, those allele terms should not be deleted, even if no longer in use.

After the gene and alleles have been entered, this gene can be used for any mouse or sample.

### 3.3.8 Rooms, Health Level, and Pens

See section 7 on Pens and Cage Cards for details on setting up room names, health levels, and pens. Section 3.5 Cage Card Setup explains the details of the different cage cards that are provided and how to create custom cards.

### 3.3.9 Mouse Use Types

Mouse use types consist of a list of terms that specify a mouse use (e.g. clinical exam). These terms are used in the Mouse Usage table and by the add and edit mouse use forms, add plug date form, and the mouse use calendar.

The screenshot shows the JAX JCMS Release 4.6.0 Administrator interface. At the top, there's a toolbar with various buttons like 'Strain', 'Gene Class', 'Gene', 'Allele', 'Life Status', 'Add Approved Mating Strains', 'Edit Approved Mating Strains', 'Owner and Secretary Accounts', 'JCMS Setup Variables', and 'Set Form Access Privileges'. Below the toolbar, a message says 'Welcome To JAX JCMS Release: 4.6.0' and 'Logged in as: mtsadmin'. A sub-menu for 'Secretaries and Owners' is open. In the center, there's a table titled 'Add or Delete Mouse Use Types (mtsadmin)' with columns for 'mouseUse', 'useDescription', 'is Active', 'd1Caption', and 'd2Caption'. The table lists several entries: 'Basic blood work' (description: 'These are required for all blood work for Project 1523'), 'Clinical exam' (description: 'for testing no records are using this'), 'dummy' (description: 'To be done once a month'), 'Necropsy results', 'Neurological exam', 'Six month complete checkup', and 'Skin graft'. The 'is Active' column contains checkboxes, some of which are checked. The 'd1Caption' and 'd2Caption' columns show standard abbreviations like 'CBC', 'D1', 'D2', etc.

Figure 3-14 Add or delete mouse use types

On the Administrator button bar, click the Mouse Use Types button. An existing mouse use term may not be edited. It may be deleted if it is not in use by any mice. All the other fields on the form may be edited.

When a mouse use is no longer needed, set it to no longer be active (uncheck “is Active”). Mouse use terms that are not active will not be shown on the drop-down list of choices to use for a term when adding. It will still be a choice for reporting and editing so older data can be found.

Captions to use for the D1, D2, ... D10 data fields can be entered for the mouse use term. Any data field with no caption (i.e. possibly not needed) will automatically be set to D1, D2, ...D10. The caption may not be blank. The captions can be edited.

## 3.4 Setup Variables Table (DbSetup) for Customizing the Installation

JCMS has a setup variables table that allows easy customization of each installation. The Administrator (mtsadmin) can change configuration variables from the **JCMS Setup Variables** button on the Administrator button bar. **Some of these variables should be initialized; those are marked below in bold print.** The others may be left at the default and changed later once the users are more familiar with the database. See section 3.5.4 on creating custom cage cards for a list and examples of the standard cage card choices. Section 7.1 Manage Pen Configuration explains the form interface used to change the settings of the setup variables used for pens, rooms, and cage cards.

JCMS Setup Variable name	Value	Description
JCMS_ACTIVATE_MATINGS_INCREMENT	false	true or false; if true the mating ID is automatically incremented after activating a mating. May be overridden on the form.

JCMS Setup Variable name	Value	Description
JCMS_ADD_AT_WEAN_INCREMENT	false	true or false; if true the mouse ID is automatically incremented after adding a mouse at weaning. May be overridden on the form.
JCMS_ADD_GENOTYPE_INCREMENT	false	true or false; if true the mouse ID is automatically incremented after adding a genotype. May be overridden on the form.
JCMS_ADD_LITTER_INCREMENT	false	true or false; if true the litter ID is automatically incremented after adding a litter. May be overridden on the form.
JCMS_ADD_LITTER_PUPS_INCREMENT	false	true or false; if true the litter ID is automatically incremented after adding a litter with pups. May be overridden on the form.
JCMS_ADD_MOUSE_INCREMENT	false	true or false; if true the mouse ID is automatically incremented after adding a mouse. May be overridden on the form.
JCMS_ADD_MOUSE_USE_INCREMENT	false	true or false; if true the mouse ID is automatically incremented after adding a mouse use. May be overridden on the form.
JCMS_ALLELE_CONF_HIGH		Character(s) displayed/printed as part of a genotype to indicate high confidence in an allele. May be blank.
JCMS_ALLELE_CONF_LOW	?	Character(s) displayed/printed as part of a genotype to indicate low confidence in an allele. May be blank. Ex: use ? To display Abc +?/+? For low confidence in both alleles.
JCMS_ALLELE_GENE_SEPARATORS		Specify separators to display around the alleles. Ex: specify [] to get Abc[+/+] or leave blank for Abc +/+
JCMS_ALLOW_USERDEFINED_GENERATIONS	false	true or false; If true user may add generations to the Generations table on the fly.
JCMS_ALLOW_USERDEFINED_STRAINS	false	true or false; If true user may add strains to the Strains table (via the Strains form) on the fly.
JCMS_AUTOINCREMENT_GENERATION	false	true or false; If true the generation for a litter will be automatically incremented when creating matings. May be overridden on the form.
JCMS_AUTO_RETIRE_MATINGS	true	true or false; If true automatically retire a mating when the sire and dam(s) all have an exit life status.
JCMS_AUTO_RETIRE_PENS	false	Automatically retire a pen when it becomes empty or all occupants are not alive.
JCMS_BILL_PARTIAL_FIRST_DAY	false	A pen is counted for billing on the day it is created or moved into a room.
JCMS_BILL_PARTIAL_LAST_DAY	false	A pen is counted for billing on the day it is retired or moved out of a room.
JCMS_CREATE_PEN_INCREMENT	false	true or false; if true the pen ID is automatically incremented when adding new pens.
JCMS_DATABASE_DBMS	MSAccess	Name of the database management system. Valid values are MSAccess or MySQL. Required for MySQL to function properly.
<b>JCMS_DATA_FILE_DIRECTORY</b>		The root directory where JCMS documents (files for images, Excel, Word, etc.) are stored
JCMS_DAYS_TO_GENOTYPE	14	The number of days from the birth date to when the pups should be genotyped.
JCMS_DEFAULT_CONTAINER_STATUS	active	Specify the default value for pen status; must match a value in the pen status table.
JCMS_DEFAULT_EXIT_TERM	E	Specify the default term used when exiting mice from the colony, usually E or K; must match a value in the life status table. May be overridden on forms.
JCMS DESIGN RETIRE MATINGS_INCREMENT	false	true or false; if true the mating ID is automatically incremented when using the design retire mating form. May be overridden on the form.

JCMS Setup Variable name	Value	Description
JCMS_EDIT_LITTER_INCREMENT	false	true or false; if true the litter ID is automatically incremented after editing a litter. May be overridden on the form.
JCMS_EDIT_MOUSE_INCREMENT	false	true or false; if true the mouse ID is automatically incremented after editing a mouse. May be overridden on the form.
JCMS_EDIT_MOUSE_USE_INCREMENT	false	true or false; if true the mouse ID is automatically incremented after editing a use. May be overridden on the form.
JCMS_ENABLE_GENOTYPE_IMPORT	true	true or false; must be set to true to allow importing genotypes.
JCMS_ENFORCE_APPROVED_MATINGS	false	true or false; if true then user is only allowed to set litter strain to pre-approved strains. May be overridden on the form.
JCMS_ENFORCE_PASSWORD_CHANGE	false	Tells application whether to remind user whether to change password after password change period has elapsed (used by JCMSWeb)
JCMS_EXT_WEAN_TIME	28	the number of days from the birth date to when a litter should be weaned - used for late weanings/"long" wean time.
JCMS FEMALES_FIRST	true	true or false; if true females are the first to be assigned mouse IDs when adding mice with a bulk add.
JCMS_GENERATION_INCREMENT_RIGHTMOST	true	true or false; if true, increment the rightmost numeric portion of the generation; if false the leftmost.
JCMS_GESTATION_PERIOD	21	The length of gestation (pregnancy) is used to determine when plug dates expire.
JCMS_IMPORT_EXP_DATA_ALLOW_MULTIPLE	false	If importing with NO experimental plan, allow more than one record for a test type/mouse combination.
JCMS_IMPORT_EXP_DATA_MICE_MUST_BE_PRESELECTED	true	true or false; if false, any mouse IDs not pre-selected will be automatically added to the plan and test.
JCMS_JAXLAB_INSTALLATION	false	Is this a Jackson Laboratory installation.
JCMS_JAX_ACCOUNT_NUMBER	12.210.3163.56048	The value to submit in every export-to-TGS file as the "JAX Account #".
JCMS_LITTERID_INCREMENT_RIGHTMOST	true	true or false; if true, increment the rightmost numeric portion of the litter ID; if false the leftmost.
JCMS_LOOP_LITTER_NUMBERS	true	true or false; if true the litter numbers recycle after 10 litters, appending a character to the number.
JCMS_MAX_IMPORT_EXP_DATA_ERRORS	10	Import Experimental data; when this number of errors is reached, verification stops and a report is printed.
JCMS_MOUSEID_INCREMENT_RIGHTMOST	true	true or false; if true, increment the rightmost numeric portion of the mouse ID; if false the leftmost.
JCMS_MTS_IMPORT_PATH	C:\	The file path to default to for non-TGS imports via the 'Import Genotype' button.
JCMS MySQL DATABASE_NAME	jcms_db	The name of your MySQL database.
JCMS MYSQL DRIVER	{MySQL ODBC 3.51 Driver}	The name of your MySQL driver.
JCMS MYSQL SERVER	localhost	The name of your MySQL server.
JCMS MYSQL USER_ID	root	The name of your MySQL user ID.
JCMS_PASSWORD_CHANGE_PERIOD	90	The number of days between a user changing and being reminded to change their password (used by JCMSWeb)
JCMS_PEN_NAMES_INCREMENT_RIGHTMOST	true	true or false; if true, increment the rightmost numeric portion of the pen name; if false the leftmost.
JCMS_PRINT_EXITED_MICE_ON_CAGE_CARDS	true	true or false; if true, mice with an exit status such as dead, euthanized, missing, shipped, etc. will print on cage cards.
JCMS_RETIRE_MATINGS_INCREMENT	false	true or false; if true the mating ID is automatically

JCMS Setup Variable name		Value	Description
			incremented after retiring a mating. May be overridden on the form.
JCMS_SAMPLE_LABEL_REPORT	PrintSampleLabels		The name of the report used to print sample labels from the Print Sample Label form
JCMS_SORT_BY_PEN_NAME	False		Causes lists with pen ID and pen name to sort alphabetically by pen name instead of pen ID.
JCMS_STANDARD_WEAN_TIME	18		the number of days from the birth date to when a litter should normally be weaned.
JCMS_STRAINNAME_FIRST	true		true or false, if true the strain name will appear first (to the left of the JR number) in all the dropdown lists.
JCMS_TGS_REQUEST_FILENAME	komp2typ.txt		The filename to store the file being submitted to TGS. Note this value should not be overridden; TGS expects this filename verbatim.
JCMS_TGS_REQUEST_PATH	\jax\jax\cs\private\appdev\src\komp\DEV		The file path to store the file being submitted to TGS. Note this should not include the actual filename or a trailing slash.
JCMS_TGS_RESPONSE_PATH	\jax\jax\cs\private\appdev\src\komp\DEV\outbox		The file path to look for response files from TGS.
JCMS_USING_HEALTH_LEVEL	true		true or false; if false, the room health level will not be displayed on most forms
JCMS_USING_PEN_COMMENTS	true		true or false; if false, the pen comment field will not be displayed on most forms
JCMS_USING_PEN_NAMES	true		true or false; if false, pen names will not be displayed on most forms
JCMS_WARN_DUPLICATE_PEN_NAME	false		true or false; if false warn if a duplicate pen name is used.
JCMS_WARN_LITTER_NOT_UPDATED	true		true or false, if true, add mice at weaning will warn if litter record was not updated.
JCMS_WRITE_FAILED_TRANSACTIONS	false		true or false; used by the Add Sample form for debugging, should be set to false.
MTS_1PEN_WEAN_CAGE_CARD	TS_1PWeanCageCard		The name of the cage card report for 1-pen wean cage cards.
MTS_2PEN_WEAN_CAGE_CARD	OS_2PWeanCageCard		The name of the cage card report for 2-pen wean cage cards.
MTS_AUTOINCR_DAMS_SIRES	false		true or false, if true the dams and sire are automatically incremented on the handheld trio/pair mating form. May be overridden on the form.
MTS_AUTO_COLOR	true		true or false; if true, then many forms get all data entry fields colored after user hits submit. Color is cleared after user visits the field.
MTS_AUTO_LITTER_NUMS	on		on or off; if on litter numbers are automatically generated for matings.
MTS_CAGE_CARD_DETAIL_NOTE	Put card note here (dbsetup)		A note that will be printed on all Detail cage cards.
MTS_DEFAULT_AUTO_INCREMENT	off		on or off; if on then auto increment is the default on forms with functions that do not have a specific auto increment setup variable.
MTS_DEFAULT_COD			Specify the default cause of death; must match a value in the cv_CauseOfDeath table.
MTS_DEFAULT_HEALTH_LEVEL	2		Specify the default room health level; must match a value in the Health Level table.
MTS_DEFAULT_MOUSE_ORIGIN			Specify the default mouse origin; must match a value in the cv_MouseOrigin table.
MTS_DEFAULT_MOUSE_ROOM			Specify the default mouse room; must match a value in the Room table.
MTS_DEFAULT_PRINTCARDS	true		true or false; if true the print cage card option on the handheld trio/pair mating form is set on. May be overridden on the form.
MTS_DEFAULT_USE_BASEMOUSE_ID	false		true or false; if true indicates using a base mouse id on

JCMS Setup Variable name		Value	Description
			the handheld add litter form. May be overridden on the form.
MTS_DETAIL_CAGE_CARD	TS_DetailCageCard		The name of the cage card report for detail cage cards.
MTS_DOB_ROLLBACK_OFFSET	7		Subtract this number of days from today's date to get the date of birth.
MTS_HELP_EMAIL	mailto:jaxcms@lists.jax.org?subject=Support issue		Specify an email address that users can send JCMS support questions to. Used as the link for "Report a problem" on the JCMS welcome window.
MTS_IMPORT_MAX_WARNING	20		Users will be warned if they try to import more than this number of mice at once (only effects bulk imports of mice).
MTS_INSTALLATION_NAME	JCMS		Name of this JCMS installation (anything you want to call it).
MTS_LITTER_ID_PREFIX	L		A short string of characters that are prefixed on litter IDs generated by JCMS (not all litter IDs are generated by JCMS).
MTS_MAIN_BUTTON_BAR	MainButtonBarJCMS		Name of the main button bar form displayed when user hits start workstation from welcome window.
MTS_MATING_CAGE_CARD	OS_MatingCageCard		The name of the cage card report for mating cage cards.
MTS_MATING_CAGE_CARD2	TS_MatingCageCardStyle2 WithBarcode		The name of the cage card report for mating cage cards style 2; handheld only, print cage card form.
MTS_MATING_ID_PREFIX	M		A short string of characters that are prefixed on mating IDs when printed on some cage cards.
MTS_MAX_MICE_PER_PEN	10		Maximum number of live mice in any pen.
MTS_MOUSE_ID_PREFIX	A		A short string of characters that are prefixed on mouse IDs generated by JCMS (not all mouse IDs are generated by JCMS).
MTS_NUM_AUTO_LITTER_NUMS	10		This variable sets the number of litter numbers that are assigned to a mating. It should be set to a value bigger than the max number of litters you ever expect. Suggested values are 10 or 100.
MTS_PEN_ID_PREFIX	P		A short string of characters that are prefixed on pen IDs when printed on some cage cards.
MTS_PI_NAME	PI Name		Name of lab PI who owns colonies tracked by JCMS, printed on some cage cards.
MTS_PI_PHONE	555-1212 (office)		Phone numbers, printed on mating card
MTS_RELAXED_PEN_NUMS	true		true or false; if false, then it is required that a cage card is printed for all pens.
MTS_THRESHOLD_MICE_BATCH_OPERATION	50		This variable will trigger a warning from the handheld wean and exit form when the number of affected mice exceeds this value.

**Table 3-1 JCMS Setup Variables**

### 3.5 Cage Card Setup

Cage cards are designed to print on a standard index card. The exact cage cards used will vary depending on the setup. Different cage card formats may be selected from the setup variables option on the Administrator's button bar. Enter into the JCMS Setup Variable value the exact name of the cage card report. When cage cards are printed from data entry forms, relevant information (such as mouse ID, strain, etc.) is printed directly on the card.

#### 3.5.1 Printing Cage Cards

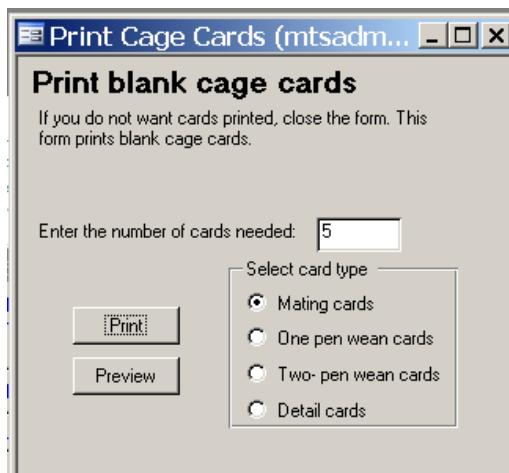
The cage cards use the default printer and default paper location. Many modern printers will use the sheet feeder as the default whenever there is paper in the sheet feeder. To print cage cards, open the sheet feeder and load the cards into the envelope feeder part of it. Also set the printer to use as straight a paper path as possible. Many printers have an option for sending sheets out

the back if it is open or have a toggle switch to change the path. The cage cards are designed to print either centered or to the far left on the paper. Adjust the envelope feeder to place the cards in the proper location. If the cage cards do not print out correctly, adjust the margins from the File/Page Setup menu (Access 2007: Print dialog box, Setup button). To solve problems with printing, see the printer notes in the Technical Guide section of this document.

Note that some wean cage cards are designed to print for one pen and some for two pens. The two-pen option is useful for side-by-side shoebox type pens.

A JCMS setup variable, JCMS\_PRINT\_EXITED\_MICE\_ON\_CAGE\_CARDS, allows configuring whether or not mice that are not in the colony print on cage cards. The default value is true, but if changed to false, mice with a status indicating that they have exited the colony (i.e. "Dead," "Missing," "Shipped") will be excluded from the cage card.

### 3.5.2 Blank Cage Cards



Periodically, blank cage cards can be printed to use in the mouse room for newly weaned mice or mice obtained from other sources. Use the **Print Cage Cards** button on the reports tab to open the Print Cage Cards form. Enter the number of cards needed. Any of the four cage card types may be printed as blank cards. JCMS will generate unique pen ID numbers for each card.

**Figure 3-15 Form: Print blank cage cards**

### 3.5.3 JCMS-Provided Cage Card Formats

JCMS defines four types of cage cards:

Pen L.	P.I.	Owner
Strain:		
B.D.      wean/bog		
#F	#M	Litter#
Mouse ID# Gen other		
Pen R.	P.I.	Owner
Strain:		
B.D.      wean/bog		
#F	#M	Litter#
Mouse ID# Gen other		

- 1) Mating cage card
- 2) One-pen wean cage card
- 3) Two-pen wean cage card
- 4) Detail cage card

#### 3.5.3.1 Two-pen Wean Cards (setup variable MTS\_2PEN\_WEAN\_CAGE\_CARD)

Two-pen wean cards can only be printed as blank cage cards. The PI phone and PI name information will be shown on the card using the values specified in the setup variables called MTS\_PI\_NAME and MTS\_PI\_PHONE (these values are set via the Administrator tool bar, *JCMS Setup Variables* button).

- Two-pen wean cards can be printed from the Print Cage Cards Form.

This card format prints on the upper left corner of the page. Make sure the envelope feeder on the printer is set to print to the far left side.

**Figure 3-16  
OS\_2PWeanCageCard**

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>PINameR</b>	Name of PI responsible for the right cage	Setup variable (MTS_PI_NAME)
<b>PINameL</b>	Name of PI responsible for the left cage	Setup variable (MTS_PI_NAME)
<b>PIPhoneR</b>	A short string of text with PI contact information for right pen	Setup variable (MTS_PI_PHONE)
<b>PIPhoneL</b>	A short string of text with PI contact information for left pen	Setup variable (MTS_PI_PHONE)
<b>PenIDR</b>	Right pen ID	generated unique by JCMS or user can generate
<b>PenIDL</b>	Left pen ID	generated unique by JCMS or user can generate

### 3.5.3.2 One-pen Wean Cards (setup variable MTS\_1PEN\_WEAN\_CAGE\_CARD)

One-pen wean cards can be printed as blank wean cards or with mouse IDs and other pertinent information on them. One-pen wean cards can be printed from the following JCMS forms:

Strain # 651	P.I. Lab name goes here xxx-xxx-xxxx	Owner nobody
Activation date 2/2/2010	Wean date 2/3/2010	Count/sex: <b>2 F</b>
Pen# <b>735</b>		
<b>BALB/cJ</b>		
born: 1/1/2010 M66 L-660		
Gen: N05		
Intended use:		
A422		
A423		
Notes:		

- Print Cage Cards
- Pen Info
- Add Pen
- Edit Pen
- Move Mice
- Add Litter With Pups
- Handheld Print Cage Cards
- Edit Pen Status/Location

These two card formats will print centered – make sure the envelope feeder sends the cards through the center of the print path.

**Figure 3-17 TS\_1PWeanCageCard or  
TS\_1PWeanCageCardWithBarcode**

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>StrainNum</b>	Strain number	Strain table
<b>PIPhone</b>	A short string of text with PI contact information	Setup variable (MTS_PI_PHONE)
<b>Owner</b>	Owner ID	mouse owner
<b>activationDate</b>	Pen activation date	Container history table
<b>weanDate</b>	Date mice were weaned or blank if mice imported from external colony	litter table
<b>PenID</b>	A unique pen number	JCMS generated or user specified
<b>countSex</b>	A string with number of mice and sex e.g. "10 M", would specify ten males	JCMS generated
<b>birthDate</b>	Date of birth of mice	litter table
<b>MatingID</b>	Mating number that produced this litter	mating table
<b>LitterID</b>	Litter number for the litter	litter table
<b>IntendedUse</b>	From CV pull down list	mouseUse
<b>generation</b>	Litter generation	litter table
<b>M1ID</b>	Mouse 1 ID (from litter)	mouse table
<b>M2ID</b>	Mouse 2 ID (from litter)	mouse table
<b>M3ID</b>	Mouse 3 ID (from litter)	mouse table
<b>M4ID</b>	Mouse 4 ID (from litter)	mouse table
<b>M5ID</b>	Mouse 5 ID (from litter)	mouse table
<b>M6ID</b>	Mouse 6 ID (from litter)	mouse table
<b>M7ID</b>	Mouse 7 ID (from litter)	mouse table
<b>M8ID</b>	Mouse 8 ID (from litter)	mouse table
<b>M9ID</b>	Mouse 9 ID (from litter)	mouse table
<b>M10ID</b>	Mouse 10 ID (from litter)	mouse table
<b>penIDBC</b>	Pen ID in barcode format. Must be of the font Code 128AB	User specified
<b>matingIDBC</b>	Mating ID in barcode format. Must be of the font Code 128AB	User specified

### 3.5.3.3 One pen wean card for designing new cards

The report named TS\_1PWeanCageCardwithXtraFields may be used for designing a new cage card; it uses the fields and format above. The following fields are added.

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>protocol1</b>	Protocol for mouse 1	Mouse table
<b>protocol2</b>	Protocol for mouse 2	Mouse table
<b>protocol3</b>	Protocol for mouse 3	Mouse table
<b>protocol4</b>	Protocol for mouse 4	Mouse table
<b>protocol5</b>	Protocol for mouse 5	Mouse table
<b>protocol6</b>	Protocol for mouse 6	Mouse table
<b>protocol7</b>	Protocol for mouse 7	Mouse table
<b>protocol8</b>	Protocol for mouse 8	Mouse table
<b>protocol9</b>	Protocol for mouse 9	Mouse table
<b>protocol10</b>	Protocol for mouse 10	Mouse table
<b>coatColor1</b>	Coat color for mouse 1	Mouse table
<b>coatColor2</b>	Coat color for mouse 2	Mouse table
<b>coatColor3</b>	Coat color for mouse 3	Mouse table
<b>coatColor4</b>	Coat color for mouse 4	Mouse table
<b>coatColor5</b>	Coat color for mouse 5	Mouse table
<b>coatColor6</b>	Coat color for mouse 6	Mouse table
<b>coatColor7</b>	Coat color for mouse 7	Mouse table

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<i>coatColor8</i>	Coat color for mouse 8	Mouse table
<i>coatColor9</i>	Coat color for mouse 9	Mouse table
<i>coatColor10</i>	Coat color for mouse 10	Mouse table
<i>comment1</i>	Comment for mouse 1	Mouse table
<i>comment2</i>	Comment for mouse 2	Mouse table
<i>comment3</i>	Comment for mouse 3	Mouse table
<i>comment4</i>	Comment for mouse 4	Mouse table
<i>comment5</i>	Comment for mouse 5	Mouse table
<i>comment6</i>	Comment for mouse 6	Mouse table
<i>comment7</i>	Comment for mouse 7	Mouse table
<i>comment8</i>	Comment for mouse 8	Mouse table
<i>comment9</i>	Comment for mouse 9	Mouse table
<i>comment10</i>	Comment for mouse 10	Mouse table
<i>room</i>	Room name	Room table
<i>penName</i>	Name of the pen	Container table
<i>penComment</i>	Comment for pen	Container table
<i>statusDate</i>	Date when pen status or room changed	ContainerHistory table
<i>statusDateWith Time</i>	Same as status date only includes the exact time of the change	ContainerHistory table
<i>penStatus</i>	Current status of pen (active, retired, proposed, etc.)	Container History table
<i>Dam1-ID</i>	Mouse ID of dam1 (from mating)	Mouse table
<i>Dam2-ID</i>	Mouse ID of dam2 (from mating)	Mouse table
<i>Sire-ID</i>	Mouse ID of sire (from mating)	Mouse table
<i>Dam1-Strain</i>	Strain of dam1 (from mating)	Strain table
<i>Dam2-Strain</i>	Strain of dam2 (from mating)	Strain table
<i>Sire-Strain</i>	Strain of sire (from mating)	Strain table
<i>Dam1-Genotype</i>	Genotype of dam1 (from mating)	Genotype table
<i>Dam2-Genotype</i>	Genotype of dam2 (from mating)	Genotype table
<i>Sire-Genotype</i>	Genotype of sire (from mating)	Genotype table
<i>PlanID</i>	Experimental plan ID	ExpPlan table (Only one plan ID for one of the mice is displayed when there are multiple possibilities)
<i>PlanName</i>	Experimental plan name	ExpPlan table
<i>M1NewTag</i>	Replacement tag for mouse 1	Mouse table
<i>M2NewTag</i>	Replacement tag for mouse 2	Mouse table
<i>M3NewTag</i>	Replacement tag for mouse 3	Mouse table
<i>M4NewTag</i>	Replacement tag for mouse 4	Mouse table
<i>M5NewTag</i>	Replacement tag for mouse 5	Mouse table
<i>M6NewTag</i>	Replacement tag for mouse 6	Mouse table
<i>M7NewTag</i>	Replacement tag for mouse 7	Mouse table
<i>M8NewTag</i>	Replacement tag for mouse 8	Mouse table
<i>M9NewTag</i>	Replacement tag for mouse 9	Mouse table
<i>M10NewTag</i>	Replacement tag for mouse 10	Mouse table

Strain # 651	P.I. xxx-xxx-xxxx	Lab name goes here Owner nobody	
Activation date 5/22/2010	Wean date 5/22/2010	Count/sex: 2 M	Pen ID 819
Room B50-22                            BALB/cJ-0045			
Status active		Status date 5/22/2010	
<b>BALB/cJ</b>			
born: 5/4/2010		M120	L-1201
Gen: N05			
Intended use:			
BALB108			
BALB109			
Notes: EXP 3			
IOP testing			

**Figure 3-18**  
**CC\_JCMS\_WeanCageCardUsingPenNames**

### 3.5.3.4 Mating Cards (setup variable MTS\_MATING\_CAGE\_CARD or MTS\_MATING\_CAGE\_CARD2)

**Figure 3-19**  
**TS\_MatingCAgeCard or**  
**TS\_MatingCageCardStyle1W**  
**ith BarCode**

**Figure 3-21 OS\_MatingCageCard**

**Figure 3-20**  
**TS\_MatingCageCardStyle2With**  
**BarCode**

**Figure 3-22 MW\_MatingCageCardLandscape**

Pen <b>L or R.</b>	P.I. Lab name goes here xxx-xxx-xxxx	Owner <b>OWN1</b>	
Mating # <b>M79</b>	Mating Date <b>1/12/2010</b>	Pen ID <b>524</b>	
Room B50-22	BALB mating 005		
Status active	Status date 2/2/2010		
Litter Strain <b>BALB/cJ</b>	Litter gen <b>N05</b>		
Dame # L340-17	B.D. 9/2/2009	Dame Strain BALB/cJ	
Sire # L340-11	B.D. 9/2/2009	Sire Strain BALB/cJ	
Note 1:			
Note 2:			
Wean note: sample wean note			
Litter # L-790	B.D. 	#born 	other 
L-791			
L-792			
L-793			
L-794			
L-795			
L-796			
L-797			
L-798			
L-799			

**Figure 3-23**  
**CC\_JCMS\_MatingCageCardUsingPenNames**

Mating cards may be printed from the following set of forms:

- Design Matings
- Activate Matings
- Edit Matings
- Add Matings
- Move Mice
- Print Cage Cards
- Handheld Print Cage Cards

The OS and CC Mating Cage Card format prints in the upper left corner of the page. The MW landscape card prints on the left of the page with a one inch left margin. The TS card formats will print centered at the top of the page. Make sure the envelope feeder sends the cards through the correct print path.

<b>Field name</b>	<b>Description</b>	<b>Data source</b>	<b>OS_Visible?</b>	<b>TS_Visible?</b>	<b>CC_Visible?</b>	<b>MW_Visible?</b>
<b>PIName</b>	Name of responsible PI	Setup variable (MTS_PI_NAME)	y	y	y	n
<b>PIPhone</b>	Contact phone for PI	Setup variable MTS_PI_PHONE	y	y	y	y
<b>owner</b>	Mating owner	mating table	y	y	y	y
<b>matingID</b>	Mating number	mating table	y	y	y	y
<b>matingDate</b>	Mating date	mating table	y	y	y	y
<b>StrainSection</b>	Strain section in mouse room	strain table	n	y	n	n
<b>PenID</b>	Pen number	(one mouse) Container table	y	y	y	y
<b>LitterStrain</b>	Strain of pups	mating table	y	y	y	y
<b>litterGeneration</b>	Generation of pups	mating table	y	y	y	y
<b>CardColor</b>	Color of cage card	strain table	n	y	y	n
<b>dam1ID</b>	Mouse ID for dam 1	mouse table	y	y	y	y
<b>dam2ID</b>	Mouse ID for dam 2	mouse table	y	y	y	y
<b>sireID</b>	Mouse ID for sire	mouse table	y	y	y	y
<b>dam1BirthDate</b>	Birth date of dam1	mouse table	y	y	y	y
<b>dam2BirthDate</b>	Birth date of dam2	mouse table	y	y	y	y
<b>sireBirthDate</b>	Birth date of sire	mouse table	y	y	y	y
<b>dam1Strain</b>	Strain of dam1	mouse table	y	y	y	y
<b>dam2Strain</b>	Strain of dam2	mouse table	y	y	y	y
<b>sireStrain</b>	Strain of sire	mouse table	y	y	y	y
<b>dam1Genotype</b>	Genotype of dam1	genotype table	n	y	n	y
<b>dam2Genotype</b>	Genotype of dam 2	genotype table	n	y	n	y
<b>sireGenotype</b>	Genotype of sire	genotype table	n	y	n	y
<b>dam1JRNum</b>	JR number of dam 1 strain	strain table	n	y	n	n
<b>dam2JRNum</b>	JR number of dam 2 strain	strain table	n	y	n	n
<b>sireJRNum</b>	JR number of sire strain	strain table	n	y	n	n
<b>dam1MatingID</b>	Mating dam1 came from	mating table via litter	n	y	n	n
<b>dam2MatingID</b>	Mating dam2 came from	mating table via litter	n	y	n	n

<b>Field name</b>	<b>Description</b>	<b>Data source</b>	<b>OS_Visible?</b>	<b>TS_Visible?</b>	<b>CC_Visible?</b>	<b>MW_Visible?</b>
<b>sireMatingID</b>	Mating sire came from	mating table via litter	n	y	n	n
<b>dam1LitterID</b>	Litter dam1 came from	litter table	n	y	n	n
<b>dam2LitterID</b>	Litter dam2 came from	litter table	n	y	n	n
<b>sireLitterID</b>	Litter sire came from	litter table	n	y	n	n
<b>dam1Gen</b>	Generation of dam1	mouse table	n	y	n	n
<b>dam2Gen</b>	Generation of dam2	mouse table	n	y	n	n
<b>sireGen</b>	Generation of sire	mouse table	n	y	n	y
<b>matingNote1</b>	A note about the mating	user selected from pick list on form	y	y	y	n
<b>matingNote2</b>	Additional mating notes	user selected from pick list on form	y	y	y	n
<b>weanNote</b>	Additional mating notes	mating table, wean note field	n	y	y	n
<b>penIDBC</b>	Pen ID in barcode format. Must be of the font Code 128AB	User specified	n	y	n	n
<b>matingIDBC</b>	Mating ID in barcode format. Must be of the font Code 128AB	User specified	n	n	n	n
<b>litter1ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter2ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter3ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter4ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter5ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter6ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter7ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter8ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter9ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>litter10ID</b>	Auto generated litter IDs	JCMS generated (optional)	y	y	y	y
<b>protocol</b>	Protocol for dam1	Mouse table	n	n	n	n
<b>room</b>	Room	Room table	n	n	y	n
<b>matingComment</b>	Comment	Mating table	n	n	n	n

<b>Field name</b>	<b>Description</b>	<b>Data source</b>	<b>OS_Visible?</b>	<b>TS_Visible?</b>	<b>CC_Visible?</b>	<b>MW_Visible?</b>
<b>penName</b>	Name of pen	Container table	n	n	y	n
<b>penStatus</b>	Current status	Container History table	n	n	y	n
<b>statusDate</b>	Date pen status or room changed	Container History table	n	n	y	n
<b>statusDateWithTime</b>	Same as above only including time	Container History table	n	n	n	n
<b>penComment</b>	Misc. information	Container table	n	n	n	n
<b>PlanID</b>	Experimental plan ID	ExpPlan table (Only one plan ID for one of the mice is displayed when there are multiple possibilities)	n	n	y	n
<b>PlanName</b>	Experimental plan name	ExpPlan table	n	n	y	n
<b>dam1NewTag</b>	Replacement tag for dam1	Mouse table	n	n	n	n
<b>dam2NewTag</b>	Replacement tag for dam2	Mouse table	n	n	n	n
<b>SireNewTag</b>	Replacement tag for sire	Mouse table	n	n	n	n

Two reports are available to use for designing new cage cards in these formats. They are named TS\_MatingCageCardStyle2WithBarcode with XtraFields and OS\_MatingCageCardwithXtraFields. The following fields are visible on both: protocol, room, statusDateWithTime, penComment, matingComment, PlanID, PlanName, dam1NewTag, dam2NewTag, and sireNewTag. Note they are located outside the normal boundaries for an index-sized card. Some of these fields are also on the MW\_MatingCageCardLandscape report, but they are not visible. Make a copy of this report before changing it.

### 3.5.3.5 Detail Cards (setup variable MTS\_DETAIL\_CAGE\_CARD)

	P.I.			Owner
Activation date:	Sec:	Count / sex	Pen#	
born:				
Gen:	genotype:			
born:				
Gen:	genotype:			
born:				
Gen:	genotype:			
born:				
Gen:	genotype:			
Intended use:				
Notes:				

Figure 3-24 TS\_DetailCageCard

	P.I.			Owner
Activation date:	Sec:	Count / sex	Pen#	
Expt #:				
#:	born:			
Gen:	genotype:			
#:	born:			
Gen:	genotype:			
#:	born:			
Gen:	genotype:			
#:	born:			
Gen:	genotype:			
Notes:				

Figure 3-25 TS\_DetailCageCardWithBarcode

	P.I. Lab name goes here xxx-xxx-xxxx			Owner nobody
Activation date:	Sec:	Count / sex:	Pen ID	
5/22/2009		1 M	721	
Room 2433	BALB8-001			
Status active	Status date 5/22/2009			
sample card note				
B002-003	M	born: 5/4/2009	M-14	L-140
BALB/cJ				
Gen: N04	genotype:			
born:				
Gen:	genotype:			
born:				
Gen:	genotype:			
born:				
Gen:	genotype:			
Intended use:				
Notes:				

Notes:	Pen#	Count / sex:		
Owner				
Phone				
Notes:				
Mouse #	DOB	Gen.	Strain	Genotype

Figure 3-27 MW\_DetailCageCardLandscape

Figure 3-26 (left)  
CC\_JCMS\_DetailCageCardUsingPenNames

Detail cards can be printed from the following forms:

- Pen Info
- Add Pen
- Edit Pen
- Bulk Add Mice
- Move Mouse
- Move Mice

- Print Cage Cards
- Handheld Print Cage Cards
- Edit Pen Status/Location

The “TS” and “CC” card formats will print centered – make sure the envelope feeder sends the cards through the center of the print path. The landscape “MW” card prints with a one inch margin, left side.

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>PIName</b>	Name of responsible PI	Setup variable (MTS_PI_NAME)
<b>PiPhone</b>	Phone number of responsible PI	Setup variable (MTS_PI_PHONE)
<b>Owner</b>	Owner of mice in pen	Mouse table (one mouse)
<b>ActivationDate</b>	Date pen was activated	Container History table
<b>Section</b>	Strain section	strain table
<b>countSex</b>	String of text with count and sex of mice on it (e.g. 4 M)	Generated by JCMS when mouse information is retrieved.
<b>PenID</b>	Pen number	Container table, from one mouse
<b>penIDBC</b>	Pen ID in barcode format. Must be of the font Code 128AB	User specified
<b>matingIDBC</b>	Mating ID in barcode format. Must be of the font Code 128AB	User specified
<b>Note</b>	A cage card note	Setup variable (MTS_CAGE_CARD_DETAIL_NOTE)
<b>M1ID</b>	Mouse 1 ID number	mouse table
<b>M1Sex</b>	Mouse 1 Sex	mouse table
<b>M1Born</b>	Mouse 1 DOB	mouse table
<b>M1MID</b>	Mouse 1 is a product of Mating ID	mating table
<b>M1LID</b>	Mouse 1's litter ID	litter table
<b>M1Strain</b>	Mouse 1's strain	mouse table
<b>M1Generation</b>	Mouse 1's generation	mouse table
<b>M1GenoType</b>	Mouse 1's genotype	genotype table
<b>M2ID</b>	Mouse 2 ID number	mouse table
<b>M2Sex</b>	Mouse 2 Sex	mouse table
<b>M2Born</b>	Mouse 2 DOB	mouse table
<b>M2MID</b>	Mouse 2 is a product of Mating ID	mating table
<b>M2LID</b>	Mouse 2's litter ID	litter table
<b>M2Strain</b>	Mouse 2's strain	mouse table
<b>M2Generation</b>	Mouse 2's generation	mouse table
<b>M2GenoType</b>	Mouse 2's genotype	genotype table
<b>M3ID</b>	Mouse 3 ID number	mouse table
<b>M3Sex</b>	Mouse 3 Sex	mouse table
<b>M3Born</b>	Mouse 3 DOB	mouse table
<b>M3MID</b>	Mouse 3 is a product of Mating ID	mating table
<b>M3LID</b>	Mouse 3's litter ID	litter table
<b>M3Strain</b>	Mouse 3's strain	mouse table
<b>M3Generation</b>	Mouse 3's generation	mouse table
<b>M3GenoType</b>	Mouse 3's genotype	genotype table
<b>M4ID</b>	Mouse 4 ID number	mouse table
<b>M4Sex</b>	Mouse 4 Sex	mouse table
<b>M4Born</b>	Mouse 4 DOB	mouse table

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>M4MID</b>	Mouse 4 is a product of Mating ID	mating table
<b>M4LID</b>	Mouse 4's litter ID	litter table
<b>M4Strain</b>	Mouse 4's strain	mouse table
<b>M4Generation</b>	Mouse 4's generation	mouse table
<b>M4GenoType</b>	Mouse 4's genotype	genotype table
<b>M5ID</b>	Mouse 5 ID number	mouse table
<b>M5Sex</b>	Mouse 5 Sex	mouse table
<b>M5Born</b>	Mouse 5 DOB	mouse table
<b>M5MID</b>	Mouse 5 is a product of Mating ID	mating table
<b>M5LID</b>	Mouse 5's litter ID	litter table
<b>M5Strain</b>	Mouse 5's strain	mouse table
<b>M5Generation</b>	Mouse 5's generation	mouse table
<b>M5GenoType</b>	Mouse 5's genotype	genotype table

### 3.5.3.6 Detail card for designing new cards

The report named TS\_DetailCageCardwithXtraFields may be used for designing a new cage card. It uses the fields and formats shown above. The following fields are added. Some of these fields are also on the MW\_DetailCageCardLandscape report, but they are not visible. Make a copy of this report before changing it. Up to 10 mice may be placed on a detail card. Fields for M6ID, M7ID up to M10ID are not provided. The user must create these. Be careful to follow the naming convention for the field name that is illustrated in these two tables.

<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>protocol1</b>	Protocol for mouse 1	Mouse table
<b>protocol2</b>	Protocol for mouse 2	Mouse table
<b>protocol3</b>	Protocol for mouse 3	Mouse table
<b>protocol4</b>	Protocol for mouse 4	Mouse table
<b>protocol5</b>	Protocol for mouse 5	Mouse table
<b>coatColor1</b>	Coat color for mouse 1	Mouse table
<b>coatColor2</b>	Coat color for mouse 2	Mouse table
<b>coatColor3</b>	Coat color for mouse 3	Mouse table
<b>coatColor4</b>	Coat color for mouse 4	Mouse table
<b>coatColor5</b>	Coat color for mouse 5	Mouse table
<b>comment1</b>	Comment for mouse 1	Mouse table
<b>comment2</b>	Comment for mouse 2	Mouse table
<b>comment3</b>	Comment for mouse 3	Mouse table
<b>comment4</b>	Comment for mouse 4	Mouse table
<b>comment5</b>	Comment for mouse 5	Mouse table
<b>room</b>	Room name	Room table
<b>weanDate1</b>	Wean date for mouse 1	Litter table
<b>weanDate2</b>	Wean date for mouse 2	Litter table
<b>weanDate3</b>	Wean date for mouse 3	Litter table
<b>weanDate4</b>	Wean date for mouse 4	Litter table
<b>weanDate5</b>	Wean date for mouse 5	Litter table
<b>penName</b>	Name of the pen	Container table
<b>penComment</b>	Comment for pen	Container table
<b>statusDate</b>	Date when pen status or room changed	ContainerHistory table
<b>statusDateWith Time</b>	Same as status date only includes the exact time of the change	ContainerHistory table

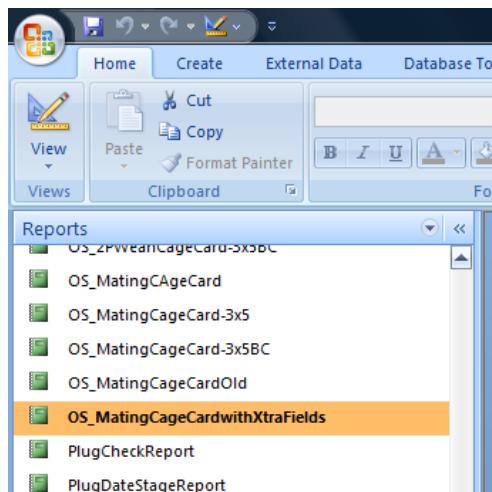
<b>Field name</b>	<b>Description</b>	<b>Data source</b>
<b>penStatus</b>	Current status of pen (active, retired, proposed, etc.)	Container History table
<b>PlanID</b>	Experimental plan ID	ExpPlan table (Only one plan ID for one of the mice is displayed when there are multiple possibilities)
<b>PlanName</b>	Experimental plan name	ExpPlan table
<b>M1NewTag</b>	Mouse 1's Replacement tag	mouse table
<b>M2NewTag</b>	Mouse 2's Replacement tag	mouse table
<b>M3NewTag</b>	Mouse 3's Replacement tag	mouse table
<b>M4NewTag</b>	Mouse 4's Replacement tag	mouse table
<b>M5NewTag</b>	Mouse 5's Replacement tag	mouse table

### 3.5.4 Creating Custom Cage Cards

JCMS is configured to allow new cage card designs to be easily created using the MS Access report design tools. An easy to follow tutorial may be found online (check out the v4.6 tutorial package found in the [JCMS Tutorials](#)).

When JCMS prints a cage card, it writes information from the database into text fields on the card. If a text box is put on a card and given a name that JCMS knows about, then JCMS will

write the associated information into that text box when the card is printed. For example, the field PIName is a text box on each cage card where JCMS prints the PI name (as specified in the setup variables).



**Figure 3-28 Office 2007 reports list shown using F11 key**

Make a backup copy of JCMS interface before making changes (Copy the interface JCMS.mdb file and save as {backup name}). To view the card designs in JCMS, logon as mtsadmin, press F11, and select the reports tab. To create a new card, simply draw the card wanted and add the fields (as specified in the tables above) that should appear on the card. Use the tables above as references showing what fields can be put on the card. JCMS is installed with a number of cards pre-configured for your use. To create a custom card, it may prove easiest to start with a card similar to the one wanted. Save a copy of the report format for this card under a new name. To make changes, open the report in "design view." Now

edit this new report format. Place the new report name into the setup variable value for the particular type of cage card. Four cage cards have been created specifically for use to help create new cage cards. These contain all possible fields. These reports have a name ending "withXtraFields".

### 3.5.5 User-designed Cage Cards

A separate JCMS cage card database, containing designs from the user community may be downloaded at the [colonymangement.jax.org](#) FAQs page.

#### Instructions for importing a cage card report (Access 2007):

- Download the JCMS Cage Card database from the JCMS website. Save.
- Open existing JCMS database (as user mtsadmin).
- Select External Data tab from the ribbon.

- Select Access Import icon.
- Select Import bullet.
- Select Browse button and navigate to the JCMS CageCard database.
- Highlight JCMS\_CageCards-1.mdb and select Open.
- Back to Get External Data window, select OK.
- Import Objects window opens. Select Reports tab.
- Choose Select All or highlight desired cage cards to import. Select OK.

**How to activate Cage Cards:**

- From the Administrator tab, select Administrator. Open JCMS Setup Variables.
- Scroll to MTS\_1PEN\_WEAN\_CAGE\_CARD. Type in the desired cage card name in the JCMS Setup Variable Value field.
- Scroll to MTS\_2PEN\_WEAN\_CAGE\_CARD. Type in the desired cage card name in the JCMS Setup Variable Value field.
- Scroll to MTS\_DETAIL\_CAGE\_CARD. Type in the desired cage card name in the JCMS Setup Variable Value field.
- Scroll to MTS\_MATING\_CAGE\_CARD. Type in the desired cage card name in the JCMS Setup Variable Value field.
- Close Setup Variables form.

### **3.6 Setting up an Experimental Plan**

In order to use the Experiments section of JCMS, the following CV tables must also have terms entered into them: **field of study** and **keywords**. Controlled vocabularies are changed from the **Administrator** button bar and may only be changed by the mtsadmin user.

See the section on [Experimental Plans](#) for more information on the details of setting up this portion of the database. Owners are able to set up all other aspects of Experimental Plans.

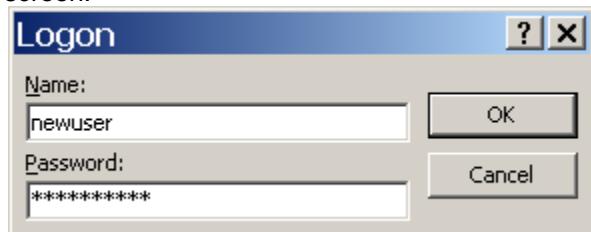
## **4 User Setup**

A new user must first be setup with an MS Access account and password logon by the Administrator. The Administrator will also do any client installation necessary on a computer used by the new user.

### **4.1 Logging On and Passwords**

To provide database security MS Access requires all users of JCMS to have a unique logon account and password. This logon is separate from any other that may be used by a user's system such as the Windows logon.

To start JCMS, double click the JCMS icon on the desktop. A dialog box will request the user name and password. The user logon name will be displayed on each form that is displayed on the screen.



**Figure 4-1 User Logon Screen**

The JCMS Welcome window displays at startup. Click the *Start (Workstation)* or *Start (Handheld Device)* button to begin work. The *Main Button Bar* is displayed next. From this button bar forms can be opened for entering and viewing JCMS data.

Check with the Administrator if entering the logon and password he/she provided does not result in the JCMS welcome screen appearing.

## 5 Basics on Using JCMS

### 5.1 What are Owners and Secretaries?

Access to the functions of the database is restricted on a form-by-form basis. That is, each form is programmed to allow access to JCMS for certain users (based on their security level). JCMS defines a security hierarchy with three levels of permission: Administrator, Owner, and Secretary. The Administrator can access all forms and all data; owners can enter mice, matings, experimental plans, samples, and experimental data and access all data entry and query forms from the Main Button Bar; secretaries can usually access only a limited subset of data entry forms.

Within the database, *mice*, *matings*, *experimental plans*, *experimental data*, and *samples* have *owners*. The owner of a mating also owns any *litters* produced by that mating. The owner of an experimental plan also owns any experimental data that is part of the plan. Owners may use all forms that have access permission to the database at an *owners* or *secretary* level. Forms for editing a mouse, mating, litter, experimental plan, samples, or experimental data usually restrict edit permission to owners only.

*Secretaries* may use only those forms with secretary level access. Additionally, secretaries are associated with owners (by being members of an owner's secretary group); thus, secretaries can only work with data associated with a specific owner.

It is possible for a user to be both an owner of their own mice and to act as the "secretary" for one or more other owners.

The Administrator will have decided the owner or secretary status of each user and set this up as part of the user's logon. The Administrator may also change the default security access level for most forms.

### 5.2 Changing Passwords

When a user logs in for the first time, he/she may not have a password or may have a special password assigned by the Administrator. To create or change this password, select from the main menu, **tools – security – user and group accounts** (Office 2007: Database Tools tab, Administer – Users and Permissions, User and Group accounts; Office 2010: File tab – Users and Permissions – User and group accounts). Click on the **Change Logon Password tab** and add or change the password. Only you can change your password.

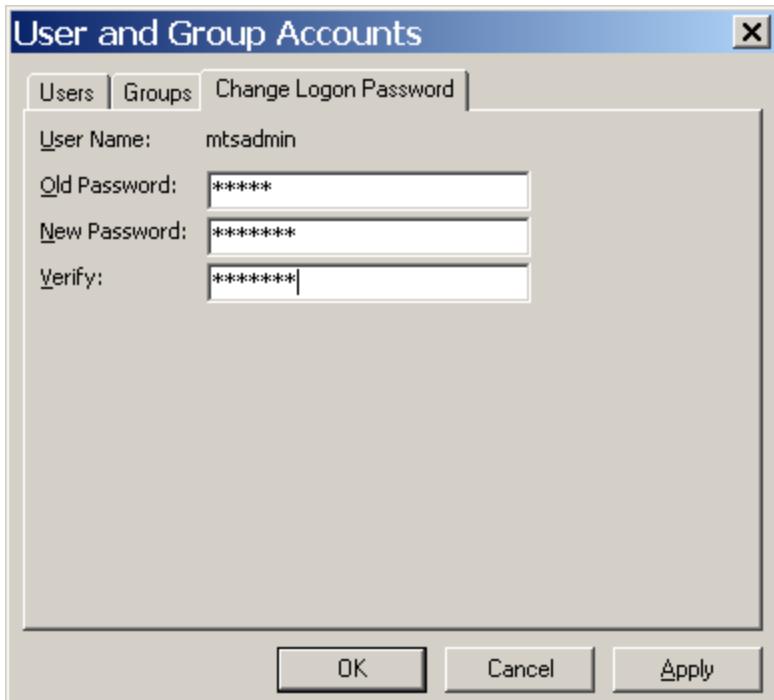


Figure 5-1 Dialog Box: Change Password

### 5.3 Button Bars

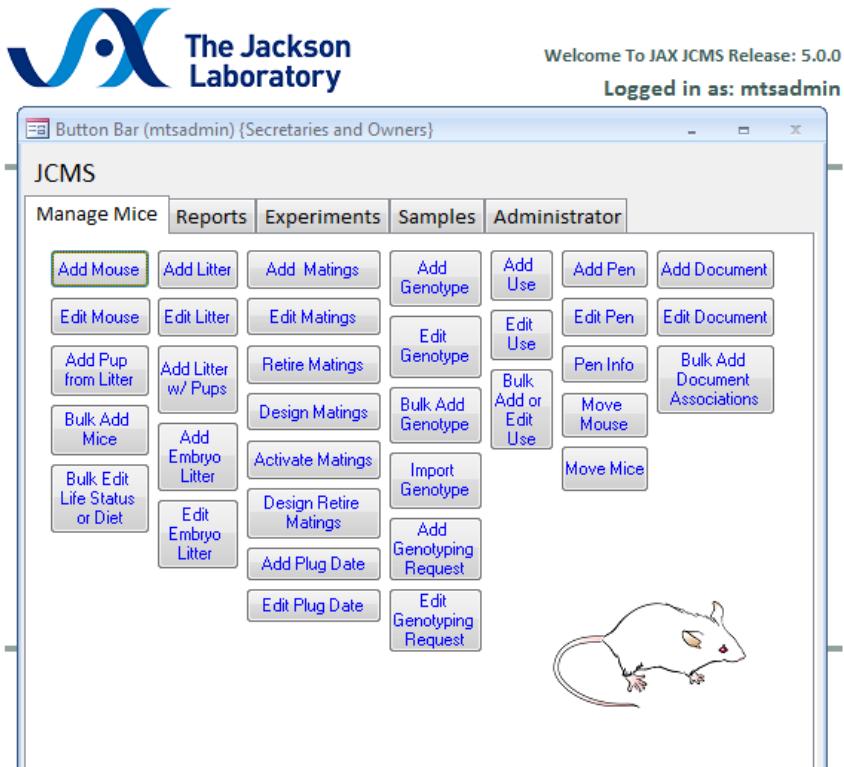


Figure 5-2 Main Button Bar

JCMS uses button bars to provide easy access to its many functions.

Clicking buttons on the button bar(s) opens JCMS forms. After pressing the start (workstation) button, the main button bar opens and is always open from then on (until JCMS is shut down). The main button bar cannot be closed, but can be minimized.

The main button bar is organized into sections with tabs and then into columns for different functions. Note that some buttons will appear to not work for "secretaries" because they do not have permission to open that form. The Administrator button will work only for the Administrator.

## 5.4 Using the Forms

Forms may be **closed** or **minimized** using the normal Windows minimize (–) button and close (x) button. Multiple forms may be open on the screen at one time, however updates submitted on one form may not be updated on another open form. Close and reopen the other form to see all updated information.

The screenshot shows a Windows-style dialog box titled 'Add a New Mouse'. At the top left is a small icon of a computer monitor. The title bar says 'Add Mouse (mtsadmin) {Secretaries and Owners}'. The main area has a black header bar with the title. Below it, there are several input fields: a text box for 'Mouse ID', a dropdown menu for 'Protocol ID', a dropdown menu for 'Litter #', and a dropdown menu for 'Strain'. The 'Strain' dropdown is currently set to 'BALB/cByJ'. There is also a radio button labeled 'All' and a text box next to it.

Figure 5-3 Example of a text box

There are several general form types: edit, add-new, request reports, and query. Some forms allow combined functionality (e.g. allow you to add new or edit). Basic data entry is done using text boxes that are white. Text boxes that are blue-green in color are "active" boxes. Entering a valid identifier in an active box followed by moving the cursor out of the box (by tabbing or clicking the mouse pointer elsewhere) will cause the database to display information associated with that item.

Data entry boxes with a drop down list (click on the down arrow ▾) allow choosing an item from a list. The item can also be typed directly into the box; as you type, the first item in the list that matches the characters typed so far is displayed. In either case, press tab to accept an item and move to the next field. Most drop down lists are limited to selecting items in the list and anything else that is typed will be rejected.

The screenshot shows a Windows-style dialog box titled 'Pen pals'. The title bar includes a small icon of a computer monitor and the text 'Pen pals'. The main area contains a table with data. The table has columns for 'ID', 'Sex', 'breedi', 'lifeSta', 'birthDate', 'generation', and 'StrainName'. Two rows of data are shown: FVB-M-101 (M, B, A, 6/1/2013, F01, FVB/NJ) and FVB-M-210 (F, B, A, 9/2/2013, F02, FVB/NJ). Below the table, there is a checkbox labeled 'Auto increment ID'. At the bottom of the form are five buttons: 'Submit', 'Clear', 'Session Report', 'Set Genotype', and 'Set Mouse Use'.

Figure 5-4 Example of regular form buttons

Gray boxes display information that cannot be changed using this form. All forms can be navigated through quickly by tabbing to move forward and shift tabbing to move backward through the fields. When tab does not work, try pressing Control and Tab simultaneously.

Any form that has a **Submit button** will not update the database until this button is clicked. If a form is closed before hitting the submit button, changes made on the form will not be saved in JCMS.

The **Clear button** is used to remove all data from the form. The clear button does not delete data from the database.

Extra button(s) on the form will often open another form showing data associated with an ID on the first form (mouse, litter, mating, pen, experimental data, etc.) For example, the **Set Genotype**

**button** on the edit mouse form opens the add genotype form showing data for the mouse ID used on the edit mouse form, all ready to work with.

#### **5.4.1 Special Features of Some Forms**

Why are adding and editing functions kept separately on different forms? In general, add and edit forms will look very similar to each other. Different forms are used because creating a new record is a separate function from changing an existing one. If an error has been made on an add form, then open the edit version of this form to correct it. Edit forms limit editing capabilities based on the *owner* or *secretary* permission status.

Color fields on submit: by default, some of the forms use auto-color. After a new record is submitted to the database, all the text fields (except checkboxes and dates) have their background colored green. When these fields are visited by tabbing or using the mouse, the colored background disappears. The purpose of this feature is to prompt visiting each field when doing a lot of data entry so that values from the previous record are not erroneously entered into the new record. This feature may be turned off by the Administrator in the setup variables by changing MTS\_AUTO\_COLOR to false.

Pen Info Button: this button allows viewing the contents of a pen related to the form. When this button is pressed the pen info form appears with a list of all mice in the pen and other pertinent information about the pen.

Auto Increment ID check box: when this box is checked, the main ID field of the form is automatically incremented after JCMS has been updated (new record added or edited for this ID). This feature is useful when entering a lot of information that uses successive ID numbers. NOTE: ID numbers may have characters and leading zeros in them. Thus, the ID number cd012 would increment to cd013, and 099 would increment to 100 (as examples). The default for this check box to be set to on or off on most forms is changed in the setup variables by the Administrator using MTS\_DEFAULT\_AUTO\_INCREMENT, JCMS\_ACTIVATE\_MATINGS\_INCREMENT, etc. Some forms always default to NOT auto increment regardless of the value of this setup variable.

#### **5.5 Navigation Buttons**

Some forms will use a special set of navigation buttons to move from one record to another.



Clicking **▶** will move forward by one record, **▶ |** will move to the last record, **▶ \*** will provide a blank space to enter a new record. The number in the box indicates which record is currently displayed (first, second, third, etc.) Records are numbered by the order they are sorted into in the database. If the sort criteria is changed, this number may change for the record on display, it only reflects the record's relative placement in the sort order.

#### **5.6 Session Reports**

Session box: session boxes give some history information about this edit session. A session lasts for the period of time the form is open on the screen. A typical session box shows the ID and one or two bits of important information about each record entered or edited. The most recent entries are listed first in the list box.

Printable session report: use the Session Report button to see a print preview of a more detailed session report than is available in the session box.

#### **5.7 Printing**

JCMS will use the default printer as set up on the client machine. Most reports will first display in print preview mode on the screen in order to help save printing unnecessary pages.

Use the *Select Default Printer* button (Administrator tab) to temporarily change the default printer for one JCMS session. The change will not affect the Windows default printer for any other program.

Many modern printers will use the sheet feeder as the default whenever there is paper in the sheet feeder. To print cage cards, open the sheet feeder and load the cards into the envelope feeder part of it. Also set the printer to use as straight a paper path as possible. Many printers have an option for sending sheets out the back if it is open or have a toggle switch to change the path.

Forms are not designed to print nicely on a page, use a report for a quality printout of information in a form. However, there are circumstances where it may be advantageous to print a copy of a form.

Use the **printer icon on the toolbar** with caution; this will often try to print the entire table, not just the values shown on the form on the screen. To determine if the form fits on one page of paper, try using the **print preview button** on the toolbar. If the screen is truncated on the right, change the page setup to landscape. This will also show if the results give all the records in the table instead of just one. Use the **close** button on the print preview window to get back to the form or the print icon to print what is shown in the preview.

To **print just a selected record** from forms that print the whole table, use the File menu. **Select Print from the File menu** and the print dialog box will open. Under the print range, pick **Selected record(s)**. Now, only the record on the form or with the cursor in a field on it will print.

Most forms will not **print with all blank fields**. When the cursor is placed into a new blank record, *print the selected record* results in only the header for the form. To print a blank copy, first create a dummy record, placing something such as a decimal point into the required fields. Then print this selected record and delete it. Also note that only the Administrator has permission to delete records from some of the tables.

## 5.8 Statistics

Click the Data Stats button on the Reports tab to get information on the total number of pens, mice, and matings in the database.

# 6 Mice

## 6.1 Mouse IDs

Each mouse must have a unique mouse ID number containing up to 16 characters. JCMS will not allow entry of duplicate ID numbers. The following setup variables may be used to help create sequences of mouse IDs when mice are added in groups.

### 6.1.1 Automatically Incrementing Mouse IDs

Many forms have a check box labeled "Auto increment ID". When the box is checked, after successfully adding a new mouse, the next mouse ID in sequence will be suggested.

The setup variable JCMS\_MOUSEID\_INCREMENT\_RIGHTMOST indicates which portion of the ID to increment if it contains more than one numeric section. For example, if the setup variable is "false" the base ID 0015BALB-600 would be incremented to 0016BALB-600 and if it is "true" it would be incremented to 0015BALB-601.

Note the numeric portion of the ID that is incremented is limited to a 10 digit number. Using numbers larger than that will return a message "Unable to increment ID". Longer numbers may be used if the mice are entered individually on forms such as Add mouse and Add mouse at wean.

### 6.1.2 Base Mouse Numbers

The forms Bulk Add Mice, Add Litter w/Pups, and Handheld Add Litter w/Mice allow entering multiple mice at once. There are two methods of determining what mouse ID to start with.

- 1) The setup variable MTS\_MOUSE\_ID\_PREFIX contains a character string used at the beginning of a mouse ID. JCMS keeps track of the last integer used in the sequence and automatically increments it when adding mice. For example, if the prefix is "A" and the last mouse ID added was A98, JCMS will use the sequence A99, A100, A101, etc. This option is the default on the forms.
- 2) The default (MTS\_MOUSE\_ID\_PREFIX) may be overridden by selecting the check box "**Use Base ID**" or "**Use base mouse number**" and then entering a starting mouse ID for the sequence. A list of the current mice is provided on the non-handheld forms to help with determining the sequences in use.

## 6.2 Adding Mice

The screenshot shows the 'Add a New Mouse' form in the JCMS application. The form is titled 'Add a New Mouse' and is part of the 'Add Mouse (mtsdadmin) {Secretaries and Owners}' module. The form contains the following fields and sections:

- Protocol ID:** A dropdown menu.
- Litter #:** A dropdown menu.
- Strain:** A dropdown menu with radio buttons for "All" and "Active only".
- Generation:** A dropdown menu.
- Date born:** A dropdown menu set to 12/16/2011.
- Cause of death:** A dropdown menu.
- COD notes:** A text input field.
- Sex:** A dropdown menu.
- Life status:** A dropdown menu.
- Breeding status:** A dropdown menu.
- Coat:** A dropdown menu.
- Diet:** A dropdown menu.
- Owner:** A dropdown menu.
- Origin:** A dropdown menu.
- Replacement Tag:** A text input field.
- Comments:** A large text area.
- Pen:** A section with fields for "ID" (dropdown menu), "Name" (dropdown menu), "Status" (dropdown menu set to "active"), and "Date" (dropdown menu set to 12/16/2011 4:44 PM). It also includes checkboxes for "Use next available ID" and "Increment Name", and buttons for "Edit Pen" and "Pen Info".
- Room:** A dropdown menu set to "unknown". Next to it is the text "H Lvl: 2 since 1/1/1990".
- Pen pals:** A section showing five empty boxes for listing pen pals.
- Buttons:** At the bottom are buttons for "Submit", "Clear", "Session Report", "Set Genotype", and "Set Mouse Use".
- Checkboxes:** A checkbox for "Auto increment ID" is located near the bottom left.

Figure 6-1 Add Mouse Form

Use the **Add Mouse** button on the manage mice tab to open a form for adding mice one at a time. If adding mice to the database when they are weaned, consider using the **Add Mice at**

**Wean** instead. It will automatically fill in the information from the litter for each mouse, saving time.

If this version of JCMS has been set up to use a prefix on the mouse ID, this is only entered if JCMS is creating the ID numbers. On this form, the prefix must be entered on the ID number by the user.

When mice are added, they must be assigned to a pen. The pen information section of these forms is used for this purpose. If the pen ID or pen name selected on the form already exists, the room, list of mice already in the pen, and other information will be retrieved and displayed in the pen information section. To add a new pen for this mouse, check "Use next available ID" to create it with the next available pen ID. Or uncheck the box and enter an unused pen ID by hand. Pens may have duplicate pen names but the pen ID is always unique.

The "Submit" button must be pressed to add the mouse into JCMS. If an error message is given, the error must be corrected and the submit button pressed again. A successful submission will result in a message stating, "Mouse xxx submitted" in red on the form. The session box will also list the mouse ID, pen ID, and pen name.

The Add Mouse form ignores the setup variables JCMS\_AUTO\_RETIRE\_PENS and JCMS\_AUTO\_RETIRE\_MATINGS.

The screenshot shows the 'Add a New Mouse' window in JCMS. The main area contains various input fields for mouse metadata such as Mouse ID, Protocol ID, Litter #, Strain, Generation, Date born, Sex, Life status, Breeding status, Coat, Diet, Owner, Origin, and Replacement Tag. There are also dropdowns for comments, room, and pen details. A 'Pen' panel on the right provides specific information about the assigned pen, including its ID, name, status, and date. A 'Mice entered this session' panel displays the message 'Mouse Oct-2011-050 submitted'. At the bottom, there are buttons for Submit, Clear, Session Report, Set Genotype, and Set Mouse Use.

**Figure 6-2 Add mouse form after a successful submit**

## 6.3 Editing Mice

Changes are made to individual mice using the **Edit Mouse button** to open the edit mouse form. This form works in the same manner as the Add mouse form. If a selected mouse has genotype information associated with it, it will be displayed on the form.

### 6.3.1 Changing the ID of a Mouse

The screenshot shows a dialog box titled "Change Mouse ID Form". At the top left is a small icon of a computer monitor. The title bar reads "Change Mouse ID (OWN1) {Owners Only}" with standard window control buttons. The main area contains the following text and controls:  
Old Mouse ID: Feb3-001  
New Mouse ID:   
Submit change      Close Form

**Figure 6-3 Change Mouse ID Form**

The ID of any mouse in JCMS may be changed as long as the new name is unique in the database. For example, if mice are ear notched so they are unique in a pen, punches may need to be added to the mouse ear when the mouse is moved to a new pen. A code could be appended at the end of the mouse ID number to indicate its current ear-punch value (e.g., 2373-R, 2373-L, 2373-RR, 2373-LL, 2373-RL, 2373-RRL, 2373-RLL, 2373-RLLL). When

the mouse moves, and the notch changes, update the mouse ID accordingly. To change a mouse ID, use the mouse edit form. Click the Change ID button on the form. Fill out the dialog box with the new ID and press submit change to change the mouse ID in JCMS. (The ID may not be changed to an already existing ID).

The Edit Mouse form also allows you to navigate through the mice in the database - a mouse browser.

**Edit a mouse**

**Apply search criteria:**

Mouse ID like	<input type="text" value="F"/>	Strain	<input type="button" value="▼"/>	<input type="button" value="▼"/>	<input type="button" value="Apply Criteria"/>
Life status equals	<input type="button" value="▼"/>	Owner ID equals	<input type="button" value="▼"/>	<input type="button" value="Clear Criteria"/>	

<Previous      Next>

**Mice edited this session**

*Mouse ID: <input type="text" value="Feb3-001"/> <input type="button" value="Change ID..."/>	Vial Position: <input type="text"/>
Protocol ID: <input type="text"/>	Vial ID: <input type="text"/>
Litter #: <input type="text" value="830"/>	
*Strain: <input type="text" value="FVB/NJ"/>	
*Generation: <input type="text" value="F01"/>	Age expressed in: <input type="radio"/> Days <input checked="" type="radio"/> Weeks <input type="radio"/> Months <input type="text" value="3"/>
*Date born: <input type="text" value="1/15/2010"/>	Date exited: <input type="text" value="2/3/2010"/>
*Sex: <input type="text" value="F"/>	Cause of death: <input type="text"/>
*Life status: <input type="text" value="A"/>	COD notes: <input type="text"/>
*Breeding status: <input type="text" value="V"/>	
Coat: <input type="text"/>	
Diet: <input type="text"/>	
*Owner: <input type="text" value="OWN1"/>	<b>Pen</b>
*Origin: <input type="text" value="internal"/>	*ID: <input type="text" value="697"/>
Comments: <input type="text"/>	Name: <input type="text" value="Section 5A-22"/>
<input type="checkbox"/> Auto increment ID	*Status: <input type="text" value="active2"/>
	*Date: <input type="text" value="2/03/2010 4:03 PM"/>
	Comments: <input type="text"/>
	*Room: <input type="text" value="LAH7756"/> H Lvl: 1 since 4/17/2009 12:00:23 PM

Add Pen      Pen Info

Submit      Clear      Session Report      Set Genotype      Set Mouse Use

**Figure 6-4 Edit Mouse With Browser Functions**

The user can set search criteria and apply them to the mice in the database. When applied, only those mice meeting the criteria will become available in the lower section for viewing or editing. The criteria are ANDed together, that is, for a mouse to be included it must meet all of the selected criteria. Some fields support substrings while others (life status and owner) accept only exact matches.

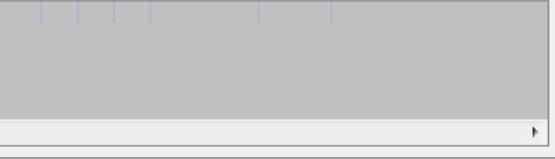
A live mouse may not be put into a retired pen. This form will retire pens and matings on the mouse exit date if the setup variables JCMS\_AUTO\_RETIRE\_PENS and JCMS\_AUTO\_RETIRE\_MATINGS are true. Note that resurrecting a mouse will not resurrect its retired matings or pen.

## 6.4 Importing or Bulk Adding Mice

This form is useful when bringing a group of mice in from outside the colony or when adding a new group of mice that are all similar. The group must all have the same information for the required fields, the fields marked on the form with an \*. JCMS will automatically generate unique mouse ID numbers and will increment by one for each additional mouse imported. Set the Base ID to the first ID number to be used. This form will not add any mouse ID Prefix specified in the setup variables; enter the prefix as part of the base ID.

**Bulk Add Mice (mtsadmin) {Owners Only}**

### Bulk Add a Group of Mice

*Number of mice:	*Mice/pen [0,10] 10	Mice entered this session	
<input checked="" type="checkbox"/> Use base ID Base ID ST-020	Current mice ST-019		
Protocol ID			
*Strain			
<input type="radio"/> All			
<input checked="" type="radio"/> Active only			
Cage card color			
*Generation			
*Date born	12/19/2011	<input checked="" type="checkbox"/> Use next available ID	
*Sex:			
*Life status:			
Exit date:	12/19/2011	<input checked="" type="checkbox"/> Increment Name	
*Breeding status:			
Coat:			
Diet:			
*Owner:			
*Origin:			
Comments:			
<b>Pen</b> *ID: <input type="text"/> Name: <input type="text"/> *Status: active *Date: 12/19/2011 1:13 PM Comments: *Room: unknown H Lvl: 2 since 1/1/1990			
<b>Pen pals</b> 			
Cage cards for last group submitted <input type="button" value="Print Cage Cards"/> Intended use <input type="checkbox"/> Preview			
<input type="button" value="Submit"/> <input type="button" value="Clear"/> <input type="button" value="Session Report"/> <input type="button" value="Undo LAST"/>			

**Figure 6-5 Bulk Add Mice Form**

The new pens will be automatically assigned and filled with the number of mice specified. "Use next available ID" must be checked. Note that live mice may not be put into a retired pen.

Once mice are added, cage cards can be printed. Any intended use entered on this form will only be printed on the cage cards; it will not be saved with the mouse.

Do not use this form if the mice have a dam and sire already in JCMS. Create a litter and use one of the litter forms instead. Otherwise, it will not be possible to create a pedigree tree.

## 6.5 Adding Pups

The Add Pup from Litter form is used to add mice into the database when they are weaned. This form adds mice one at a time and must be used if the litter record has already been created. If not, consider using the Add Litter w/Pups form instead. That form will add the litter record plus all the pups as weaned mice in one step. Which form will work best is dependent upon the workflow.

Once the litter # is entered, the form will display the strain, generation, date born, wean date, and tag date from the litter record. It automatically sets the life status to alive and breeding status to virgin.

Any change made to the wean date or tag date on this form will also update the litter record when the mouse is submitted.

Add Pup from Litter (mtsadmin) {Owners Only}

### Add Pup from Litter

**Mouse Oct-2011-054 submitted**

*Mouse ID	Oct-2011-055	Protocol ID	Oct-2011-054	Current mice	<input checked="" type="radio"/> Litter strain only <input type="radio"/> All
*Litter #	50		Oct-2011-050		
*Strain	<input type="radio"/> All <input checked="" type="radio"/> Active only B6.129P2-Apo<tmlUnc> /J				
Generation	N02	Date Born	10/21/2011		
Weaning date	11/8/2011	Tagging date	11/8/2011		
*Sex	M	Name:	Section J-052	<input type="checkbox"/> Use next available ID <input checked="" type="checkbox"/> Increment Name	
Status	A	*Status:	active		
Breeding Status	V	*Date:	11/15/2011 12:44 PM		
Coat	Grey/White	Comments:			
Diet	4%				
*Owner	nobody				
Origin					
Comments					

**Mice entered this session**

Oct-2011-054/71 Section J-052
Oct-2011-053/71 Section J-052
Oct-2011-052/70 Section J-051
Oct-2011-051/70 Section J-051
Litter #50 updated. Born=5 F=3 M=2

**Pen**

*ID:	71	<input type="checkbox"/> Use next available ID
Name:	Section J-052	<input checked="" type="checkbox"/> Increment Name
*Status:	active	
*Date:	11/15/2011 12:44 PM	
Comments:		

**Pen pals**

ID	Sex	breed	lifeSta	birthDate	generation	StrainName
Oct-2011-053	M	V	A	10/21/2011	N02	B6.129P2-Apo<tmlUr
Oct-2011-054	M	V	A	10/21/2011	N02	B6.129P2-Apo<tmlUr

Add to experimental plan(s)

<input checked="" type="radio"/> None	3	Test Plan 3
<input type="radio"/> Plan only	2	Clinical tests
<input type="radio"/> Plan + all tests	1	Use for testing documents

Auto increment ID

**Litter Record**

# Born	5	# Females	3	# Males	2
Actual Mouse Records	5	3	2	<b>Update Litter Rec. to</b>	
<b>Update Litter Rec. to</b>					

**Buttons**

- Submit
- Clear
- Session Report
- Set Genotype
- Set Mouse Use

**Figure 6-6 Add Pup from Litter Form**

The current mice box (top center) is an aid for determining the next mouse ID to use if a sequence of ID numbers should be maintained. It shows the existing mouse IDs.

As mice are added, the litter record box (bottom right of form) is updated to show the actual number of mice in JCMS for this litter. Once all pups have been added, click the Update Litter button to change the litter record to the numbers shown in the red “update” boxes. The “update” boxes may be changed by the user to any desired number (or blank). If pups have died, the number born or females/males in the litter record may be different from the actual mouse records.

As mice are added, they may also be added to experimental plan(s) and all the experimental tests in those plans. The session box will list the plan and test IDs.

When the setup variable MTS\_RELAXED\_PEN\_NUMS is false a “Next ID” button is displayed instead of the “Use Next Available ID” check box. This button will provide a pen ID. Make sure to use the Pen Info button and print the cage card.

Note that a live mouse may not be put into a retired pen.

## 6.6 Changing Life Status or Diet of a Group of Mice

The screenshot shows the 'Bulk Edit Life Status or Diet' window. At the top, there are two radio buttons: 'Edit Life Status' (selected) and 'Edit Diet'. Below this, there are three sections for selecting a group type: 'By mouse ID' (selected), 'By litter number', and 'By pen ID'. In the 'Mouse ID' section, there are three options: 'Range' (with min and max fields), 'Selected mice from list' (which is selected and shows a list of mice IDs: A338, A339, A337, A336, A335, A334, A333), and 'Litter #' (with range and selected litter fields). In the 'Pen ID' section, there are also three options: 'Range' (with min and max fields), 'Selected pens by Name from List' (selected and shows pens AB-01 through AB-06), and 'Selected pens from list'. On the left, there are fields for 'New diet' (dropdown menu), 'New life status' (dropdown menu with 'E' selected), 'Exit date' (set to 2/9/2010), 'Cause of death' (dropdown menu with 'E' selected), and 'COD notes' (text input field). Below these fields is a note: 'The "Show-me" button will display a list of the mice that may be edited. This list can then be printed. To make changes you must press "Submit." Any mice that you do not own or mice that already have their exit date set will not be changed.' At the bottom are three buttons: 'Show me', 'Show me print', and 'Submit'. A message at the bottom left says 'Show me list: 5 mice selected.' On the right side of the window, there is a table showing the current status of the selected mice:

Mouse ID	Strain	Gen	Sex	Birth Date	Exit Date	LS	BS	Diet	Pen	Owner
A338	C3H/HeJ	F01	F	1/12/2010		A	V	4%	630	OWN1
A337	C3H/HeJ	F01	F	1/12/2010		A	V	4%	629	OWN1
A336	C3H/HeJ	F01	F	1/12/2010		A	V	4%	629	OWN1
A335	C3H/HeJ	F01	F	1/12/2010		A	V	4%	629	OWN1
A334	C3H/HeJ	F01	F	1/12/2010		A	V	4%	628	OWN1

**Figure 6-7 Form: Bulk Change Life Status or Diet**

JCMS allows users to do a few edit operations en-mass. Select the Bulk Change Life Status or Diet button. First select change the life status or change the diet. Mice may be picked for the edit operation by litter, pen, or mouse ID. Use the “Show me” button to display a list of mice that may be affected by the operation when submit is pressed. No changes are made to the mice until submit is pressed. Any mice not owned by the current user or that already have their exit date set will not be changed. Use the “Show me print” button for a printable list of the mouse changes. If the setup variable JCMS\_AUTO\_RETIRE\_PENS is true, pens now containing only exited mice will be retired as of the exit date. If the setup variable JCMS\_AUTO\_RETIRE\_MATINGS is true, all mice that are exited have their matings checked. Those active matings now consisting of only exited mice will be retired as of the exit date.

## 6.7 Changing Life Status using the Handheld Forms

The Bulk Exit handheld form allows changing the life status of all mice in the selected pens at once. All the mice must be changed to the same new life status, exit date, and cause of death. This form does not allow changing a mouse to a non-exit life status. It also will not make any changes to mice that already are exited. If the setup variable JCMS\_AUTO\_RETIRE\_PENS is true, pens now containing only exited mice will be retired as of the exit date. If the setup variable JCMS\_AUTO\_RETIRE\_MATINGS is true, all mice that are exited have their matings checked. Those active matings now consisting of only exited mice will be retired as of the exit date.

## 6.8 Change Life Status of an Individual Mouse using Handheld

Use these handheld forms to change the mouse life status. Select pens and use **Bulk Exit** to simultaneously exit and wean all mice in the pens. Select one pen and use **Individual Mouse Change Life Status** to change the life status and exit information for selected mice in that pen. The new life status, exit date, and cause of death will be the same for all mice. Using the **Change** form, a mouse may be changed from an exit to a non-exit life status. However, if the pen is

retired, the mice in the pen may not have their life status changed to a non-exit life status (retired pens are expected to be empty, i.e. the mice left in them are no longer alive). Either move the mice into an active pen or change the pen to no longer be retired.

If the setup variable JCMS\_AUTO\_RETIRE\_PENS is true, after changing the life status of the mice, pens now containing only exited mice will be retired as of the mouse exit date. If the setup variable JCMS\_AUTO\_RETIRE\_MATINGS is true, all mice that are exited have their matings checked. Those active matings now consisting of only exited mice will be retired as of the exit date. Note that resurrecting a mouse will not resurrect its retired matings.

ID	Status	ExitDate
A432	E	2/9/2010
A433	E	2/9/2010

**Figure 6-9 Bulk Exit Handheld Form**

ID	Status	ExitDate
A432	E	2/9/2010

**Figure 6-8 Individual Mouse Change Life Status Handheld Form**

## 7 Pens and Cage Cards

In JCMS we use the terms pen and cage interchangeably.

All mice reside in a cage or pen. In JCMS each pen has a unique pen ID number. A pen may have an optional pen name that does not have to be unique. A pen may be empty. A pen has a pen status, such as proposed, active, or retired, and a history of when the pen's status changed and when it moved between rooms.

### 7.1 Manage Pen Configuration Form

The Manage Pen Configuration form, accessed from the Administrator button bar, provides an interface for changing the settings of setup variables related to pens. See section 3.4 for more details.

The screenshot shows the 'Manage Pen Configuration' window with the title 'Manage Pen, Room, and Cage Card Setup Variables'. The window is divided into three main sections: 'Defaults', 'Pen Billing', and 'Cage Cards'.

- Defaults:** Contains fields for 'Pen Status' (active), 'Room' (Annex 6), 'Health Level' (unknown), and 'Maximum number of live mice in a pen' (10). It also includes several checkboxes:
  - Use Pen Names
  - Use Pen Comments
  - Use Health Level
  - Sort by Pen Name
  - Warn about duplicate Pen Name
  - When incrementing a pen name, increase the rightmost number
  - "Use next available Pen ID" option is selected when adding pens
  - Automatically retire pens if empty or containing only exited mice
  - Handheld: Print cage cards option is selected
- Pen Billing:** Contains radio buttons for 'Pen Billing' options: Full Days (selected), Partial First Day, Partial Last Day, and Any Day.
- Cage Cards:** Contains settings for 'Formats':
  - Detail: CC\_JCMS\_DetailCageCardUsingPenNames
  - Mating Format 1: CC\_JCMS\_MatingCageCardUsingPenNames
  - Mating Format 2 (For Handheld): TS\_MatingCageCardStyle2WithBarcodeIt also includes fields for 'Print prefixes before:' (Pen ID, Mating ID M, Litter ID L), 'Detail card note' (sample card note), 'Phone number' (xxx-xxxx-xxxx), 'Lab name' (Lab name goes here), and checkboxes for 'Print exited mice' and 'Relax restrictions on pen IDs'.

A 'Submit' button is located at the bottom left of the form.

**Figure 7-1 Manage Pen Configuration Form**

The default *pen status* and *room name* are used to automatically pre-set these values on forms where new pens are added. Most forms (except some of the handheld forms) provide an option for overriding the default. Similarly, the default *health level* is used on the form where new rooms are added. The health level is always associated with a room.

Many add forms limit the number of live mice in a pen to the number specified in the *maximum number of mice in a pen* which may be overridden on the form. This number is used to separate the mice into different pens when adding a large litter or group of mice at once.

*Pen names, pen comments, and room health level* are optional and do not have to be used.

When using pen names, it may be convenient to “sort by pen name”, which will cause some drop-down lists to be ordered by the pen name instead of pen ID. It is possible to have *duplicate pen names* (the pen ID will always be unique). When duplicate pen names are not desired, have JCMS warn before adding a duplicate pen name.

When adding pens, if “*increment name*” is checked on the form, the pen name shown will be incremented in the same fashion as mouse IDs. *Increment rightmost* will add one to the rightmost integer. If not selected, one will be added to the leftmost integer. The numeric portion of the pen name that is incremented is limited to a 10 digit number. Using numbers larger than that will return an “Unable to increment” error message. Longer numbers may be used if the pen names are entered individually by hand.

Pen ID numbers are unique integers that are either generated by JCMS (and therefore assured to be unique) or provided by JCMS users. When “*Use next available ID*” is checked, JCMS will assign the next largest pen ID. If the user prefers to generate pen IDs, they may hand enter them instead. JCMS will never allow duplicate IDs to be entered into the database. JCMS will also warn if a user tries to enter a pen ID much larger than any previous pen ID used (it keeps track of the largest pen ID used). The largest pen ID that can be entered into JCMS is 2,000,000,000. Very large pen IDs may not print out correctly on all cage cards due to lack of space. If this is a problem, cage cards can be configured to have more space for display of the pen IDs. A short string of characters can be prefixed on Pen IDs generated by JCMS when they are printed. The Administrator must specify this string in the *PEN\_ID prefix* setup variable.

*Mating ID* and *Litter ID* prefixes may also be specified for printing on cage cards.

When *automatically retire pens* is selected, any time a mouse is moved, pup is weaned and moved, or a mouse’s life status is changed to an exit life status the pen is checked to determine if it is empty or only contains exited mice. If yes, then the pen is retired using the date of the change (exit date, wean date, mating date, or today’s date).

If the *handheld: print cage cards option* is checked then certain handheld forms open with the print cage card check box selected.

Pens have a *history* that records the date and time a pen was established, retired, had its status changed, or was moved into a different room. JCMS provides the default *pen status* terms proposed, active, and retired. The user may add other terms. A pen status may be either billable or non-billable. For example, a proposed pen may not contain mice and, therefore, is not billed; a retired pen also contains no live mice and is not billed.

*Pen billing* is used by the Cage Use and Cage Use Summary reports to determine how many pens to count each day. For reporting purposes, a pen is either “billable” (counted) or “non-billable” (not counted) on a specific date. Use the Administrator *pen status* form to change the billable attribute of a pen status term. When counting, a “partial” day is a day when a pen was moved into or out of a room or a day when a pen’s status changed from a billable to a non-billable status. The date a pen was established is a “partial” day. The date a pen was retired is a “partial” day.

JCMS supports several *cage card formats*. Each type of format must have a cage card *report* selected. For example, a detail cage card could use the report CC\_JCMS\_DetailCageCardUsingPenNames or MW\_DetailCageCardLandscape.

The *Detail card note*, *Phone number*, and *Lab name* are printed on many cage card formats.

*Print exited mice* specifies if a cage card should list all mice in that cage, including those with an exit life status, or show only those still alive.

*Relax restrictions on pen IDs* indicates if the user should have an unrestricted ability (checked) to create pens or if some forms will only create pens that already have had a cage card (possibly blank) printed with the pen ID on it.

## 7.2 Rooms and Health Level

*Health level* is associated with a *room* and a *history* of health level changes is maintained for each room. The Administrator has several forms used for manipulating the health level and room terms.

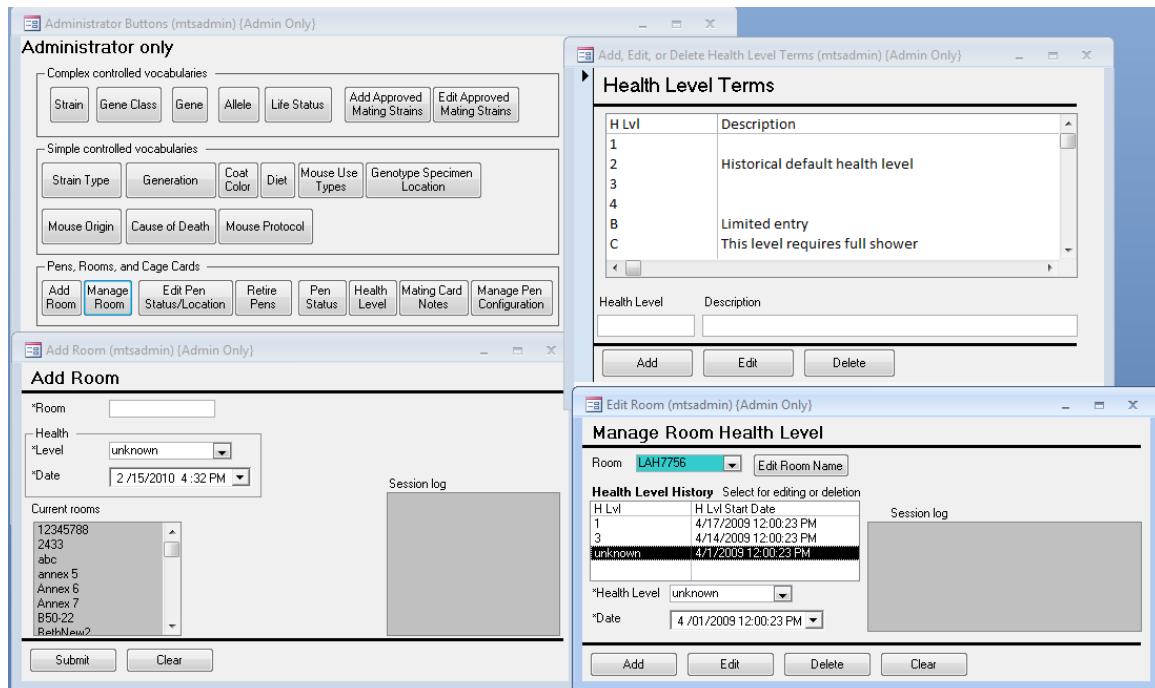


Figure 7-2 Administrator Forms for Managing Room and Health Level

Health level terms may not be deleted unless the term has no association (past or present) with a room. An edited health level term will be changed in all places where it was used. An edited room name will be changed in all places where it was used. If the date/time of a health level change is incorrect it may be edited using the *Manage Room* form. If a health level has been incorrectly associated with a room it may be deleted using the *Manage Room* form, however the room must have one health level associated with it.

## 7.3 Adding Pens

New pens are added when new mice are created in the database using the **Add Mouse**, **Bulk Add Mice**, **Add Mice at Wean**, or **Add Litter w/ Pups** forms. A new pen can be created when changes are made to a mouse using the **Edit Mouse** form. The **Add Matings** form will automatically generate new pens. The **Design Matings** form will assign pen IDs to matings and the pen is created in the default room with a status of proposed. When the mating is activated using the **Activate Matings** form the pen's status will also be changed to active as of the mating date. The handheld forms **Add Litter w/Mice** and **Make a Trio or Pair Mating** will create pens using the default room name and default pen status term.

### 7.3.1 Pen Info Form

The pen information form can be accessed from the manage mice tab and from many other forms that contain a *Pen Info* button.

The screenshot shows the 'Pen Info' form. In the 'Pen' section, the ID is 613, Name is FVB-01, Status is active, Date is 1/22/2010 10:33 AM, and Comments are Project 512. The 'Room' is B50-22. In the 'Pen pals' section, there are two entries: A303 and A304, both female (F), virgin (V), and alive (A). They were born on 12/1/2009 and are F01 generation, strain FVB/NJ. In the 'Pen history' section, there are two entries: one on 1/22/2010 at 10:33:09 AM in room B50-22 with status active, and another on 1/11/2010 at 10:18:29 AM in room annex 5 with status proposed. On the right, there is a 'Cage card printing' section with options to print detail card, print wean card, or preview, and an intended use dropdown set to Exp. 7354.

Select a pen by pen ID or pen name. The mice currently in the pen are listed (“pen pals”) and the history of the status and room changes for the pen are displayed.

Cage cards may be printed or previewed using the detail format or wean format if the pen contains mice all from the same litter. An optional *intended use* may be printed on the card. This use is not saved in JCMS or associated with the pen. It simply provides a way of printing a short note on the card.

Figure 7-3 Pen Info Form

### 7.3.2 Add Pen Form

The screenshot shows the 'Add Pen' form. In the 'Pen' section, the ID is 610, Name is FVB-01, Status is proposed, Date is 1/11/2010 10:18 AM, and Comments are Experiment 7354. The 'Room' is annex 5. On the right, there is a 'Cage card printing' section with options to print detail card, print wean card, or preview, and an intended use dropdown set to Exp. 7354. At the bottom are buttons for Submit, Close, and Clear.

Figure 7-4 Add Pen Form

The add pen form can be accessed from the manage mice tab and from many other forms that contain an *Add Pen* button. When *Use next available ID* is checked JCMS will assign the pen ID number. When it is unchecked, a number may be hand entered. An existing pen name may be selected for incrementing (*Increment name* must be checked). The pen that is added will have a name that is one larger than what is shown in the name field (for example, FVB-01 will be added

as FBV-02). Note that if increment name is not checked or the name has no numeric portion to increment, a duplicate pen name can be added with a unique pen ID. After submitting the new pen, a blank cage card for it may be printed.

## 7.4 Editing and Retiring Pens

The edit pen form can be accessed from the manage mice tab and from many other forms that contain an **Edit Pen** button. Select a pen ID or pen name and the current information for that pen will appear.

**Edit Pen**

**Pen**

- \*ID: 613
- Name: FVB-01
- \*Status: retired
- \*Date: 2/16/2010 1:17 PM
- New Name: FVB-01
- Comments: Project 512
- \*Room: B50-22
- H Lvl: 2 since 7/8/2009 4:38:23 PM

**Pen pals**

Mouse ID	Sex	Breed	Status	Date born	Generation	Strain
A303	F	V	A	12/1/2009	F01	FVB/NJ
A304	F	V	A	12/1/2009	F01	FVB/NJ

**Pen history**

Date	Room	Pen Status
1/22/2010 10:33:09 AM	B50-22	active
1/11/2010 10:18:29 AM	annex 5	proposed

**Cage card printing**

- Print detail card
- Intended use:
- Print wean card
- Preview

**Buttons:** Submit, Clear, Delete

Figure 7-5 Edit Pen Form

This form may be used to move a pen into a different room or to change the pen status. Enter the date/time of the change before submitting it. A successful change will be displayed in the pen history list. To change the pen name, enter the change in the *New Name* field. To remove the name, enter blanks in the *New Name* field. The pen ID may not be changed. The pen comments may be edited.

When retiring a number of pens, check the *Retire* box and the pen status field will be displayed as “retired” instead of the current status, saving having to change the status box before submitting.

A pen containing live mice (mice with a non-exit life status) may not be retired.

## 7.5 Moving Mice between Pens

**Move Mouse (OWN1) (Owners Only)**

**Transfer a mouse from one pen into another**

**Current Pen ID** 613      **Name** FVB-01

**New Pen ID** 616      **Name** FVB-02

**Health Lvl** 2      **Room** B50-22      **Status** active      **Established** 1/11/2010

**Cage Cards**

- Print Cage Card
- Intended use:
- Preview

**Buttons:** Move Mouse, Create New Pen

Figure 7-6 Move Mouse Form

There are two forms for moving mice. The **Move Mouse** button on the manage mice tab is used for moving one mouse at a time. Select the mouse and then the pen to put it into.

A detail cage card may be printed for the selected pen.

A live mouse (non-exit life status) cannot be moved into a retired pen.

When mice are moved around they may need to be moved into pens that do not exist yet

in JCMS. In this case, use the **Create New Pen** button to add the pen.

The session box will indicate the mouse ID moved and it's old and new pen ID number.

The **Move Mice** button on the manage mice tab will open the handheld move mice form. See below for a description.

## 7.6 Moving Mice Using Handheld

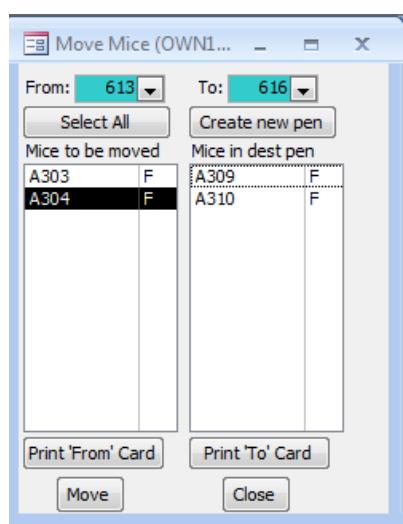


Figure 7-7 Handheld Move Mice Form

When opening from the handheld main menu, select the two pen ID numbers and choose *Move Mice*. When opening from the manage mice tab the form will show the two highest pen ID numbers. Select one or more mice to move. A new pen may also be created to move the mice into.

The Move mice form allows optional printing of new cage cards for both the *from* and *to* pens.

## 7.7 Printing Blank Cage Cards

Most forms that allow the creation of a pen also provide a method of printing cage cards. Blank cage cards may be printed using the **Print Cage Cards** button on the reports tab. No pen records are put into the JCMS database. These blank pre-numbered cards are created to prevent the accidental use of duplicate pen IDs in the mouse room. Blank cards that are not used (because they are lost, damaged, etc.) have numbers that may never be used in the database. If the setup variable `MTS_RELAXED_PEN_NUMS` is set to false, this form may be used to print pen IDs on cage cards for later use.

## 7.8 Retiring Pens Automatically

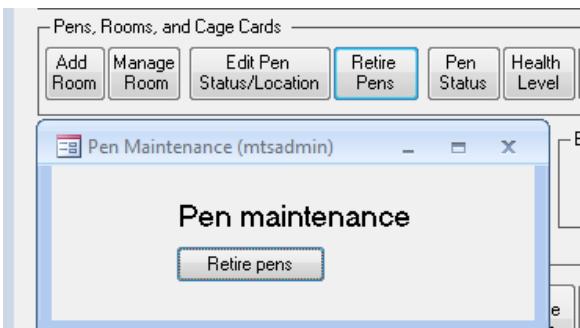
JCMS may be configured to attempt to retire pens automatically when the setup variable `JCMS_AUTO_RETIRE_PENS` is set to true. Every time a mouse's life status is changed from a non-exit life status to an exit life status JCMS will check if the pen it is in should be retired. Every time a mouse is moved into a new pen, JCMS will check if the old pen should be retired. The retired date will be the date of the action (exit date of the mouse or today's date if the mouse is moved). A pen will be retired if it is empty (contains no mice) or if it contains only mice that have an exit life status.

Note the following exceptions:

- A pen will not be retired twice in a row.
- A pen will not be retired if the retire date would be prior to some action that is already recorded in the pen history. (For example: the mouse's exit date might be last week and this pen's status was changed this week.)
- Changes to a pen's status are not just date sensitive, they are also time sensitive. Wean dates are not time sensitive and are considered to occur at midnight on a particular day. Midnight (12:00:00 AM) is the earliest time something can occur.
- The retire date used will be the date that pertains to the mouse that is changed. This means that if there are exited mice in the pen whose exit dates are more recent than the exit date entered for the last mouse, the pen retire date could end up being prior to those exit dates. Suggestion – watch the order that mice are exited so the most recent one is done last.
- The bulk forms for changing life status will attempt to also retire all pens. However, if an exception occurs in the middle of retiring pens, JCMS will stop retiring pens and not try to

continue. Correct the exception and use the Administrator's bulk retire function to complete retiring the pens.

## 7.9 Bulk Retire Pens



Bulk retiring of pens is an Administrator function, found on the Administrator button bar. Use the **Pen Maintenance** button to open this form and click the **Retire Pens** button. This will search the database for pens that are empty (have no mice in them or have only mice in them that have an exit life status). These pens will have their pen status set to retired on today's date.

NOTE: the pen maintenance function can take a long time to complete. Trial runs show that on a Pentium III 600MHz computer running

JCMS with 2000 pens, the pen maintenance functions take about 3 minutes each to run. The compute time will probably grow as  $n(\log n)$  (but could be as bad as  $n^2$ , depending on how MS Access optimizes its queries), where  $n$  is the number of cages in the database. If you have more than 10,000 pens, you should consider running the pen maintenance functions when you have several hours available in which the database will not be used. You should also back up your database (JCMS\_db.mdb or MySQL) before running these functions just in case they take too long to complete. Whether or not this precaution is necessary will not be known until we have more experience with JCMS performance with very large data sets.

## 7.10 Correcting Pen Status and Date Errors

Figure 7-9 Edit Pen Status/Location Form

The Administrator uses the Edit pen status/location form for correcting errors to pen history records. Select a pen history row and edit it using the fields below the pen history list. A pen must always have one pen history record, others can be deleted.

## 7.11 Cage Use Reports

The two cage use reports are designed to provide information on the number of cages (pens) that should be paid for during a particular time period. Only the cages that have a "billable" pen status on a date are counted. Pen status terms are defined as either billable or non-billable. Use the Administrator *pen status* form to change the billable attribute of a pen status term.

If the pen status terms are set up to track a cage "use", then by filtering the report results for a specific billable pen status, the number of cages for that "use" are listed. For example, a pen status term "breeding" could be used for all pens containing a mating pair. By setting the filter option for status to "breeding" only those pens are counted. Additional filter options may be selected. All the filter options are combined (ANDed) together and further limit the results. For example, the report could count cages in a specific room, with a specific owner and strain.

A pen's status may change in the middle of a day. *Pen billing* is used to determine if a pen is counted on a half day (partial day) during which its status changed between billable and non-billable. For example, the date a pen was established is a partial first day. The date a pen was retired is a partial last day. Full days count only the "non-partial" days. Any day counts all partial days.

### 7.11.1 Cage Use Report

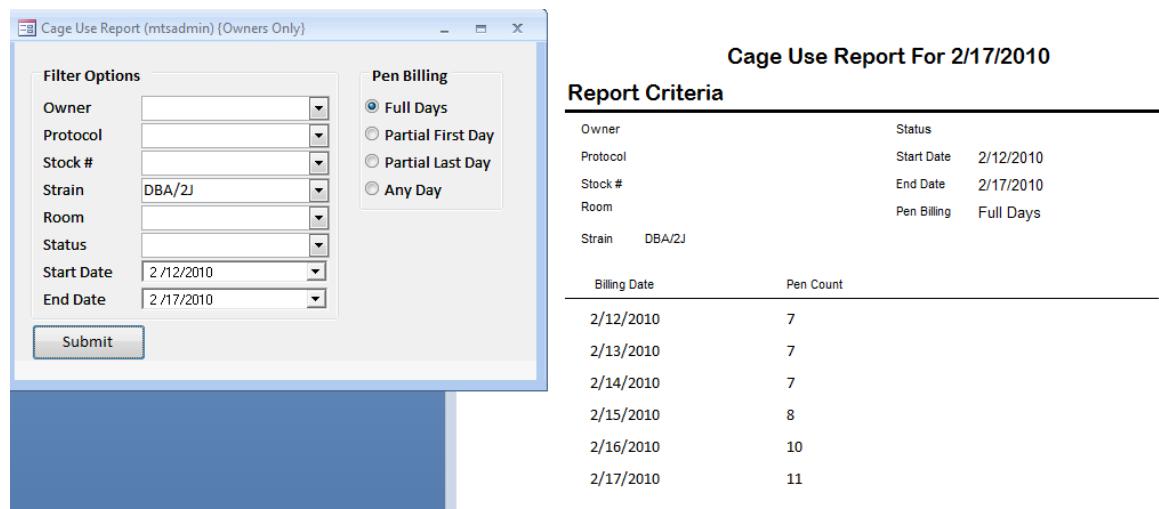


Figure 7-10 Cage Use Report

This report may be printed or exported to Excel. Several filter options may be chosen at once.

#### **Important Notes:**

If JCMS has a large number of cages and a long date range is selected, the report may run noticeably slow. It may be best to run this report overnight in this case.

If a status that is non-billable is selected as a filter option, the pen count will always be zero.

All strains have an associated stock # (JR #). It is not possible to select a stock number for one strain and a different strain as simultaneous criteria and receive a correct pen count. Use either stock # or strain as the criteria.

## 7.11.2 Cage Use Summary

This report is formatted as a datasheet (spreadsheet-like columns) and may be printed or exported to Excel. The resulting columns are limited to only those in the selected "Group" that have pens that match the filter options.

If a column contains all zeros, then there are an unknown number of pens that match the criteria, but all those pens have a non-billable status. For example, in Figure 7-11, owner "nobody" has C57BL/6J mice in pens in 4 rooms, but the pens in Big Jims and LAH7756 happen to all be retired and, therefore, the count is zero for those rooms. "Nobody" has never had any C57BL/6J mice in the room "B-52" so there is no column for B-52 in the report.

**Figure 7-11 Cage Use Summary Report**

The screenshot shows two windows side-by-side. The top window is titled 'Cage Use Report (mtsadmin) (Owners Only)' and displays a 'Report Criteria' section with fields for Owner, Protocol, Stock #, Room, Status, Start Date, and End Date. Below this is a table titled 'BillingReportQuery62265' with columns for Bill Date, New mutant 1, and New mutant 2. The bottom window is titled 'Cage Use Report Summary (mtsadmin) {Owners Only}' and also displays a 'Report Criteria' section with the same fields. It includes a 'Group By' section with radio buttons for Protocol, Strain, Room, Status, and Stock #. Below this is a table titled 'BillingReportQuery43530' with columns for Bill Date, annex 5, Big Jims, LAH7756, and unknown. Both windows have a 'Submit' button at the bottom.

Bill Date	annex 5	Big Jims	LAH7756	unknown
2/17/2010	1	0	0	1
2/18/2010	1	0	0	1
2/19/2010	1	0	0	1

Billing Date	New mutant 1	New mutant 2
2/12/2010	0	0
2/13/2010	0	0
2/14/2010	0	0
2/15/2010	1	0
2/16/2010	1	0
2/17/2010	2	1

**Figure 7-12 Cage Use Summary (bottom) vs. Cage Use Report (top)**

The important notes listed above for the cage use report also apply to the summary report with the following exception.

If a stock number that applies to several strains is chosen as the filter criteria and strain is chosen for grouping, the results can be seen by strain. For example, in Figure 7-12 the strains New Mutant and New Mutant 2 both have stock # zero.

There are some counting difficulties in the case where a cage contains mice with different characteristics (protocol or strain/stock#). A cage is counted only once, not multiple times. Which protocol or strain/stock # it is counted for is selected by random. The following combinations of criteria and grouping exhibit this problem.

Filter by protocol --- group by strain  
Filter by strain --- group by protocol  
Filter by stock # --- group by protocol  
No filter --- group by protocol  
No filter --- group by strain  
No filter --- group by stock #

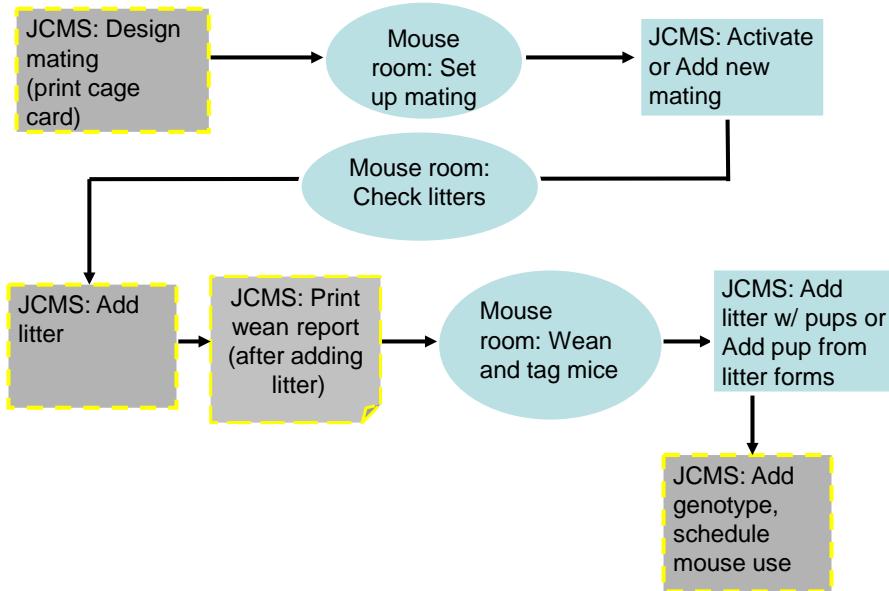
To see accurate results, filter by both criteria (example: filter by protocol and strain).

NOTE: It is possible to assign the same stock # to more than one strain (for example, if none is assigned, the stock number will be zero or blank). If a stock number associated with several strains is selected, the cage count will indicate the total number of cages for all those strains. If several strains have no stock number or a stock # of zero, they will all be counted and reported in one column with a header of 0.

## 8 Matings

JCMS is flexible in that you can choose to either set up matings on the computer (design mating) or do the mating work in the mouse room or laboratory and later enter the information into JCMS (add mating). If the design first method is used, then the mating must be *activated* on the computer after the mating is implemented in the mouse room.

### Matings – Litters - Weaning



**Figure 8-1 Mating to Weaning Diagram**

This diagram shows a *typical* flow of information from JCMS to the mouse room and back to JCMS. Dashed boxes are optional. JCMS also has a special add litter form for use for those who prefer to wean mice before entering the litter information into JCMS.

#### 8.1.1 Automatic Litter Number Generation

JCMS can generate litter numbers automatically. To use auto litter numbering, set the setup variable MTS\_AUTO\_LITTER\_NUMS to "on." When this variable is set to on, JCMS generates litter numbers and associates the litter numbers with a mating.

Litter numbers are generated by JCMS in batches. The size of the litter number batches is a function of the setup variable called MTS\_NUM\_AUTO\_LITTER\_NUMS (the number of automatic litter numbers). It is recommended that the value of this variable be set to 10 or 100. By using a round number (like 10 or 100) it is easy to identify a litter as being the first, second, third etc., by simply looking at the litter number.

#### 8.2 Technical Matings

In addition to a Natural (N) mating between one or two dams and a sire, JCMS supports Embryo Transfer, Ovary Transfer, and In vitro Fertilization matings.

- Embryo Transfer (ET) involves an embryo sample and host dam
- Ovary Transfer (OT) involves an ovary or ovarian tissue sample, host dam, and sire
- In vitro Fertilization (IVF) involves an egg sample and sperm sample

All matings except IVF may have a natural (live birth) litter. All mating types may have an embryo litter. An embryo litter consists of harvested embryos or embryonic stem cell samples. To obtain a live birth from an IVF mating, create embryo sample(s) that are then used in an embryo transfer mating.

### 8.3 Which Mating Forms to Use?

Design Mating is used by those who want to set up their matings on paper by using the computer ahead of time. Activate Matings must then be used after the mating has been performed in the mouse room. These two forms are only used for natural matings. Others prefer to first do the matings in the mouse room or laboratory, then return to the computer and use the Add Mating form to enter the information. The Add Mating form may also be used in the mouse room on a laptop computer. If handheld computers are used in the mouse room, there is a special Do Pair or Do Trio mating form to use.

The Edit Mating form is provided for correcting any errors that were made in previous data entry.

Eventually, the mating will be ended or “retired”. Some plan the ending of natural matings ahead of time using the Design Retire Matings form. All matings use the Retire Matings form when ended. When the Design Retire matings is used first, information given in the design stage is used for the defaults on the Retire mating form. This information may still be changed to reflect last minute changes made in the mouse room.

Sires are often used in more than one mating simultaneously. JCMS will warn that a mouse is already in use in another unretired mating.

Cage card note choices are set up by the Administrator by using the Administrator button bar.

### 8.4 Approved Mating Strains

Dam strain	Sire strain	Litter strain
BALB/cJ	DBA/2J	New mutant
C57BL/6J	C3H/HeJ	New mutant

Figure 8-2 Add Approved Strains Form

The Administrator has the optional ability to create a table of approved litter strains. An approved litter strain field is a function of the dam strain and the sire strain. A sire strain and dam strain combination is **not** unique. That is, a sire and dam strain combination may result in multiple litter strains. The special case of when the sire and dam strains are the same will result in that strain always being ‘approved’ and need not be entered in the database. Approved strain records can be active or inactive. Inactive records are ignored as candidates for litter strains. They are kept in the database for documentation purposes and possible future use.

The “approved matings” radio button on the mating forms will be initialized to the value of JCMS\_ENFORCE\_APPROVED\_MATINGS in the setup variables. When this is set to true, only approved litter strains may be selected. The user may change the radio button setting to active or all and enter the unapproved strain.

The forms for adding and editing approved litter strains are invoked from the Administrator tab by clicking the **Administrator** button. Use the two buttons: **Add Approved Strains** and **Edit Approved Strains**.

## 8.5 Designing a New Mating

**Design Matings**

Mating Type: Natural  
 Select only live mice

Strain/Stock # criteria			
FVB/NJ	FVB/NJ	FVB/NJ	
<b>*Litter Strain</b> FVB/NJ <input type="radio"/> All <input type="radio"/> Active only <input checked="" type="radio"/> Approved only Cage card color	<b>*Dam 1</b> FVB-M-214 FVB/NJ F02 Birth date Sex Pen ID 9/2/2013 F 88 Pen Name Life Status A Breeding V Diet Genotype	<b>Dam 2</b> FVB-M-213 FVB/NJ F02 Birth date Sex Pen ID 9/2/2013 F 88 Pen Name Life Status A Breeding V Diet	<b>*Sire</b> FVB-M-205 FVB/NJ F02 Birth date Sex Pen ID 9/1/2013 M 87 Pen Name Life Status A Breeding V Diet

\*Owner nobody  
Wean Note:  
Comments:  
Proposed Diet 6%

Matings / pen IDs [litter IDs] submitted in this session

Print Cage Card  
 Preview  
Cage Card Note 1 Check daily  
Cage Card Note 2

Submit    Clear    Session Report

**Figure 8-3 Design Mating Form**

Matings can be set up “on paper” before they are implemented in the mouse room. The mating information may be printed on a cage card to be used by the mouse room workers. Once the mating is set up in the mouse room, JCMS must be updated using the *Activate mating* form described below. This form is limited to designing natural matings.

Use the **Design Matings** button to open the form. The choices of mice may be limited to only live mice (use the select only live mice check box) and to one strain (use the strain/stock # criteria choice above the dam or sire).

Use the blue-green drop down boxes to select the dam(s) and sire. Dam2 is optional. It is not possible to enter the mating date at this point. When “approved only” is selected, only approved litter strains will be provided as a choice.

There are a number of “proposed” data items that can be entered on the design mating form (such as proposed diet). These items are stored in the database but not associated with the mice until the mating is activated (using the Activate mating form). The “proposed” data items allow printing out the information to bring to the mouse room. When the mating is selected for activation, the “proposed” items become the defaults so they don’t have to be reentered unless a change has been made.

**Click on the Submit button** to add the mating into the database. If there are no error messages, JCMS will automatically generate a mating number and a pen in the default room with a pen ID, pen status of proposed, and today’s date as the pen established date. The mating number and pen ID will be listed in the session box.

After the submit is successful, the **Print Cage Card button** will be activated. Printing cage cards is an optional step. The card printed will show the mating that was just submitted and the proposed pen ID number.

The **Clear** button is used to clear all the fields at once.

## 8.6 Activating a Mating

The screenshot shows the 'Activate Matings' window with the following details:

- Mating Type:** Natural
- Mating ID:** 59
- Dam 1:** FVB-M-214 (FVB/NJ, F02, Birth Date: 9/2/2013, Sex: F, Pen ID: 88)
- Dam 2:** FVB-M-213 (FVB/NJ, F02, Birth Date: 9/2/2013, Sex: F, Pen ID: 88)
- Sire:** FVB-M-205 (FVB/NJ, F02, Birth Date: 9/1/2013, Sex: M, Pen ID: 87)
- Pen Settings:**
  - ID: 102 (checkboxes: Use next available ID, Increment Name)
  - Name: (text input)
  - Status: active (dropdown)
  - Date: 9/25/2013 12:00:01 AM (dropdown)
  - Comments: (text input)
- Changes to all mice:**
  - New Diet: 6%
  - Suggested First litter #: 590
  - Auto increment ID: checked
- Cage Card Notes:** Cage card notes are printed on cage cards but are not stored in the database with each mating. NOT ALL CAGE CARDS SUPPORT CAGE CARD NOTES.
- Buttons:** Print Cage Card, Preview, Submit, Clear, Session Report.

**Figure 8-4 Activate Mating Form**

Once the matings have been set up in the mouse room, use the **Activate Mating** button to enter the mating date and any changes. First select the mating ID number and the mating as it was set

up will appear. If a different dam or sire was used, they can be changed using the drop down boxes. When “approved only” is selected, only approved litter strains will be provided as a choice.

When a mating is activated, the diets for each mouse in the mating will be automatically updated according to the new diet combo box. The default **new diet** will be the **proposed diet** specified with the design mating form. The diets for each mouse can be changed by selecting a new diet from the combo box if the proposed diet is not the actual diet.

The suggested pen ID will be used unless a different pen is selected and the three mice will be moved into it. The pen status is changed to active on the mating date. All fields that were changed will be displayed in red after a successful submit. The mating ID and pen ID will also be added to the session box.

There is no way to delete a mating. If three completely different mice were used, use the Add Mating form to create a new mating instead.

When the setup variable JCMS\_AUTO\_RETIRE\_PENS is true, if the pens vacated by moving the dams and sire are now empty, the pens will be retired on the mating date.

## 8.7 Add a Natural Mating

**Add Matings**

Mating Type: N

Select only live mice

\*Litter Strain: FVB/NJ  
All Active only Approved only Cage card color  
 Autoincrement Generation  
\*Litter Generation: F02  
\*Mating Date: 9/23/2013  
Standard Wean Time:   
Needs Genotyping:   
\*Owner: nobody  
Wean Note:   
Comments:   
Changes to all mice:  
New Diet: 6%

Strain/Stock # criteria			
Dam 1: FVB-F-020 FVB/NJ F01 Birth Date: 6/1/2013 Sex: F Pen ID: 78 Pen Name: Cage-00014 Life Status: A Breeding: V Cur. Diet: 4%	Dam 2: FVB-F-019 FVB/NJ F01 Birth Date: 6/1/2013 Sex: F Pen ID: 78 Pen Name: Cage-00014 Life Status: A Breeding: V Cur. Diet: 4%	Sire: FVB-M-113 FVB/NJ F01 Birth Date: 6/1/2013 Sex: M Pen ID: 81 Pen Name: Cage-00017 Life Status: A Breeding: V Cur. Diet: 4%	

Print Cage Card  
 Preview  
Cage Card Note 1: Cage Card Note 2:   
Matings / pen IDs [litter IDS] submitted in this session

Pen  
ID:  Use next available ID  
Name: Cage-00022 Increment Name  
Status: active  
Date: 9/23/2013 4:01:25 PM  
Comments:   
Room: Room 26 H Lvl: 3 since 9/6/2013 3:14:47 PM

Submit Clear Session Report Add Sample

**Figure 8-5 Add Mating Form**

The **Add Matings** form allows designing and activating a mating on one form. This form is useful for those who implement the matings in the mouse room or laboratory and then want to come

back to the computer to enter the data or for those who use laptops or other computers in the mouse room.

The choices of mice may be limited to only live mice (use the select only live mice check box) and to one strain (use the strain/stock # criteria choice above the dam or sire). Use the drop down boxes to select the dam(s) and sire. Dam 2 is optional. When “Approved only” is selected, only approved litter strains will be provided as a choice.

On this form, there is a check-box on the lower right hand side that, when checked, tells JCMS to put the mice in this mating in the next available pen. If the check box is un-checked, then a pen ID number must be manually entered in the pen ID box or an existing pen must be selected.

When the setup variable JCMS\_AUTO\_RETIRE\_PENS is true, if the pens vacated by moving the dams and sire are now empty, the pens will be retired on the mating date.

Hint for tab key users: In order to tab out of the dam1, dam2, sire, host dam, and sample drop down boxes, hold down both the Control key and Tab key simultaneously.

## 8.8 Add a Technical Mating

Use the same Add Mating form that is provided for adding a natural mating. In the upper left, select the mating type. The form’s columns will change to allow selection of the appropriate mating components. The choice of samples are limited to sample class “Live” and sample types embryo, egg, ovary, ovarian tissue, and sperm. The select live mice check box and strain/stock # criteria will also limit the choices for sample, host dam, and sire.

**Add Matings - Ovary Transfer**

Mating Type: DT

Select only live mice

Strain/Stock # criteria	
FVB/NJ	C57BL/6J
FVB/NJ	FVB/NJ

**Maternal Sample:** FVB-Ovary-203

**Host Dam:** A77

**Sire:** FVB-M-112

**Changes to all mice:** New Diet: 6%

**Pen:**

- ID: Cage-00022
- Status: active
- Date: 9/23/2013 4:01:25 PM
- Comments:

**Print Cage Card**   **Preview**

Cage Card Note 1:

Cage Card Note 2:

Matings / pen IDs [litter IDs] submitted in this session:

Submit   Clear   Session Report   Add Sample

Figure 8-6 Add Ovary Transfer Mating

**Add Matings - Embryo Transfer**

Mating Type		ET	Strain/Stock # criteria	
<input checked="" type="checkbox"/> Select only live mice		FVB/NJ	C57BL/6J	FVB/NJ
<input checked="" type="radio"/> All <input checked="" type="radio"/> Active only <input type="radio"/> Approved only Cage card color <input type="checkbox"/> Autoincrement Generation Litter Generation F02 Mating Date 9/23/2013 Standard Wean Time Needs Genotyping Owner nobody Wean Note		<b>*Litter</b> FVB/NJ <b>Strain</b> <input checked="" type="radio"/> All <input checked="" type="radio"/> Active only <input type="radio"/> Approved only Cage card color <input type="checkbox"/> Autoincrement Generation Litter Generation F02 Mating Date 9/23/2013 Standard Wean Time Needs Genotyping Owner nobody Wean Note	<b>*Embryo Sample</b> FVB-Embryos-010 FVB/NJ Sample Class Live Sample Type Embryo Preservation Type Frozen Harvest Date 9/24/2013 Age 1 Status Unprocessed Location Storage Facility\Building1\Floor1\Freezer1	<b>*Host Dam</b> A79 C57BL/6J F03 Birth Date Sex Pen ID 7/1/2013 F 90 Pen Name Cage-00021 Life Status A Breeding V Cur. Diet 4%

**Figure 8-7 Add Embryo Transfer Mating**

**Add Matings - In vitro Fertilization**

Mating Type		IVF	Strain/Stock # criteria	
<input checked="" type="checkbox"/> Select only live mice		FVB/NJ	C57BL/6J	FVB/NJ
<input checked="" type="radio"/> All <input checked="" type="radio"/> Active only <input type="radio"/> Approved only Cage card color <input type="checkbox"/> Autoincrement Generation Litter Generation F02 Mating Date 9/23/2013 Standard Wean Time Needs Genotyping Owner nobody Wean Note Comments		<b>*Litter</b> FVB/NJ <b>Strain</b> <input checked="" type="radio"/> All <input checked="" type="radio"/> Active only <input type="radio"/> Approved only Cage card color <input type="checkbox"/> Autoincrement Generation Litter Generation F02 Mating Date 9/23/2013 Standard Wean Time Needs Genotyping Owner nobody Wean Note Comments	<b>*Maternal Sample</b> FVB-EGGS-005 FVB/NJ Sample Class Live Sample Type Egg Preservation Type Frozen Harvest Date 8/5/2013 Age 0 Status Processed Location Storage Facility\Building1\Floor2\Room 10\Freezer	<b>*Paternal Sample</b> FVB-SPERM-003 FVB/NJ Sample Class Live Sample Type Sperm Preservation Type Frozen Harvest Date 8/5/2013 Age 0 Status Processed Location Storage Facility\Building1\Floor3\Holding tank
Changes to all mice: New Diet 6%		<div style="border: 1px solid black; padding: 5px;"> Print Cage Card  <input type="checkbox"/> Preview  Cage Card Note 1  Cage Card Note 2 </div> <div style="border: 1px solid black; padding: 5px; margin-top: 10px;"> Pen  ID: 104 <input checked="" type="checkbox"/> Use next available ID  Name: Cage-00024 <input checked="" type="checkbox"/> Increment Name  Status: active  Date: 9/23/2013 4:01:25 PM  Comments:  Room: Room 26 H Lvl: 3 since 9/6/2013 3:14:47 PM </div>		
<input type="button" value="Submit"/> <input type="button" value="Clear"/> <input type="button" value="Session Report"/> <input type="button" value="Add Sample"/>				

**Figure 8-8 Add In vitro Fertilization Mating**

Once a sample becomes part of a mating, its status is changed to "Processed". Once an IVF mating is created, the Add Sample button on the bottom of the form is enabled. It will open the Add sample form allowing creation of an embryo or other type of sample. Note that IVF matings are not associated with a cage.

Creating pups from an IVF mating is a multi-step process. First create the mating, next create embryo samples, and finally, use the embryo sample in an embryo transfer mating.

## 8.9 Edit a Mating

Edit an existing mating using the **Edit matings** form. With this form it is possible to select new mice (dam 1, dam 2, host dam, and sire) or samples and change the basic mating information. A duplicate or new cage card can be printed from this form.

**NOTE:** If a new dam 1, dam 2, host dam, or sire for a mating is selected, it may be necessary to do other cleanup work. For example, the mouse that was in the mating and the new dam, host dam, and/or sire, may need to have their breeding status and diets changed (use the edit mouse form). It may also be necessary to use the move mice form to move the replaced mice to appropriate pens. If a pregnant dam is moved out of a mating, the pedigree information for her litter may be lost.

The edit matings form does not automatically make changes to any of the mice in the mating (unlike the activate mating form which changes breeding status, diet, and pen automatically); however, a warning will display, telling exactly which mice to make changes to.

## 8.10 Design Matings to be Retired

The screenshot shows the 'Design Retire Matings' form window. The left side contains general mating details: Mating Type (Natural), Mating ID (44), Litter (New Mutant 4), Strain (New Mutant 4), Generation (N03), Mating Date (8/8/2013), Date to Retire (9/24/2013), Standard Wean Time (checked), Needs Genotyping (unchecked), Owner (nobody), and a Comments field (testing). An 'Auto increment ID' checkbox is checked. The right side is divided into three sections: Dam1, Dam2, and Sire, each showing detailed information for a mouse (e.g., New-010, New-011, New-012; N02, N02, N02; and New Mutant 4). Below these sections is a summary table for the three mice. At the bottom, there is a note about matings slated for retirement and a list of buttons: Submit, Clear, Session Report, Move Mice, and Pen Info.

**Figure 8-9 Design Retire Matings Form**

Natural matings can be retired in two steps. First, on the computer decide which matings to retire (design step). Next go to the mouse room and do the work, then come back to the computer and actually retire the matings on the computer. Or, the *design* step can be skipped.

The design retire matings form allows setting up a work list of matings to retire without actually making changes to the primary mating information in the database. The database stores **proposed** information in the mating table. When this form is used, it only updates the **proposed** fields in the database. Thus, no changes will be seen when a query for mating or mouse information is made. The idea is to set up a work report (by printing out copies of the Design retire matings form as you work). Then, after the work has been completed in the mouse room, use the Retire matings form to actually update the primary information in the database. When selecting a mating to retire, the proposed information (entered previously in the Design Retire matings form) will appear as the defaults in the Retire matings form. Any changes can be made at this point (to reflect what really happened in the mouse rooms) before submitting the changes to the primary information in the database.

## 8.11 Retire a Mating

**Figure 8-10 Retire Matings Form**

Once the matings have been retired in the mouse room, the date retired must be entered into JCMS. Technical matings also should be retired using this same form. Use the **Retire Matings** button. Select the mating type and mating ID from the blue/green box. Only matings that have not already been retired are available for selection. All information about this mating will display as soon as the cursor is moved out of the drop down box by tabbing or clicking elsewhere. Enter the date retired. The dam and sire life statuses can be changed to an exit status by using the drop down menus. Similarly, the breeding status can be changed to R for retired breeder and a new diet can be selected. No changes occur until the **Submit** button is clicked.

The retire mating form also has a *Set Mice to Retire and Euthanized* button which sets the mouse life status to the value of the setup variable JCMS\_DEFAULT\_EXIT\_TERM and the mouse's breeding status to retired (R). Using this button also sets the date exited to the date retired. Note if the default term is not euthanized, the button will display the default term on it.

## 8.12 Working with Matings using a Handheld

Mating ID: 6      Pen ID: 177

Owner: OVS      Mating date: 11/6/2005

Dam1: 1  
Strain: ABC/xyzJ  
Gen.: N01

Dam2:  
Gen.:

Sire: 4  
Strain: ABC/xyzJ  
Gen.: N01

Litter: ABC/xyzJ  
Strain: ABC/xyzJ  
Gen.: N03

Card color: none  
 Print cage cards on 'Submit'  
 Autoincrement dams/sires

Submit

**Figure 8-11 Handheld Form: Pair Mating**

Natural matings can be created using handheld computers by scanning or entering pen IDs. Two types of matings are available, pair or trio. A cage card can optionally be printed showing the pen ID, Mating ID, and suggested litter numbers. The automatically assigned pen ID and Mating ID will also be shown at the top of the form. Only active strains are shown as choices for the litter strain.

## 8.13 Automatic Retiring of Matings

It is possible to have JCMS attempt to retire matings as a background function if the setup variable JCMS\_AUTO\_RETIRE\_MATINGS is set to "True". When a mouse's life status is changed to an exit life status, JCMS will check the following:

- Is the mouse in any matings?
- If so, check each active mating and retire those where all dams, host dams, and sires have an exit life status.
- The retire date used will correspond to the exit date for this mouse.

Note: This check is done only for mice whose current life status is a non-exit status. It will not be done if the life status is changed from one exit status to another or if the exit date is changed.

## **9 Litters**

JCMS associates litters with matings. Each litter must have a unique litter ID number associated with it. JCMS can generate litter ID numbers or you can use your own litter number scheme.

All litter forms except those for handhelds show a list of all litters currently associated with the selected mating.

### **9.1 Automatic Litter Number Generation**

JCMS may be set up by the Administrator to automatically generate litter numbers. Once this function is set on, all JCMS users will have automatic litter numbers generated. Note, however, JCMS does not enforce which litter numbers are used (even if it generates a set of numbers); it is up to you when you enter a new litter in JCMS to decide which number to use. If you prefer to use your own litter number series, then turn off the JCMS litter number generator.

#### **9.1.1 Turning Automatic Litter Numbering On or Off**

The Administrator controls automatic litter numbering from the Administrator button bar, using the button for *JCMS Setup Variables*. The variable called MTS\_AUTO\_LITTER\_NUMS can take the values "on" or "off".

#### **9.1.2 Setting the Number of Litter Numbers that are Generated for Each Mating**

JCMS increments its internal litter number counter by a value specified in the *JCMS Setup Variables* table. Set how many litter numbers should be allocated to litters by changing the value of this variable. NOTE: to prevent any confusion, it is best to set this number when first starting to use JCMS and then not change it. We recommend that this number be set to 10 unless you really have long breeding pairs that you track. The number of litter numbers that are generated is set by the JCMS Setup Variable called MTS\_NUM\_LITTER\_NUMS (from the Administrator button bar, click the JCMS Setup Variables button).

Changing the number of litter numbers can create minor confusion (but no date problems) because the add and edit litter forms show the range of litter numbers associated with each mating. The range of numbers is calculated from the first suggested number plus the number of numbers generated (as specified in the MTS\_NUM\_LITTER\_NUMS variable). The only point of confusion will be on the litter forms that may show an incorrect range for matings that were allocated a different number of litter numbers (if this is confusing, don't worry; just don't change the MTS\_NUM\_LITTER\_NUMS variable very often).

## **9.2 Which Litter Forms to Use?**

### **Multiple step process:**

Using this approach, first identify the litters from the mating and record them in JCMS (Add Litter button). When the pups are sexed and/or tagged, add the number of male and female pups to the litter record (Edit Litter button). Next use the Wean report (Wean Work Report button) to get a listing of litters that should be weaned during a particular time period. Finally, add the weaned pups and their wean and tagging dates to JCMS (Add Pup from Litter button).

### **One step process:**

Add the litter to the database when the mice are weaned by using the Add Litter w/ Pups button. This form allows adding all of the litter information AND adding the pups as individual mice in one step.

### Handheld forms:

Use the Add Litter w/Mice choice from the handheld forms to enter a new litter at weaning. This form will create a litter record and then create the newly weaned mice in pens. It will not enter the wean date in the litter record. Use the Handheld Wean or Wean and Exit form to enter the wean date into the litter record.

## 9.3 Adding Litters

Litter strain:	Litters	Litter #	Births #	Harvested #	Date
FVB/NJ	Min/Max allocated numbers	450	8	5	9/1/2013
	451	8	5	9/2/2013	
	452	0	5	9/23/2013	
	453	0	5	9/24/2013	

**Figure 9-1 Add Litter Form**

To add a new litter, specify the Mating ID number. The mating information will show on the screen. Type the litter number into the litter number box or use the *Next Litter Num* button. Litters that were successfully added will be listed in the session box along with their mating ID number. Embryo litters are listed in the # harvested and date columns. Live birth litters may be added to natural, embryo transfer, and ovary transfer matings. Use the Add Embryo litter form to create litters that are composed of harvested samples.

## 9.4 Wean Report

Report date: Tuesday, February 16, 2010  
Wean start date: 3/22/2010

owner	date born	projected wean date	litter ID	# born	# F	# M	wean date	tag date	mating ID	gen.	needs typing	pen ID	pen name
Room: B50-22	H Lvl: 2 since 7/8/2009 4:38:23 PM						Strain: BALB/cJ						
OWN1	1/5/2010	1/23/2010	771	8	8	0		77	N02		<input type="checkbox"/>	524	BALB mating 005
Room: B50-22	H Lvl: 2 since 7/8/2009 4:38:23 PM						Strain: FVB/NJ						
OWN1	3/4/2010	3/22/2010	1170	6				117	F02		<input checked="" type="checkbox"/>	801	check early
Room: Big Jims	H Lvl: B since 9/29/2009 7:22:00 AM						Strain: C3H/HeJ						
OWN1	2/8/2010	2/26/2010	930	2	1	1		93	F02		<input type="checkbox"/>	748	

**Figure 9-2 Wean Work Report**

The **Wean Work Report** button on the reports tab will open the request wean report form. It is used to request a listing of litters that have not yet been weaned by returning all litters with a status of "A" and no recorded wean date. A start date is entered to eliminate from the list all litters that will be too young on that date for weaning. Litters will be selected that are 18 or more days old (for the standard wean time) or 28 or more days old (for long weans) on the start date. The resulting list may also be limited by selecting only certain mouse owners. Note the defaults for the standard and extended wean times are setup variables (JCMS\_STANDARD\_WEAN\_TIME and JCMS\_EXT\_WEAN\_TIME) that may be changed.

The report is available in three formats: sorted by strain, sorted by room and strain, and ready to export to Excel.

## 9.5 Editing Litters

Litters	Litter #	Births	Harvested
Min/Max allocated numbers	# Date	# Date	
500/509	500	6 8/17/2013	

**Figure 9-3 Edit Litter Form**

Only litters that have the litter type "live birth" are shown on this form. However, the mating could be natural, embryo transfer, or ovary transfer. Click on the **Edit Litter** button to open the form. Select the litter number from the drop down list. When the cursor leaves the litter number box (either via tabbing out of the box or by using the mouse to go to another box,) the litter data will appear on the form.

No changes are made to the database until the **Submit button** is clicked.

The **Clear button** clears all edit fields in order to start over. However, if the clear button is pressed by accident then recover simply by clicking on the litter number box and then moving the cursor back out of the box (to redo the query).

When pups are weaned, this form is used to enter the number of males and females, weaning date, and tagging date. Only an owner or owner's secretary has permission to use this edit form.

Note that it is possible to use the Add Litter w/Pups form to create mice in proposed pens prior to the pups being weaned (possibly on the tagging date). These pens can be activated (pen status changed to active on the wean date) by entering the wean date on this form and checking the *activate pens* box.

If the Add Litter w/Pups form was used to give the pups mouse IDs and the pups were left in the mating pen, after entering the wean date for the litter, use the *Move Pups* button to open the Handheld Move Mice form and transfer the pups into the weaning pens.

If a sample ID is underlined, it may be clicked to open the Edit Sample form. The purpose of this feature is to allow viewing the sample information. If a change to the sample strain is made, it will not be shown on the litter form. Select the litter again on the litter form in order to refresh this information.

## 9.6 Wean Litters

Use the **Add Pup from Litter** button to open this form. Each newly weaned mouse is entered individually. The wean and tag dates for the litter may also be entered on this form. See Adding Pups section 6.5 for a description of this form.

## 9.7 Adding Litters with Pups at Weaning

**Mating Information**

- Litter #: 531
- Next Litter Num:
- Mating ID: 53
- Mating Pen ID: 94
- Pen Name: Cage-00022
- Mating Type: Ovary Transfer
- Generation:
- Strain Name:
- Maternal Sample: FVB-Ovaru-200
- FVB/NJ

**Host Dam:** A80      F03      C57BL/6J

**Sire:** FVB-M-115      F01      FVB/NJ

**Litter:** FVB/NJ strain: **Litters**      **Litter #**      **Births**      **Harvested**

Min/Max allocated numbers	Litter #	Date	#	Date
530 / 539	530	8/20/2013	6	

**Comments:**

Auto increment litter ID

**Put pups into database**

- Auto generate mouse records
- Females first
- Use base mouse number->
- \*Mouse origin: internal
- Protocol ID: P-0902
- Leave pups in mating pen
- Num females/pen [1..10]: 10
- Num males/pen [1..10]: 10
- Add to experimental plan(s):
  - None
  - Plan only
  - Plan + all tests

**Pen**

- ID:
- Name: Cage-00024
- Use next available ID
- Increment Name
- Status: active
- Date: 10/12/2013 9:07:18 AM
- Comments:

**Room**: Room 26      H Lvl: 3 since 9/6/2013 3:14:47 PM

**Cage cards for last group submitted**

Intended use: Blood work

Print Cage Cards

Preview

Submit    Clear    Session Report

Figure 9-4 Add Litter w/Pups Form

This form is used to both add the litter into JCMS and add all pups as mice with individual mouse IDs in one action. Natural, embryo transfer, and ovary transfer matings may have live birth litters. Use the **Add Litter w/Pups** button. First select the mating and then enter the litter number by typing it or using the *Next Litter Num* button. Enter the rest of the litter information. The current mice drop down box is there to help determine the last mouse ID number used. When the base mouse number is blank, selecting a current mouse number will cause the next number in the sequence to be placed in the base mouse number box.

As mice are added, they may also be added to experimental plan(s) and all the experimental tests in those plans. The session box will list the plan and test IDs.

Several workflows are supported by this form.

- 1) Create the weaned mice and litter all at once. Pups will be placed into new pens.
  - a. Enter the wean date and optional tag date
  - b. Select auto generate mouse records
  - c. Enter a starting pen ID or use next available ID
- 2) Create the litter with tagged pups remaining in the mating pen
  - a. Enter the tag date but no wean date for the litter
  - b. Select auto generate mouse records
  - c. Select leave pups in mating pen
  - d. When it is time to wean the pups, use the Edit litter form to enter the wean date for the litter. Then click on the *move pups* button, which opens the *Move mice* form. Transfer the mice into their new pens.
- 3) Create the litter with tagged pups placed into "proposed" pens. This allows printing the cage cards for the new pens even though they are not yet in use. The cards might later be used in the mouse room to indicate where to put the pups when they are weaned.
  - a. Enter the tag date but no wean date for the litter
  - b. Select auto generate mouse records
  - c. Enter a starting pen ID or use next available ID
  - d. Use the pen status "proposed" and the tag date
  - e. Note that JCMS will now show the pups located in the proposed pens, even though they physically are still in the mating pen.
  - f. When the pups are weaned enter the wean date for the litter and change the pen(s) status to active using either
    - i. Edit litter form: make sure activate pens is checked when the wean date is submitted
    - ii. Handheld forms: scan the pen IDs on all the proposed cards for the litter, select the wean option, make sure activate pens is checked, submit the wean date.

This form normally creates pens starting with the next available pen ID. However, a starting pen ID may be specified so long as the pen sequence created does not already exist and there are enough sequential pen IDs available to hold all the pups in the litter. Note that empty or retired pens may not be used. The starting pen ID must be typed into the pen ID box. For example, if there is a gap between 68 and 74, then 69, 70, 71, 72, and 73 are available.

Pen ID
DBA_016
DBA_015
DBA_012
DBA_011
DBA_010
DBA_009
DBA_008
DBA_007

Figure 9-5 Example of a pen gap

## 9.8 Weaning Mice using a Handheld

The figure displays three handheld computer screens side-by-side, each showing a different form for managing mouse litters:

- Add Litter & Mice (...):** This form is used to add a new litter. It includes fields for 'Mice born (female/male)' (Total 5, F 3, M 2), 'Litter ID' (1190), 'Date of birth' (2/12/2010), 'Wean date' (7/12/2006), 'Life status' (A), 'PenID' (808), 'Use base ID' (unchecked), 'Mating ID' (119), 'Strain' (C3H/HeJ), 'Generation' (N02), and a checkbox for 'Print cage cards on 'Submit''. Buttons for 'Submit' and 'Close' are at the bottom.
- Wean (OWN1) {Secretaries and ...}:** This form is used to wean mice. It shows 'Pens' (813, 812) and a 'Wean date' (2/19/2010). A checkbox 'Activate pens' is checked. Below is a table for 'Litter ID: 1190' with 'Total Mice' (5) and a list of mice (A455-A459) each marked with an 'A'. Buttons for 'Submit' and 'Done' are at the bottom.
- Bulk Change Life Status (OWN1...):** This form is used to bulk change life status. It shows 'Pens' (813, 812) and an 'Exit date' (2/19/2010). Below is a table for 'Litter ID: 1190' with 'Total Mice' (5) and a list of mice (A455-A459) each marked with an 'A'. A dropdown 'Life status' is set to 'E' and a dropdown 'C.O.D' is set to 'Euthanasia'. Buttons for 'Submit' and 'Done' are at the bottom.

Figure 9-6 Handheld Forms: Weaning and Add litter and mice

Handheld Forms: Weaning and Add litter and mice

These three forms are provided for using handheld computers in the mouse room for weaning mice. Scan the pen ID of a mating pen and select **Add litter w/mice** from the handheld main menu. This will open the Add litter and mice form. Use the **Next Avail** button to select the litter ID. Enter the litter information and a Base ID number (mouse ID number to start with) for the new mouse records. Printing cage cards will provide the proper labeling for the new pens. Submit will create a new litter record for this mating, new mouse records for each member of the litter, and new pen records with a pen status of *proposed*. This form does not enter the wean date into the litter record. Setup variable defaults are used for all values that cannot be entered (room name, mouse ID prefix if no base ID is specified, maximum number of mice in a pen).

To enter the wean date into the litter record, scan one of the new pens and select **Wean mice** from the handheld main menu. Enter the wean date and click submit. This will enter the wean date for the whole litter. If proposed pens should be activated (pen status changed to active on the wean date) make sure to scan all the pen IDs for the litter. If one is missed, repeat using the form, the pens will be activated even though a message will indicate the wean date is already entered.

A second method of entering the wean date is to enter it when the mice in the pen are given their exit date. Scan the pen IDs and select **Bulk Exit** from the handheld main menu. Enter the wean date, exit date, life status, and cause of death (C.O.D.). If the mice in the pen have already had a wean date entered, it will not be changed, so this form can also be used just to exit mice. If the setup variable JCMS\_AUTO\_RETIRE\_PENS is set to true, if all the mice in a pen are given an exit life status, the pen will be retired on the exit date.

The setup variable MTS\_THRESHOLD\_MICE\_BATCH\_OPERATION should be set to a maximum number of mice to be weaned or exited at once. If the number of mice listed is greater than this limit, a warning message is displayed, which offers the choice to continue or cancel.

# 10 Embryo Litters

An embryo litter is created by harvesting embryos or embryonic stem cells from a dam in a mating. The harvested material can optionally be recorded and tracked as a Sample. For specific information about how JCMS tracks samples, see Section 20.

## 10.1 Add Embryo Litter w/Samples

Add Embryo Litter w/Samples (mtsadmin) {Secretaries and Owners}

### Add Embryo Litter w/Samples

Mating Information

Mating ID:	59	Mating Pen ID: 88	Pen Name:
Mating type:	Natural	Generation	Strain Name
Dam 1:	FVB-M-214	F02	FVB/NJ

Plug date: 9/22/2013

Dam 2:	FVB-M-213	F02	FVB/NJ
--------	-----------	-----	--------

Plug date: 9/22/2013

Sire:	FVB-M-205	F02	FVB/NJ
-------	-----------	-----	--------

Litter strain: FVB/NJ

Litters	Litter #	Births	Harvested
Min/Max allocated numbers	#	Date	# Date
590/ 599	590	0	7 9/24/2013

DPC

Calc from Dam 1 plug 2

Calc from Dam 2 plug

Enter manually  Add Dam 1 plug  Add Dam 2 plug

Status: H

Comments

Auto increment litter ID

**Put samples into database**

Auto generate sample records

**\* Sample ID**

Individual Samples FVB-Embryos-011

Add to Embryo Pool

Create Embryo Pool

**\* Sample Class** Live

**\* Sample Type** Embryo

Harvest Method Protocol 77

Preservation Type Frozen

Preservation Method -30 C

Preservation Detail

**\* Sample Status** Unprocessed

Current samples with litter source:  Litter strain only  All FVB-Embryos-010

**\* Location**

Storage Facility

- Building1
  - Floor1
    - Freezer1
  - Floor2
  - Floor3

Litters / Mating ID (Sample ID) List

Plug date 9/22/2013 added to Dam2  
Plug date 9/22/2013 added to Dam1  
590/59 (FVB-Embryos-011::FVB-Embryos-012::FVB-Embryos-013::FVB-Embryos-014::FVB-Embryos-015::FVB-Embryos-016)

Submit Clear Session Report Print Sample Labels Set Genotype

Figure 10-1 Add Embryo Litter with Sample records

Start by selecting the mating. Any type of mating (natural or technical matings) may create an embryo litter. Any previous litters and existing plug dates will be displayed. Use the next litter button to obtain a litter # from the allocated litter number series.

The days post conception (DPC) is required and, if samples are created, this number is saved as the age in days from conception in the sample record. A table showing the relationship between

the DPC and Theiler Stage is provided to aid in determination of the proper number. This table is derived from e-Mouse Atlas (EMA):

[http://www.emouseatlas.org/emap/ema/theiler\\_stages/StageDefinition/stagedefinition.html](http://www.emouseatlas.org/emap/ema/theiler_stages/StageDefinition/stagedefinition.html). The EMA Anatomy Atlas of Mouse Development uses embryological mouse models to provide a digital atlas of mouse development. It is based on the definitive books of mouse embryonic development by Theiler (1989) and Kaufman (1992).

The DPC may be entered using one of these methods:

- Manually by selecting from the table
- Calculated from either Dam1 or Dam2 plug date
  - This requires that a plug date has previously been entered (see Section 11 on Plug Dates). JCMS will calculate the DPC using the plug date selected in the mating information section of the form and the harvest date entered on the form. It will update the DPC displayed on the form when the harvest date or plug date selections are changed.

Checking “Add Dam 1 plug” and/or “Add Dam 2 plug” will update the plug date records with a new plug date for the specified dam(s) when the litter is added. The plug date is added as obsolete. It is determined from the harvest date minus DPC.

If sample records are created, there are three options:

- Individual samples: Create a sample for each embryo harvested. The number of samples created is equal to the # harvested shown on the form.
- Create embryo pool: Create only one sample record.
- Add to embryo pool: Select an existing sample with a litter source. The litter record will be added to that sample. Samples are able to have multiple sources. In this case, the pool of embryos comes from several litters.

The sample ID must be unique. If several samples are created, the ID must contain a number that can be incremented in the same fashion as mouse IDs are incremented. Use the “Current samples with a litter source” combo box to select an ID to use for starting the incrementing. If the sample ID box is blank, the next ID in the sequence will be placed in it, unless the choice is to “Add to embryo pool”. In that case, select the ID of the embryo pool.

See Section 20 on Samples for information on the other sample fields. Note that no weight or description is saved in the sample record(s). The owner is set to the mating owner.

## 10.2 Edit Embryo Litter

This form is provided for correcting errors in the basic litter information. Only litter types 'E' (harvested embryos) or "ES" (embryonic stem cells) may be edited using this form.

The screenshot shows the 'Edit Embryo Litter' form. Key fields include:

- Litter #: 452
- Litter type: E
- Number harvested: 5
- Harvest date: 9/23/2013
- DPC: 1
- Status: H
- Comments: (empty)
- Auto increment litter ID: checked

Mating Information:

Mating ID: 45	Mating Pen ID: 83	Pen Name: Cage-00019
Mating Type: Natural	Generation	Strain Name
Dam 1: FVB-F-001	F01	FVB/NJ

Nearby plug date: 9/22/2013

Dam 2: FVB-F-002	F01	FVB/NJ
------------------	-----	--------

Nearby plug date: 9/23/2013

Sire: FVB-M-100	F01	FVB/NJ
-----------------	-----	--------

Litter strain: FVB/NJ

Litters	Litter #	Births #	Harvested #	Date
Min/Max allocated numbers	450	8	5	9/1/2013
	451	8	5	9/2/2013
	452	0		9/23/2013
	453	0		9/24/2013
Litter generation: F02	450 / 459			
Owner: nobody				

Samples:

Sample ID	Owner	Sample Status	Storage Facility	Sample Date	Harvest Method
FVB-Embryos-001	nobody	Unprocessed	Building1	9/23/2013	Protocol 77
FVB-Embryos-002	nobody	Unprocessed	Building1	9/23/2013	Protocol 77
FVB-Embryos-003	nobody	Unprocessed	Building1	9/23/2013	Protocol 77
FVB-Embryos-004	nobody	Unprocessed	Building1	9/23/2013	Protocol 77
FVB-Embryos-005	nobody	Unprocessed	Building1	9/23/2013	Protocol 77

Session Log (Litter/Mating): (Empty)

Buttons: Submit, Clear, Session Report, Edit Sample

**Figure 10-2 Edit Embryo Litter form**

When there are two dams, JCMS is unable to tell which dam was the mother of this litter. If this information is important, we suggest you place it into the comments field or create separate matings for each dam.

The "nearby plug dates" displayed on the form are a best guess selected as a plug date within the JCMS\_GESTATION\_PERIOD (number of days) just prior to the harvest date.

A change made to the harvest date or DPC value will NOT update any associated sample or plug date records.

To make a change to one of the samples, click on the sample ID and then press the "Edit Sample" button, which will open the Edit Samples form.

# 11 Plug Dates and Pregnancy Checking

Vaginal plugs form following copulation. The plug date table is used to record the date a plug is observed and associate it with a dam and mating. Later, the plug date may be marked obsolete to indicate the dam has given birth or that conception did not take place. Over time, a dam may accumulate a series of plug dates. Only natural matings may have plug dates entered.

Plug dates are used to generate several reports containing the following types of information:

- Stage of pregnancy (lists dams by strain and date of plug).
- Work report: dams to check for plugs (lists dams by strain that currently have no plug)
- Plug date history for mice or matings (lists all the dam's plug dates)

Pregnancy checks:

- Plug Date work report: dams to check for conception/pregnancy (lists dams by strain and date of plug).
- Pregnancy Check work report: for those not using plug dates, this report lists all matings with no litter in the pen and ignores any plug dates.

## 11.1 Add Plug Date

The screenshot shows the 'Add Plug Date' window. Key fields include:  
- \*Mating ID: 28 (selected from a dropdown)  
- \*Plug Date: 9/7/2012  
- \*Apply to: Dam 1 (checkbox checked), Dam 2 (checkbox checked)  
- \*Obsolete: checkbox (unchecked)  
- Comments: A large text area.  
- Session Box: A scrollable list of previous entries.  
Detailed Dam Information:  
Dam 1:  
ID-0039, FVB/NJ, F03, Room unknown, Birth date 4/25/2012, Sex F, Pen ID 70254, Life Status A, Pen Name Mating001.  
Previous plug dates:  
Date: 9/5/2012 (Obsolete N), 6/4/2012 (Obsolete Y).  
Projected Date - Use:  
9/25/2012: Basic blood work; 9/15/2012: Neurological exam; 9/9/2012: Basic blood work; 6/24/2012: Basic blood work; 6/14/2012: Neurological exam; 6/8/2012: Basic blood work.  
Dam 2:  
ID-0038, FVB/NJ, F03, Room unknown, Birth date 4/25/2012, Sex F, Pen ID 70254, Life Status A, Pen Name Mating001.  
Previous plug dates:  
Date: 6/4/2012 (Obsolete Y).  
Projected Date - Use:  
6/24/2012: Basic blood work; 6/14/2012: Neurological exam; 6/8/2012: Basic blood work.  
Sire:  
ID-0036, C3H/HaJ, F01, Room unknown, Birth date 4/6/2012, Sex M, Pen ID 70254, Life Status A, Pen Name Mating001.  
At the bottom, the 'Add Mouse Uses' section includes:  
- A grid showing Mouse Use (Basic blood work, Neurological exam), DPC (4.5, 10.5, 20.5), and Projected Date (9/11/2012, 9/17/2012, 9/27/2012).  
- Buttons: >, Clear Grid, and a red X button.  
- Buttons at the bottom: Submit, Clear, Session Report.

Figure 11-1 Add Plug Date Form

The plug date is associated with a particular dam in a specific mating. Only matings that are active (not proposed or retired) are listed as choices. To add the same plug date to both dam 1 and dam 2 at the same time, check both *apply to* boxes. If a dam has previous plug dates that are associated with a different mating, they will not be displayed on this form.

Mouse uses that are associated with a date post conception (DPC) may be entered at the same time the plug date is entered. Fill the grid at the bottom of the form with the uses and DPC. The projected date will be calculated based on the plug date. The projected date is updated whenever the plug date is changed on the form. Previously entered uses are shown for each mouse to help prevent accidental duplication. The maximum choice for the DPC value is set by default to 21. To set it to a different value, change the setup variable JCMS\_GESTATION\_PERIOD.

## 11.2 Edit Plug Date

Only plug dates that are associated with an active mating (not proposed or retired) may be edited or deleted. Select the plug date to edit from the list below the dam by double clicking on it. Once the specific plug date record is displayed on the left side of the form, the comments, obsolete check box, and plug date may be changed. Mark plug dates obsolete to keep the dam from being listed on the plug date and pregnancy stage report. A dam's plug dates must all be obsolete for her to be listed on the plug check report.

The screenshot shows the 'Edit Plug Date' window with the following details:

- Mating ID:** 28 (selected)
- Owner:** nobody
- Plug Date:** 6/4/2012
- Applies to:** Dam 1 (checkbox checked), Dam 2 (checkbox unchecked)
- Obsolete:** checked
- Comments:** A large text area for notes.
- Session Box:** A large gray area for session logs.
- Double click Plug Date to Edit:** A table showing existing plug dates:
 

Date	Obsolete
9/5/2012	N
6/4/2012	Y
- Buttons at the bottom:** Submit, Delete, Clear, Session Report, Expire Plug Dates (highlighted with a yellow circle).

Figure 11-2 Edit Plug Date Form

To delete a particular plug date, double click it on the dam's plug date list. The delete button will become an available choice. If any editable field is changed on the form (plug date, obsolete, or comments), the delete button will no longer be a choice. This is to prevent accidentally deleting instead of editing a plug date.

### 11.2.1 Automatically mark plug dates as obsolete

The **Expire Plug Dates** button found on the Edit Plug Date form is used to mark **all** plug dates prior to a certain date as obsolete. The date is calculated by subtracting the value of the setup variable JCMS\_GESTATION\_PERIOD from today's date. The default value is 21 days. For

example, if today is November 12, 2012 minus 21 yields the date October 22, 2012. This action must be confirmed before any changes are made.

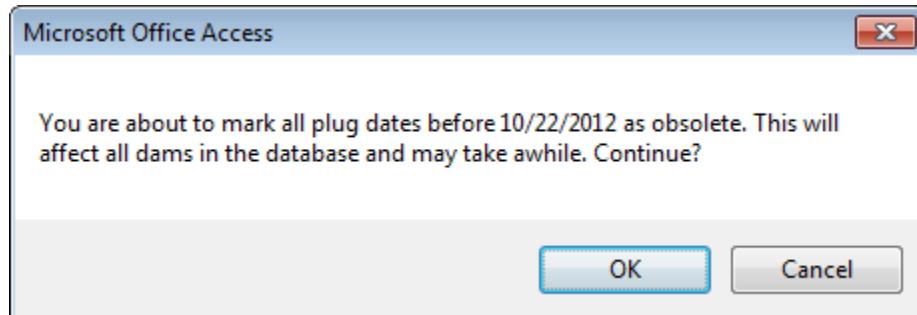


Figure 11-3 Confirm plug date expiration

### 11.3 Plug Date and Pregnancy Check Work Reports

A Microsoft Access form titled "Request Plug Date and Pregnancy Work Reports". It has three main sections: "Choose report" (radio buttons for "Plug date / Pregnancy stage" (selected), "Check for plugs", and "Pregnancy or conception check / not using plugs"), "Choose owner(s)" (list box showing "nobody" and "OWN1"), and "Format" (radio buttons for "Print" (selected) and "Tabular Spreadsheet"). A "Preview" button is located at the bottom left.

Figure 11-4 Form: Request plug date or pregnancy reports

Three different reports are selected using this form. Any report may be filtered by owner. The tabular spreadsheet format may be exported to Microsoft Excel. If Microsoft Office 2007 gives the message "Excel found unreadable content..." answer "yes" to recover the contents. No data will be lost.

### 11.3.1 Plug Date / Pregnancy Stage Report

Plug Date / Pregnancy Stage							23-Oct-09	
Mouse ID	DPC	Generation	Life Status	Owner	Mating ID	Mating date	Room	Pen ID
<b>Strain: B6D2F1/J</b>								
Plug Date:	10/23/2009							
F0001	0.5	F02	A	nobody	8	10/23/2009	113	
F0002	0.5	F02	A	nobody	9	10/23/2009	114	
F0003	0.5	F02	A	nobody	10	10/23/2009	115	
F0004	0.5	F02	A	nobody	11	10/23/2009	116	
Plug Date:	10/9/2009							
F99-03	14.5	F01	A	nobody	5	10/9/2009	107	
F99-04	14.5	F01	A	nobody	4	10/6/2009	106	
<b>Strain: BALB/cByJ</b>								
Plug Date:	10/6/2009							
Litter106	17.5	F02	A	nobody	6	10/9/2009	109	
Litter106	17.5	F02	A	nobody	7	10/9/2009	109	
Litter106	17.5	F02	A	nobody	7	10/9/2009	109	

Figure 11-5 Sample plug date / pregnancy stage report

This report is sorted by strain, plug date, room, and pen ID. It lists for active matings, all plug dates that are not marked obsolete. In the sample report shown above, dams A2, F3, and F1 are listed several times because they have multiple plug dates that are not marked obsolete. For this report to be accurate, all old plug dates must be marked obsolete.

Use this report to determine the pregnancy stage by calculating the number of days between today and the plug date. Use this report to list dams that should be checked for conception, pregnancy, or litters based on their plug date.

Plug Date / Pregnancy Stage										
	Strain	Plug Date	Mouse ID	Mating #	Mating Date	Room #	Pen #	Generation	Status	Owner
BALB/cByJ	12/5/2008	A2	7	10/8/2008		15	F01	A	nobody	
BALB/cByJ	12/1/2008	A3	7	10/8/2008		15	F01	A	nobody	
BALB/cByJ	10/14/2008	A2	7	10/8/2008		15	F01	A	nobody	
BALB/cJ	12/6/2008	F3	4	8/25/2008	B102 R55	10	F02	A	nobody	
BALB/cJ	12/3/2008	F4	6	10/10/2008		12	F02	A	nobody	
BALB/cJ	12/3/2008	F3	4	8/25/2008	B102 R55	10	F02	A	nobody	
BALB/cJ	12/2/2008	F1	5	8/27/2008	B99 R273	11	F02	A	nobody	
BALB/cJ	11/30/2008	F1	5	8/27/2008	B99 R273	11	F02	A	nobody	
BALB/cJ	10/6/2008	F3	4	8/25/2008	B102 R55	10	F02	A	nobody	
FVB/NJ	7/14/2009	Old2	9	3/1/2008		19	N/D	A	nobody	

Figure 11-6 Sample tabular spreadsheet format plug date / pregnancy stage report

### 11.3.2 Plug Check Work Report

This report lists all active matings where one (or more) dams have no plug (i.e. all the dam's plug dates are marked obsolete or it has no plug dates). The report is sorted by strain, room, and pen ID.

#### Plug Check - Dams with no Plug

Pen Name	Pen ID	Dam Generation	Mating #	Mating Date	Owner	Dam ID
Dam Strain	BALB/cJ					
<b>Room abc</b>						
BALB_0000	259	F02	27	10/20/2009	OWN1	Test001903
BALB_0000	259	F02	27	10/20/2009	OWN1	Test001904

Figure 11-7 Plug Check Report

### **11.3.3 Pregnancy Check Work Report**

This report ignores plug dates. It lists active matings that have no litter in the pen. A mating has no litter if all litters associated with the mating have a wean date or it has no litter records associated with it. A litter that has a status other than "A" for alive will be ignored since a litter with a status such as born dead might not have a wean date.

#### **Pregnancy Check - Matings with no Litter**

Pen Name	Pen ID	Mating	Mating Date	Litter Generation	Owner	Dam 1
<hr/>						
Litter Strain BALB/cJ						
<hr/>						
Room abc						
	476	58	12/2/2009	N03	OWN1	A113
<hr/>						
Room Annex 6						
	256	24	10/17/2009	N02	OWN1	Test001907
<hr/>						
Room B50-22						

**Figure 11-8 Sample pregnancy check work report**

### **11.3.4 Plug Date History**

Use the Mouse Query or Mating Query to obtain a list of all plug dates recorded for selected dams or matings.

## 12 Breeding Performance Reports

There are six variations of the Breeding Performance Report. Access them from the Reports tab on the main button bar.

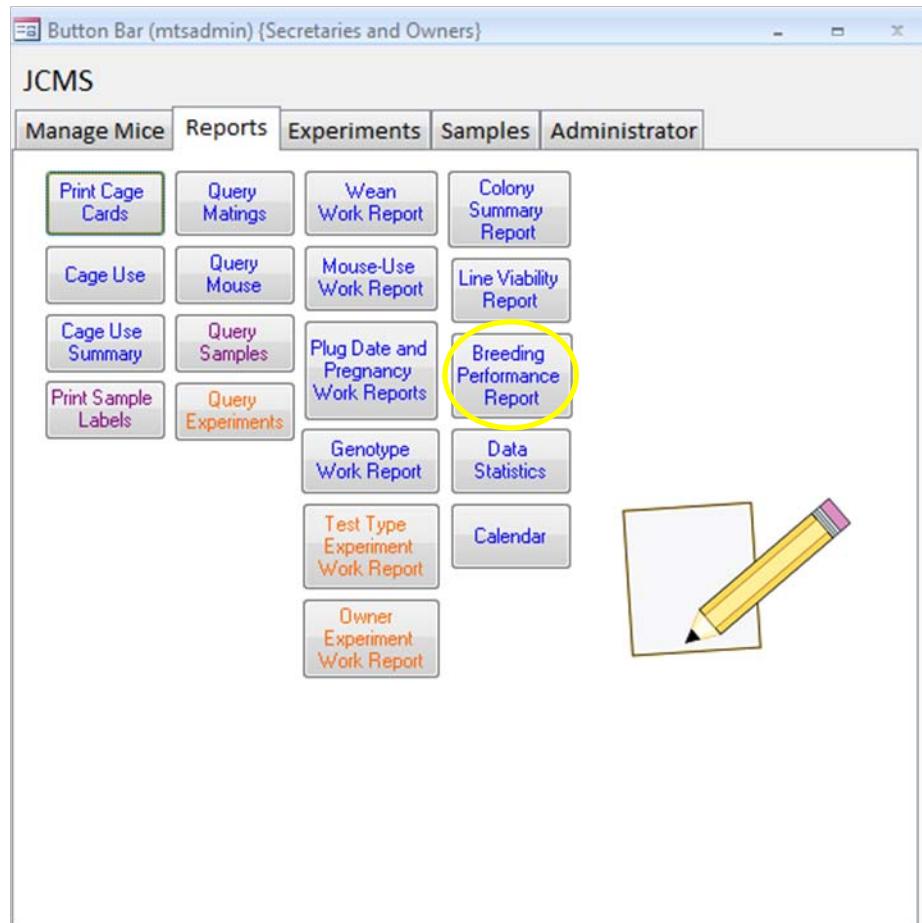


Figure 12-1 Select the Breeding Performance Report

A screenshot of the 'Breeding Performance Report' configuration form. The form has four main sections: 1) 'Choose report' with radio buttons for 'Full Report' (selected), 'Summary', and 'Tabular Spreadsheet'. 2) 'Choose strain name/stock #' with radio buttons for 'All' (selected) and 'Selected from List'. 3) 'Choose owner(s)' with a dropdown list containing 'nobody' and 'OWN1'. 4) 'Report time frame' with dropdown menus for 'Past 6 months', 'Mating Date', 'Start Date' (set to 12/13/2011), 'Past year', 'Past year and a half', 'End Date' (set to 6/13/2012), and 'Report time frame' (which is also labeled 'Report time frame'). At the bottom left of the form is a 'Preview' button.

Figure 12-2 Form: Request a breeding performance report

Only natural matings and their litters are shown on these reports.

The criteria for the report include the following:

- Time frame
  - The data included will be for all matings with a mating date within the start and end date range. Litters associated with the selected matings, whose birth date is

within the date range are included. Use the buttons to set a 6, 12, or 18 month time span or enter the exact dates for the range.

- Strain
  - Include all strains or one selected from the list
- Owner
  - Select one or more from the list

## 12.1 Full Breeding Performance Report

The complete report includes details on all matings and associated litters, including the mating ID; mating and retired dates; dam1, dam2, and sire; current pen ID and pen name for dam 1; litter ID; birth, wean, and tag dates; # born, # females, and # males. There are summary rows for the total number of matings, litters, # born, # females, # males, and matings with no litters for each mating, strain, and owner. The litter status (A for alive, H for harvested, B for born dead, D for dead at weaning, K for killed, and M for missing) is listed for each litter. Note that harvested litters will normally have a # born of zero.

Breeding Performance Report						Tuesday, September 24, 2013 10:08:45 AM
<hr/>						
Owner: nobody						
Stock # Strain						
1800 FVB/NJ						
Mating ID: 45	Mating Date: 8/5/2013	Retired:	Dam 1 Pen ID: 83	Cage: 00019		
Dam 1 ID: FVB-F-001	Dam 2 ID: FVB-F-002		Sire ID: FVB-M-100			
Litter ID	Birth Date	Wean Date	Tag Date	# Born	# Female	# Male
453				0		H
452				0		H
450	9/1/2013	9/19/2013	9/19/2013	8	4	A
451	9/2/2013	9/19/2013	9/19/2013	8	8	A
<hr/>			Totals for Mating 45	# Litters	# Born	# Female
				4	16	12
					4	
Mating ID: 46	Mating Date: 8/5/2013	Retired:	Dam 1 Pen ID: 84	Cage: 00020		
Dam 1 ID: FVB-F-003	Dam 2 ID:		Sire ID: FVB-M-101			
Litter ID	Birth Date	Wean Date	Tag Date	# Born	# Female	# Male
<hr/>			Totals for Mating 46	# Litters	# Born	# Female
				0		
					0	
Mating ID: 47	Mating Date: 8/5/2013	Retired:	Dam 1 Pen ID: 85	Cage: 00021		
Dam 1 ID: FVB-F-004	Dam 2 ID: FVB-F-005		Sire ID: FVB-M-101			
Litter ID	Birth Date	Wean Date	Tag Date	# Born	# Female	# Male
<hr/>			Totals for Mating 47	# Litters	# Born	# Female
				0		
					0	
Mating ID: 56	Mating Date: 9/20/2013	Retired:	Dam 1 Pen ID: 99	Cage: 00022		
Dam 1 ID: FVB-M-210	Dam 2 ID:		Sire ID: FVB-M-101			
Litter ID	Birth Date	Wean Date	Tag Date	# Born	# Female	# Male
<hr/>			Totals for Mating 56	# Litters	# Born	# Female
				0		
					0	
<hr/>						
Totals for Stock # / Strain						
1800 FVB/NJ						
<hr/>						
# Matings	# Litters	# Born	# Female	# Male	# Matings w/no Litters	
4	4	16	12	4		3

Figure 12-3 Detail section of the report

=====						
Totals for Stock # / Strain						
656 CBA/J	# Matings	# Litters	# Born	# Female	# Male	# Matings w/no Litters
	5	11	63	31	22	0

Figure 12-4 Strain total section of the report

=====						
Totals for Owner	# Matings	# Litters	# Born	# Female	# Male	# Matings w/no Litters
OWN1	11	31	195	81	55	1
=====						
Summary: Grand Totals	# Matings	# Litters	# Born	# Female	# Male	# Matings w/no Litters
	16	35	230	97	70	4

Figure 12-5 Owner and grand total section of the report for all strains

## 12.2 Summary Breeding Performance Report

The summary report provides the same totals as the full report but eliminates the detail section that shows each litter.

Breeding Performance Report							Wednesday, June 13, 2012 4:39:02 PM			
<hr/>										
Owner: OWN1										
Stock # Strain										
656 CBA/J										
<hr/>										
Totals for Mating 9	# Litters	# Born	# Female	# Male						
	4	18	11	7						
<hr/>										
Totals for Mating 10	# Litters	# Born	# Female	# Male						
	1	1	1	0						
<hr/>										
Totals for Mating 11	# Litters	# Born	# Female	# Male						
	2	14	9	5						
<hr/>										
Totals for Mating 12	# Litters	# Born	# Female	# Male						
	2	8	2	0						
<hr/>										
Totals for Mating 13	# Litters	# Born	# Female	# Male						
	2	22	8	10						
<hr/>										
Totals for Stock # / Strain	# Matings	# Litters	# Born	# Female	# Male	# Matings w/no Litters				
656 CBA/J	5	11	63	31	22	0				

Figure 12-6 Summary shown for one strain

## 12.3 Breeding Performance Report Tabular Spreadsheet

The tabular spreadsheet provides the data in a format that can be easily exported to Excel. There is one row for each litter or, if the mating has no litters, one row for the mating.

BreedingPerformanceQuery60386														
Owner	Stock #	Strain	Mating #	Mouse ID	Mouse ID	Mating Date	Date Retire	Litter #	Number Born	Date Born	# F	# M	Weaning	
nobody	659	C3H/HeJ	1 A5		A9	5/7/2012								
nobody	659	C3H/HeJ	3 A3		A7	5/15/2012								
nobody	659	C3H/HeJ	14 C3H-F-060	C3H-F-059	C3H-M-050	3/26/2012		140		11	4/16/2012	4	5	5/4/2012
nobody	659	C3H/HeJ	14 C3H-F-060	C3H-F-059	C3H-M-050	3/26/2012		141		8	4/15/2012	4	3	5/4/2012
nobody	659	C3H/HeJ	15 C3H-F-058	C3H-F-057	C3H-M-049	4/2/2012		150		8	4/27/2012	4	3	5/15/2012
nobody	659	C3H/HeJ	15 C3H-F-058	C3H-F-057	C3H-M-049	4/2/2012		151		8	4/28/2012	4	4	5/15/2012
nobody	659	C3H/HeJ	16 C3H-F-056	C3H-F-055	C3H-M-048	5/7/2012								
OWN1	659	C3H/HeJ	2 A4		A8	5/7/2012								
OWN1	656	CBA/J	9 CBA-F-020	CBA-F-019	CBA-010	3/19/2012		90		8	4/16/2012	4	4	5/4/2012
OWN1	656	CBA/J	9 CBA-F-020	CBA-F-019	CBA-010	3/19/2012		93		6	5/30/2012	3	3	
OWN1	656	CBA/J	9 CBA-F-020	CBA-F-019	CBA-010	3/19/2012		91		2	4/17/2012	2	0	5/4/2012
OWN1	656	CBA/J	9 CBA-F-020	CBA-F-019	CBA-010	3/19/2012		92		2	5/29/2012	2	0	
OWN1	656	CBA/J	10 CBA-F-018	CBA-F-017	CBA-009	3/19/2012		100		1	4/9/2012	1	0	4/27/2012
OWN1	656	CBA/J	11 CBA-F-016	CBA-F-015	CBA-008	4/5/2012		110		10	4/26/2012	5	5	5/14/2012
OWN1	656	CBA/J	11 CBA-F-016	CBA-F-015	CBA-008	4/5/2012		111		4	4/27/2012	4	0	5/15/2012
OWN1	656	CBA/J	12 CBA-F-014	CBA-F-013	CBA-007	4/18/2012		120		4	4/27/2012	0	0	5/15/2012
OWN1	656	CBA/J	12 CBA-F-014	CBA-F-013	CBA-007	4/18/2012		121		4	5/9/2012	2	0	5/27/2012
OWN1	656	CBA/J	13 CBA-F-012	CBA-F-011	CBA-006	4/18/2012		131		11	5/9/2012	4	5	5/27/2012
OWN1	656	CBA/J	13 CBA-F-012	CBA-F-011	CBA-006	4/18/2012		130		11	5/9/2012	4	5	5/27/2012
OWN1	671	DBA/2J	4 DBA-F-040	DBA-F-039	DBA-M-030	3/2/2012		40		6	3/21/2012	4	2	4/8/2012
OWN1	671	DBA/2J	4 DBA-F-040	DBA-F-039	DBA-M-030	3/2/2012		42		9	4/27/2012	6	3	5/15/2012
OWN1	671	DBA/2J	4 DBA-F-040	DBA-F-039	DBA-M-030	3/2/2012		43		8	4/28/2012	4	3	5/16/2012

Figure 12-7 Tabular spreadsheet format

## 12.4 Breeding Performance Report Litter Detail Pivot Table

The litter detail pivot table provides a dynamic table showing the # born, # females, and # males for each mating and litter broken out by year/quarter/month.

Form1																									
owner	strainName	StartDate	EndDate	Years ▾ Quarters Months																					
OWNER1	CBA/J	All	All	Grand Total																					
<input type="checkbox"/> 2012																									
<input type="checkbox"/> Qtr2																									
<input type="checkbox"/> Apr																									
<input type="checkbox"/> May																									
<input type="checkbox"/> Total																									
<input type="checkbox"/> matingID																									
<input type="checkbox"/> 90																									
<input type="checkbox"/> 91																									
<input type="checkbox"/> 92																									
<input type="checkbox"/> 93																									
<input type="checkbox"/> Total																									
<input type="checkbox"/> 100																									
<input type="checkbox"/> Total																									
<input type="checkbox"/> 110																									
<input type="checkbox"/> 111																									
<input type="checkbox"/> Total																									
<input type="checkbox"/> 120																									
<input type="checkbox"/> 121																									
<input type="checkbox"/> Total																									
<input type="checkbox"/> 130																									
<input type="checkbox"/> 131																									
<input type="checkbox"/> Total																									
<input type="checkbox"/> Grand Total																									

Figure 12-8 Litter detail pivot table

## 12.5 Breeding Performance Report Litter Summary Pivot Chart

The litter summary pivot chart provides a dynamic column graph showing the # litters, # born, # females, and # males by year/quarter/month.

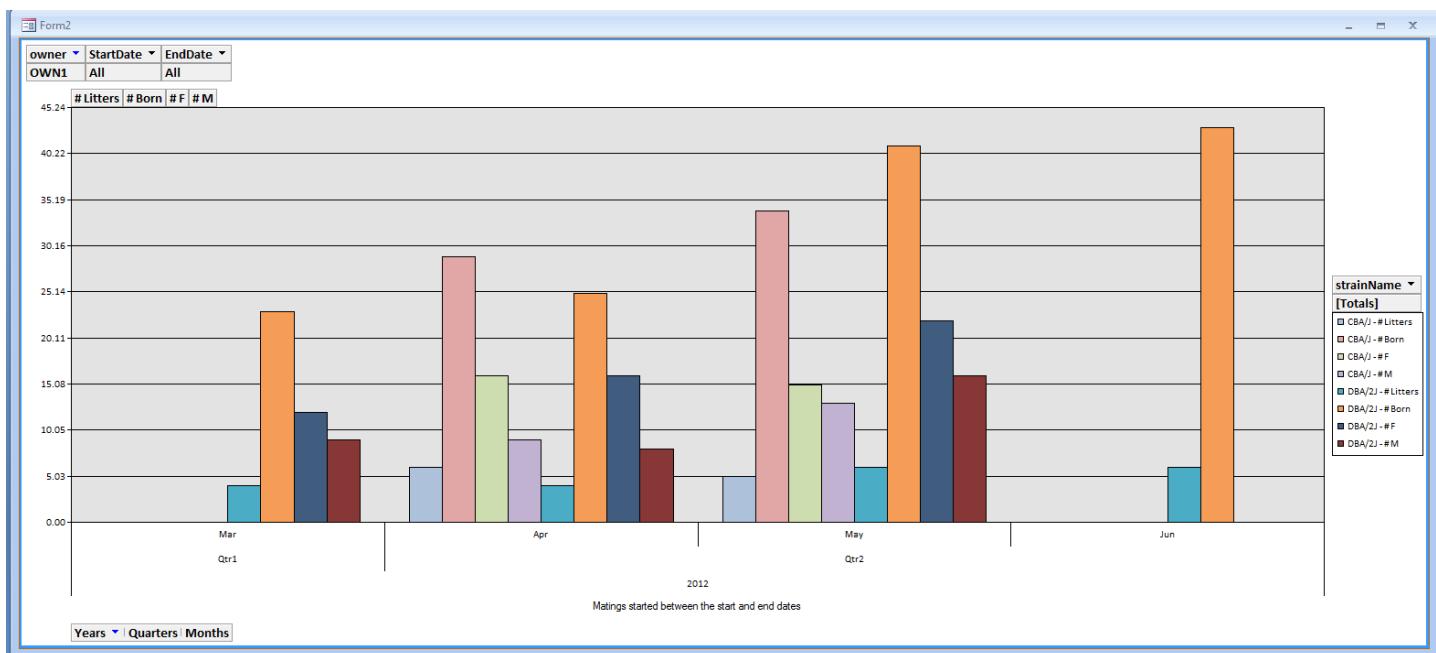


Figure 12-9 Litter summary pivot chart

## 12.6 Breeding Performance Report Mating Count Pivot Chart

The mating count pivot chart shows the number of matings started by year/quarter/month/week.

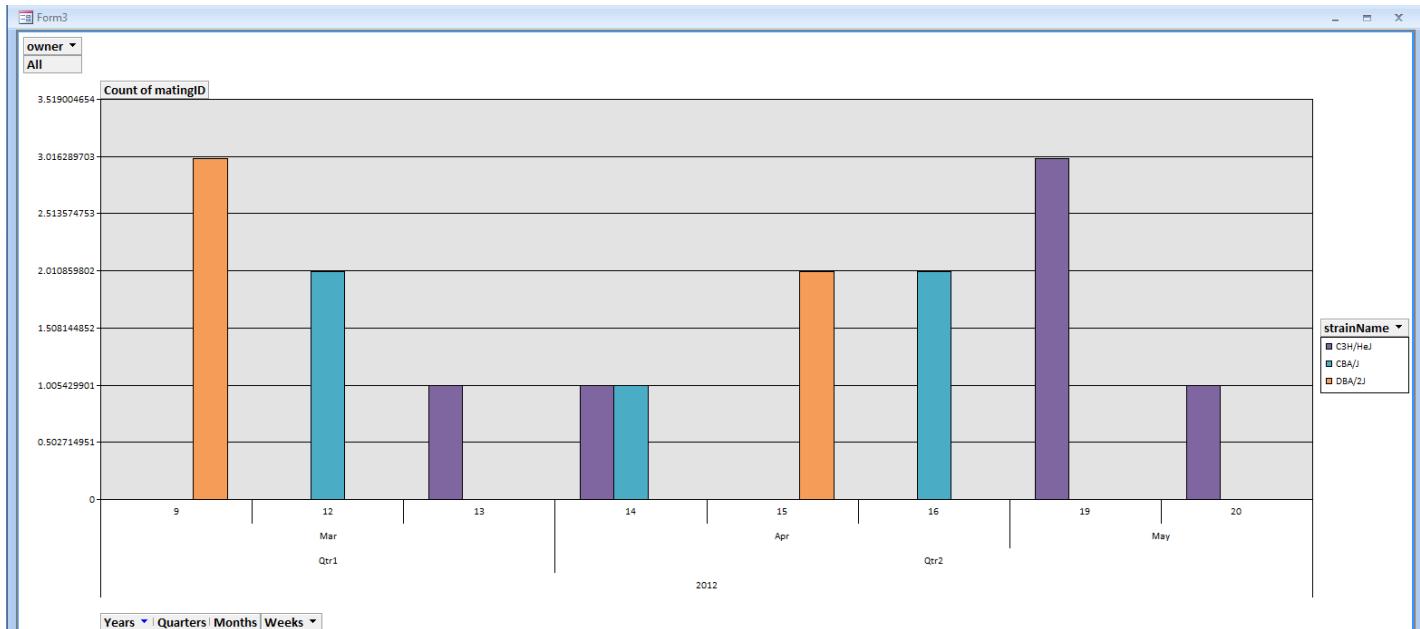


Figure 12-10 Mating counts by week

## 12.7 How to Use the Dynamic Features of a Pivot Table or Chart

Microsoft Access pivot charts and tables contain a number of customizable elements. JCMS has saved these charts and tables as a temporary form associated with the requested data (matings and litters for the time frame, strain, and owner criteria). These forms will be removed by JCMS when the application is closed. The forms are named sequentially as Form1, Form2, Form3, etc. We encourage manipulation of the forms to show the data in different ways. These changes are only displayed on the temporary form and do not change the underlying data. After changing a table or chart you can't get it back to a previous format, simply request the pivot table or chart again, creating a new temporary form.

Pivot tables and charts may be printed once the desired format is achieved. The size of the form can be adjusted by dragging the corners. Below are some tips about how to make changes.

### 12.7.1 Changing the Chart Type

Below is an example of what the mating count pivot chart looks like if it is changed to use the 3-D chart format. Changing the chart type is on the PivotChart Tools – Design tab (yellow circle).

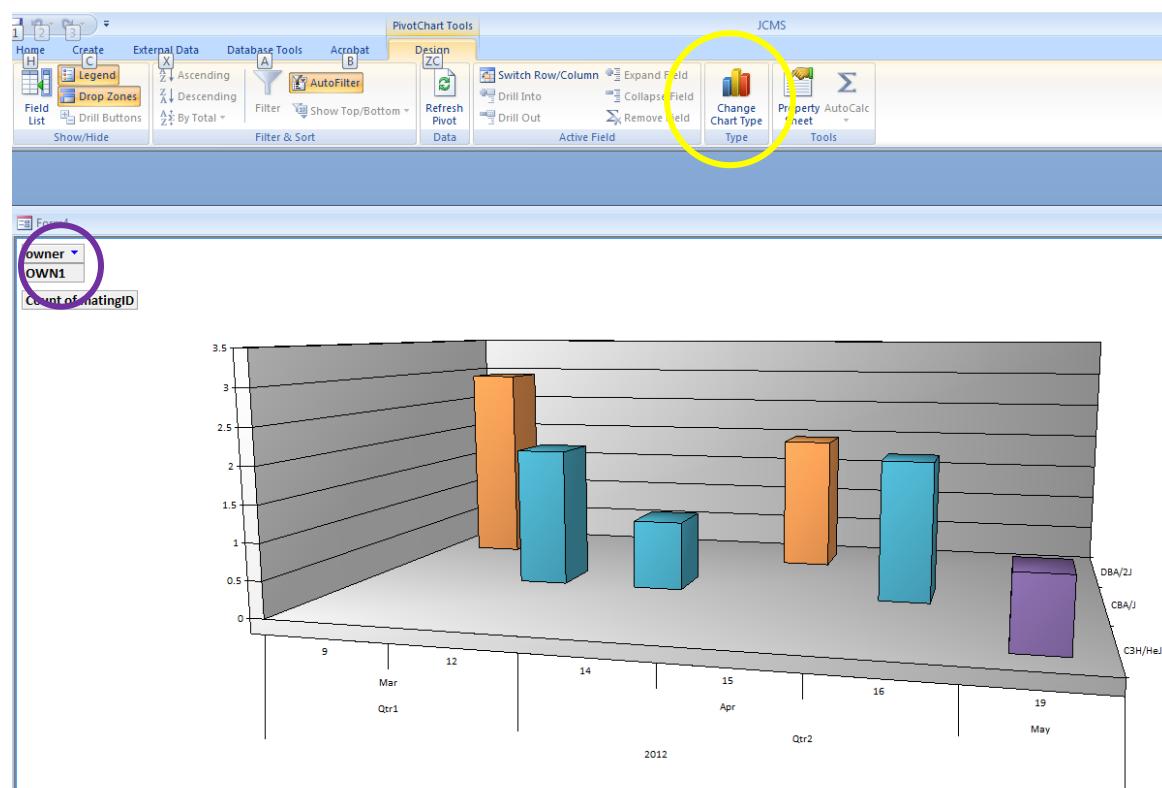


Figure 12-11 Chart changed to 3-D and only one owner

### 12.7.2 Showing Only Some of the Criteria

The charts and tables contain sections called “drop zones” (an example is shown in the purple circle above). Click on these to refine the possible choices. In the chart above, only “OWN1” has been selected instead of all owners.

The tables contain “drill buttons” that may be used to show or not show details. In the table below Qtr2 has been summarized (+) and March has been expanded (-) to show the birth dates of the litters.

owner	strainName	StartDate	EndDate																				
All	All	All	All																				
Years   Quarters   Months   Days																							
⊖ 2012																							
⊖ Qtr1																							
⊖ Mar																							
⊕ 21-Mar    ⊕ 23-Mar    ⊕ 29-Mar    ⊕ 30-Mar    Total																							
+ - + - + - + - + - + - + - + - + -																							
matingID	litterID	# Born	# F	# M	# Born	# F	# M	# Born	# F	# M	# Born	# F	# M	# Born									
⊖ 4	40	+ -	6	4	2									6 4 2									
	41	+ -				6	2	3						6 2 3									
	42	+ -												9 6 3									
	43	+ -												8 4 3									
	44	+ -												8 0 0									
	45	+ -												8 0 0									
	Total	+ -		6	4	2		6	2	3				12 6 5 12 6 5 33 10 6 4									

Figure 12-12 Changed “drill buttons”

### 12.7.3 Adjusting the Axis Scale

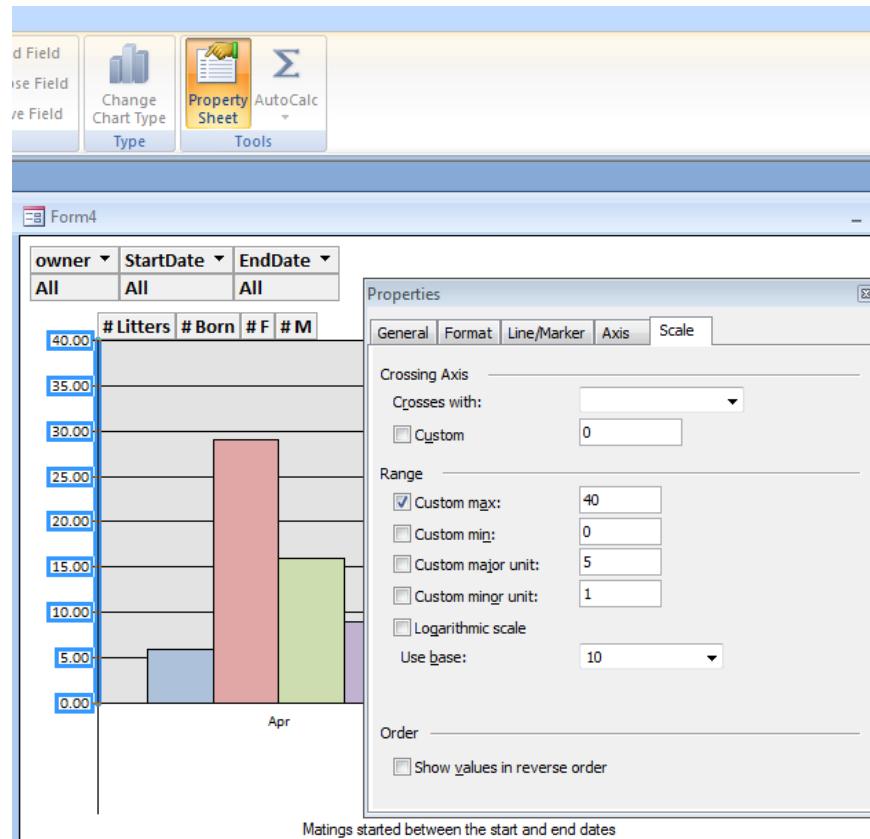


Figure 12-13 Properties dialog box

#### 12.7.4 Switch from Pivot Chart to Pivot Table



It is possible to change any pivot table into a pivot chart or vice versa. Click on the View button on the ribbon to select either PivotTable View or PivotChart View.

Figure 12-14 View button

#### 12.7.5 Viewing Values for Portions of a Pivot Chart or Table

Hover the mouse over a column on a pivot chart or cell in a pivot table to see a box containing the values summarized by that element.

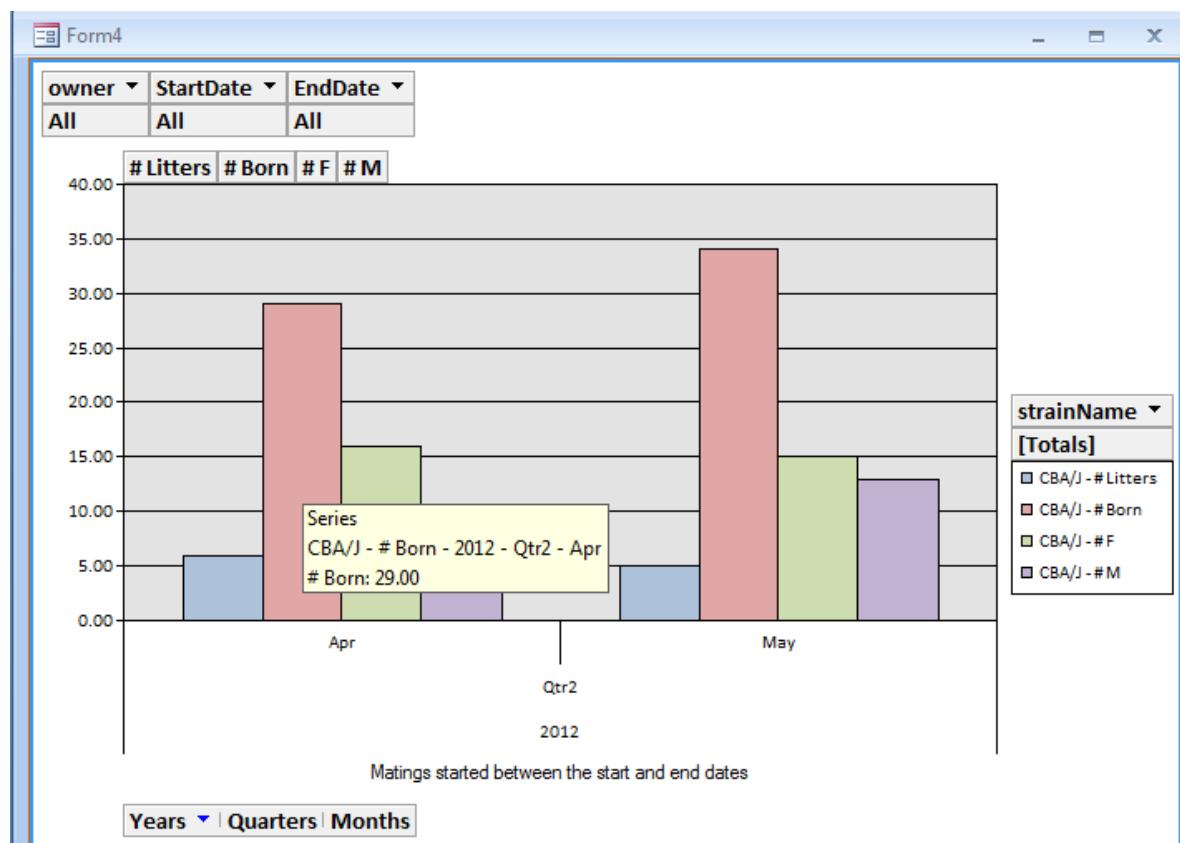


Figure 12-15 Hover over a column with the mouse to see the values

# 13 Genotyping

## 13.1 How Does Genotyping Work?

Any mouse or sample may have several genotypes. Each genotype consists of a gene and the two alleles that were found for that gene. In JCMS, alleles are associated with specific genes or gene classes. The JCMS forms that display alleles that may be associated with a gene use both criteria (association to the gene and association to the gene class) to come up with a list of alleles that may be associated with a given gene.

The Administrator is responsible for setting up the available genes, alleles and gene classes. See section 3.3.7 on initializing controlled value (CV) tables for genes, alleles, and gene classes for more information on how these are set up.

The following Gene Class values are preset, but may have been changed or added to by the Administrator.

Class name	Comments
E	endogenous
MKO	multi allele knock out
MTG	multi allele transgene
TG	trans gene
KO	knock out
KI	knock in
Floxed	tissue specific knock out
CTK	Combination transgene + KO

## 13.2 Adding a Genotype to a Mouse or Sample

The screenshot shows a Windows application window titled "Add Genotype (mtsadmin) {Owners Only}". The main title bar has standard window controls (minimize, maximize, close). Below the title bar is a horizontal toolbar with a "New" icon, a "Save" icon, and a "Cancel" icon. The main content area is divided into several sections:

- Strain / Stock # criteria:** A dropdown menu with a "Clear criteria" button.
- \*Mouse or Sample:** A group of radio buttons:
  - Mouse: Mouse ID:
  - Sample: Sample ID:
- Owner:** A text input field.
- \*Gene:** A dropdown menu.
- \*First Allele:** A dropdown menu. **Second Allele:** A dropdown menu. **Allele 2 Confidence:** A dropdown menu.
- \*Page No.:** A text input field. **Genotype Specimen Location:** A dropdown menu. **Genotype Date:** A date picker showing "7/5/2013".
- Comments:** A text area.
- Current Genotype:** A large empty text area.
- Genotype Session Log:** A large empty text area.
- Buttons at the bottom:** "Submit" and "Clear".
- Checkboxes at the bottom:** "Auto increment ID", "Include documents", and "Clear document list after submit".

Figure 13-1 Form: Add Genotype

Click the **Add Genotype** button on the manage mice tab to open this form. To add a new genotype, specify the mouse ID or sample ID. The current genotype will show on the screen. Next select the gene. Now the alleles that have been set up for the chosen gene will appear as choices in the first and second allele drop down boxes. The first allele and page number are required fields. If there is no page number, enter "None" in this field.

If the allele drop down boxes are blank, make sure a gene has been chosen. If they are still blank, have the Administrator add the alleles to the controlled value (CV) tables.

**Strain/Stock # criteria:** Select a value in the strain/stock # combo box to limit the mouse ID and sample ID choices. Use the *Clear criteria* button to show all the mouse and sample ID choices.

### 13.2.1 Mouse-Genotype Documents

Documents (computer files) may be associated with a mouse-genotype, but not a sample genotype. First check "include documents" to show the document list on the form.

The screenshot shows the 'Add a Genotype' form with the following details:

- Strain / Stock # criteria:** A dropdown menu with a 'Clear criteria' button.
- Mouse or Sample:**
  - Mouse: Mouse ID: DOE-Live-013
  - Sample: Sample ID: (empty)
  - Owner: OWN1
- Gene:** SKID
- First Allele:** Allele 1 Confidence: +, Allele 2 Confidence: -
- Page No.:** None, Genotype Specimen Location: Freezer A, Genotype Date: 7/5/2013
- Comments:** (empty text area)
- Associated Documents:**

Title_Description	FileName	DateUpload	Owner
Genotype information	A Gene protocol.docx	2/18/2013 1:22:10	OWN1

  - Buttons: ... Upload New, Associate Existing, Delete Association
  - Links: Double click to view
- Buttons:** Submit, Clear

**Figure 13-2 Add genotype documents**

This portion of the form is used to add document associations to a specific mouse-genotype combination. First create a list of one or more documents using the Upload New, Associate Existing, and Delete Association buttons. The document associations are created at the same

time as the genotype record, when the submit button is clicked. The associated documents are copied into a special “genotype folder”. Upload new is used to copy the file into the folder and also add it to the list of associated documents for this genotype. The Associate Existing button is used to select a file that has already been uploaded into the genotype folder and optionally associated with other mouse-genotype combinations. The “Delete association” button is used to remove a document from the document list. It does not delete the file or remove it from the genotype folder. See Documents (section 17) for more information on document and file handling.

Double clicking on a row in the document list will treat the file as a hyperlink and open it in the default browser. Use the “clear document list after submit” check box to save time selecting documents to associate if they should be associated with multiple mouse-genotype combinations.

### 13.3 Editing a Genotype

Gene:	allele1	conf	allele2	conf	g-page	location	date
SKID	+	True	-	True	None	Freezer A	7/5/2013

**Figure 13-3 Form: Edit Genotype**

Click the **Edit Genotype** button on the manage mice tab to open this form. To edit a genotype, first select the Mouse ID or Sample ID. Then all the current genotypes for that mouse will show in the list box on the right side of the form. Double click on the genotype to be edited or deleted and that information will appear in the boxes on the left side of the form. Click the Delete genotype button to remove this genotype. Make changes to the information in the boxes and then click Update Genotype to edit it.

Documents that are associated with this mouse-genotype combination are also edited when the update genotype button is clicked. Use the Upload new, associate existing, and delete association buttons to make changes to the list of associated documents. These changes are not finalized until the update genotype button is clicked.

Delete association does not remove the document from the genotype folder. Delete genotype will automatically remove all document associations for this mouse-genotype but does not remove the files from the genotype folder. For more information on document handling see section 17.

## 13.4 Adding a Genotype to a Group of Mice

The screenshot shows the 'Bulk Add Genotype (OWN1) (Owners Only)' window. At the top, there's a header 'Use this form to add a genotype to a group of mice'. On the left, under 'Select Group Type', there are three buttons: 'By mouse ID' (highlighted in blue), 'By litter number', and 'By pen'. Below this, a dropdown menu shows 'Gene: FSN'. Underneath, there are fields for 'First Allele: Allele1 Confidence' and 'Second Allele: Allele2 Confidence', both with dropdown menus. There are also fields for 'Page No.', 'Genotype Specimen', 'Genotype Date' (set to 1/15/2010), and 'Location'. A 'Comments:' text area is present. In the center, there are three sections: 'Mouse ID' (with 'Range' and 'Selected mice from list' options), 'Litter #' (with 'Range' and 'Selected litters from list' options), and 'Pen ID' (with 'Range' and 'Selected pens from list' options). Below these sections is a note: 'The "Show-me" button will display a list of the mice that may have genotype information added. This list can then be printed. To make changes you must select rows with your computer mouse and press "Submit."'. Three buttons are available: 'Show me', 'Show me print', and 'Submit'. At the bottom, a table displays the details of the selected mice:

Mouse ID	Strain	Gen	Sex	Birth Date	LS	BS	Diet	Pen	Owner	Already typed for gene?
A75	BALB/cJ	F03	M	9/2/2009	A	V		265	nobody	N
A74	BALB/cJ	F03	M	9/2/2009	A	V		265	nobody	N
A73	BALB/cJ	F03	M	9/2/2009	A	V		265	nobody	N
A72	BALB/cJ	F03	M	9/2/2009	A	V		265	nobody	N
A71	BALB/cJ	F03	M	9/2/2009	A	V		265	nobody	Y

**Figure 13-4 Form: Bulk Add Genotype**

Use this form for mice with an identical genotype. Click the **Bulk Add Genotype** button on the manage mice tab to open this form. Select the group of mice by mouse ID, Litter ID, or Pen. Click the **Show Me** button to see the list of possible mice in the group. Enter the genotype information on the left side of the form in the same fashion as on the Add Genotype form. Now, select the specific mice to have the genotype added by clicking on them in the show me box. Hold down the shift key to select a range or the ctrl key to select one at a time. Any mouse that already has this genotype will have "Y" listed in the "Already typed for gene?" column. If they are included in the group that is submitted, they will not have it added again or changed.

## 13.5 Genotype String Format

The administrator can configure how genotype strings are displayed and if the confidence level is shown or not. Three setup variables are used to indicate the string's appearance.

The default display of a genotype is: gene AB/CD? where nothing is displayed for high confidence and ? represents low confidence and the allele names in this example are AB and CD.

The setup variables for the confidence level: JCMS\_ALLELE\_CONF\_HIGH and JCMS\_ALLELE\_CONF\_LOW may be set to a value that is 8 characters or less.

The setup variable JCMS\_ALLELE\_GENE\_SEPARATORS specifies one character that is used before the alleles and one at the end. The default is no entry. If no characters are given, one space will be placed between the gene name and the alleles.

Note that the characters ' (single quote) " (double quotes) ; (semicolon) and , (comma) are not allowed. Cage cards do not print the allele confidence levels.

**Table 13-1 Examples of genotype strings**

JCMS_ALLELE_CONF_HIGH	JCMS_ALLELE_CONF_LOW	JCMS_ALLELE_GENE_SEPARATORS	RESULT
-Y	-N	[]	gene[AB-Y/CD-N]
blank	(?)	blank	gene AB/CD(?)
blank	blank	:	gene:AB/CD
(y)	(n)	<>	gene<AB(y)/CD(n)>

## 13.6 Genotype Work Report

The genotype work report is used to generate a list of litters due for genotyping during a particular time span.

Genotype Work Report (mtsadmin) {Secretaries and Owners}

### Request Genotype Work Report

List litters based on the litter birth date and the days to genotyping setup variable. The mating record must specify "needs genotyping".

Report time frame

\*Start date: 6 / 4 /2010

\*End date: 6 / 4 /2010

Owners: nobody, OWN1

Preview

**Figure 13-5 Request a genotype work report**

Only litters whose matings have “needs genotyping” set will be on the report. The setup variable JCMS\_DAYS\_TO\_GENOTYPE is used to determine how many days after birth genotyping is expected to be done. The default value is 14 days. JCMS is unable to determine if the genotyping has been done, therefore, all litters whose expected genotype date is within the selected time span are listed.

Genotype Work Report												
Owner	Mating #	Dam1 ID	Litter #	Strain	Room	Pen ID	Pen Name	Number Born	Date Born	Genotype Date	Age	
OWN1	95	A438	950	C3H/HeJ	B50-22	766		4	2/15/2010	3/1/2010	14	
nobody	110	Chuck004	1100	DBA/2J	BLDG29	786		5	2/18/2010	3/4/2010	14	
nobody	111	A436	1110	New mutant	BLDG29	787		2	3/2/2010	3/16/2010	14	
OWN1	117	A303	1171	FVB/NJ	B50-22	801		8	3/6/2010	3/20/2010	14	
OWN1	117	A303	1170	FVB/NJ	B50-22	801		6	3/4/2010	3/18/2010	14	

**Figure 13-6 Genotype Work Report**

## 14 Genotype Loader

There are two methods for importing mouse genotype data into JCMS.

- Use a comma separated value (CSV) file of genes, gene markers, or SNP data.
- Use a special file provided by the Jackson Laboratory Transgenic Genotyping Service (TGS).

Additionally, for Jackson Laboratory users, a genotyping request may be sent to the Transgenic Genotyping Service (TGS).

Details on the TGS options are included in Appendix 1 at the end of this document.

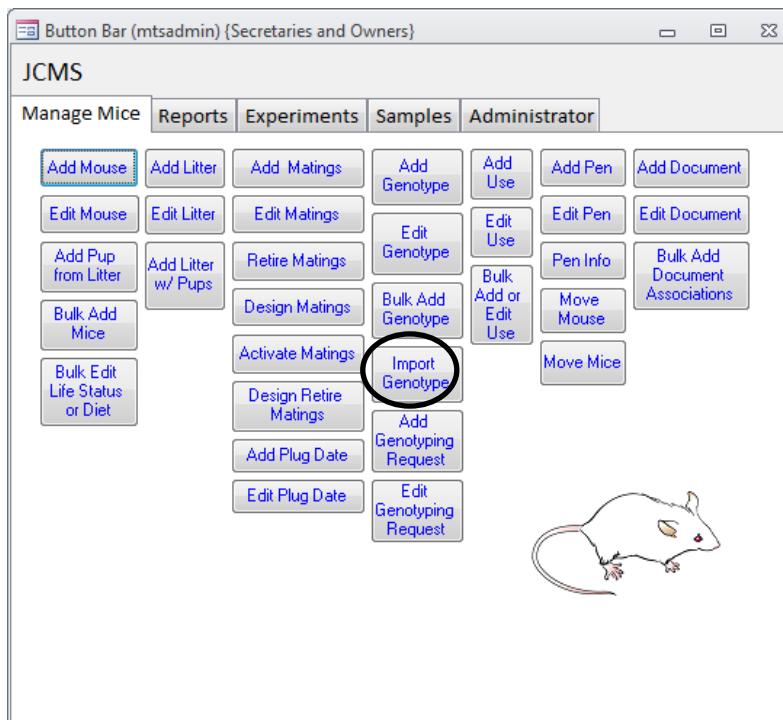


Figure 14-1 Form: The main button bar with the "Import Genotype" button visible

### 14.1 Import genotypes from a comma separated value (CSV) file

The genotype loader utility imports genotyping information and maps the rows and columns from a comma separated value (CSV) file to tables in the JCMS database. The result of the importation will be a set of new genotyping records for a given group of mice.

The software reads a formatted input file, parses it, validates it, and automatically populates the genotype records for the mice identified in the file.

Each file can be successfully loaded only once. If the user tries to load the file a second time it will generate error messages.

Note: This utility was originally written to import SNP (single nucleotide polymorphism) genotypes. Therefore, the acronym SNP may appear occasionally. The user can justifiably substitute the word "gene marker" or "gene". The resulting records are the same.

**The JCMS setup variable JCMS\_ENABLE\_GENOTYPE\_IMPORT must be set to true in order to bulk import genotype information from a file.**

### 14.1.1 Input File Format

There are a number of rules concerning the format of the input file. These are described fully in this section of the user guide.

The input file is a comma separated value (CSV) file. A CSV file has one row per line and columns separated by commas. The file can be created by any text editor or can be generated from Microsoft Excel. Only CSV files are accepted by the genotype loader.

The contents of **row one, column one** must contain the string:

JAX-CMS SNP Genotype Import vers 1.0

If this value is not present then JCMS will abort the importation.

**Row one, column four** starts the *gene* identifiers. If these do not appear in the Gene table in JCMS then first the user will need to add them.

In **row two, column four** begin the *gene class* identifiers. These are optional.

Beginning in **row three, columns one through three** are the *Mouse ID*, a *Vial ID*, and a *Position id*, respectively. These values must be unique throughout the file. Stated another way, they can only appear in an input file once.

**Columns four to the end of the line** contain the allele identifiers associated with the genes from row one of the same column.

Alleles have a confidence level associated with them. Please note that when loading genotype data from a CSV file the confidence level is automatically set to high. They can be modified later by using the **Edit Genotype** form.

Parsing of the input file stops when:

1. the mouse ID column is empty OR
2. the gene marker column is empty OR
3. the allele column is empty.

### Example import file:

The screenshot shows an Excel spreadsheet titled "JAX-CMS SNP Genotype Import vers 1.0". The first row contains headers: "JAX-CMS SNP Genotype Import", "01-167008", "01-173221", "01-178047", "01-182325", "01-186311", "01-190055", "01-19173", "07-124090", "07-135024", and "12". Rows 2 through 22 contain data for individual mice. Column A lists "Mouse ID"s, column B lists "Vial ID"s, and column C lists "Position"s. Columns D through L show "KO" (homozygous), "het" (heterozygous), and "?" (untyped) status for various genes. Column F contains "Gene designations" like "16-1-A01" and "16-1-A10". Column G contains "Gene class" codes like "A/C" and "A/G". Column H contains "Allele designations" like "G/A" and "T/C". Column I contains "Vial ID (optional)" values like "35330173" and "35330160". Column J contains "Position in the tray" values like "16-1-B01" and "16-1-B02". Column K contains "Allele designations" like "G/A" and "T/C". Column L contains "Vial ID (optional)" values like "35330173" and "35330160".

	A	B	C	D	E	F	G	H	I	J	K	L
1	JAX-CMS SNP Genotype Import	01-167008	01-173221	01-178047	01-182325	01-186311	01-190055	01-19173	07-124090	07-135024	12	
2	Mouse ID	Vial ID	Position	KO	KO		KI		TG			
3	200001	35330173	16-1-A01	het/wt	het	het	?	het	het	het	het	
4	200002	35330172	16-1-A02	A/T	het	het	het	het	het	A	C	
5	200003		16-1-A03	+	C/T	G	T	het/het	het	het/het	A/C	
6	200004	35330176	16-1-A04	/A	T	C/G	T	G	?	G	het	
7	200005	35330498		A	T	G	C/T	G	T	G/T	A	
8	200006	35330497	16-1-A06	A/A	T	C/G	T	G	T	G	A	
9	200007	35330496	16-1-A07	het/het	het	het	het	het/wt	het	het	het	
10	200008	35330495		/het	het	het	het	het	het	A/G	C	
11	200009		16-1-A09	het	het	het	het	het	het	A	C/G	
12	200010	35330493	16-1-A10	A	T	/G	T	C	/T	G	het	
13	200011	35330160		A	T	G	T	G	T	G	het	
14	200012	35330161	16-1-A12	A/C	T	G	T	G	T	G	A	
15	200013	35330479	16-1-B01	A	T	G	T	het	het	het	C	
16	200014	35330480		A	T					G		
17	200015		16-1-B03	het	het					het		
18	200016	35330482	16-1-B04	het	het					G		
19	200017	35330483	16-1-B05	A	C					A		
20	200018	35330484		A	G/					C	T	
21	200019	35330485	16-1-B07	het	het	het	het	het	het	het	C	
22	200020					?	?	T	het	het	?	

Figure 14-2 Spreadsheet: the input file as seen in MS Excel

The table below lists the components of the input file and whether they are mandatory or not. The column values are case insensitive. It also provides more details about the values in the columns, their meaning, and how they map to the JCMS database tables.

Table 14-1 The mapping of the input fields to the data tables

Field	Mandatory	Properties	Description [Table.column]
Mouse ID	Yes	Alphanumeric, 16 characters maximum	Unique identifier. Mouse IDs must exist in the database prior to importing the file. [Mouse.ID]
Vial id	No	Alphanumeric, 16 characters maximum	Maps to the sampleVialID tag field. Must be unique. [Mouse.sampleVialID]
Position	No	Alphanumeric, 16 characters maximum	Maps to the sampleVialTagPosition. [Mouse.sampleVialTagPosition]
Gene class ID	No	Alphanumeric, 16 characters maximum	Optional field. If present identifies a class of genes. [cv_GeneClass.GeneClass and Gene.geneClass fields]

Field	Mandatory	Properties	Description [Table.column]
Gene markers	Yes	Alphanumeric, 32 characters maximum	Treated as gene names. [Gene.labSymbol]
Allele designations	Yes	Alphanumeric, 8 characters maximum.	Associated with the gene marker (Gene) not the gene class. [Allele.allele, Genotype.allele1, and Genotype.allele2]

### 14.1.2 Alleles

Alleles can be represented four ways in the input file. The table below lists the four ways along with examples and the results produced by the example.

**Table 14-2 Genotype loader allele representations**

Allele representation	Example	Result
Two string separated by a slash	"abc/xyz"	Allele1 = "abc", allele2 = "xyz"
A single string	"hom"	Allele1 = "hom", allele2 = ""
A string followed by a slash	"cre1 /"	Allele1 = "cre1", allele2 = ""
A slash followed by a string	"/cre1"	Allele1 = "", allele2 = "cre1"

### 14.1.3 Import File Integrity Checks

The following table lists the tests performed and exceptions caught.

**Table 14-3 Genotype loader error messages**

Rule	Exception	Action
Row one, column one does not contain a valid genotype file identifier.	File is not an import file.	Popup message box: "The file <filename> does not appear to be a valid JAX-CMS formatted genotype importation file." <ok> After clicking OK the file dialog exits.
All mice IDs from the input file must exist in the database.	A mouse ID is not found	Record the error as "Mouse Id doesn't exist in the database, no allele and genotype records are created for that mouse id' in the error log generated at the end and skip this mouse record in the file.
Mouse record must have an empty vialID or it must match the current value.	Mouse record has a sampleVialID that is not NULL and is different from the value we are trying to write.	Record the error as "Vial id doesn't match the value in the input file' in the error log generated at the end and don't update the mouse record.
If a gene name (marker) exists in the database then the gene class in the input file must match.	Gene and Gene class tuple from input file do not match what is in the database.	Record the error as 'Gene already exists in the database but geneClass doesn't match the value in the input file" in the error log generated at the end and do not add the gene / gene class tuple to the database.

Rule	Exception	Action
Input file must be in a valid format (con't)	File is a valid file but does not conform to the expected format.	Popup message box: "input file <filename> is not in a valid format. <additional info>" (ok)  The validation dialog remains visible with the error message and any additional info entered into the 'details' box but the only option is the Cancel button.
Mouse ID(s) not unique	Mouse ID (recipient ID) <id> already exists in database at line	Popup message box: "Mouse ID (recipient ID) <id> already exists in database at line <n>." and abort the operation
Not a CSV file	Invalid file	Popup message box: "Invalid file" and abort the operation
Missing mouse id field in the input file	Missing Mouse ID at line	Popup message box: "Missing Mouse ID at line <n> in the input file" and abort the operation
Missing Gene in the input file	Missing Gene at line	Popup message box: "Missing Gene at line <n>" and abort the operation
Missing Allele in the input file	Invalid Allele at line	Popup message box: "Invalid Allele at line <n>"

#### 14.1.4 Select the type of Import

The "Import Genotype" button appears in the 'genotype column' on the manage mice tab. It opens a dialog box used to indicate which import to use. Select "Import JCMS \*.csv format file".

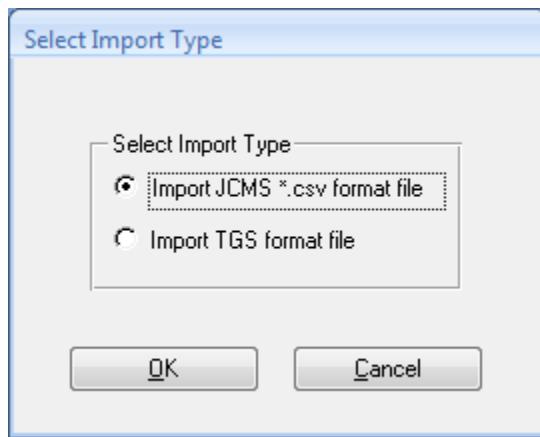


Figure 14-3 Select import type dialog box

The JCMS setup variable `JCMS_ENABLE_GENOTYPE_IMPORT` must be set to true in order to bulk import genotype information from a file. This is the default.

When functional, clicking OK brings up a standard file open dialog. See Figure 14-4. The user can browse to files or type in a pathname, including a network (UNC) pathname. It accepts only comma separated value (.csv) files.

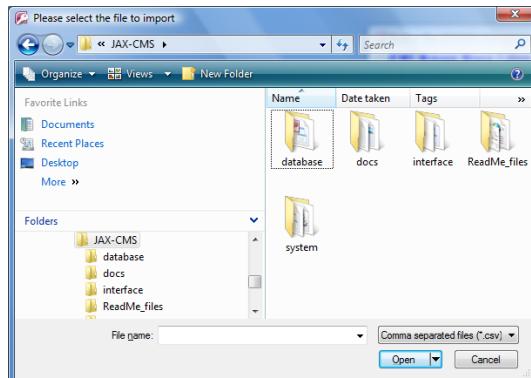


Figure 14-4 The file open dialog

If the user clicks cancel, then the message “Request to upload the file was cancelled” is displayed and the operation is aborted.

If the input file is not a comma separated value file the error message “The file xxx.csv is not in a valid format” is displayed and the operation is aborted.

### 14.1.5 Verification

Once a file is selected and verified, the software runs a series of validations against it. If any of the validations fail, the user will not be able to import the genotypes. If all the validation tests pass, the user may still elect to cancel.

Figure 14-5, Figure 14-6, and Figure 14-7 show the user notification dialog that appears during validation, after a successful validation, and after the import, respectively.

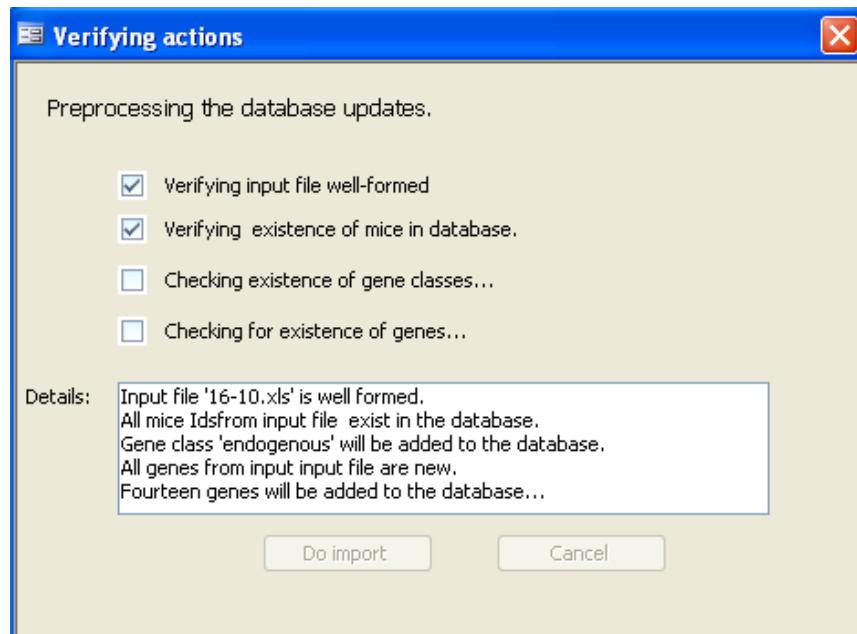
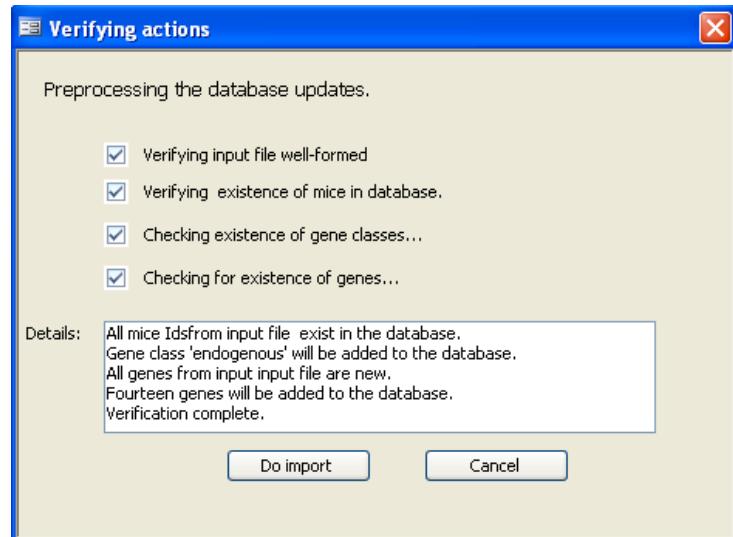


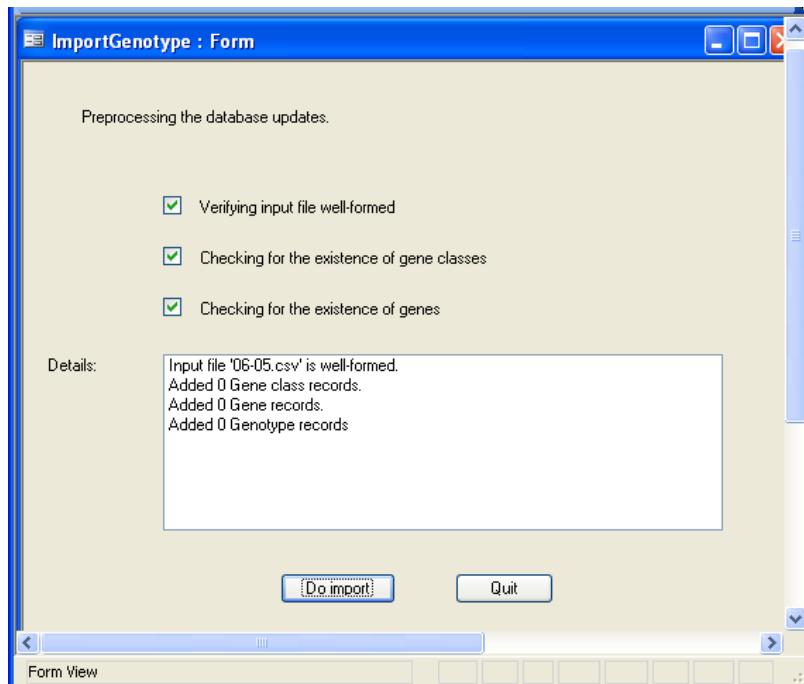
Figure 14-5 The user notification screen during validation



**Figure 14-6 The user notification after validation**

The validation process:

1. Checks the input file contents of row 1, column 1 to be the string “JAX-CMS SNP Genotype Import vers 1.0”, If this value is not present then error message “The file xxx.csv does not appear to be a valid JAX-CMS formatted genotype importation file” is displayed and the operation is aborted.
2. Once the file is validated, the software checks if the gene class from the input file exists in the database, if not it is added to the database.
3. Tests if the gene markers exist in the input file, if not then error message “Gene doesn’t exist” is displayed and aborts the operation. If the input file contains gene markers then the software checks that those genes exist in the database, if not the genes are added to the database.



**Figure 14-7 The user notification after import**

When the user clicks the **Do Import** button, the software:

1. Checks if mouse ID from the input file exists in the database. If not it is recorded in the load report and the row is skipped. If it does exist then it checks if the user has permissions to edit the mouse.
2. Updates the mouse record with new vial Id and vial tag position values.
3. Checks if the allele(s) from the input file exist in the database; if not, it adds all the new alleles to the database.
4. Checks if the genotype record for each mouse, gene, and allele exists in the database; if not, it adds all the genotype records with respective mouse, gene, and allele to the database

#### 14.1.6 Genotype Load Report

The screenshot shows a Windows application window titled "SNPLoadReport : Form". The main title is "SNP Genotype Load Report". Below the title, there is a section titled "Summary" which contains a message box stating: "The file 06-05.csv is a valid JAX-CMS formatted genotype importation file" followed by a list of statistics: "Added 0 geneclass records", "Added 0 gene records", "Added 0 Allele records", "Added 0 Genotype records", and "Updated 0 Mouse records". Below the summary is a section titled "Details of the Errors" containing a scrollable list of error messages. The errors listed are all related to invalid alleles, such as "Invalid Allele at mouseId 200022 and gene 07-135024189-M", "Invalid Allele b at mouseId 200001 and gene 01-182325478-N", and many others. At the bottom left of the window, there is a "Form View" button.

**Figure 14-8 Genotype Load Report**

A load report is generated at the end of the import process.

This form is invoked from the Import Genotype form after the importation process is done. It gives the summary of the input file verification, count of gene classes added, genes added, alleles added, genotypes added, and mice updated in the database. It also gives the list of errors encountered during the importation process.

# 15 Scheduling Procedures (Uses)

## 15.1 How do Mouse Uses Work?

Mice may be assigned various uses over time. JCMS uses the term "use" to refer to a procedure, protocol, test, experiment, examination, assessment, etc. that was done with a mouse at a particular point in time. Some "uses", such as taking a weight measurement, may be repeated several times over the life of a mouse.

Uses may have a use age that provides a projected date for the use. The projected date can be used to generate a report of work that needs to be done during a particular time frame. When the use is complete the actual date is entered, and up to 10 fields are available for entering data results (text format). The mouse use Calendar (Section 16) is used to display the projected and/or actual dates by the month, week, or day. Uses that are associated with plug dates and a date post conception (DPC) may be entered at the same time as the plug date (Section 11.1 Add Plug Date).

If the "mouse uses" system does not provide enough flexibility or data collection, use the Experimental Plan portion of JCMS instead. Experimental plans allow for user-designed experimental tests and metadata plus more complex scheduling.

## 15.2 Adding a Use to a Mouse

The screenshot shows the 'Add Mouse Use' window. At the top, it displays the mouse's identification information: Mouse ID (ST-007), Mouse Owner (OWN1), Birth Date (7/1/2011), and Sex (M). The 'Use' dropdown is set to 'Basic blood work'. Under 'Use Age Determination', the 'Manually enter' option is selected, and the use age is listed as 400.00 days. The 'Projected Date' is 8/4/2012, and the 'Actual Date' is 11/5/2012. The 'Done' checkbox is unchecked. The 'Comments' text area is empty. There are two checkboxes at the bottom left: 'Auto increment ID' and 'On Submit, don't clear data'. On the right, there is a large grid table with columns for 'Use', 'Use Age', 'Proj Date', 'Act Date', 'Done?', and 'Comments'. One row in the grid is visible, showing 'Skin graft', '366.00', '7/1/2012', '9/10/2012', 'Done', and an empty comments field. Below the grid, there are sections for laboratory results: 'CBC' (with 10 rows), 'WBC' (with 10 rows), 'RBC' (with 10 rows), 'Lipids' (with 10 rows), and 'Cholesterol' (with 10 rows). To the right of these sections is a 'Session log' area, which is currently empty. At the bottom of the form are 'Submit' and 'Clear' buttons.

Figure 15-1 Form: Add Use

Click the **Add Use** button on the manage mice tab. Enter the mouse ID. Any current uses for this mouse will display in the gray list box. The use age is stored in days. Use the “Days/Weeks/Months” radio buttons to indicate how the use age is displayed. JCMS will convert months or weeks into days by using 30.4375 days in a month and 7 days in a week. With each use, some information may be stored in the comment field. The ten fields at the bottom of the form are provided to store results or other information associated with this use. It is the responsibility of the user to keep track of what the data in each field means. A suggestion is to include a meaningful caption for each field when the use term is added (by the administrator). If there is no caption for the field it is labeled as D1, D2, ...D10 on the form.

The projected date is always calculated. Select the “manually enter” radio button and the projected date will be calculated based on the mouse’s birth date. Select the “calculate from plug date and DPC” radio button and the projected date will be set to the plug date plus DPC (rounded down). The use age is then calculated as the difference between the mouse’s birth date and the projected date. It is assumed that the mouse was not exited prior to the projected date.

**Figure 15-2 Adding a use based on a plug date**

### 15.3 Mouse Use Work Report

Use the **Mouse-Use Work Report** button on the reports tab to request a listing of mouse uses. There are multiple criteria including projected or actual date, use status (done or not done), owner, and life status. The date may be within a range or open-ended by un-checking either the “from” or “to” date. It is possible for a mouse-use to have no projected or actual date. Include these mouse-uses by checking “include if no date.”

This report is designed to be printed in landscape format. If the print preview shows the report in portrait format, change the page layout to landscape.

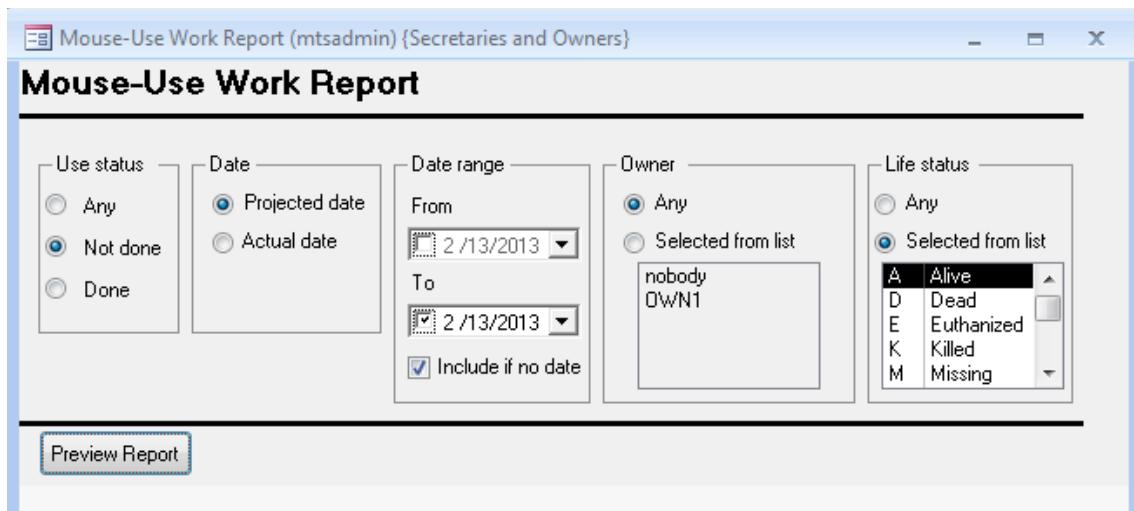


Figure 15-3 Request mouse use work report

## 15.4 Editing a Mouse Use

Use	Use Age	Proj Date	Act Date	Done?	Comments
Neurological exam	2.96	9/29/2011	9/28/2012	Done	
Skin graft	12.02	7/1/2012	9/5/2012	Done	

Figure 15-4 Form: Edit Use

Click the **Edit Use** button on the manage mice tab. Enter the mouse ID. Any current uses for this mouse will display in the gray list box. To edit a use, double click on the use in the list box. The values for this use will appear in the white editing boxes. Press the Submit button for the changes.

to be saved or the Delete button to remove the selected use from JCMS. New uses cannot be added on the edit use form, only changes to existing ones.

Often data results are repetitive. When using the auto increment ID function, it is not possible to repeat the data for the next mouse as this is an edit form. To make repeat entry of the same data easier, there are two buttons on the form. Click on **Copy Data to JCMS Clipboard** before clicking the Submit button. Select the proper use from the next mouse. Click the **Paste Data from JCMS Clipboard** and the data fields will have the values from that previous mouse pasted into them. No other fields will be affected by this special paste. The clipboard will continue to contain these values so they may be pasted into a third, fourth, etc. mouse use. This clipboard cannot be used to paste these values into any other application.

## 15.5 Adding or Editing a Mouse Use for a Group of Mice

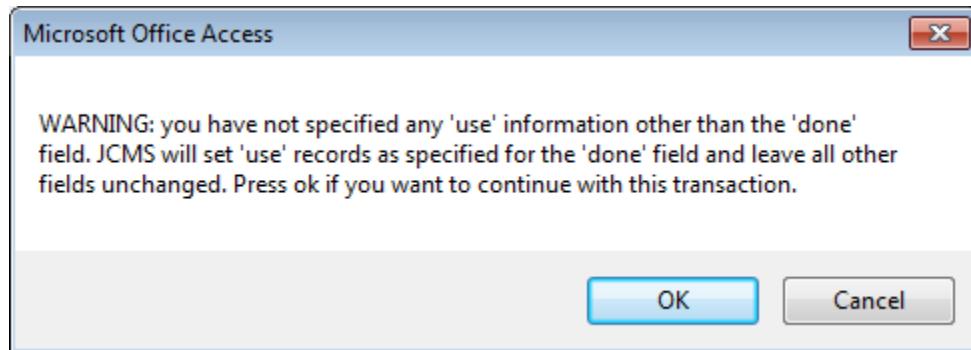
**Figure 15-5 Bulk Add or Edit Use Form**

Click the **Bulk Add or Edit Use** button on the manage mice tab. This form allows both adding and editing mouse uses for a set of mice. Both functions cannot be done at once, first choose add or edit. The group may be selected by mouse ID, Litter #, or Pen ID. Click the **Show Me** button to see the list of possible mice.

Edit use: specify an “old-use” (the use to be edited). Only mice that have the “old-use” will be displayed in the “show-me” list. Thus, it is possible to select mice to work on from one of the selection criteria list boxes, and potentially none will be displayed in the show-me list because they do not have the old-use.

One mouse can be scheduled for the same use multiple times. Thus it is possible to see the same mouse listed multiple times in the show-me box. Use records can be distinguished from each other by looking at the old-use age. This form requires first selecting the mice to be modified and then choosing which of possibly many *use-records* to change.

Only the new use info fields that contain information in them will be changed. Blank fields are ignored. The exception to this is the *Done* field. All selected use records will be changed. If all fields in the new use section are blank, JCMS will verify that the done field should be changed.



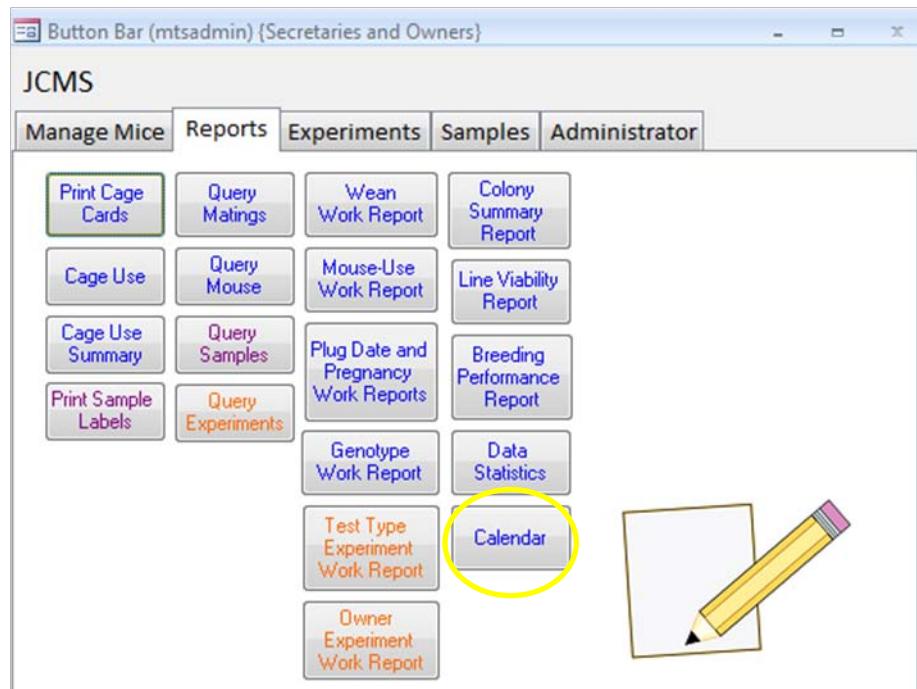
**Figure 15-6 Warning about making a change to the "done" field**

NOTE: When editing a use, the contents of the comment field will replace any existing comments associated with the specific set of uses being edited. This may seem counter intuitive when thinking in terms of editing the comments. However, this edit function is designed to copy over any existing information unless the comments field is blank. It is not possible to use this form to completely remove the comments.

Add new use: select the “add a mouse use” choice and specify the mice to be operated on via one of the three selection criteria (mouse ID, pen, or litter). It is valid to add the same use many times for one mouse. This form will not calculate the projected use date; it must be entered on the form and will be the same for all selected mice. The birth date, use age, and projected date will not be synchronized unless the user carefully determines them.

## 16 Calendar

The calendar is used to display the mouse uses scheduled for a particular date. To open the calendar, click on the Calendar button on the Reports tab.



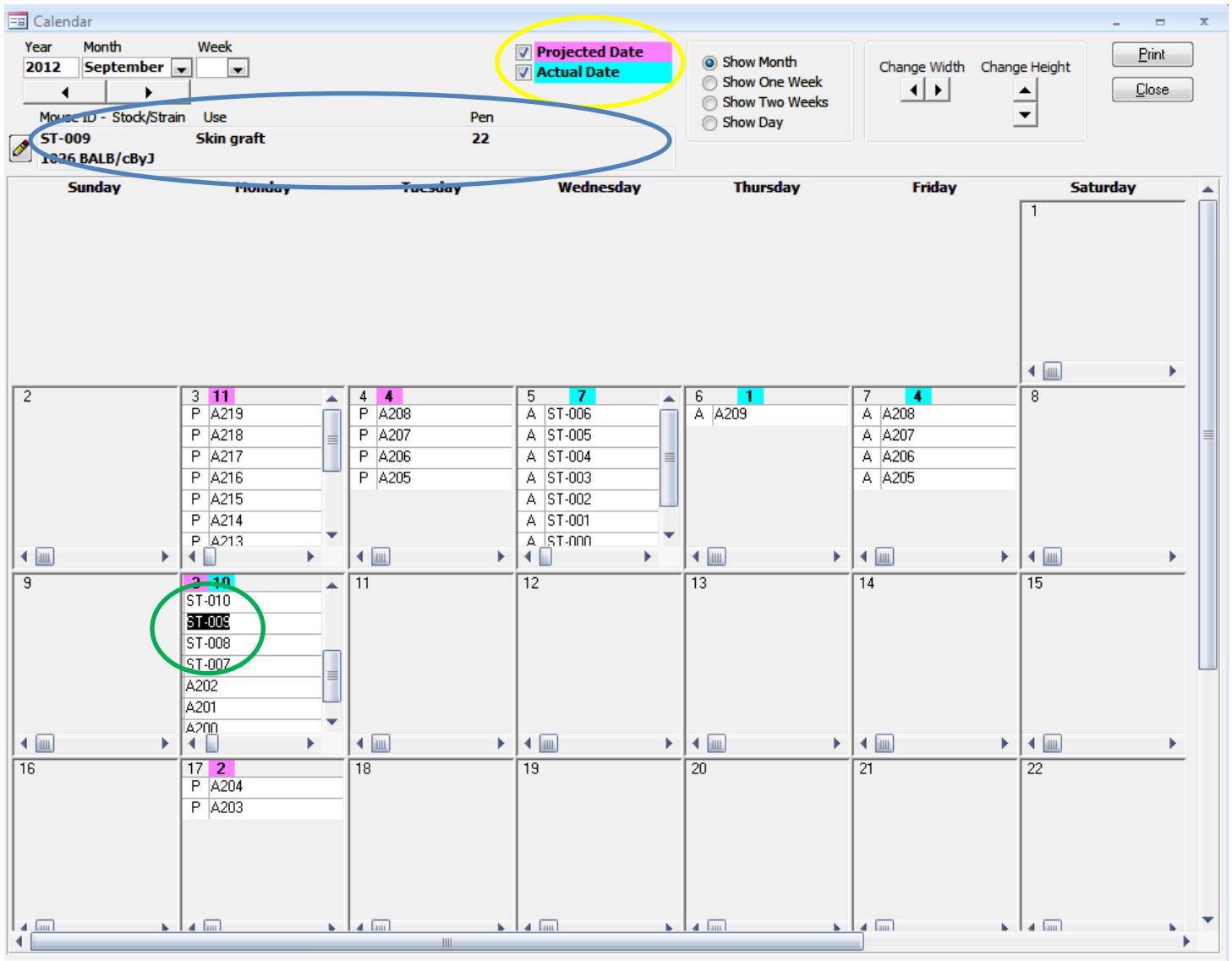
**Figure 16-1 Open the calendar**

The request mouse use calendar form will open. Specify one or more mouse uses, strains, and owners to be shown on the calendar. Mouse uses have a status of done or not done. The calendar can show only those that are not done yet or can be used to show a history of all those that were done on a particular date.

Click the Display Calendar button.

A screenshot of the 'Request Mouse Use Calendar' form. The title bar says 'Request Mouse Use Calendar (mtsadmin) (Owners Only)'. The form has several sections: 1) 'Calendar Criteria' with a dropdown for 'Choose mouse use(s)' containing items like 'Necropsy results', 'Six month complete checkup', etc.; a radio button group for 'Use status' ('Any', 'Done', 'Not Done'); and a dropdown for 'Choose owner(s)' containing 'nobody' and 'OWN1'. 2) 'Choose stock # / strain name(s)' with a radio button group ('Any', 'Selected from list') and a dropdown list showing stock numbers and strain names: 2052, 100006, 1026, 651, 659, 664, 656, followed by strain names: B6.129P2-Apoe<tm1Unc>/J, B6D2F1/J, BALB/cByJ, BALB/cJ, C3H/HeJ, C57BL/6J, CBA/J. 3) A 'Display Calendar' button at the bottom left.

**Figure 16-2 Form: Request mouse use calendar**



**Figure 16-3 Example of a mouse use calendar**

The calendar displays two dates, the projected date (date the use is planned for) and the actual date the use was completed. Check and/or uncheck the projected and actual dates (yellow oval) to add or remove them from the calendar. Projected dates are marked with a "P" and actual dates with an "A".

Click on a row in the calendar (green circle) and the details of that mouse use are displayed (blue oval). Also the scroll bars on the bottom of the date box may be used to see the same details [mouse ID, use, pen ID/pen name, and stock #/strain].

The size of the date boxes can be adjusted by using the Change Width and Change Height buttons. The year, month, and week can also be changed.

The format of the calendar can be changed from month to one week, two weeks, or one day.

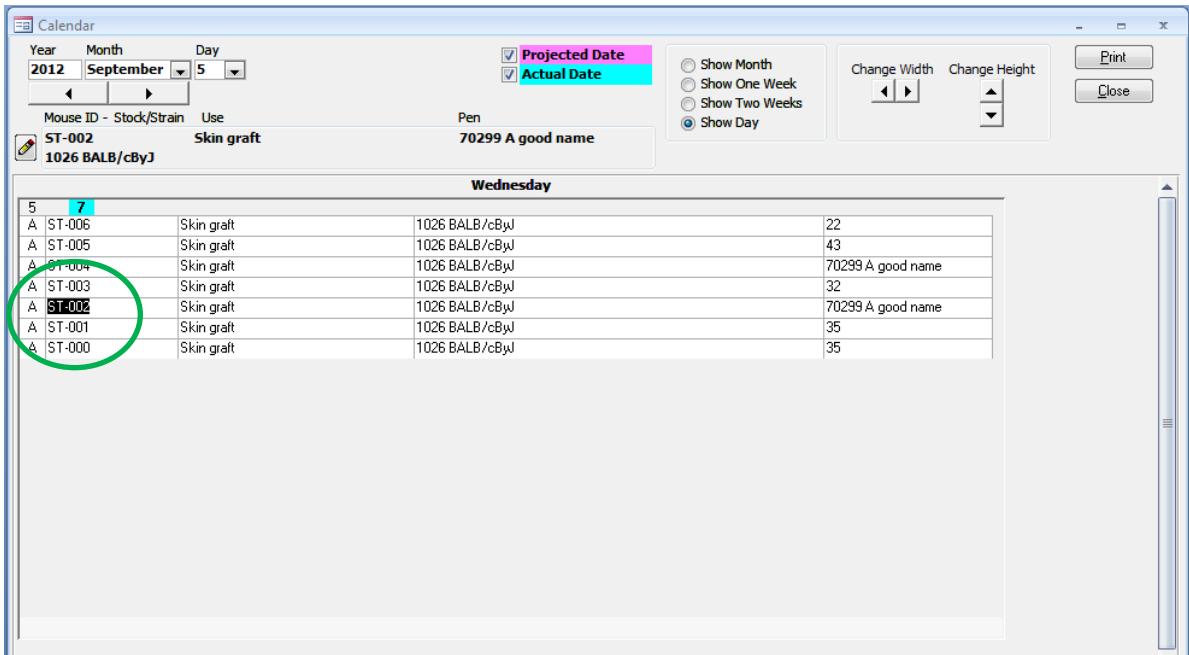


Figure 16-4 Single day format

## 16.1 Edit a Mouse Use Shown on the Calendar

To open a mouse use for editing, click on the pencil icon button or on the use (black circles). This will open the Edit Mouse Uses form as shown below. Please note that the calendar is not automatically refreshed to show changes made to the mouse use. To see the changes, close the calendar and reopen it.

Year	Month	Week		
2012	October		<input checked="" type="checkbox"/> Projected Date	<input checked="" type="checkbox"/> Actual Date
			<input type="radio"/> Show Month	<input type="radio"/> Show One Week
			<input type="radio"/> Show Two Weeks	<input checked="" type="radio"/> Show Day
			Change Width	Change Height
			<input type="button" value="Print"/>	<input type="button" value="Close"/>

Mouse ID - Stock/Strain: A93  
Use: Neurological exam

Pen: 70300

**Edit Mouse Use**

*Mouse ID: A93	Mouse Owner: <b>nobody</b>	Birth Date: 4/1/2012	Sex: F
*Use: Neurological exam	Strain: C3H/HeJ		
Use Age in: <input type="radio"/> Days <input type="radio"/> Weeks <input checked="" type="radio"/> Months: 6.24	Select a use to edit or delete from the list below by double clicking on the item. When you press the 'Submit' or 'Delete' button, the item selected will be updated or deleted.		
Projected Date: 10/8/2012	<b>NOTE: Any changes to a mouse use will not display on the calendar. Close the calendar and re-open it to see changes.</b>		
Actual Date: 11/9/2012			
<input type="button" value="Submit"/>	<input type="button" value="Delete"/>	<input type="button" value="Cancel"/>	<input type="button" value="Comments"/>

Figure 16-5 Opening the edit mouse uses form

## 16.2 Mouse Use Details Form

Double click on a row on the calendar or on the mouse ID, pen, or stock#/strain (orange circles) to open the Mouse use details form showing information about that specific mouse use.

The screenshot shows two windows side-by-side. On the left is a 'Calendar' window for September 2012. A mouse ID ('ST-016') and strain ('1800 FVB/NJ') are highlighted with orange circles. On the right is a 'Mouse Use Details' form window. The 'Actual Date' checkbox is checked. The 'Pen' field shows '24'. The 'Mouse Use Details' form contains the following data:

Mouse ID:	ST-016	Owner:	nobody
Use:	Skin graft	Pen ID:	24
Use Age in months:	12.00	Pen Name:	
Projected Date:	8/1/2012	<input checked="" type="checkbox"/> Done	Actual Date: 9/10/2012
Comment:			
D1	D6		
D2	D7		
D3	D8		
D4	D9		
D5	D10		
Strain:	FVB/NJ		
Stock #:	1800	Generation:	F02
Sex:	M	Life Status:	S
Birth Date:	8/1/2011	Exit Date:	5/14/2012

Figure 16-6 Mouse use details form

### **16.3 Print a Calendar Report**

Use the Print button to obtain a report formatted for printing that shows all the rows on the calendar.

#### **Mouse Use Calendar**

Monday, September 10, 2012

September, 2012		
Mouse ID	Cage ID / Name	Stock # / Strain Name
<b>September 5, 2012</b>		
<b>Actual Date</b>		
<b>Skin graft</b>		
ST-005	43	1026 BALB/cByJ
ST-004	70299 A good name	1026 BALB/cByJ
ST-003	32	1026 BALB/cByJ
ST-002	70299 A good name	1026 BALB/cByJ
ST-001	35	1026 BALB/cByJ
ST-000	35	1026 BALB/cByJ
ST-006	22	1026 BALB/cByJ
<b>September 6, 2012</b>		
<b>Actual Date</b>		
<b>Skin graft</b>		
A209	70323	656 CBA/J
<b>September 7, 2012</b>		
<b>Actual Date</b>		

**Figure 16-7 Example of a calendar printout**

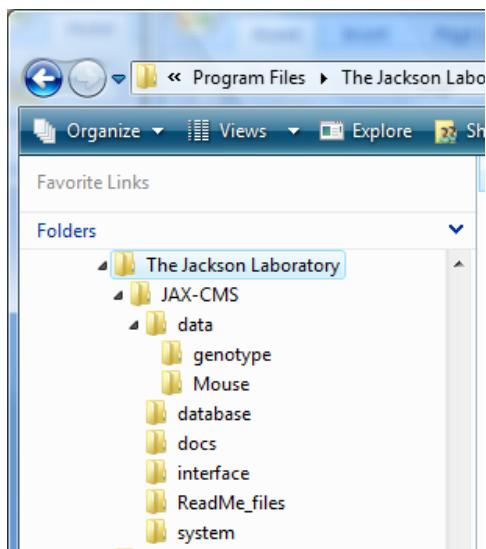
#### **16.3.1 Suggestions for speeding up the calendar form**

When the database is large, the calendar may take a long time to open. An active hourglass is visible during the wait for the calendar to display. To reduce the waiting time, try to limit the criteria to only one or two owners. The more owners included, the slower the search is to find all the mouse uses. Limit the criteria to only one or two mouse uses. Allow "any" strain instead of selecting certain ones.

# 17 Documents

A document is a computer file, which may be associated with one or more IDs in certain JCMS categories, such as a mouse, sample, mouse genotype, experimental data, etc. In the first implementation of this feature, one or more documents may be associated with a mouse-genotype combination (a specific mouse ID and one of its genotypes). The description of this feature uses mouse genotype in the examples. Associations to other categories are now available.

## 17.1 Document Storage



**Figure 17-1 Possible data file setup for documents**

Any file stored on the computer may become a JCMS document. JCMS does not incorporate the documents into its database. It moves a copy of the original file into a special "data" folder and maintains a link to the file in the JCMS database. JCMS allows the user to open the file by treating this link as a standard hyperlink, opening the file in the default browser or program depending on the defaults set on the user computer. This link will be broken if the user manually moves or deletes the file from this folder.

The JCMS Setup variable

JCMS\_DATA\_FILE\_DIRECTORY specifies the path to the data folder. This is the root directory where all JCMS document files are stored and it needs to be set before using this feature. Typically, the value of this variable is set to the directory where JCMS is installed, for example:

C:\Program Files\The Jackson Laboratory\JAX-CMS\data

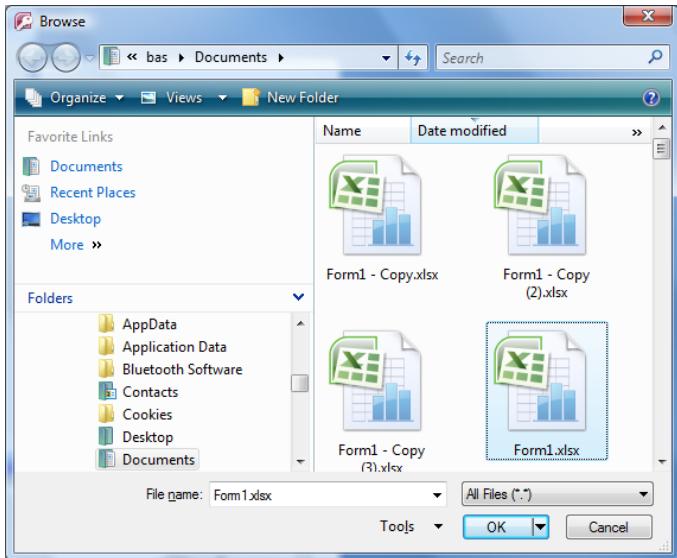
The subfolders, such as *genotype* and *Mouse* shown in Figure 17-1, will be created by JCMS when needed. All mouse-genotype documents are copied to the genotype folder, all documents associated with a mouse ID are copied to the Mouse folder, all documents associated with a sample ID are copied to the Sample folder, etc.

## 17.2 Adding (Uploading) Documents

A screenshot of a web-based application window titled 'Add Document (Upload Files)'.

- Category:** Radio buttons for 'Mouse' (selected), 'Mouse Genotype', 'Mating', 'Litter', 'Mouse Use', 'Sample', 'Experimental Plan', 'Test Type', and 'Experimental Data'.
- File Title / Description:** Text input field containing 'Example of a document'.
- File Name:** Text input field containing 'C:\Program Files\The Jackson Laboratory\JAX-CMS\docs\ReadMe\_files\image001.png' with a 'Browse...' button to its right.
- Owner:** A dropdown menu currently set to 'nobody'.
- Buttons:** 'Submit', 'Clear', and 'Close'.

**Figure 17-2 Add document form**



**Figure 17-3 File browser**

When a document is added (uploaded) to a JCMS document folder, the file will be copied to the folder, given a title/description and assigned an owner. A document may be added from the Manage Mice tab by using the *Add Document* button. Documents added this way will not be associated yet to an ID.

The *Browse...* button is used to open a file browser window for selecting the file to copy. The *View Document* button is used to open the file hyperlink prior to copying it. The file will not be copied into a JCMS data folder until the *Submit* button is clicked.

Title_Description	FileName	DateUpload	Owner
Example 1	Report1.xls	7/5/2013 2:25:56	nobody
Example 2	Form1.xlsx	7/5/2013 2:27:15	nobody

**Figure 17-4 Upload New button**

An alternative method for adding a document and associating it with an ID is to use the *... Upload New* button on the Add or Edit form for the ID. This button will open the Add Document form,

where one or more documents may be added. When the Add Document form is closed, those documents will be displayed on the Associated Documents list (example: Report1.xls and Form1.xls in Figure 17-4.) All documents in the list will be associated with the ID (mouse-genotype in this case) when it is added using the *Submit* button.

### 17.3 Selecting Documents to Associate

Once a document has been added to a document folder it may be associated to many different IDs by using the *Associate Existing* button on an add or edit form. This button opens the Associate Documents form. Several criteria may be used on this form to search for the appropriate document(s), including upload date, owner, and a text string in the file name or title/description. The *Search* button will fill the documents list with the matches. Select one or more documents in the list and click the *Associate* button. (If the search button does not work and MySQL is used for the database management system, make sure the setup variable JCMS\_DATABASE\_DBMS is set to MySQL.)

The screenshot shows the 'Associate Documents' dialog box. At the top, there are three search criteria groups: 'Select Documents by' (radio buttons for 'Any' or 'Title / Description or File Name like'), 'Date Uploaded' (radio buttons for 'Any' or 'Range', with dropdowns for dates), and 'Owner(s)' (radio buttons for 'Any' or 'Selected', with a list box showing 'nobody' and 'OWN1'). A 'Search' button is located at the top right. Below these is a table titled 'Documents' with columns: Title\_Description, FileName, Owner, and DateUpload. The table contains three rows: 'Example two' (FileName: Form2.xlsx, Owner: nobody, DateUpload: 6/6/2011 10:10:24 AM), 'Example Three' (FileName: Form3.xlsx, Owner: nobody, DateUpload: 6/6/2011 10:11:02 AM), and 'Example document' (FileName: Form1.xlsx, Owner: nobody, DateUpload: 6/6/2011 10:24:35 AM). To the right of the table are buttons for 'Select All' and 'Clear All'. At the bottom left are 'Associate', 'Clear', and 'Close' buttons. A status message 'Documents have been added to the list to associate.' is displayed at the bottom right.

**Figure 17-5 Associate Documents form**

This process may be repeated. When the Associate Documents form is closed, all documents that were selected will appear on the original add or edit form in the associated documents list.

The screenshot shows a portion of an edit form with a table titled 'Associated Documents' containing the same three rows as the previous screenshot: 'Example two', 'Example Three', and 'Example document'. To the right of the table are three buttons: '... Upload New', 'Associate Existing' (which is highlighted with a blue border), and 'Delete Association'. At the bottom left are 'Submit' and 'Clear' buttons.

**Figure 17-6 Associate Existing button**

The *Delete Association* button is used to remove documents from the list of associated documents. It does not delete the document.

Any changes to the associated documents list are not saved until the add or edit is submitted.

## 17.4 Edit Document

Use the *Edit Document* button on the Manage Mice tab to open this form. It is used to change the document owner or file title/description. A *Search* button is provided at the top of the form to use in selecting the proper document by a combination of date uploaded, owner, category, or a text string in the file name or title/description. From the list of documents, choose one to edit by highlighting it and clicking the *Select* button. The *Submit* button must be used to save any changes.

The screenshot shows the 'Edit Document Information' window. At the top, there are search filters for 'Select Documents by': 'Any' (radio button), 'Title / Description or File Name like' (radio button selected), and a text input field containing 'form'. There are also filters for 'Date Uploaded': 'Any' (radio button) and 'Range' (radio button selected), with dropdown menus for '≥' and '≤' set to '6/6/2011'. For 'Owner(s)', there are 'Any' (radio button) and 'Selected' (radio button selected), with a dropdown menu showing 'nobody' and 'OWN1'. For 'Category', there are 'Any' (radio button) and 'Selected' (radio button selected), with a dropdown menu showing 'Exp Data', 'Genotype' (selected), and 'Litter'. A 'Search' button is located in the top right corner. Below the filters is a table titled 'Documents' with columns: FilePath, Title\_Description, FileName, Owner, and DateUpload. The table contains three rows:

FilePath	Title_Description	FileName	Owner	DateUpload
\genotype	Example two	Form2.xlsx	nobody	6/6/2011 10:10
\genotype	Example Three	Form3.xlsx	nobody	6/6/2011 10:11
\genotype	Example document	Form1.xlsx	nobody	6/6/2011 10:24

A 'Select' button is located to the right of the table. Below the table, there is a section for editing a selected document: 'File Title / Description:' with a text input field containing 'Example Three - Change the description here', and 'Owner:' with a dropdown menu showing 'nobody'. At the bottom are 'Submit' and 'Clear' buttons.

**Figure 17-7 Edit Document form**

If the search button does not work and MySQL is used for the database management system, make sure the setup variable JCMS\_DATABASE\_DBMS is set to MySQL.

## 17.5 Bulk Add Document Associations

The bulk add documents form allows adding associations between multiple documents and multiple IDs at once. It is opened from the Manage Mice tab.

The screenshot shows the 'Bulk Add Document Associations' window. In the top left, there's a radio button group for 'Category': Mouse (selected), Mouse Genotype, Mating, Litter, Use, Sample, Experimental Plan, and Test Type. Below this is a table titled 'Associated Documents' with columns: Title\_Description, FileName, DateUpload, Owner, and a 'Double click to view' column. A 'Session Box' is on the right. In the 'Select IDs to Associate' section, there's a search bar for 'ID like' (containing 'test'), dropdowns for 'Gene' (FSN) and 'Alleles' (Allele-specific search), and a list of items: Test6, Test7, GenoTest. To the right is a grid table with columns: Mouse ID, Allele 1, Allele 2, Strain, Generation, Sex, Birth Date, and Mouse ID. Rows in the grid include Test6 (DBA/2J, F?, M, 2/18/2013), Test7 (DBA/2J, F02, F, 3/23/2013), and a bottom row with 'New 1'. Buttons at the bottom include 'Submit', 'Session Report', and 'Clear'.

**Figure 17-8 Bulk Add Document Associations form**

Documents may only be added to one category at a time. The select documents section works as described above for the add genotype form.

The select IDs to associate section will change depending on the selected category. It is used for entering criteria to fill the search list (bottom left of the form) when the **Search** button is clicked.

Select IDs to associate to the document(s) and use the > button to move them into the grid on the right side of the form. The red X button is used to remove the selected row from the grid. If no row is selected, the bottom row is removed.

Two categories, experimental plan and test type have drop down boxes that may be used to move the item directly into the grid. In the example below, selecting plan ID 2 has moved this plan into the grid.

The screenshot shows the 'Select IDs to Associate' section for 'Experimental Plan'. A dropdown menu is open, showing 'Experimental Plan' and '2 Clinical tests'. To the right is a grid table with columns: Experimental Plan ID, Experimental Plan Name, Plan Status, Field of Study, and Plan Owner. The single row in the grid is '2 Clinical tests' (active, clinical, OW/N1).

**Figure 17-9 Experimental plan category**

Once all the IDs have been moved into the grid, use the **Submit** button to create the document associations. The session box and report will list associations that have been created and a second list of those that already existed.

Note that if a document will be associated with items from different categories, it will need to be uploaded once for each category.

# 18 Queries

## 18.1 Colony Summary Report

Some useful queries within JCMS are pre-packaged into reports. The Colony Summary Report is one of these.

Colony Summary Report For 2/19/2010															
<b>Summary Statistics</b>															
Number of live* mice for all owners:	567														
Number of active strains for all owners:	16														
Number of active pens* for all owners:	294														
Number of active matings for all owners:	76														
Number of active plans for all owners:	5														
Number of active tests for all owners:	8														
Number of mouse tests scheduled for all owners:	47														
<small>*live mice have a life status that is not an exit status; active pens are those containing live mice</small>															
<b>Detail Reports</b>															
<b>Number of live mice by owner and strain</b>															
Owner: nobody															
<table border="1"><thead><tr><th>Strain name</th><th># live mice</th></tr></thead><tbody><tr><td>B6.129P2-Apo&lt;tm1Unc&gt;J</td><td>4</td></tr><tr><td>B6D2F 1/J</td><td>24</td></tr><tr><td>BALB/dByJ</td><td>13</td></tr><tr><td>BALB/cJ</td><td>102</td></tr><tr><td>C3H/HeJ</td><td>6</td></tr><tr><td>C57BL/6J</td><td>3</td></tr></tbody></table>		Strain name	# live mice	B6.129P2-Apo<tm1Unc>J	4	B6D2F 1/J	24	BALB/dByJ	13	BALB/cJ	102	C3H/HeJ	6	C57BL/6J	3
Strain name	# live mice														
B6.129P2-Apo<tm1Unc>J	4														
B6D2F 1/J	24														
BALB/dByJ	13														
BALB/cJ	102														
C3H/HeJ	6														
C57BL/6J	3														

**Figure 18-1 Colony Summary Report**

Note: If mice in a pen have different owners, the pen will be counted more than once in the summary, once for each owner. Therefore, the total number of pens will agree with the sum of the number of pens each owner has.

## 18.2 What are Queries used for? – or How to Search the Database

One of the most important functions of JCMS is to provide methods of searching for answers to specific questions about the data. The best way to obtain copies of the data entered into JCMS is by using one of the special query forms to set up a **search**. All searches have two parts, the question (criteria) and a description of the data to return (result fields to show).

To invoke it click on the **Colony Summary Report** button on the reports tab.

The Colony Summary Report function produces a report that summarizes the state of the colony in a number of important areas. The report presents the information at a global level and additionally breaks it down in a detailed view by owner.

The report displays:

- total number of live mice
- number of active strains
- number of active pens
- number of active matings
- number of active experimental test plans
- number of active experimental tests
- number of mouse tests scheduled

## 18.2.1 Basics on using the Query Forms

The screenshot shows the 'Query Mouse' interface. On the left, there are several search criteria groups:

- Mouse IDs:** Any, Range, Like, Selected from list. Sub-options include ST-019, ST-018, ST-017, ST-016, ST-015, ST-014.
- Strain or Stock/JR #:** Any, Selected strains / stock #'s from list. A dropdown shows strain names like B6.129P2-Apoe<tm1Unc>J, B6D2F1/J, BALB/cByJ, BALB/cJ, C3H/HeJ, C57BL/6J, CBA/J, DBA/2J, FVB/NJ, and Mixed.
- Experimental plans:** Any, In active plan, Not in active plan, In any plan, Not in any plan. Sub-options include Owner (Any, Selected owners from list), Origin (Any, Selected origins from list), and Mouse protocol ID (Any, Selected from list).
- Pen:** Any, Selected pen IDs from list, ID Range (Min: 29, Max: 23), Selected pen names from list.
- Generation:** Any, Selected litter #'s from list. Sub-options include Generation (F01, F02, F03, NU2, LNU2), Date of birth (Any, Range, ≥ Min value: 1, ≤ Max value: 10), and Exit date (Any, Range, ≥ Min value: 10/21/2011, ≤ Max value: 10/21/2011).
- Room:** Any, Selected from list. Sub-options include Health Level (Any, Selected from list, 1, 2, 3, 4).
- Sex:** Any, Male, Female, Unknown.
- Genotype date:** Mice typed on specified dates. Sub-options include Genotype date (Any, Range, ≥ Min value: 1, ≤ Max value: 10), Necropsy results (Any, Selected from list, Six month complete, Skin graft, Three month comp, Twelve month comp, Two year complete), and Mouse uses (Any, Selected from list, Necropsy results, Six month complete, Skin graft, Three month comp, Twelve month comp, Two year complete).
- Check this box if only selected genotypes are desired. Up to 10 may be selected.** Select AND/OR logic for GT constraints (AND, OR). Restrict output to show selected genotypes only, Show genotype details.

On the right side, there is a list of 'Check off the result fields to show' with checkboxes for various mouse attributes. Some checkboxes are checked by default, such as 'Mouse ID', 'Any', 'Selected from list', and 'Any' under 'Cause of death'.

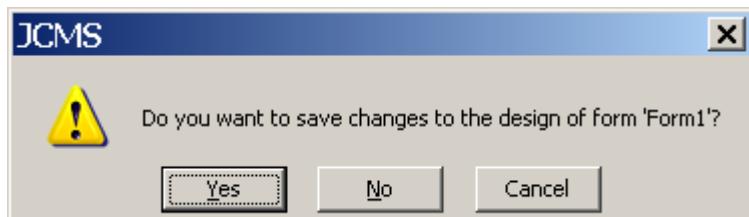
Figure 18-2 Mouse Query Form

The above sample query form shows the general layout of the query forms. Each form provides many choices for the **criteria**. The criteria specify how to decide the data to return. Criteria that are not used are set to **Any**. These possible criteria are ignored. The second part of a query is to indicate what **result fields** to return. The query forms all have a set of check boxes on the right side listing choices of result fields.

Form1 : Form						
	Mouse ID	Strain	Generation	DOB	Mouse Uses	Genotype
▶	1	ABC/xyzJ	N01	5/5/2005		
	10	ABC/xyzJ	F01	5/5/2005	abc	cpdm[+Y/+Y]
	11	ABC/xyzJ	F01	5/5/2005		
	12	ABC/xyzJ	F01	5/5/2005		
	16	ABC/xyzJ	N03	11/6/2005		
	17	ABC/xyzJ	F02	11/6/2005		
	18	ABC/xyzJ	F02	11/6/2005		
	19	ABC/xyzJ	F02	11/6/2005		
	2	ABC/xyzJ	N01	5/5/2005		
	9	ABC/xyzJ	F01	5/5/2005	abc, abc, testing	cpdm[+Y/+Y], fsn[−Y/−Y]

Figure 18-3 Datasheet: Mouse Query Results

The Query forms allow extraction of information from JCMS into a datasheet form by clicking the **Run Query** button. This output of a query looks like a spreadsheet with a column for each result field, but it cannot be edited. The example above shows the results of a query asking for a particular strain. The datasheet can be easily **exported into Microsoft Excel** by selecting on the menu bar Tools – Office Links – Analyze it with Microsoft Office Excel (Office 2003) or selecting on the ribbon External Data – Export – Excel (Office 2007).



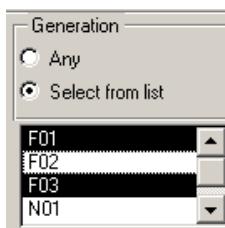
**Figure 18-4 Query: Do you want to save changes to Form?**

the option to save the query form that has been created. However, saving the form in JCMS will not preserve the query information between JCMS sessions. It is best to NEVER SAVE A QUERY FORM IN JCMS. Always answer **No** to this dialog box.

The contents of a query form in JCMS cannot be saved between sessions. To keep the information from a query, be sure to export it to Excel before leaving the JCMS session.

When the datasheet form is closed, a prompt appears with

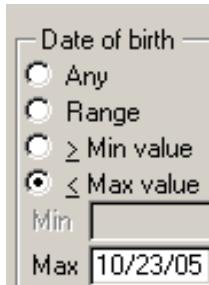
### 18.2.2 How to Select Query Criteria



**Figure 18-5**  
**Query Form: List Box Criteria**

For many of the criteria choices there is a list box and some push buttons. Push buttons allow specifying to select specific criteria items from the list box, or if you don't care about a particular criteria item, select the "Any" button. Select individual items in the list box by holding down the control key (ctrl) on the keyboard and, while holding down the key, use the mouse to select individual items by clicking on them. A range of items may be selected from the list box by holding down the shift key while using the mouse to select the first and last in the range. If the ctrl or shift key is not held down, selecting an item from the list box clears all other selections.

The example criteria are: show all mice with generation equal to F01 or F03.



**Figure 18-6**  
**Query Form:**  
**Range Criteria**

Another push button option is "Range." Using this option allows specifying a minimum and maximum value or range of values for the query. Only mice between the minimum and maximum range, including the minimum and maximum values, will be included in the query results.

The example criterion is: show all mice with a date of birth less than or equal to 10/23/2005.

Another type of criteria choice provides multiple check boxes. If the "Any" button is selected, no check boxes are available (they are all gray and cannot be clicked on). Click the any button to unselect it to make the check boxes white (available to choose). Multiple check boxes may be chosen or unselected by clicking on them with the mouse.



**Figure 18-7 Query**  
**Form: Check Box**  
**Criteria**

The example criteria are: show all mice that are breeders OR virgins.

When more than one criterion is set to something other than "Any", the criteria is put together using "AND" Boolean logic. If the three examples above were used the criteria becomes: show all mice with (generation equal to F01 OR F03) AND (a date of birth less than or equal to 10/23/2005) AND (are breeders OR virgins).

Click the **Run Query** button to see the results. The Clear all and Select all buttons will only clear the choices of result fields. The criteria choices must all be cleared individually by clicking the any buttons. Output result fields must be specified for the query. If no fields are selected, an error message will be displayed.

### 18.2.3 Like Criteria for Mouse ID

**Figure 18-8 Query Form: Like Criteria**

The Like criteria will return all mouse IDs that contain the string entered in the text box. In the example shown in the figure, only M0008-A0032 will be returned. Entering "00" would return all of the mouse IDs shown in the list.

JCMS has converted the string entered in the Like box to \*A\*. Otherwise the criteria of A would return only one mouse whose ID was exactly A. If a leading or ending \* is entered by the user, no \* will be added by JCMS. This allows entering \*A to return all mouse IDs ending with A and A\* to return all mouse IDs beginning with A. Standard wildcards as described below will work in the search. The information below is from Microsoft Access Help.

Built-in pattern matching provides a versatile tool for making string comparisons. The following table shows the wildcard characters you can use with the **Like** operator and the number of digits or strings they match.

Character(s) in pattern	Matches in expression
? or _ (underscore)	Any single character
* or %	Zero or more characters
#	Any single digit (0— 9)
[charlist]	Any single character in charlist
[!charlist]	Any single character not in charlist

You can use a group of one or more characters (*charlist*) enclosed in brackets ([ ]) to match any single character in *expression*, and *charlist* can include almost any characters in the ANSI character set, including digits. You can use the special characters opening bracket ([ ), question mark (?), number sign (#), and asterisk (\*) to match themselves directly only if enclosed in brackets. You cannot use the closing bracket ( ]) within a group to match itself, but you can use it outside a group as an individual character.

In addition to a simple list of characters enclosed in brackets, *charlist* can specify a range of characters by using a hyphen (-) to separate the upper and lower bounds of the range. For example, using [A-Z] in *pattern* results in a match if the corresponding character position in *expression* contains any of the uppercase letters in the range A through Z. You can include multiple ranges within the brackets without delimiting the ranges. For example, [a-zA-Z0-9] matches any alphanumeric character.

### 18.2.4 Mouse Age

**Figure 18-9 Mouse age result choice**

The mouse age may be returned as days, weeks, or months depending on the selection. Two columns are returned in the results, one giving the mouse's age today for non-exited mice and the second giving the mouse's age on the exit date.

## 18.2.5 Filtering Strains or Stock #'s

Run Query		Add Mice to Experiment	
Strain or Stock/JR #		Stock # filter:	
<input type="radio"/> Any	<input checked="" type="radio"/> Selected strains / stock #'s from list	660	to 2000
stockNum	strainName		
564	C57BL/6J		
671	DBA/2J		
1026	BALB/cByJ		
1303	NOD.CB17-Prkdc<scid>/J		
1800	FVB/NJ		
1976	NOD/ShiLtJ		

Figure 18-10 Strain or Stock # Filter

A filter is available for the strain name/stock # choices. When the setup variable JCMS\_STRAINNAME\_FIRST is set to false, a stock # filter is displayed. A range of numbers may be entered. If only one number is entered, the list will be displayed starting

with it as the minimum stock # or ending with it as the maximum stock number. The strains are displayed in stock number order. When the setup variable is set to true, the filter works similar to the like criteria described above for the mouse ID. The strain names will be displayed in alphabetical order.

## 18.3 Mouse Query

The [basics](#) of using this query form are described above. Below is an explanation of how to use the genotype, documents, and mouse use portions of the mouse query form.

### 18.3.1 Query by Genotype (QGT)

<input checked="" type="checkbox"/> Check this box if only selected genotypes are desired. Up to 10 may be selected.																															
Select AND/OR logic for GT constraints																															
<input type="radio"/> AND	<input checked="" type="radio"/> OR																														
<input checked="" type="checkbox"/> Restrict output to show selected genotypes only <input type="checkbox"/> Show genotype details																															
<table border="1"> <tr> <td>fsn</td> <td>cpdm</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>?</td> <td>?</td> <td>-</td> <td>-</td> <td></td> <td></td> </tr> <tr> <td>+</td> <td>+</td> <td>+</td> <td>+</td> <td></td> <td></td> </tr> <tr> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> </tr> </table>		fsn	cpdm					?	?	-	-			+	+	+	+														
fsn	cpdm																														
?	?	-	-																												
+	+	+	+																												
Genotype date Mice typed on specified dates Genotype date <input type="radio"/> Any <input checked="" type="radio"/> Range <input type="radio"/> ≥ Min value <input type="radio"/> ≤ Max value Min 1/1/05 Max 12/12/05																															

Figure 18-11 Query Form: Genotype

As part of the mouse query, it is possible to query for mice by genotype. That is, select only those mice with a particular set of genotypes (up to 10 may be specified). To use the QGT option, first select the check box near the bottom of the form that enables this feature.

It is important to understand that QGT constraints have two levels: the Gene level, and the allele level. For each Gene, it is possible (but not required) to select one or two alleles that must be matched in order for the query to return results.

There is a choice of using AND logic or OR logic (use the push button just above the QGT selection boxes to choose a logic type) when combining constraints at the Gene level. But at the allele level AND logic is always used.

For example, suppose there are three genes X, Y, and Z. And for each of these genes there are allele possibilities x1, x2, y1, y2, y3, and z1, z2 where the letter "x" associates with gene X etc.

Using OR logic you could select all mice with genotype X(x1,x2) OR Y(-,-). The “-“ means you don’t care what alleles are associated with the gene. This selection would return all mice that have the specific genotype of X(x1,x2) and all mice that have been genotyped for gene Y. There can be overlap in the sets since a mouse could have genotype X(x1,x2) and also Y(y3,y2).

Using AND logic you could select for mice with genotype X(x1,x2) AND Y(-,-). Only mice that have both the specific genotype X(x1,x2) and have also been genotyped for gene Y will be found.

Mice shown may also be restricted by the date they were genotyped. The genotype date selection criteria will further limit the mice that are found to only those mice genotyped on a specific date or date range.

### 18.3.2 Interpreting the Genotype Output

Any mouse may have zero or more genotypes. Each genotype is reported in the following format:

labSymbol[allele1-conf/allele2-conf]

Example: bax[1-Y/0-N].

The allele confidence is reported as “Y” for “yes,” we have confidence or “N” for “no,” we do not have confidence. The use of [] as gene separators and Y or N for gene confidence may be customized using the setup variables JCMS\_ALLELE\_GENE\_SEPARATORS, JCMS\_ALLELE\_CONF\_HIGH, and JCMS\_ALLELE\_CONF\_LOW.

### 18.3.3 Restricting Genotypes in the Query Output

fsn	cpdm	More GTs
	cpdm[+Y/+Y] (pg=None) (sl=shelf 22) (dt=11/7/2005)	No More
	cpdm[+Y/+Y] (pg=None) (sl=shelf 22) (dt=11/7/2005)	No More
	cpdm[+Y/+Y] (pg=None) (sl=shelf 22) (dt=11/7/2005)	No More
fsn[--Y/-Y] (pg=None) (sl=shelf 55) (dt=11/7/2005)	cpdm[+Y/+Y] (pg=None) (sl=shelf 22) (dt=11/7/2005)	No More

**Figure 18-12 Query Datasheet: Restricted Genotype Output**

If the box for restricted genotype output is checked, only the genotypes for the genes selected in the genotype constraints will be output. When this option is used, each genotype is reported in a separate column in the output form. Genotype details, view the sample location, and page number for the sample may also be requested. Since a mouse may have been typed for more genes than those selected, a column is added to the output form that indicates if there are “More” genotypes for this mouse or “No More” genotypes for this mouse in the database.

Below is an example of a genotype with the genotype details; **pg** is page number and **sl** is sample location. **sl=NONE** implies that there is no information about sample location in the database. Also **dt** is the genotype date field. In this case the date was not stored in JCMS so it is listed a no date.

Sod1[1-Y/1-Y] (pg=g7-84) (sl=NONE) (dt=no date)

### 18.3.4 Listing Documents Associated with Genotypes

A screenshot of a software interface showing a list of checkboxes for selecting document types. The checkboxes are labeled: 'Genotype' (checked), 'Associated documents, Max of 1' (checked), 'Vial ID' (unchecked), and 'Vial position' (unchecked). The 'Associated documents' checkbox has a tooltip indicating it selects up to 10 columns of documents.

Documents associated with mouse-genotype combinations may be requested in the output only when genotype is also requested. Up to 10 columns of documents associated with a genotype may be requested. An extra column is added that will indicate if there are more

**Figure 18-13 Requesting documents in the query output**

documents than columns. For a search that lists multiple genotypes in one output column it is not possible to recognize that the same document is associated with several of those genotypes, possibly causing the document to be listed in more than one column.

### 18.3.5 Listing Other Associated Documents

Documents associated with a mouse ID, litter #, mouse use, or mating may also be requested, up to a maximum of 10 per category. In order to list documents, the associated category must be checked. An extra “More Documents?” column is added that will indicate if there are documents that were not listed for a category. The full path to the document is returned as an active hyperlink, which may be clicked to open the document file.

### 18.3.6 Query by Mouse Use

As part of the Mouse Query, it is possible to set up criteria for mouse uses. If multiple uses are selected the results will include all mice that have one of the selected uses (i.e. the uses are ORed together in the query). If the check box to restrict output to only the selected uses is checked, JCMS will put each selected use in a separate column (instead of putting all of the uses in one comma delimited list). Use details (such as comments) can also be shown this way. If a mouse has more uses than those displayed, the *more uses* column will have the word *more* in it. To be sure to see all uses for all mice selected in the query, the restricted output checkbox must be unchecked. Mouse uses must be checked in the results fields in order for the “restrict output to show selected uses only” check box to be enabled (see green circle). The “show use details” checkbox will not be enabled unless the “restrict output to show selected uses” box is checked.

The output can also be restricted to show only mice that have selected uses that are *not done* or *are done*. For example, this feature can be used to show all mice that have not yet been tested for a specific use.

The use details use the following codes: UA for use age, PD for proposed date, comments, and DATA: D1=, D2=, etc for the data field values.

User note: use the Mouse-Use Work Report button on the Reports tab to

**Figure 18-14 Query Form: Mouse Use** generate a report of all mice that are scheduled for use.

Form1					
Mouse -	MouseUse-Skin graft	Documents: Skin graft1	More Uses	More Documents?	
A7	Skin graft[ UA-2.5 (not done) PD-8/15/2011](DATA: )	C:\Copy of zip\JCMS\DocTest\MouseUsage\Document5.docx	No More		
A9	Skin graft[ UA-2.5 (not done) PD-8/15/2011](DATA: )	C:\Copy of zip\JCMS\DocTest\MouseUsage\Document5.docx	No More		
ST-014	Skin graft[ UA-3 (not done) PD-10/30/2011](DATA: )		No More		

**Figure 18-15 Query Datasheet: Mouse Uses**

#### 18.3.6.1 Mouse query failure created by certain special characters in a mouse use type term

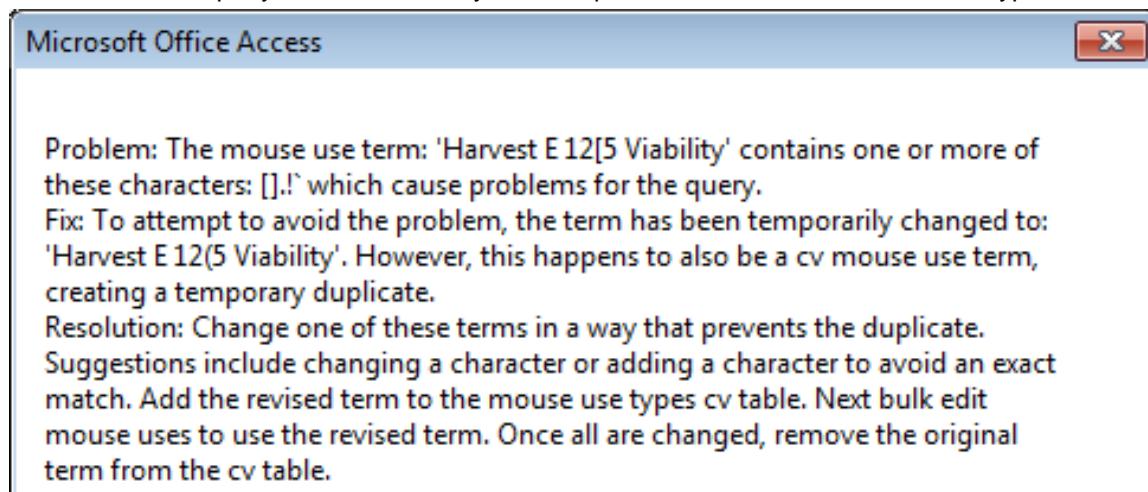


Figure 18-16 Mouse query - use term error message

When a mouse use term contains one or more of these special characters: period “.”, left bracket “[”, right bracket “]”, exclamation point “!”, or accent grave “`”, JCMS replaces them in the column header of the output. No changes are made to the data. The message above outlines a rare situation where the character substitution will not work.

## 18.4 Mating Query

The screenshot shows the 'Query Matings' form with several sections for filtering data:

- Mating ID:** Set to 'Any'. Options include 'Any', 'Range' (Min: 60, Max: 9), and 'Selected matings from list'.
- Litter strain:** Set to 'Any'. Options include 'Any' and 'Selected strains or stock/JR #'s from list'. A dropdown shows strain names like B6.129P2-ApoecJU, BALB/cByJ, C3H/HeJ, C57BL/6J, CBA/J, DBA/2J, FVB/NJ, etc., with a 'jrNum' column.
- Litter generation:** Set to 'Any'. Options include 'Any' and 'Selected generations from list'. A dropdown shows F01 through N07.
- Mating owner:** Set to 'Any'. Options include 'Any' and 'Select from list'. A dropdown shows 'nobody' and 'OWN1'.
- Pens:** Set to 'Any'. Options include 'Any', 'Range' (Min: 104, Max: 99), and 'Select Pen IDs from List'. A dropdown shows cage numbers like Cage-00001 through Cage-00005.
- Check off the mating results to show:** A large list of checkboxes for various mating and litter details, such as 'Mating ID', 'Associated documents, Max of [1]', 'Mating type', 'Pen Info', 'Dam 1 ID', 'Dam 1 strain', 'Dam 1 stock#', 'Dam 1 generation', 'Dam 1 up to three genotypes', 'Dam 1 date of birth', 'Dam 1 plug dates', 'Dam 2 ID', 'Dam 2 strain', 'Dam 2 stock#', 'Dam 2 generation', 'Dam 2 up to three genotypes', 'Dam 2 date of birth', 'Dam 2 plug dates', 'Sire ID', 'Sire strain', 'Sire stock#', 'Sire generation', 'Sire up to three genotypes', 'Sire date of birth', 'Mating date', 'Date retired', 'Litter strain', 'Litter stock#', 'Litter generation', 'Wean time', 'Wean note', 'Needs typing', 'Mating comments', and 'Mating owner'.
- Show summary or detailed litter information:** Options include 'Summarize litter information' (selected) and 'Show litter details on [ ] litters'.
- Check off the litter results to show:** A list of checkboxes for litter details like 'Total number of litters', 'Total number harvested', 'Litter # sort @ ascending / most recent', 'Associated documents, Max of [1]', 'Number born', 'Wearing date', 'Total number of males weaned', 'Total number of females weaned', 'Total number of litters born dead', 'Total number of females harvested', 'Date of birth', 'Tagging date', '# Females', 'Litter status', '# Males', 'DPC', 'First and last birth dates', 'Number harvested', 'Harvest date', 'Litter comments', and 'Litter Type'.
- Buttons:** 'Run Query', 'Clear All', 'Select All'.

Figure 18-17 Query Mating Form

The [basics](#) of using this query form are described above. The query is limited to showing only matings from one of the four mating types at a time: Natural (N), Embryo Transfer (ET), Ovary Transfer (ET), or In vitro Fertilization (IVF).

The screenshot shows the 'Query Matings' form with the 'Mating type' dropdown set to 'OT'.

Figure 18-18 Mating type selection

The results section column on the right will change so that only the pertinent result choices may be selected.

In addition to the normal results field column on the right of the form, the mating query has a special results section at the bottom of the form for checking off the litter results to show. Choose either summary litter information or detailed litter information.

The screenshot shows a 'Form1 : Form' table with columns for matingID, Dam 1 ID, Dam 2 ID, Sire ID, Mating Date, litter ID1, DOB1, # Female1, # Male1, Wean Date1, litter ID2, and DOB2. The data includes rows for entries 1, 5, 6, and 7.

matingID	Dam 1 ID	Dam 2 ID	Sire ID	Mating Date	litter ID1	DOB1	# Female1	# Male1	Wean Date1	litter ID2	DOB2
1	12	10	6	11/6/2005	41	11/6/2005	3	3	11/24/2005		
5	9	8	6	9/29/2005	51	11/6/2005			12:00:00 AM		
6	1		4	11/6/2005	61	11/6/2005			12:00:00 AM		
7	18	19	22	9/29/2005							

Figure 18-19 Query Datasheet: Mating

## 18.5 Experimental Data Query

The [basics](#) of using this query form are described above.

The screenshot shows the 'Query Experiments' interface with the following sections:

- Search Criteria:** Includes 'Any Data' (selected), 'Mice in Experimental Plan(s)' (selected), 'With results' (selected), and a 'Run Query' button.
- Experimental Plans:** 'Any' (selected) or 'Selected from list'.
- Test Types:** 'Any' (selected), 'One from list', or 'Multiple from list'. A dropdown menu shows 'Example'.
- Keywords:** 'Any' (selected) or 'Selected from list'.
- Mouse ID:** 'Any' (selected) or 'Range'. Input fields for Min and Max values are present.
- Mouse owner:** 'Any' (selected) or 'Selected from list'. A dropdown menu shows 'nobody', 'OWN1', 'TestWG1', and 'USR1'.
- Strain:** 'Any' (selected) or 'Selected from list'. A dropdown menu lists strain numbers and names, including:
 

stockNum	strainName
651	BALB/cJ
656	CBA/J
659	C3H/HaJ
664	C57BL/6J
671	DBA/2J
1026	BALB/cByJ
1303	NOD.CB17-Prkdc<scid>/J
1800	FVB/NJ
1976	NOD/ShiLtJ
2052	B6.129P2-Apo<tm1Unc>/J
100006	B6D2F1/J
- Experimental Tests:** 'Any' (selected), 'One from list', or 'Multiple from list'. A dropdown menu shows 'testName' with items 'Example 1', 'Proposed for Dec 1', 'Proposed for Jan 1', and 'Proposed for Feb 1' (the last one is highlighted).
- Keywords:** 'Any' (selected) or 'Selected from list'.
- Proposed test date:** 'Any' (selected), 'Range', '≥ Min value' (2/26/2014), '≤ Max value' (2/26/2014).
- Data collection date:** 'Any' (selected), 'Range', '≥ Min value' (2/26/2014), '≤ Max value' (2/26/2014).
- Age on data collection date:** 'Expressed in' (Days selected), 'Any' (selected), 'Range', '≥ Min. value' (2/26/2014), '≤ Max. value' (2/26/2014).
- Abnormal data:** 'Any' (selected), 'Abnormal only', 'Normal only'.
- Data owner:** 'Any' (selected) or 'Selected from list'. A dropdown menu shows 'nobody', 'OWN1', 'TestWG1', and 'USR1'.
- Output fields:** A large list of checkboxes grouped under 'All Mouse' (selected), 'All Experimental Plan' (selected), and 'All Experimental Data' (selected). Examples include 'Mouse ID', 'Strain', 'Stock/JR #', 'Generation', 'Birth Date', etc.

**Figure 18-20 Query Experiment Form**

This form offers three choices when selecting the criteria:

1. Any data. Results will be returned for all mice that have experimental data records associated with them. This can result in a mixture of results that include data from experimental plans and data records that are not part of experimental plans.
2. Mice in experimental plan(s) with results. The results will only include mice that have experimental data records AND are part of an experimental test. The rest of the criteria should specify the experimental test(s) and other choices. The possible choices have been limited to eliminate those that would cause a conflict. For example: an experimental test is always part of only one plan, therefore it is not possible to choose a plan name as this plan might not include the test that was chosen, a situation that would yield no results. If experimental test is left set to "any" then other criteria such as a range of mouse IDs or test status may be specified. This will yield all mice with data records that fit the criteria AND which are part of some experimental test.

3. Mice in experimental plan(s) not tested. This will give results for all mice that have been pre-selected for the experimental test, but have NO DATA RECORD for that test. No criteria except the one test may be specified. The output result field choices will not include the "All Experimental Data" section.

When only one test type is specified in the criteria, it is possible to choose output results using a data caption list instead of the generic D1, D2, D3, etc. list. To see the data result captions as a column header make sure to use the "Data Fields" option. If the query criteria options create a mixture of test types, the results may have different types of values in the D1, D2, D3, etc. list. In this situation, check "show data captions in column" so the caption will appear in the results in a column next to the data value.

### **18.5.1 Documents Associated with Experimental Data**

An experimental plan and test type may have documents associated with them. These documents may be requested as output fields, up to a maximum of 10 per experimental plan or test type. In order to list documents, the associated category (plan ID or test type) must be checked. An extra "More Documents?" column is added that will indicate if there are documents that were not listed for a category. The full path to the document is returned as an active hyperlink, which may be clicked to open the document file.

A test type may have a result field defined with the format of "file" indicating that the information stored in that field is a document. All data results are displayed formatted with the standard blue underline used for a hyperlink. However, only actual file paths will act as a hyperlink. If the color is annoying, the underline and color of a column in the datasheet may be changed by using conditional formatting. In Excel they may be changed by changing the cell formatting.

Mou:	Data	Plan	Documents:	More Documents?	WBC	Excel output
ST-002	42	2	C:\Copy of zip\JCMS\DocTest\ExpPlan\Test word doc for ExpData.docx	More Exp Plan	10	C:\Copy of zip\JCMS\DocTest\ExpData\ST-MiceBloodValues.xlsx

**Figure 18-21 Experiment Query Datasheet**

In the figure above, one document associated with the experimental plan is shown. The "More Documents?" column indicates this plan has other documents associated with it. Two data results columns are shown. "WBC" and "Excel output" are the captions for these data results.

### **18.6 Sample Query**

This query is described in the Samples section 20.6.

### **18.7 Microsoft Query**

Microsoft Query (MS Query) may be used to select data from an "outside source" such as JCMS and bring it into MS Excel. In this case, MS Excel must be set up to ask for data from a Microsoft Access database named JCMS.mdb. See Microsoft's documentation for information on using MS Query.

## 19 Experimental Plans

JCMS provides a method for defining experiments conducted using the mice within the database, setting up definitions for the various experimental tests, data, and data defaults, and recording experimental results. It can also be used to setup, track, and schedule mice for use in the experiments.

The following tables are used:

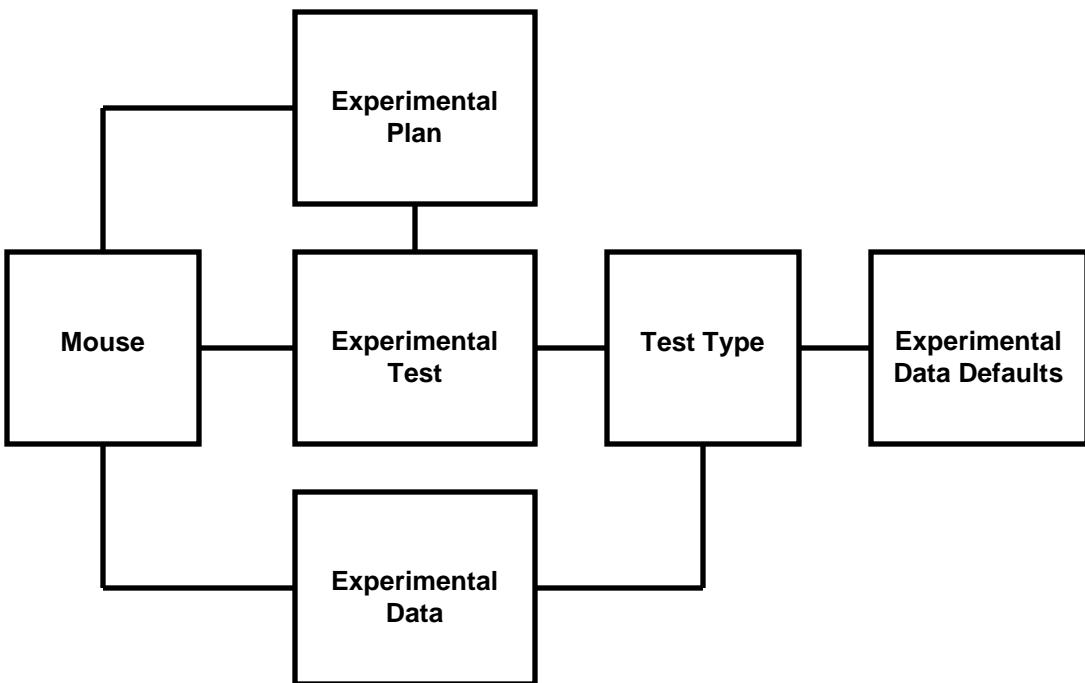


Figure 19-1 Diagram: Experimental Plan Tables

**Experimental Data:** One experimental data record consists of a set of data results for a mouse. A mouse may have many experimental data records, each one for a different test. Each experimental data record has to have an associated test type record that defines the format of the data results. Usually, each data record is associated with a test within a plan. However, it is possible to create data records that are not associated with a plan/test.

**Experimental Plan:** A definition of one experiment or project, referred to as the plan. The plan is used to coordinate the tests, mice planned for use in a test, and results. Each plan may have many mice scheduled to be part of it. Conversely, a mouse may be scheduled into many plans.

**Experimental Test:** One plan usually consists of several different tests (procedures). Some tests will be repeated multiple times perhaps using different sets of mice or with changes in protocol. JCMS considers each repetition a separate experimental test. Each test may have many mice scheduled to be part of it. Conversely, a mouse may be scheduled into many tests. A test must have one plan associated with it and one test type. The test type defines the format of the data results for this test. A test may have many experimental data records (results) associated with it, one for each mouse scheduled for the test.

**Test Type:** To make the repetition of tests easier, JCMS allows the user to define each type of experimental test used. This test type and data description includes a specific definition of the data collected. Up to 30 different result fields may be collected per mouse as part of one test.

Each result field may have specified a meaningful caption, maximum value, minimum value, format (date, numeric, text, or file), and whether or not it is required.

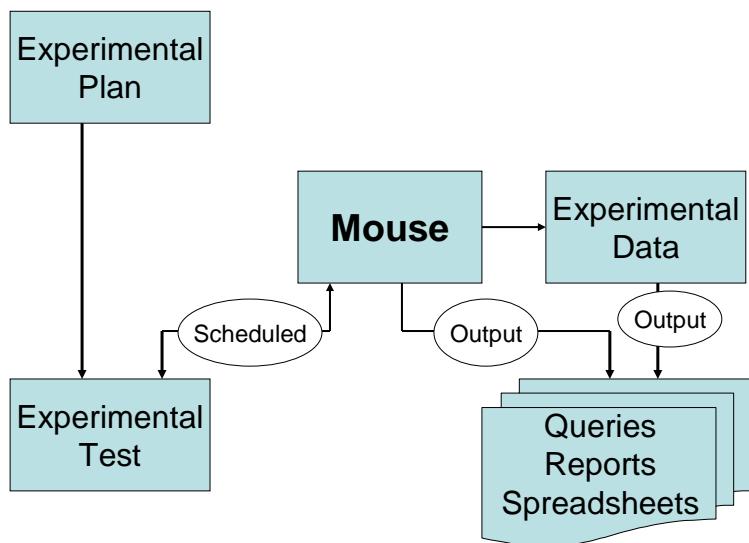
**Experimental Data Defaults:** Sets of default values for particular test types may optionally be defined to help with data entry of the experimental data results. Data defaults are associated with only one test type.

Experimental data may be recorded without an association to a plan and test. However, all experimental data must be associated with an experimental test type in order to define the data result fields.

An experimental plan may have mice pre-selected for it for planning purposes. An experimental test may also have pre-selected for it a subset of the mice selected for the plan. One mouse may be selected for multiple tests within a plan and may also be selected for other plans.

## 19.1 How to use an Experimental Plan

# Experimental Plan



**Figure 19-2 Diagram: Experimental Plan**

The experimental plan flow shown above is what the user will normally work with. The plan and tests will be defined. Mice will be scheduled for the various tests. Experimental data is collected and entered for the mice. Finally, reports and spreadsheets are output based on the mice and data collected.

Behind the scenes, some things have to be defined to set up the flow above. Each step is defined in detail below. The general plan definition and execution process includes:

- Define the **test types** that will be used for the tests in the plan. Once a test type is created it may be used over and over again within this and other plans. The test type defines the data results fields to be collected.
- Define the **data default** values for the test type. These values are sets of expected or standard results for some or all of the data result fields. They are used to help with data entry. These may be created at any time during the life of a plan. There may be several choices of data defaults created for a test type.
- Create the **experimental plan**.

- Create **experimental tests** for this experimental plan using the pre-defined test types. Each experimental test includes information about the proposed date, projected number of mice and ages of the mice. A specific test type may be used repetitively for different groups of mice, different dates, and/or different ages.
- **Schedule** mice into the plan and various tests. Scheduling mice is optional, if mice are not pre-scheduled, they will be automatically entered into the plan and test when the data for the mouse is entered.
- Print the **experiment work report**. This report is used to list what tests are scheduled for a particular time period.
- Enter the **experimental data** results.
- Use the **experiment query** to export results into MS Excel, determine mice that have not had the data results entered yet, and answer any other questions about the status of experimental plans, tests, or data.

## **19.2 How to Create Experimental Data without using an Experimental Plan**

It is possible to enter experimental data for mice into JCMS without going through the process of creating experimental plans and tests or scheduling mice. This process includes:

- Define the **test types** that will be used for the data results. Once a test type is created it may be used over and over again. The test type defines the data results fields to be collected. Pre-defined test types for a plan may also be used to create experimental data records outside of a plan. However, it is not possible to use default values for the experimental data without having an experimental test record.
- Enter the **experimental data** results.
- Use the **experiment query** to export data results into MS Excel and answer questions about the experimental data.

### 19.3 Setting up Test Types (Data Descriptions)

The test type and data description must be created before experimental tests or experimental data results can be added to an experimental plan. Even if an experimental plan is not used, the test type must be created before experimental data results may be entered. Once the test type is created, it may be reused for this or other experiments and data results.

Test Type	Blood values	Notes: Sample for documentation			
Caption	Field description	Format	Required	Min value	Max value
D1	Blood Glucose	dec	<input checked="" type="checkbox"/>	70	100
D2	Body Weight	dec	<input checked="" type="checkbox"/>		
D3	Comments	text	<input type="checkbox"/>		
D4	CBC	dec	<input type="checkbox"/>		
D5	WBC	dec	<input type="checkbox"/>		
D6	RBC	dec	<input type="checkbox"/>		
D7		text	<input type="checkbox"/>		
D8		text	<input type="checkbox"/>		
D9		text	<input type="checkbox"/>		
D10		text	<input type="checkbox"/>		
D11		text	<input type="checkbox"/>		
D12		text	<input type="checkbox"/>		
D13		text	<input type="checkbox"/>		
D14		text	<input type="checkbox"/>		
D15		text	<input type="checkbox"/>		
D16		text	<input type="checkbox"/>		
D17		text	<input type="checkbox"/>		
D18		text	<input type="checkbox"/>		
D19		text	<input type="checkbox"/>		
D20		text	<input type="checkbox"/>		
D21		text	<input type="checkbox"/>		
D22		text	<input type="checkbox"/>		
D23		text	<input type="checkbox"/>		
D24		text	<input type="checkbox"/>		
D25		text	<input type="checkbox"/>		
D26		text	<input type="checkbox"/>		
D27		text	<input type="checkbox"/>		
D28		text	<input type="checkbox"/>		
D29		text	<input type="checkbox"/>		
D30		text	<input type="checkbox"/>		

Double click to repeat data description

test type	D1	D2	D3	D4
Blood Work and Body Weights	Blood Glucose	Body Weight	Comments	CBC
Body Weights	WEIGHT			

Session Box

Submit    Clear    Session Report    Add default data values

**Figure 19-3 Form: Add Test Type**

Click the **Add Test Type – Data Description** button to open the form. This form may also be opened by clicking a button on the Add an experimental test form. Only owners or an Administrator may create test types.

Use a brief but meaningful name for the test type, up to 32 characters long. This name must be unique (not used by any other test type). Use the notes field to enter a description of the test, protocols, etc. The description may be as long as needed. Up to 30 data result fields may be defined. These are labeled D1, D2, D3, etc. on the form. The results will later be entered into an experimental data record using this description to validate them. For each data result field, enter the following:

- Caption: Used on the forms as the “name” of the data to enter. Use a short but meaningful caption, up to 32 characters long.
- Field description: An optional short description of the result field. The description might indicate the units such as cm, gm, etc. or the expected text entries such as yes/no or mutant/control. It may be up to 32 characters long.
- Format: Select text, date, file, integer, or decimal value. The format will be used during experimental data entry to check for errors. For example, when the data results are entered, if a numeric format is chosen no text or special characters will be allowed except for the normal ones associated with numbers (- +, .). The “file” format is used to associate this data result field with a document (computer file such as an image, spreadsheet, word document, etc.)

- Required: Indicates if the field is required or not during data entry. By default, fields are not required.
- Min value: Optional field. When experimental data is entered, numeric data may not be less than this minimum value.
- Max value: Optional field. When experimental data is entered, numeric data may not be greater than this maximum value.

A list box at the bottom of the form shows test types that have already been defined. If the test type to add is similar to an existing one, double click it in the list box. The values for that test type will be repeated on the form. These can then be changed and submitted for the new test type.

After successfully submitting a new test type, click the “Add default data values” button to set up any defaults that would help with experimental data entry.

## 19.4 Editing a Test Type

Click the **Edit Test Type** button to open the form. This form looks and works in the same manner as the Add test type form. New data result fields may be added to an existing test type and changes to fields may be made only if no experimental data or test type defaults exist for this test type. A test type may be deleted only if there is no experimental data in the database for this test type, no experimental tests that are using this test type, and no default data records for it.

## 19.5 Setting up Default Data for Experiments

**Figure 19-4 Form: Add Test Type Defaults**

Click the **Add Test Type Defaults** button to open the form. The Test Type defines the different data fields used for an experimental test. The test type data default values define a set of standard results that might be expected. Several data default records can be created for one test type. Each will contain one set of possible default results. There does not need to be a default value for each data value field. One of these sets of standard results (data defaults) can be selected at the creation of an experimental test. These defaults are then used when experimental data is entered to speed up the data entry process. The values that are specified have to fit within any minimum, maximum, or format that is defined for the data field. No data field is required to have a default value.

When experimental data is entered, the data defaults are displayed on the data entry form, where they may be edited to change any values that differ from the defaults.

Only the plan owner or the Administrator may create the data defaults.

## 19.6 Editing Default Data

Click the **Edit Test Type Defaults** button to open the form. Only the Administrator may edit or delete a test type data default record. Any edit changes made to a data default will not automatically change any experimental test that is already using this data default. That experimental test will still contain the old defaults. To change these in an existing experimental test, edit that experimental test and select the defaults again from the list of possible data defaults.

## 19.7 Adding an Experimental Plan

The screenshot shows a Windows-style dialog box titled "Add Experimental Plan (mtsadmin) {Owners Only}". The main title is "Add an Experimental Plan". The fields include:

- Plan ID: 3
- \*Owner: nobody
- \*Exp plan name: Sample plan for documentation
- Program and/or Field of Study: Dermatology
- \*Plan status: active
- A plan may have up to five keywords associated with it:
  - Keyword 1: Skin
  - Keyword 2
  - Keyword 3
  - Keyword 4
  - Keyword 5
- Notes:

This is for documentation only.
- Buttons at the bottom: Submit, Clear, Add tests to plan. A message "Plan submitted successfully." is displayed in red text next to the "Submit" button.

**Figure 19-5 Form: Add Experimental Plan**

Click the **Add Exp Plan** button on the experiments tab to open the form. The plan ID number will be assigned by JCMS. All experimental plans must have an owner. All experimental data generated for this plan will be assigned this same owner. All plans must be given a name and status. The program and/or field of study and keyword fields are provided so the user may later query for plans using this information. The Administrator sets up the choices for these fields.

## 19.8 Adding an Experimental Test

The screenshot shows the 'Add Experimental Test' form. At the top, there are dropdown menus for 'Default data values' and 'Caption / Format / Minimum / Maximum value'. Below these are fields for 'Plan ID' (set to 3), 'Name' (Sample plan for documentation), 'Test ID' (Documentation test), 'Test Type' (set to 'Blood values'), 'Proposed date' (set to 9/1/2008), 'Age entered as' (radio buttons for Days, Weeks, Months, with Days selected), 'Projected # mice' (set to 25), and 'Test status' (set to active). To the right, there are several data value fields with their captions and default values: 'Blood Glucose / dec / 70 / 100' (value 80.5), 'Body Weight / dec / /', 'Comments / text / /', 'Sample' (highlighted in green), 'CBC / dec / /', 'WBC / dec / /', and 'RBC / dec / /'. A 'Repeat Test' section displays a previous entry: 'BW test cycle 1' (Type: Body Weights, Prop. Date: 9/15/2008, Min. age: 20, Max. age: 100, # mice: 1000, Status: active). At the bottom, there are buttons for 'Submit', 'Clear', 'Session Report', and 'Add test type / data description'.

**Figure 19-6 Form: Add Experimental Test**

Click the **Add Exp Test** button to open the form. This form may also be opened from the Add an experimental plan form or Manage an experimental plan form. Only owners and the Administrator may add experimental tests.

An experimental test is always associated with only one experimental plan. Start by **selecting the plan**.

In order to re-use existing tests (even from other plans), you can select a test from the white list box on the lower left hand side of the form and click **Repeat Test**.

Then enter a unique test name of up to 32 characters. JCMS will assign a test ID. Choose the test type. JCMS will then display the captions for the data value fields and any choices for data defaults.

Several fields are used to track the progress of the experiment including the current test status, a proposed date for the test, projected number of mice to use, and a suggested minimum and maximum age for the mice. This proposed age range may be entered in days, weeks (7 days/week), or months (30.4375 days/ month).

The caption, format, min and max values used to define the data results to be collected for this test type are displayed on the form. To help with data entry, default values for these data result fields may be entered. These default values may be chosen from a list of previously created data defaults: select the default name and click the "Use these data defaults" button on the form. Changes may then be made to the default data values to customize them for this test. A default value may not be specified for the "file" format.

## **19.9 Editing an Experimental Test**

Click the **Edit Exp Test** button to open the Edit Experimental Test form. Only owners and the Administrator may edit experimental tests. This form looks and works in the same manner as the Add Experimental Test form. Select the experimental test by its name and ID number. The experimental plan the test is associated with may not be changed. The test type may not be changed once experimental data exists for this test. The delete button to remove a test from an experimental plan will only function if there is no experimental data for this test and no mice have been selected for the test.

## **19.10 Selecting Mice for an Experimental Plan**

Three different methods may be used to select mice for experimental plans and tests.

- 1) The simplest method is to add the mice to the plan when the data is added. This method does not allow scheduling to be used.
- 2) Add the mice using the Mouse Query form. This method takes advantage of the query form criteria for selecting possible mice for the plan. A second step in the process uses the Bulk add mice from query form to add only a subset of the mice in the query result set.
- 3) The Manage an experimental plan form is used to remove mice from a plan and to move pre-selected mice into and out of experimental tests that are part of the plan.

## 19.11 Adding Mice to a Plan using the Mouse Complex Query Form

A group of mice may be **pre-selected** (chosen for planning purposes), for experimental plans and tests. The **Query Mouse** form is used to add mice to experimental plans and tests. Click the **Query Mouse** button to open the form and use it as described in the Basics on using the query forms section to select a set of mice. No results fields need to be chosen. Click the “**Add mice to Experiment**” button. The **Bulk add mice from Query** form will open. It will show the mice selected by the query in the “mice in query results list” on the left side of the form.

ID	DOB	Age	Sex	Owner	Life	stat	plan	test	plans	Strain
CBA-W-104	4/26/2012	50	M	OWN1	A	N	N	N	0	CBAA/J
CBA-W-096	4/26/2012	50	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-097	4/26/2012	50	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-098	4/26/2012	50	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-099	4/26/2012	50	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-100	4/26/2012	50	M	OWN1	A	N	N	N	0	CBAA/J
CBA-W-101	4/26/2012	50	M	OWN1	A	N	N	N	0	CBAA/J
CBA-W-102	4/26/2012	50	M	OWN1	A	N	N	N	0	CBAA/J
CBA-W-095	4/26/2012	50	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-103	4/26/2012	50	M	OWN1	A	N	N	N	0	CBAA/J
CBA-W-106	4/27/2012	49	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-107	4/27/2012	49	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-108	4/27/2012	49	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-105	4/27/2012	49	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-114	5/9/2012	37	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-109	5/9/2012	37	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-110	5/9/2012	37	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-111	5/9/2012	37	F	OWN1	A	N	N	N	0	CBAA/J
CBA-W-112	5/9/2012	37	F	OWN1	A	N	N	N	0	CBAA/J

ID	DOB	Age	Sex	Owner	Life	Strain
CBA-W-103	4/26/2012	50	M	OWN1	A	CBAA/J
CBA-W-102	4/26/2012	50	M	OWN1	A	CBAA/J
CBA-W-101	4/26/2012	50	M	OWN1	A	CBAA/J
CBA-W-100	4/26/2012	50	M	OWN1	A	CBAA/J
CBA-W-104	4/26/2012	50	M	OWN1	A	CBAA/J

Figure 19-7 Form: Bulk Add Mice from Query

Choose the plan to add the mice to. Any experimental tests that are part of this plan will now be listed in the “Tests in the plan” box at the top right of the form. It is possible to add the mice to some (or all) of the tests in this plan by selecting the tests in this box. Be sure to **select** the “**add to test(s) selected at right also**” radio button if also adding mice to tests. Otherwise, the mice will not be added to the selected tests, only to the plan. Note that a mouse cannot be added to a test unless it is also added to the plan. Mice may be added to a plan that does not have any tests. It will not be possible to add experimental data records for these mice until a test describing that data has been added to the plan. This may be done later.

Select mice to be added and click the “> Add mice” button to move them into the list at the right. Mice may be removed from the “mice proposed to add” list box by selecting them and clicking the “< Remove mice” button.

The mice in the query results list box will indicate with a Y/N if each mouse is already in (pre-selected for) the selected plan and selected test. The number of other plans the mouse is pre-selected for will also be displayed. If more than one test is selected, the Y/N will be only for the first (or top) test in the list as shown in the tests in this plan list box.

The “**Show/print query results**” button will show a print preview giving the contents of the Mice in query results list.

Once the list of mice proposed to add is ready, click the “**Submit mice**” button to make the changes to the database. A print preview of the submit report will appear on the screen. This report will indicate what actions were taken. It lists each mouse ID, the owner, and then a set of codes indicating if the mouse was added or not. The header will list the plan ID and any selected test IDs. The codes will appear in the order given in the header. The code “A” indicates the mouse was added to the plan or test and the code “E” indicates the mouse already existed in the database (previously pre-selected) for that plan or test. An additional code ‘C’ will appear at the end of the listing if the mouse has an owner different from the owner of the plan. This is a reminder to check with the owner of the mouse before using it.

### 19.11.1 Example of the report format:

Plan owner = ABC, Plan = 6, Test(s) = 12, 15

MOUSE ID, OWNER, PLAN, TEST(S)

123, ABC, A, A, A  
456, ABC, E, A, E  
789, XYZ, E, A, A, C  
123, ABC, E, E, E

The above sample of a printout indicates that mouse 123 (owned by ABC) was added to plan 6 and tests 12 and 15. Mouse 456 was already selected for plan 6 and test 15. It was not already selected for test 12, so it was added to that test only. Mouse 789 was already selected for plan 6. It was not pre-selected for either test, so it was added to both. The code “C” is a reminder to check with owner XYZ for permission to use this mouse. Mouse 123 already existed for both the tests and plan (note: it may have been accidentally repeated in the list of mice to add). No more action was taken for that mouse.

```
Bulk Add Mice from Query - Submit report for mtsadmin (11/18/2005 3:42:13 PM)
Plan Owner = OVS, Plan = 1, Test(s) = 1
A=Added; E=already existed; C=Check w/owner for permission to use
MOUSE ID, OWNER, PLAN, TEST(S)

-----
63, OVS, A, A
64, OVS, A, A
65, OVS, A, A
66, OVS, A, A
67, OVS, A, A
68, OVS, A, A
69, OVS, A, A
70, OVS, A, A
71, OVS, A, A
72, OVS, A, A

-----
Printed 11/18/2005 3:45:36 PM
```

**Figure 19-8 Report: Add Mice to Plan**

### 19.11.2 Error message: “XX mice were selected by the query, more than can be held in the Query Results box.”

“The list has been shortened to show only xx mice. Use these or close the form and re-do the query.” It is possible to select a very large number of mice with the Mouse complex query form. The Bulk Add Mice from Query form arbitrarily selects only about 50 of these mice for use with

the form. Otherwise, the dataset can become awkward to deal with. If the mice displayed are not those desired, close the form and change the query criteria.

## 19.12 Managing an Experimental Plan

Click the **Manage/Edit Exp Plan** button to open the form. It is used to move mice in and out of tests and for deletion of mice from tests and plans. This form is also used for editing the experimental plan fields and deletion of tests from a plan. **Mice are considered “in” a test and/or plan** if they have been pre-selected for it or if they have an experimental data record containing the results for the test. Mice do not have to be pre-selected; they will be automatically added to the plan/test when the data results are entered.

### 19.12.1 Choosing Mice for Experimental Tests

The Manage/edit Experiment Plan form is used for moving mice pre-selected for a plan into and out of experimental tests associated with that plan.

ID	DOB	Sex	Owner	Life stat	Gen	Genotype	In this test?	# of tests	Strain
A5	6/1/2008	F	nobody	A	F01	none	N	0	C3H/He
A3	6/1/2008	M	nobody	A	F01	none	N	0	C3H/He
A2	6/1/2008	F	nobody	A	F01	none	N	0	C3H/He
A1	6/1/2008	M	nobody	A	F01	none	N	0	C3H/He
A4	6/1/2008	F	nobody	A	F01	none	N	0	C3H/He
Lt2	7/27/2008	F	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt10	7/27/2008	M	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt1	7/27/2008	F	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt3	7/27/2008	F	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt4	7/27/2008	F	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt5	7/27/2008	M	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt6	7/27/2008	M	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt7	7/27/2008	M	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt8	7/27/2008	M	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
Lt9	7/27/2008	M	nobody	A	F01	none	N	0	ABCD <sup>E</sup>
A10	8/1/2008	-	nobody	A	F01	none	N	0	C3H/He
A6	8/1/2008	-	nobody	A	F01	none	N	0	C3H/He
A7	8/1/2008	M	nobody	A	F01	none	N	0	C3H/He
A9	8/1/2008	M	nobody	A	F01	none	N	0	C3H/He
A8	8/1/2008	M	nobody	A	F01	none	N	0	C3H/He

**Figure 19-9 Form: Manage Experimental Plan**

First choose the plan ID. Then select one experimental test by clicking on it in the “Tests in this plan” list box (upper right of the form). All mice in the plan will be listed in the bottom left box and all mice currently in this test will be listed in the bottom right box. Select mice and move them back and forth between the boxes with the “Add mice to test >” and “< Remove mice from test” buttons. Multiple mice may be chosen by holding down the Control (Ctrl) key while clicking on each mouse. The “Select all” and “Unselect” buttons may be used to pick all the mice at once or to remove the selection from all the mice.

No changes will be made to the database until the “Submit changes to the list of mice in this test” button is clicked. A report is displayed on the screen showing the changes that were made to the database. If no mice are listed, then all mice in the “Proposed mice for the test above” list

box (right side) were already in the test. Otherwise, one of the following actions will be listed for each mouse: **D** (mouse was removed from those pre-selected for the test); **A** (mouse was added to those pre-selected for the test); or **X** (mouse was not removed because it already has an experimental data record for this test).

The “**Show/print mice in test**” button will display a report listing all mice in the test and the current action that has been specified for each mouse. This action may be: **A** for add this mouse, **R** for remove this mouse, or **N** for no action.

Edit/Manage Exp. Plans: Submit results for changes made to mice in a plan/test. Report for mtsadmin (11/18/2005 3:58:49 PM)

Listing of changes to mice for Plan ID 1 and Test ID 3

ID, DOB, Sex, Owner, Life Status, Gen, Genotype, Strain, Action taken (D=removed, A=added, X=not removed, mouse has exp. data), Warnings

---

63, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
64, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
65, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
66, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
67, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
68, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
69, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
70, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
71, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A  
72, 10/20/2005, M, OVS, A, N01, none, XYZxJ, A

---

Printed 11/18/2005 4:00:47 PM

**Figure 19-10 Report: Edit/Mange Experimental Plan**

### 19.12.2 Removing Mice from an Experimental Plan

The Manage/edit Experiment Plan form is used to remove mice from the list of those pre-selected for an experimental plan.

First **choose the plan ID**. Select the mice to be removed from the plan in the “**Mice in this plan**” list box (left side). Multiple mice may be chosen by holding down the Control (Ctrl) key while clicking on each mouse. The “Select all” and “Unselect” buttons may be used to pick all the mice at once or to remove the selection from all the mice. Click the “**Remove mice selected in the left box**” button. A report is generated showing the changes that were made for each mouse selected. “Y” indicates the mouse was removed from those pre-selected for the plan. “**Unable to delete from plan**” indicates that the mouse is also pre-selected for an experimental test or has experimental data records for one or more tests. Mice that have data may not be removed from the plan. Mice that are only pre-selected for an experimental test must be removed first from the test. Then the mouse may be removed from the plan.

The “**Show/print mice in plan**” button will display a report listing all mice in the plan with a Y/N indicating if each mouse may be deleted or not. This report also indicates if the mouse is in the currently selected test and how many tests in this plan that each mouse is in.

### 19.12.3 Editing Experimental Plan Fields

The Manage/edit Experiment Plan form is used for changing the fields describing the experimental plan. These are the fields shown in the upper left section of the form. Select the Plan ID. The “**Clear Plan fields**” button will blank all fields including the required ones. The “**Submit changes to Plan fields**” button must be clicked to make the changes permanent in the database. The owner and plan ID may not be changed. The plan name and status are required.

#### **19.12.4 Deleting an Experimental Plan**

The Manage/edit Experiment Plan form is used for deleting experimental plans. Select the plan ID. Click the “**Delete Plan**” button. A plan cannot be deleted if it has any experimental tests, any pre-selected mice, or any experimental data records associated with it.

#### **19.12.5 Deleting an Experimental Test**

The Manage/edit Experiment Plan form is used for deleting experimental tests. Select the test from the list box at the upper right side of the form. The delete test button will only function if there is no experimental data for this test and no mice have been pre-selected for the test.

## 19.13 Adding or Editing Experimental Data for a Mouse

The screenshot shows the 'Add Experimental Data' form. At the top left, there's a checkbox for 'This data is associated with an experimental plan'. Below it, a dropdown for 'Test type' is set to 'Blood Values'. The 'Plan ID' dropdown is set to '2' and the 'Test ID' dropdown is set to '9'. A 'Specimen' dropdown shows 'Mouse ST-002' and a 'Data owner' dropdown shows 'OWN1'. In the 'Data collection' section, there are buttons for 'Auto-calculate age' (selected) and 'Manually enter age', with a date field showing '10/3/2011'. Below this, a radio button group for 'Age expressed in' has 'Days' selected. A 'Data values' table lists several items: 'Blood Glucose/70 / 100' with a value of '80.5', 'Body Weight/ /' with a value of '30', 'Comments/ / text' with a value of 'Sample', 'CBC/ /' (empty), 'WBC/ /' (empty), 'RBC/ /' (empty), and 'Excel output/ / file' set to 'ST-MiceBloodValues.xls'. At the bottom, there are buttons for 'Submit', 'Clear all', 'Session Report', 'Next', 'Clear data value fields', 'Reset data value fields to defaults', and 'Test complete for this Exp Plan'.

**Figure 19-11 Form: Add Experimental Data**

Click the **Add Exp Data** button from the experiments tab to open the form. Experimental data does not have to be associated with an experimental plan and test. To add data for mice not in a plan, remove the check mark next to “**This data is associated with an experimental plan**”. The unnecessary fields will now be grayed out and all mice will be listed in the mouse drop down box.

Begin adding experimental data by choosing the **test type**. The captions for the data values will then appear on the form. Any minimum or maximum values will be displayed and the required box will be checked if the field must have data entered. Choosing the test type will narrow down the choices of experimental plans and tests to only those for this test type.

Choose the **Plan ID** to narrow down the choices of experimental tests. Then choose the **Test ID**. The test ID may be chosen without first picking the plan ID. Once the test is chosen, any default data values that have been set up for it will be displayed. These are to aid in data entry and should be changed to the actual value for each specimen.

Once the test ID is chosen, only mice that need to have a data record added will be listed in the **mouse** drop down box. These are mice that are pre-selected for this test. See the section on managing an experimental plan to learn how to pre-select mice.

To enter data for a mouse that is not pre-selected for a test, type the mouse ID into the mouse drop down box. A confirmation box will appear to verify that the mouse should be added to the list of pre-selected mice for this test. If YES is selected, the mouse will be added to the list of those pre-selected **even if the data record is not later successfully submitted**. The mouse is added

to the pre-selection list first, before the experimental data submit occurs. See the section on managing an experimental plan to learn how to remove mice from those pre-selected for a test.

JCMS always sets the owner of the data to be the same as the owner of the plan. If no experimental plan and test is used, then the owner of the data may be set to any valid owner.

The age field indicates the **age of the specimen at the time of data collection**. JCMS will calculate this from the data collection date and birth date of the mouse if the **Auto-calculate age** button is selected. A warning message will appear if the mouse's life status is not "alive" and the experiment date is after the mouse's exit date. JCMS will store the age field value as number of days old. Age may be entered in weeks or months by selecting the appropriate radio button.

If one or more data values in the test results for this specimen are abnormal, then check the Data is abnormal box. Later, this value may be used to easily locate all experimental data records with unusual results.

If the "**Change the life status of this mouse and set the exit date to the data collection date**" box is checked, then the mouse record will be updated at the same time as the experimental data record is submitted. The mouse will not be updated unless the submit is successful for the experimental data record.

If "**Auto Increment ID**" is checked, then the next pre-selected mouse ID (alphabetically) will be displayed in the mouse drop down box.

The "file" data format is used to add an association between this data record and a document. Documents are computer files (such as an image, spreadsheet, word document, etc.) It is not possible to directly enter the file name into the data result box. Use the ...(upload), assoc., or X(delete) buttons to select the file. See Section 17 on documents for more information on the forms used for uploading and associating documents. More than one data result field may be defined with the file format; however, a specific file may only be associated once with a specific experimental data record.

The **Data ID** for this experimental data record is assigned by JCMS and will be displayed in the "Data added this session" box.

The **Test complete for this Exp Plan** button will change the test status in the experimental test record for the test currently indicated in the Test ID box to "done".

## **19.14 Editing Experimental Data**

Click the **Edit Exp Data** button to open the form. Select either **Data ID** or **Mouse** as the method to use for selecting the experimental data record to edit.

Selecting a **Plan ID** or **Test ID** will limit the choices in the other drop down boxes. If the desired drop down boxes are grayed out (not available to choose from), then click the Data ID or Mouse buttons again. This will start the selection process over. When no test ID is selected, a particular mouse may be listed multiple times in the mouse drop down box, once for each experimental test the mouse has data results for.

The Data ID and Mouse drop down boxes will show all experimental data records that exist for the selected test ID. These boxes will not show pre-selected mice that have no experimental data record yet. Use the Add experimental data form to add the data for the pre-selected mice.

The rest of the fields on the form work the same as on the Add experimental data form described above.

## 19.15 Adding Experimental Data to Several Mice at Once

The screenshot shows the 'Add Data to a Specific Experimental Test (mtsadmin) (Secretaries and Owners)' interface. The main title is 'Bulk Add Data to an Experimental Test'. Key fields include 'Test type' (Blood values), 'Data owner' (nobody), 'Plan ID' (3), 'Plan name' (Sample plan for documentation), and 'Test ID' (3). A note says 'All mice pre-selected for this test are listed with Y/N indicating if data is already added. Select multiple mice to submit.' Below this is a table of mice with columns: Added, ID, birthDate, sex, owner, lifeSt, strain. The table contains rows for mice A1-A5. To the right of the table are 'Data values' for various parameters like Blood Glucose, Body Weight, and CBC, each with a 'dec' (decimal) field and a 'Req' (Required) checkbox. A 'Comments' field is also present. At the bottom are buttons for 'Submit', 'Clear all', 'Session Report', 'Clear data value fields', 'Reset data value fields to defaults', and 'Test complete for this plan'.

**Figure 19-12 Form: Bulk Add Experimental Data**

Click the **Bulk Add Data to Specific Test** button to open the form. This form is used to enter new data values for mice that have already been selected as part of a particular plan and test. If several mice should have all the same data values, this form may be used to speed up data entry. Or, it could be used to create data records for a group of mice where many of the data values are the same, then use the Edit Experimental Data form to change or add specific values for each individual mouse.

Select the experimental **plan ID** to limit the choices in the test ID box to only those for the specific plan.

Once a **test ID** is chosen, the list box will show all mice that have been pre-selected for the test. The first column indicates if a data record has already been added (Y) or not (N) for that mouse ID.

**Select one or more mice in the list box.** WARNING: The whole batch will be rejected if any of the selected mice already have a data record (Added = Y).

The rest of the form works similarly to the Add Experimental Data form described above.

Note: This form does not allow adding information in a data value field using the "file" format. If a "file" format field is required the form may not be used for that experimental test.

## 19.16 Experiment Work Report for Scheduling Procedures

Click the **Experiment Work Report** button on the Reports tab to open the form for requesting the report. This report is designed to provide lists of mice for a work "to-do" schedule. There are many criteria available for limiting the mice shown on the report, such as a date range,

experimental plan owner, mouse life status, experimental test status, test type, or specific experimental tests. Entering selections for experimental plan owner, experimental test status, and/or test types will cause the list of experimental tests to change to show only those that satisfy the criteria. There are two forms of output, a formatted report or a datasheet for exporting to Microsoft Excel or other spreadsheet.

**Experiment Work Report (mtsadmin) {Secretaries and Owners}**

### Request Experiment Work Report

<b>Report Time Frame</b>	<b>Limit By</b>	<b>Mouse Life Status</b>																												
<input checked="" type="radio"/> This week <input type="radio"/> This month <input type="radio"/> Next month	*Start date: 10/9/2013 *End date: 10/16/2013	<input checked="" type="checkbox"/> No Data Collected Yet <input checked="" type="radio"/> Proposed Test Date in Time Frame <input type="radio"/> Mouse Age in the Proposed Min/Max Age Range																												
<b>Experimental Test Criteria</b> <table border="1"> <tr> <td><b>Experimental Plan Owner</b></td> <td><b>Experimental Test Status</b></td> <td><b>Test Types</b></td> </tr> <tr> <td> <input checked="" type="radio"/> Any  <input type="radio"/> Selected            nobody            0wN1         </td> <td> <input type="checkbox"/> Proposed  <input checked="" type="checkbox"/> Active  <input type="checkbox"/> Done         </td> <td> <input checked="" type="radio"/> Any  <input type="radio"/> Selected            Blood tests            Clinical Exam            Eye exam            IOP            Macular degeneration exam            Optional extra genotyping            Primary genotyping            Secondary genotyping         </td> </tr> </table>			<b>Experimental Plan Owner</b>	<b>Experimental Test Status</b>	<b>Test Types</b>	<input checked="" type="radio"/> Any <input type="radio"/> Selected nobody 0wN1	<input type="checkbox"/> Proposed <input checked="" type="checkbox"/> Active <input type="checkbox"/> Done	<input checked="" type="radio"/> Any <input type="radio"/> Selected Blood tests Clinical Exam Eye exam IOP Macular degeneration exam Optional extra genotyping Primary genotyping Secondary genotyping																						
<b>Experimental Plan Owner</b>	<b>Experimental Test Status</b>	<b>Test Types</b>																												
<input checked="" type="radio"/> Any <input type="radio"/> Selected nobody 0wN1	<input type="checkbox"/> Proposed <input checked="" type="checkbox"/> Active <input type="checkbox"/> Done	<input checked="" type="radio"/> Any <input type="radio"/> Selected Blood tests Clinical Exam Eye exam IOP Macular degeneration exam Optional extra genotyping Primary genotyping Secondary genotyping																												
<b>Experimental Tests</b> <table border="1"> <tr> <td><input checked="" type="radio"/> Any</td> <td><input type="radio"/> Selected</td> </tr> <tr> <td colspan="2"> <table border="1"> <thead> <tr> <th>testID</th> <th>testName</th> </tr> </thead> <tbody> <tr><td>5</td><td>10 week exam</td></tr> <tr><td>4</td><td>5 week exam</td></tr> <tr><td>13</td><td>AMD</td></tr> <tr><td>16</td><td>Early genotyping</td></tr> <tr><td>12</td><td>Eye Exam</td></tr> <tr><td>14</td><td>FSN bloodwork</td></tr> <tr><td>17</td><td>Late genotyping</td></tr> <tr><td>1</td><td>One month blood work</td></tr> <tr><td>9</td><td>One month IOP</td></tr> <tr><td>6</td><td>One month IOP</td></tr> <tr><td>10</td><td>...</td></tr> </tbody> </table> </td> </tr> </table>			<input checked="" type="radio"/> Any	<input type="radio"/> Selected	<table border="1"> <thead> <tr> <th>testID</th> <th>testName</th> </tr> </thead> <tbody> <tr><td>5</td><td>10 week exam</td></tr> <tr><td>4</td><td>5 week exam</td></tr> <tr><td>13</td><td>AMD</td></tr> <tr><td>16</td><td>Early genotyping</td></tr> <tr><td>12</td><td>Eye Exam</td></tr> <tr><td>14</td><td>FSN bloodwork</td></tr> <tr><td>17</td><td>Late genotyping</td></tr> <tr><td>1</td><td>One month blood work</td></tr> <tr><td>9</td><td>One month IOP</td></tr> <tr><td>6</td><td>One month IOP</td></tr> <tr><td>10</td><td>...</td></tr> </tbody> </table>		testID	testName	5	10 week exam	4	5 week exam	13	AMD	16	Early genotyping	12	Eye Exam	14	FSN bloodwork	17	Late genotyping	1	One month blood work	9	One month IOP	6	One month IOP	10	...
<input checked="" type="radio"/> Any	<input type="radio"/> Selected																													
<table border="1"> <thead> <tr> <th>testID</th> <th>testName</th> </tr> </thead> <tbody> <tr><td>5</td><td>10 week exam</td></tr> <tr><td>4</td><td>5 week exam</td></tr> <tr><td>13</td><td>AMD</td></tr> <tr><td>16</td><td>Early genotyping</td></tr> <tr><td>12</td><td>Eye Exam</td></tr> <tr><td>14</td><td>FSN bloodwork</td></tr> <tr><td>17</td><td>Late genotyping</td></tr> <tr><td>1</td><td>One month blood work</td></tr> <tr><td>9</td><td>One month IOP</td></tr> <tr><td>6</td><td>One month IOP</td></tr> <tr><td>10</td><td>...</td></tr> </tbody> </table>		testID	testName	5	10 week exam	4	5 week exam	13	AMD	16	Early genotyping	12	Eye Exam	14	FSN bloodwork	17	Late genotyping	1	One month blood work	9	One month IOP	6	One month IOP	10	...					
testID	testName																													
5	10 week exam																													
4	5 week exam																													
13	AMD																													
16	Early genotyping																													
12	Eye Exam																													
14	FSN bloodwork																													
17	Late genotyping																													
1	One month blood work																													
9	One month IOP																													
6	One month IOP																													
10	...																													
<input type="button" value="Print Format"/> <input type="button" value="Spreadsheet Format"/>																														

**Figure 19-13 Request Experiment Work Report**

The formatted report gives a listing sorted by experimental plan owner, experimental plan, experimental test, and mouse ID. Basic information about the mice will be shown (ID, pen, strain, generation, date of birth, age today, sex, and life status). In addition it shows the plan ID, test ID, and the test's proposed values for minimum age, maximum age, and to-do date. When a minimum and/or maximum age is available, an age window is shown for each mouse (date the

mouse reaches the minimum age and date the mouse reaches the maximum age). When there is a proposed date for the test, the mouse's age on the proposed date is shown.

Experiment Work To Do Report: Untested Mice								Report date	Wednesday, October 09, 2013
Mouse ID	Age window from - to	Age on proposed date	Age today	Birth date	Sex	Life status	Gen. Pen ID	Pen name	Strain
<b>Plan ID 1 Flaky skin</b>								Plan Owner	nobody
Test ID 2	Two month blood work		Proposed min age 61			Proposed max age: 88	Proposed date	Test status: active	
NOD-001	10/12/2013	11/8/2013	58	8/12/2013	F	A	F02	101	NOD.CB17-Prkdc<scid>J
NOD-002	10/12/2013	11/8/2013	58	8/12/2013	M	A	F02	52 Cage-00005	NOD.CB17-Prkdc<scid>J
NOD-003	10/12/2013	11/8/2013	58	8/12/2013	M	A	F02	52 Cage-00005	NOD.CB17-Prkdc<scid>J
Test ID 3	Three month blood work		Proposed min age 91			Proposed max age: 119	Proposed date	Test status: active	
A31	9/30/2013	10/28/2013	100	7/1/2013	M	A	F01	33	NOD.CB17-Prkdc<scid>J
A32	9/30/2013	10/28/2013	100	7/1/2013	M	A	F01	35	NOD.CB17-Prkdc<scid>J
A41	9/30/2013	10/28/2013	100	7/1/2013	F	A	F01	36	NOD.CB17-Prkdc<scid>J
A42	9/30/2013	10/28/2013	100	7/1/2013	F	A	F01	34	NOD.CB17-Prkdc<scid>J
A43	9/30/2013	10/28/2013	100	7/1/2013	F	A	F01	32	NOD.CB17-Prkdc<scid>J
Test ID 4	5 week exam		Proposed min age 35			Proposed max age: 41	Proposed date	Test status: active	
FVB-M-200	10/6/2013	10/12/2013	38	9/1/2013	F	A	F02	86	FVB.NU
FVB-M-201	10/6/2013	10/12/2013	38	9/1/2013	F	A	F02	86	FVB.NU
FVB-M-202	10/6/2013	10/12/2013	38	9/1/2013	F	A	F02	86	FVB.NU
FVB-M-203	10/6/2013	10/12/2013	38	9/1/2013	F	A	F02	86	FVB.NU
FVB-M-204	10/6/2013	10/12/2013	38	9/1/2013	M	A	F02	87	FVB.NU

**Figure 19-14 Experiment Work To Do Report Based on the Mouse Age Range**

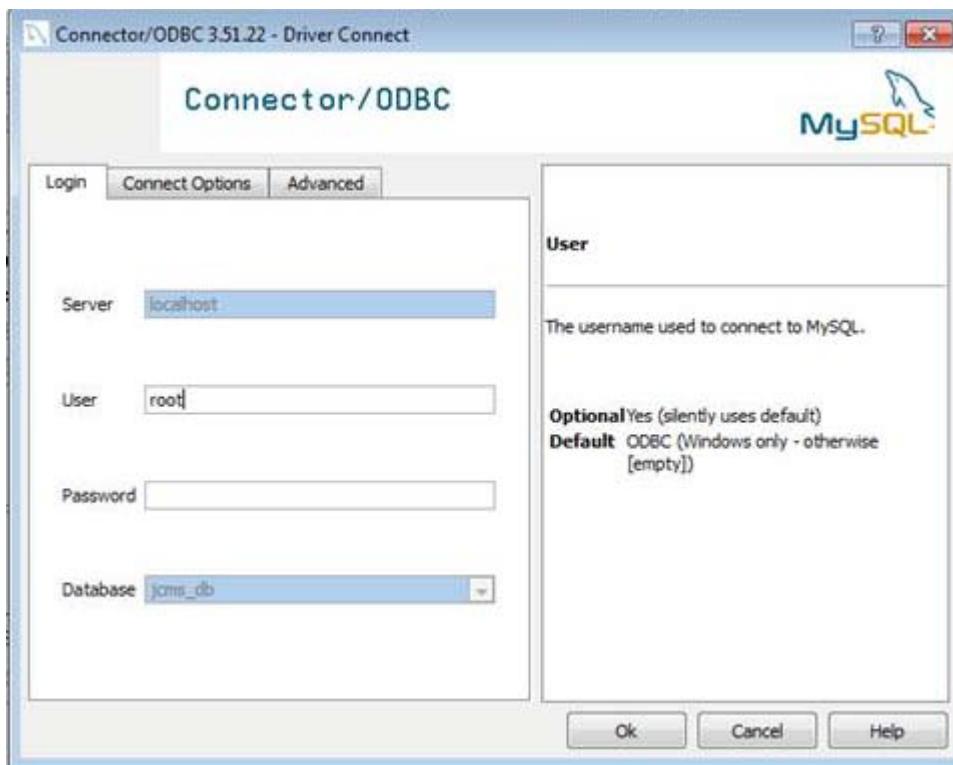
The spreadsheet version of the report contains all the same information and includes an extra column for data collected (Y/N). If the check box "No data collected yet" is not checked, then all mice selected for the test will be listed.

Plan ID	Exp plan	Owner	testType	Test ID	Test name	Mouse ID	DataCollected	Proposed min age	AgeWindowBegins	Proposed max age	AgeWindowEnd	AgeToday
1	Flaky skin	nobody	Blood tests	3	Three month blood work A31	N		91	9/30/2013	119	10/28/2013	100
1	Flaky skin	nobody	Blood tests	3	Three month blood work A32	N		91	9/30/2013	119	10/28/2013	100
1	Flaky skin	nobody	Blood tests	3	Three month blood work A41	N		91	9/30/2013	119	10/28/2013	100
1	Flaky skin	nobody	Blood tests	3	Three month blood work A42	N		91	9/30/2013	119	10/28/2013	100
1	Flaky skin	nobody	Blood tests	3	Three month blood work A43	N		91	9/30/2013	119	10/28/2013	100
1	Flaky skin	nobody	Blood tests	2	Two month blood work NOD-001	N		61	10/12/2013	88	11/8/2013	58
1	Flaky skin	nobody	Blood tests	2	Two month blood work NOD-002	N		61	10/12/2013	88	11/8/2013	58
1	Flaky skin	nobody	Blood tests	2	Two month blood work NOD-003	N		61	10/12/2013	88	11/8/2013	58
1	Flaky skin	nobody	Clinical Exam	4	5 week exam	FVB-M-200	N	35	10/6/2013	41	10/12/2013	38
1	Flaky skin	nobody	Clinical Exam	4	5 week exam	FVB-M-201	N	35	10/6/2013	41	10/12/2013	38
1	Flaky skin	nobody	Clinical Exam	4	5 week exam	FVB-M-202	N	35	10/6/2013	41	10/12/2013	38
1	Flaky skin	nobody	Clinical Exam	4	5 week exam	FVB-M-203	N	35	10/6/2013	41	10/12/2013	38

**Figure 19-15 Experiment Work Report Spreadsheet**

### 19.16.1 Connector/ODBC Window Pops-up When Using the Experiment Work Report

This window will appear if some of the parameters for the MYSQL database have changed.



**Figure 19-16 Connector/ODBC window**

To eliminate the problem update the MySQL setup variables shown below.

## 19.16.2 Setup Variables Used by the Experiment Work Report

**Table 19-1 MySQL setup variables**

JCMS(MySQL_DATABASE_NAME	jcms_db	The name of your MySQL database.
JCMS(MySQL_DRIVER	{MySQL ODBC 3.51 Driver}	The name of your MySQL driver.
JCMS(MySQL_SERVER	localhost	The name of your MySQL server.
JCMS(MySQL_USER_ID	root	The name of your MySQL user ID.

These four setup variables are necessary for the query used by the report to work efficiently. The defaults are shown. The upgrader/installer will initialize these; however, if later on changes are made to one or more of these values, it may be necessary to also manually change them in the setup variable table.

## 19.17 Experimental Plan Query

This query is described in the Queries section under Experimental Plan Query.

## 19.18 Importing Experimental Data

The Import Experimental Data function of JCMS loads data from files and automatically populates the *experimental data* records for the mice identified in the file. The data may be associated with *experimental tests* that are part of an *experimental plan* or just be associated with mice.

Many research programs use machines to generate data, such as blood work. It is impractical to try to support individual file formats for the many different machines that already exist and new ones being built each year. Therefore, imported data must be in a specific file format known as CSV (comma separated values). Many applications, such as MS Excel, are capable of creating a

file using this standard format. Some user-manipulation of the file may be required prior to importing it.

The data format must match a user-created and defined JCMS *experimental data definition (test type)*. This test type may be set up to closely match the output file from a machine or application. The JCMS identifier (*mouse ID*) is used to match the data to the mice.

#### Process Diagram

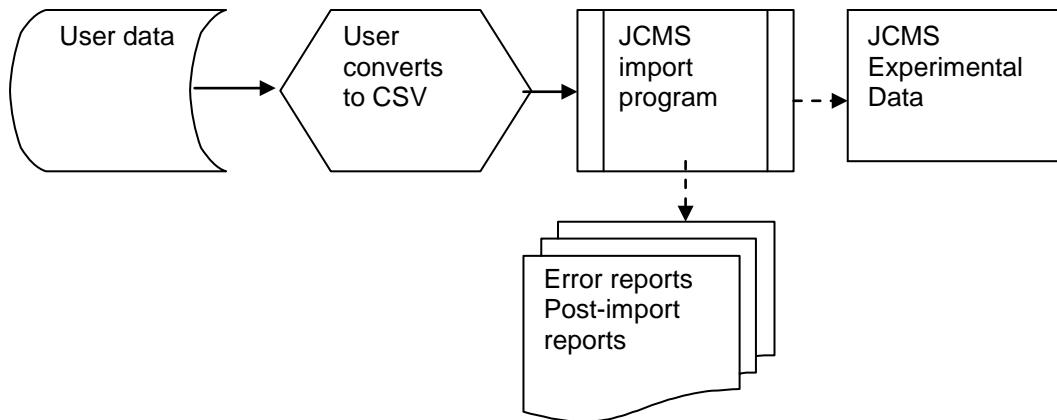


Figure 19-17 Process for importing experimental data from a user file

#### 19.18.1 Input File Format

A file using the CSV (comma separated values) format consists of multiple lines or rows with the data values separated with commas. If the file is created using MS Excel, then each line is the same as a row in a spreadsheet. Each data value consists of the contents of one cell in a column. MS Excel will insert the comma used to separate the data values when the file is saved. These commas are not visible in the cells. CAUTION NOTE: data values, such as comments, cannot contain comma characters because commas are used as data value separators.

##### JCMS format requirements

Token row: The input file may contain rows that are ignored at the beginning. It must start with a row containing “JCMS\_DATA”. This is a token used by the import program to determine where to start processing the file.

Header row: The header row must be the next row following the JCMS\_DATA token. The header row contains the captions indicating what data value is in each column.

- There are three special captions: Mouse ID, Abnormal Data Flag, and Data Collection Date. The first two are required and the Data Collection Date is optional.
- All data values that are imported must be in a column with a caption that matches one defined in the JCMS *Test Type*.
- Columns that do not match a *Test Type* caption or the three special captions are ignored.
- If the *Test Type* has required data, a column for that data must be present.
- The columns may be in any order.
- Duplicate captions are not allowed and will generate an error message.

##### Example of an input file

Blood, body weight test  
Tested on 1/2/08

Chuck Donnelly

#### JCMS\_DATA

Mouse ID,	Abnormal Data Flag,	Data Collection Date,	Blood Glucose,	Body Weight,	Comments
CJD-002,	T,	,	0,	45.0,	found dead
CJD-003,	F,	1/2/08,	89.2,	34	
CJD-010,	F,	1/3/08,	76.95,	30	
CJD-005,	T,	1/2/08,	,	36.4,	malfunction

Data rows: All rows after the header are assumed to contain data.

- Data collection date is an optional field and may be left blank. If all the data has the same collection date, then the date may be entered once on the import form instead of being present in the input file.
- Blank rows within the file and at the end are ignored.

#### 19.18.2 Create a Test Type

This is an example of the Add test type form showing creation of a test type that matches the input file header above.

Test Type   Blood Work and Body Weights						
	Caption	Field description	Format	Required	Min value	Max value
D1	Blood Glucose		dec	<input checked="" type="checkbox"/>	70	100
D2	Body Weight	grams	dec	<input checked="" type="checkbox"/>		
D3	Comments		text	<input type="checkbox"/>		
D4	CBC		dec	<input type="checkbox"/>		
D5	WBC		dec	<input type="checkbox"/>		
D6	RBC		dec	<input type="checkbox"/>		
D7			dec	<input type="checkbox"/>		
D8			dec	<input type="checkbox"/>		
D9			dec	<input type="checkbox"/>		
D10			dec	<input type="checkbox"/>		
D11			dec	<input type="checkbox"/>		
D12			dec	<input type="checkbox"/>		
D13			dec	<input type="checkbox"/>		
D14			dec	<input type="checkbox"/>		
D15			dec	<input type="checkbox"/>		

	Caption	Field description	Format	Required	Min value	Max value
D16			dec	<input checked="" type="checkbox"/>		
D17			dec	<input checked="" type="checkbox"/>		
D18			dec	<input checked="" type="checkbox"/>		
D19			dec	<input checked="" type="checkbox"/>		
D20			dec	<input checked="" type="checkbox"/>		
D21			dec	<input checked="" type="checkbox"/>		
D22			dec	<input checked="" type="checkbox"/>		
D23			dec	<input checked="" type="checkbox"/>		
D24			dec	<input checked="" type="checkbox"/>		
D25			dec	<input checked="" type="checkbox"/>		
D26			dec	<input checked="" type="checkbox"/>		
D27			dec	<input checked="" type="checkbox"/>		
D28			dec	<input checked="" type="checkbox"/>		
D29			dec	<input checked="" type="checkbox"/>		
D30			dec	<input checked="" type="checkbox"/>		

Double click to repeat data description

test type	D1	D2	D3	D4
Body Weights	WEIGHT			

Session Box

Session Report

Add default data values

Submit Clear Session Report Add default data values

**Figure 19-18 Create a test type to match the input fields**

Note there can be captions in the test type that are not used in the importation process.

### 19.18.3 User Interface

The *Import Exp Data* button on the experiments tab is used to initiate the import process.

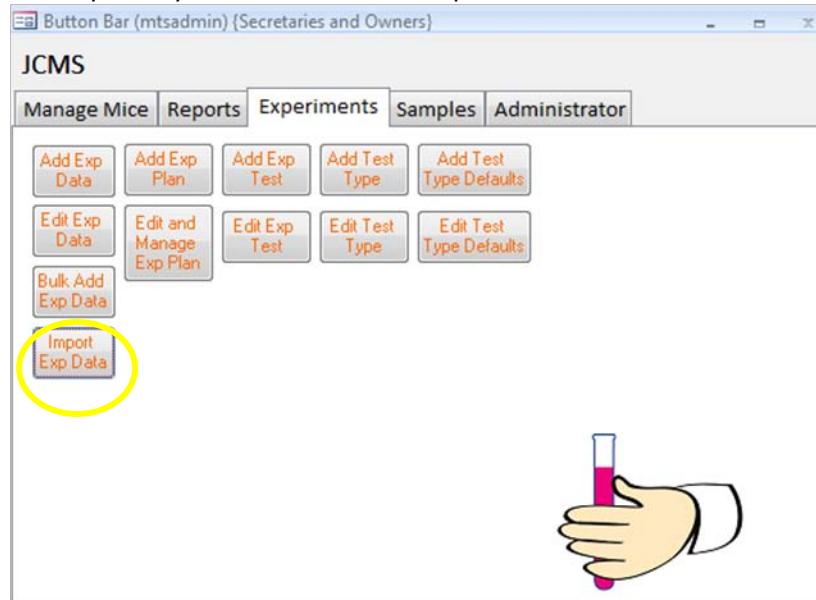


Figure 19-19 Import Exp Data button

Filling in the Import Experimental Data form.

A screenshot of the "Import Experimental Data" form. At the top, it says "Import Experimental Data". Below that is a table titled "No Plan" with one row: "no plan". Under "Test type" is "Body Weight" and "1". Under "PlanOwner" is "nobody". Below the table is a note: "Blood Work and Body Weights". At the bottom of the form are input fields: "\* Input file" with a "Browse..." button, "\* Data collection date" (set to 9/22/2008), "\* Data collection age" (radio buttons for "Auto-calculate age on collection date" and "Leave age blank"), and "\* owner" (a dropdown menu). At the very bottom are buttons for "Validate Input", "Load Data", and "Clear".

Figure 19-20 Import Experimental Data with no experimental plan

To add experimental data to mice without using an experimental plan, select “no plan” from the choices in the Plan ID combo box.

Select the test type from the list of choices. Browse to find and select the input file. The data collection date may be entered with the same value for all the data records by selecting it using the calendar. Otherwise, use the radio button to indicate it is present in a column in the input data file. The data collection age is an optional field. It may be auto-calculated (by using the mouse's birth date and the data collection date) or left blank. When an experimental plan is not used, the owner of the data must be specified. This may be different from the owner of the mice.

The screenshot shows the 'Import Experimental Data' window. At the top, there is a dropdown menu labeled 'Plan ID' with 'No Plan' selected. Below it, a section for 'Test choice' lists 'Test type' as 'Body Weights' and 'Blood Work and Body Weights' (which is highlighted). An 'Input file' field contains the path 'C:\JCMS\TestForDocumentation.csv'. A 'Browse...' button is next to it. Below the input file field are three groups of options: 'Data collection date' (radio button selected for '9/22/2008'), 'Data collection age' (radio button selected for 'Auto-calculate age on collection date'), and 'owner' (dropdown menu set to 'nobody'). A red arrow points to the 'Validate Input' button at the bottom left of the form. Other buttons at the bottom include 'Load Data' and 'Clear'.

**Figure 19-21 Validate input**

Use the Validate Input button to initiate the validation process. The data file will be checked to determine if there are any problems and a report will be displayed indicating if the data may be imported.

Import Experimental Data — Summary Report

---

Loading data from: C:\JCMS\TestForDocumentation.csv

No plan is used, data is added for test type: Blood Work and Body Weights

Data Owner: nobody Data collection date entered from a column in the input file. Age will be calculated.

\*\*\*\*\*

JCMS\_DATA token found in row5

Header row has 8 columns.

Columns with these headers will be processed: MOUSE ID, ABNORMAL DATA FLAG, DATA COLLECTION DATE, BLOOD GLUCOSE, BODY WEIGHT, COMMENTS

---

Errors found in the input file are listed by row number.

Row7: BLOOD GLUCOSE: Range error, the value is less than the minimum allowed.

---

Total number of errors: 1

Total number of valid rows found: 3

---

The following is a listing of the data that will be imported.

If the data collection date (expDate) is entered from the form instead of the input file, that column will display '-'.

Any blank values will be displayed as '--'.

\*\*\*\*\*

BLOOD GLUCOSE; BODY WEIGHT; COMMENTS; Mouse ID; abnormalData; expDate; age

\*\*\*\*\*

89.2; 34; --; CJD-003; False; 1/2/2008; 32

76.95; 30; --; CJD-010; False; 1/3/2008; 30

--; 36.4; malfunction; CJD-005; True; 1/2/2008; 2

---

Validation is complete. To start the LOAD, click the LOAD DATA button on the Import Experimental Data form.

No changes have been made to JCMS during the validation process.

How to proceed

**Figure 19-22 Sample validation report**

In the case above, one input file row has been rejected with an error. The rest of the input file may be imported or the input file can be corrected and the validation run again.

The number of errors that are acceptable is determined from a setup variable called JCMS\_MAX\_IMPORT\_EXP\_DATA\_ERRORS. Once the error limit is reached validation will stop and errors must be corrected in the input file. The default is 10 errors.

Setup Variables		
JCMS Setup Variable name	JCMS Setup Variable value	Description
► JCMS_MAX_IMPORT_EXP_DATA_ERRORS	10	When maximum is reached verification

**Figure 19-23 Maximum number of validation errors**

Import Experimental Data (mtsadmin) {Owners Only}

## Import Experimental Data

\* Plan ID 1 Clinical data on body weights and blood work

\* Test choice

Test ID	Test name	Test type
1	BW test cycle 1	Body Weights
2	Blood work	Blood Work and Body Weights

\* Input file C:\JCMS\TestForDocumentation.csv

\* Data collection date   Column in the input data  Auto-calculate age on collection date  Leave age blank

\* Data collection age  Auto-calculate age on collection date  Leave age blank

\* owner

**Figure 19-24 Import Experimental Data using an Experimental Test**

To record experimental data that is associated with a particular experimental test, first choose the experimental plan ID. The list of choices will change to show all active tests for that experimental plan. The owner is automatically set to be the experimental plan owner. Browse to select the input file. Select the experimental test and data collection date and age settings. Click the Validate Input button.

Import Experimental Data — Summary Report

---

Loading data from: C:\JCMS\TestForDocumentation.csv

Plan ID: 1 Plan name: Clinical data on body weights and blood work Test ID: 2 Test name: Blood work

Data Owner: nobody Data collection date: 2/1/2008 Age will be calculated.

+++++  
JCMS\_DATA token found in row5  
Header row has 8 columns.  
Warning: Data collection date column is present in the input data file but NOT USED. Date of 2/1/2008 is used instead.  
These columns will be ignored: DATA COLLECTION DATE  
Columns with these headers will be processed: MOUSE ID, ABNORMAL DATA FLAG, BLOOD GLUCOSE, BODY WEIGHT, COMMENTS

---

Errors found in the input file are listed by row number.

Row7: MOUSE ID: CJD-002 is not pre-selected for this plan and test. BLOOD GLUCOSE: Range error, the value is less than the minimum allowed.  
Row8: MOUSE ID: CJD-003 is not pre-selected for this plan and test.  
Row9: MOUSE ID: CJD-010 is not pre-selected for this plan and test.  
Row10: MOUSE ID: CJD-005 is not pre-selected for this plan and test.

---

Total number of errors: 5  
Total number of valid rows found: 0  
No data rows to import.  
No changes have been made to JCMS.

DATA VALIDATION FOR IMPORT HAS FAILED

Validation failure

---

Printed 9/22/2008 12:03:14 PM

**Figure 19-25 Data validation failure**

In the case above, no rows have passed the validation step and the input file or other parameters must be changed. The mouse IDs need to be added to this test using the Manage Exp Plan form or the setup variable shown below must be changed to “false”, allowing the mice to be added to the experimental test at the same time as the data is imported.

Setup Variables		
JCMS Setup Variable name	JCMS Setup Variable value	Description
JCMS_IMPORT_EXP_DATA_MICE_MUST_BE_PRESELECTED	true	If false, any mouse IDs not pre-selected will be

**Figure 19-26 Allow experimental data to be imported to an experimental test without preselecting the mice**

Set the setup variable JCMS\_IMPORT\_EXP\_DATA\_ALLOW\_MULTIPLE to true if importing with NO experimental plan and you want to allow more than one record for a test type/mouse combination.

When some or all of the entire input file has passed the validation step, the Load Data button will be enabled. Click it to proceed. The load report will be displayed listing the rows added and those rejected with errors.

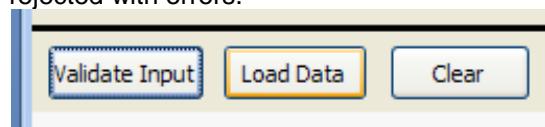


Figure 19-27 Enabled load data button

Import Experimental Data — Summary Report

Information about the request.

Loading data from: C:\JCM S\TestForDocumentation.csv

Plan ID: 1 Plan name: Clinical data on body weights and blood work Test ID: 2 Test name: Blood work

Data Owner: nobody Data collection date: 2/1/2008 Age will be calculated.

\*\*\*\*\*

JCMS\_DATA token found in row5

Header row has 8 columns.

Information about the input file

Warning: Data collection date column is present in the input data file but NOT USED. Date of 2/1/2008 is used instead.

These columns will be ignored: DATA COLLECTION DATE

Columns with these headers will be processed: MOUSE ID, ABNORMAL DATA FLAG, BLOOD GLUCOSE, BODY WEIGHT, COMMENTS

Errors found in the input file are listed by row number.

Errors found

Row7: BLOOD GLUCOSE: Range error, the value is less than the minimum allowed.

Total number of errors: 1

Total number of valid rows found: 3

Experimental Data records for valid rows have been imported and are listed below.

Any blank values will be displayed as '-'. Input file data rows that have been imported

BLOOD GLUCOSE; BODY WEIGHT; COMMENTS; Mouse ID; abnormalData; expDate; age

=====

89.2; 34; -; CJD-003; False; 2/1/2008; 62

76.95; 30; -; CJD-010; False; 2/1/2008; 59

-; 36.4; malfunction; CJD-005; True; 2/1/2008; 32

All mice pre-selected for this experimental test now have data.

Printed 9/22/2008 12:29:13 PM

Figure 19-28 Final load report

When a load to an experimental test is completed, if all the preselected mice have a data record, the option to change the experimental test status from “active” to “done” is offered.

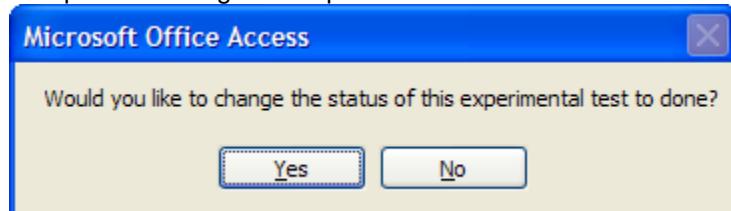
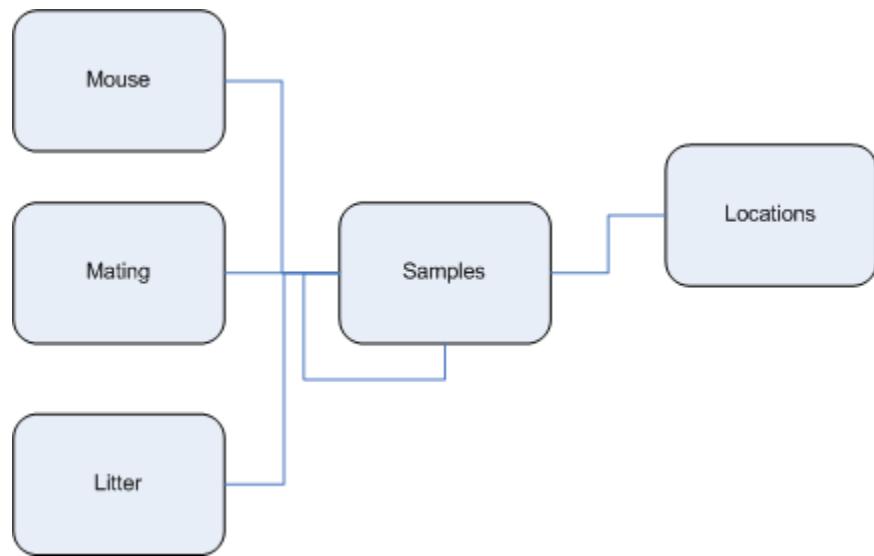


Figure 19-29 Changing the experimental test status

#### 19.18.4 Troubleshooting Notes

- Data values, such as comments, cannot contain comma characters because commas are used as data value separators.
- The *test type* selected directly (no plan) or associated with the experimental test when using an experimental plan may not contain data value fields with the captions: Mouse ID, Data Collection Date, or Abnormal Data Flag. These are special captions used by the importation process.
- Do not insert spaces between the commas used to separate missing values. These can be accidentally interpreted as a value. A line of data containing A,B,,,C is not interpreted the same as A, B, , , C. There is no need to follow a comma with a space.
- Some applications insert characters at the beginning of a file that provide information to the application. If the file has the JCMS\_DATA token on the first line these special characters may cause it to not find the token. The solution is to insert a blank line at the beginning of the file.

## 20 Samples



## 20.1 Set up Controlled Vocabulary for Sample Tracking

The first step in using the sample tracking features is to set up the controlled vocabulary for your lab. Sample tracking has both simple controlled vocabulary (see section 3.3.1) and several more complex controlled vocabularies.

The simple controlled vocabularies for sample tracking are as follows:

- 1) Epoch: Used to qualify the sample's age
- 2) Harvest Method: Indicates how the sample was harvested
- 3) Weight Unit: Units the weight is measured in
- 4) Sample Date Type: Used to qualify the sample date
- 5) Sample Status: All potential statuses a sample may be identified with
- 6) Time Unit: Used for indicating sample age
- 7) Sample Class: Top level category for a sample

Some values have already been added to these tables. Examine each one and add or remove values as necessary. These values are accessed by clicking on the corresponding button on the Administrative button bar:

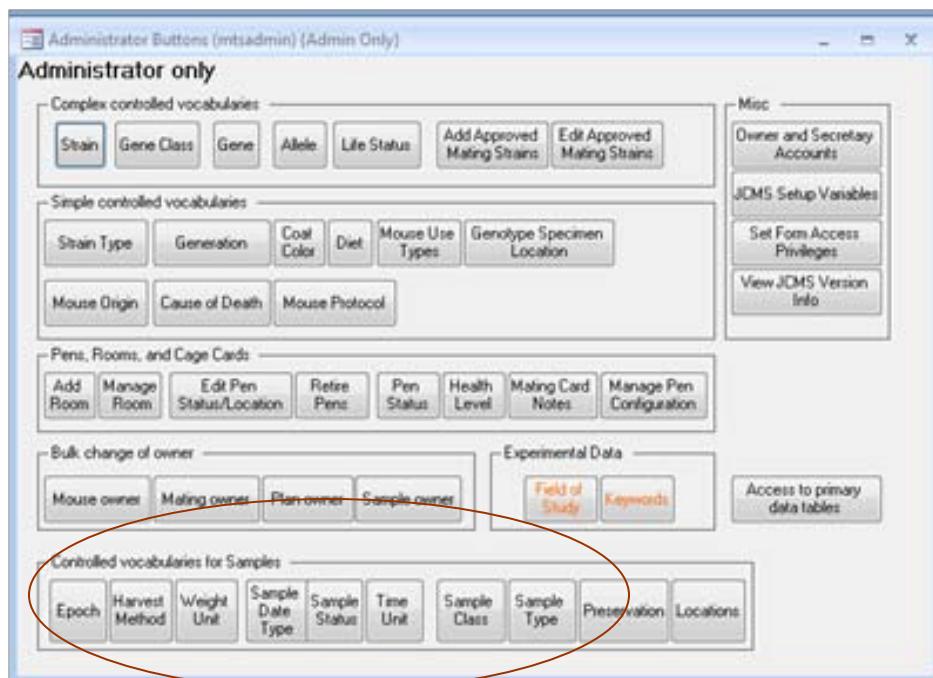


Figure 20-1 Simple controlled vocabulary buttons for sample tracking

After clicking the button, the values are displayed in table format:

AddOrDelete cv_WeightUnit	
	weightUnit
►	grams
	ounces
	milligrams
	picolitres
*	

Figure 20-2 Example of editing simple controlled vocabulary

**To add a new value**, place the cursor in the bottom blank row and type in the new value. Move the cursor out of that row for the new value to save.

**To remove a value** from the table, click on the box at the left side of the row in the table. The whole row will be highlighted. Press the delete key. A dialog box will ask for confirmation that the record should be deleted.

See section 3.1.1 for more details regarding simple controlled vocabularies.

In addition, there are three complex controlled vocabularies for sample tracking. They are:

- 1) Sample Type: Within class, identifies the sample type
- 2) Preservation (Type, Method, Detail): Values for indicating how the sample is preserved
- 3) Location: Where the sample resides

As with simple controlled vocabularies, add and remove values as necessary. Do this by clicking the corresponding buttons on the Administrator button bar:



**Figure 20-3 Complex controlled vocabulary buttons for sample tracking**

To administer sample types, click the Sample Type button. The following screen is displayed:

A screenshot of a Windows-style application window titled "Manage Sample Type". The window has a title bar with standard minimize, maximize, and close buttons. Inside, there are two main sections. On the left, a "Sample Class" dropdown menu is set to "Live". Below it is a "Sample Type" list box containing several items: Sperm, Embryo, Cell Line, Egg, Ovary, and Ovarian tissue. The "Sperm" item is currently selected and highlighted with a black background. At the bottom of the window are three buttons: "Add", "Rename", and "Delete". A note at the bottom states: "NOTE: Technical matings require the sample type terms Embryo, Egg, Ovary, Ovarian tissue, and Sperm to be in the Live sample class. They should not be deleted." The entire window is enclosed in a light blue border.

To add a sample type, first select the sample class it should belong to from the drop down at the top of the screen. Then enter the value in the Sample Type text box and click Add. To rename one, double click it, update the name, then click Rename. To remove one, highlight it and click Delete. A sample type may not be removed after it has been used for samples.

Note that technical matings require the "Live" sample class to contain the following Sample Type terms: Egg, Embryo, Ovary, Ovarian tissue, and Sperm. These are initially added by either the installation or upgrade of JCMS. Deleting any of them will cause there to be no sample choices for technical matings

**Figure 20-4 Manage sample types screen**

To administer preservation vocabularies, click the Preservation button on the Administrator button bar. This window is displayed:

The screenshot shows the 'ManagePreservation : Form' window. At the top, there is a dropdown menu labeled 'Sample Class' with the value 'Live'. Below this, there are three main sections: 'Preservation Type' (containing 'Liquid Nitrogen'), 'Preservation Method' (containing 'Protocol-231' and 'Protocol-1972'), and 'Preservation Detail' (containing 'Version 18.01' and 'Version 18.02'). Each section has a corresponding text input field below it, and a group of buttons ('Add', 'Rename', 'Delete') to its right.

**Figure 20-5 Manage preservation vocabularies screen**

It is important to remember that, on this screen, the values in a list are relevant only to the selection made in the list to its left, and all values pertain to the selected sample class at the top of the screen. The preservation methods listed pertain to the selected preservation type, and the preservation details pertain to the selected method. To add any value, type the value in the text box below the list and click the Add button. To rename a value, double click it, update the value in the text box and click Rename. To remove a value, highlight it and click Delete. As with other vocabularies, an item may not be removed if it has already been used in a sample record.

Finally, to administer locations in your facility, click the Locations button on the Administrator button bar. Following is the location administration screen:

The screenshot shows the 'ManageLocations : Form' window. On the left, there is a 'Storage Locations' tree view with nodes like 'Storage Facility', 'Lab', and 'Refrigerator'. Above the tree view are buttons for 'Add', 'Rename', and 'Delete', and a text input field labeled 'New Field'. On the right, there is a large text area labeled 'Location Description' with a 'Update Description' button at the bottom.

**Figure 20-6 Manage sample locations screen**

This screen is organized as a “tree view” with “nodes” representing locations. You may expand and collapse the nodes in the tree, and add and remove nodes at any level. For example, you

may have a building node under the root level, and beneath that node are room nodes, beneath that may be shelf nodes. This allows you to be as general or specific about locations as necessary. To add a new value, highlight the parent value, type the name in the New Field text box, and click Add. Following the previous example, to add a new room to a building, highlight the building, enter the name of the room, and click Add. To rename a value, highlight it, enter the new name in the New Field text box, and click Rename. To remove a value, highlight it and click Delete. Enter text descriptions for any node by typing in the Location description box on the right hand side of the screen and click Update Description.

## 20.2 Pooled Samples

A *pooled sample* is one from multiple sources. These sources may all have the same strain or have differing strains (*Relax Single Strain Constraint*). When pooling samples from sources from multiple strains, it is suggested to set up a mechanism to help readily identify and search for such pooled samples. Set up a dummy Strain. The dummy Strain is not associated with any actual mice or matings; it is used only as the Sample's assigned Strain. This enables easily identifying all multi-strain pooled samples using the Sample Query form.

The screenshot shows the 'Add or Edit Strain' dialog box with the title bar 'Add or Edit Strain (mtsadmin)'. The main area contains the following fields:

- \*Strain:** Mixed
- \*Frozen Embryos:** 0
- FE Max Generation:** (dropdown menu)
- \*Frozen Sperm:** 0
- FS Max Generation:** (dropdown menu)
- \*Frozen Ovaries:** 0
- FO Max Generation:** (dropdown menu)
- Strain Status:** A
- Section:** (dropdown menu)
- Min. Tag:** (text box)
- Max. Tag:** (text box)
- Last Tag:** (text box)
- JR # / Stock #:** 0
- Card color:** (dropdown menu)
- Strain type:** (dropdown menu)
- Comments:** Used only for multiple-strain pooled samples.

Below this section is a heading 'Line Viability Constraints' followed by two groups of constraints:

- YELLOW** group:
  - Minimum Number of Males: (text box)
  - Minimum Number of Females: (text box)
  - Maximum Age in days for Males: (text box)
  - Maximum Age in days for Females: (text box)
- RED** group:
  - Minimum Number of Males: (text box)
  - Minimum Number of Females: (text box)
  - Maximum Age in days for Males: (text box)
  - Maximum Age in days for Females: (text box)

At the bottom of the dialog box are navigation buttons: Record: 14 < 12 of 12 > No Filter Search.

Figure 20-7 Dummy strain for a pooled sample

## 20.3 Add Samples

**Figure 20-8 Add Sample form**

The Add Sample form is accessed by clicking the Add Sample button from the Samples tab. Use the Add Sample form to add as many samples during a session as necessary. No sample information is saved to the database until you click the Submit button at the bottom left of the screen.

This screen is divided into two sections, the top part is for identifying the source of the sample and the bottom part is for the sample data. Each section contains a grid that can hold one or more items. It is important to understand that a sample or samples may have one or more sources, and a source or sources may have one or more samples associated with it. You may view samples previously entered for the source(s) by clicking the Show Existing Samples button.

The source(s) of a sample may be another sample, a mouse, mating, or litter. There is also an *Other* option; if this is selected, only the strain is saved as the source of the sample. Except for the *Other* option, when you make a selection for the source type, the ID of Source drop down list is populated with ID's relevant to the source type. For example, if you select a source type of Mouse, the ID's of Source drop down list will contain mouse ID's.

By default, the *Sample Strains* list is populated with strains relevant to the *Source Type*. Selecting a strain from the *Sample Strain* list in turn filters the list of *Source IDs* for those associated with the specified *Sample Strain*.

A check box is provided (below the *ID of Source* selection list) that relaxes this constraint, allowing Users to create samples that reference multiple sources from different strains. When checked, the *Sample Strain* list shows all defined strains and the selection of one does not restrict the IDs in the *ID of Source* list. The *Sample Strain* value becomes the Strain value for the Sample. The *Source* record's Strain value may differ from the Sample's Strain value. See section 20.2 Pooled Samples for more information about pooled samples.

To select an item from the list as a source, select it in the *ID's of Source* drop down list and click the ">" button to move it into the source grid. The source grid indicates the source(s) that have been selected for this sample. If multiple sources are indicated, they must be of the same type. For example, both a mouse and a litter may not be indicated as the sources for a sample. In addition, the sources selected must be of the same strain when the *Relax Single Strain Constraint* is not checked (the default). To remove a source from the selected sources grid, highlight it and click the red "X" button.

After identifying the source(s), enter the sample information. Each sample is identified by a Sample ID and these must be unique. Enable the system to generate these by checking Auto-generate ID. When using this, indicate the format to use with the first sample entered, and the system will then use that format for subsequent samples. For example, to enter five samples in the format S\_01, S\_02, S\_03, S\_04, and S\_05, enter the first one (S\_01) and the system will increment the numeric digits for the remaining samples. Required data entry fields are identified with asterisks. Indicate the sample's class and type (i.e. Live, Embryo) and the values for the sample type will be based on the selected sample class. Values in the drop down lists for Preservation Type, Method, and Detail are dependent on the selection for Sample Type.

Note that the sample date, weight, and age have both a data entry box and one or more drop down lists to qualify the value. For example, enter a value for age, and then use the drop down lists to indicate the time units and how the age is measured. The Calculate Age button may be used to automatically generate the age based on the relevant date of the source. Relevant dates are birth date for mouse, mating date for mating, and litter born date for litter. If more than one source is indicated, and the relevant dates are different (i.e. two mice with different born dates,) the Calculate Age button will be disabled.

The sample's location is specified by using the location "Tree" at the bottom left side of the screen. Expand the tree by clicking on the "+" to the left of any branch of the tree, allowing you to drill down to whatever level of detail is necessary. Select any branch of the tree at any level, which allows saving the location from something as general as a building, to something as specific as a box on a shelf in a room in a building.

After entering the sample information, use the ">" button next to the sample grid to move the sample into the grid. Then repeat the process to add additional samples to the grid. All samples added to the grid will be associated with the source(s) identified in the first step. To remove a sample from the grid, highlight it and click the red "X" button. Note that both the source and samples grids have a Clear Grid button that will remove all entries from that grid. Edit a sample in the grid by highlighting it and clicking the "<" button, updating the data on the left, and then clicking the ">" button.

When the samples grid is complete and you are ready to save, click the Submit button. This enters the samples into the database and the rows in the grid will turn green to indicate that they were saved successfully. Then either close the form or continue entering additional samples. The Clear button at the bottom of the screen will clear all values so that you can start entering a new set of samples. Everything entered will appear in the Samples Entered this Session list.

## 20.4 Edit Samples

The screenshot shows the 'Edit Sample' window with the following details:

- Sample ID:** DOE-embryo-016 (dropdown menu with 'Change ID' button)
- Owner:** nobody
- Sample Class:** Live
- Sample Type:** Embryo
- Description:** (empty text area)
- Sample Date:** 3/20/2013 (dropdown menu) / Date Harvested (dropdown menu)
- Harvest Method:** Extraction protocol 354
- Weight:** 0 gram (text input) / Calculate Age (button)
- Age:** 2 Days (text input) / From Conception (dropdown menu)
- Preservation Type:** Frozen
- Preservation Method:** Liquid nitrogen
- Preservation Detail:** (dropdown menu)
- Sample Status:** Unprocessed
- \* Location:** Storage Facility > Building 51 > First Floor > 1000 > Freezer B > 1010 > Second Floor
- Samples Entered this Session:** (Large grayed-out area)
- Source of Sample:**
  - \*Source Type:** Parent Sample (radio button), Mouse (radio button), Litter (radio button, selected), Mating (radio button), Other (radio button). Buttons: >, X.
  - Sample Strain:** DBA/2J
  - \*ID of Source:** 103

Litter	Birthday	Strain	Total Born	Status	Harvest Date	Number Harv
103		DBA/2J	0	H	3/20/2013	
- Buttons at the bottom:** Submit, Clear, Show/Print Session Report, Print Sample Labels, Set Genotype.

**Figure 20-9 Edit Sample Form**

The Edit Sample form is accessed by clicking the Edit Sample button on the Samples tab. All data about a sample, including its source(s), may be updated on the edit sample screen. This screen is used to update one sample at a time. The first step is to select the Sample ID of the sample to be updated from the Sample ID drop down list at the top of the screen. After selecting a sample, make whatever updates are necessary. The data entry fields on this screen function the same way as their corresponding data entry fields on the Add Sample screen. After the information has been updated, click the Submit button to save the changes.

## 20.5 Bulk Change Samples

Sample ID	Owner	Sample Status	Storage Facility	Sample Date	Harvest Method
SS002	nobody	Unprocessed	Building 5	12/10/2008	

**Figure 20-10 Bulk Sample Update Form**

JCMS Sample Tracking provides a specialized screen for bulk updates to a group of samples to change their status, location, and/or owner. To get to this screen, click the Bulk Sample Update button on the Samples tab.

The top half of this screen is used for indicating the group of samples to be updated. This is done by searching for samples by Sample ID, selecting one or more samples from the search results, and moving them into the selected samples grid. You may enter a full or partial sample ID in the Sample ID box and then click the Search button. Samples matching the search criteria will appear in the search results list. Select one or more samples by highlighting them and click the ">" button to move the samples into the selected samples grid. You may perform as many searches as necessary to fill the selected samples grid. As on the Add Sample screen, you may remove a sample by highlighting it and clicking the red "X" button.

On the bottom half of the screen, select one of the update options, and then select the new value to be applied. When the submit button is clicked, the updated value will be applied to all samples in the grid. You may make additional updates if necessary. For example, to change a status and location for a group of samples, you would first select Sample Status and pick the new status and click Submit. Then you would select Sample Location and pick the new location value and click Submit.

Only samples belonging to the logged on user may be transferred to a new owner using this form. If ownership transfers by an administrator from one owner to another are desired, the Bulk Change of Ownership form, under Admin Functions, is used.

## 20.6 Query Samples

The screenshot shows the 'Query Samples' form with the following details:

- Source Type:** Mouse
- Source ID's:**
  - Any (radio button selected)
  - Range (radio button)
  - Min: JANE-001
  - Max: JANE-001
  - ID Like: (text input)
- Sample ID's:**
  - Any (radio button selected)
  - Range (radio button)
  - Min: New-100
  - Max: New-100
  - ID Like: (text input)
- Harvest Method:**
  - Any (radio button selected)
  - Select from List (radio button)
- Sample Class:**
  - Any (radio button selected)
  - Select from List (radio button)
  - Histology (dropdown menu: Histology)
- Genotype:**
  - Any (radio button selected)
  - Select from list (radio button)
  - AND from second list (radio button)
  - Clear allele selections (button)
- Source Age:**
  - Any (radio button selected)
  - Range (radio button)
  - ≥ (text input)
  - ≤ (text input)
- Source Sex:**
  - Any (radio button selected)
  - Select from List (radio button)
  - F (checkbox)
  - M (checkbox)
  - (checkbox)
- Sample Strain:**
  - Any (radio button selected)
  - Select from List (radio button)
  - SampleStrain (dropdown menu: Blia.ENU23XD2Gpnmb(wt))
- Weight:**
  - Any (radio button selected)
  - Range (radio button)
  - ≥ (text input)
  - ≤ (text input)
  - gram (dropdown menu: gram)
- Age:**
  - Any (radio button selected)
  - Range (radio button)
  - ≥ (text input)
  - ≤ (text input)
  - Hours (dropdown menu: Hours)
  - From Birth (dropdown menu: From Birth)
- Sample Date:**
  - Any (radio button selected)
  - Range (radio button)
  - ≥ (text input)
  - ≤ (text input)
  - Hours (dropdown menu: Hours)
  - From Birth (dropdown menu: From Birth)
- Sample Status:**
  - Any (radio button selected)
  - Select from List (radio button)
  - Unprocessed (dropdown menu: Unprocessed)
- Sample Type:**
  - Any (radio button selected)
  - Select from List (radio button)
  - Brain (dropdown menu: Brain)
- Preservation Type:**
  - Any (radio button selected)
  - Select from List (radio button)
- Preservation Method:**
  - Any (radio button selected)
  - Select from List (radio button)
- Location:**
  - Any (radio button selected)
  - Select from Tree (radio button)
  - Storage Facility (checkbox)
    - Building 51 (checkbox)
      - First Floor (checkbox)
      - Second Floor (checkbox)
- Preservation Detail:**
  - Any (radio button selected)
  - Select from List (radio button)

Figure 20-11 Query Samples Form

This screen is accessed by clicking the Query Samples button on the Reports tab. It is used for producing a query and obtaining results for samples. It allows maximum flexibility, as any combination of sample information may be selected to filter the results, and there is control over which output values are viewed. To see all results for a specific search criterion, leave the selection at the default "Any." To narrow results, select a value either by typing it in or selecting it from the drop down list where applicable. Where possible, a range of values may also be entered. Use the check box list along the right hand side of the screen to select which values to include in the results.

Search criteria on this screen may be enabled or disabled based on a selection made in another area. This happens if the two search criteria are dependent on each other. For example, sample type and preservation information are dependent on the sample class. When "Histology" is selected for sample class, the sample status, sample type, and preservation type areas become enabled. When a specific preservation type is selected, rather than "Any," the preservation method area becomes enabled, and so forth. This allows "drilling down" into the dependent information.

When the query selections have been made, click the Run Query button to see the results in tabular format.

Note: the Strain criterion for Sample searches is matched **only** to the Sample's Strain, not to the strain associated with individual sample sources. When pooled sources span multiple strains, the search does not include the source's strains, i.e., it does not answer the question "show me samples where any constituent source is from strain X." See section 20.2 Pooled Samples for more information about pooled samples.

### 20.6.1 Searching for sample genotypes

**Genotype**

- Any
- Select from list
- AND from second list

**Clear allele selections**

**FSN**

+	+
-	-

**SKID**

+	+
-	-

Up to two genes with allele combinations may be selected as criteria. When two genes are selected, results are returned only if the sample is typed for both genes. When no alleles are highlighted, the results will contain any allele combination for the gene. Use the *Clear allele selections* button to remove the allele criteria. In this example, samples are returned that have the genotype FSN +/- AND any SKID genotype.

Figure 20-12 Sample genotype criteria

**Select Query**

**Results to Show**

**Clear All** **Select All**

Sample ID  
 Associated documents, Max of 1  
 Sample Class  
 Sample Type  
 Sample Date  
 Harvest Method  
 Weight  
 Age  
 Preservation Type  
 Preservation Method  
 Preservation Detail  
 Sample Status  
 Sample Source(s)  
 Location  
 Genotype  
 Restrict output to show only genotypes in the criteria  
 Strain

**sampleQuery35281**

SampleID	'FSN'	'SKID'	MoreGTs
Test-001	FSN +/-	SKID -/-	More GTs

The genotype does not have to be included in the results. When it is, there are two options. Checking "Restrict output to show only genotypes in the criteria" will create a separate output column for each gene, plus a third column called "MoreGTs" that indicates if the sample has been typed for other genes.

Figure 20-13 Sample genotype results in multiple columns

When "Restrict output to show only genotypes in the criteria" is not checked, all genotypes are displayed in one column separated with commas. Genotype results are always returned this way if there are no genotype criteria.

SampleID	Genotype
Test-002	FSN +?/-, new.one new.alle/, SKID +/-, T +/-
Test-003	SKID +/-
Test-004	SKID +/-
Test-005	FSN +/?+, SKID +/+
Test-100	
Test-101	
Test-102	
Test-103	

**Figure 20-14 Genotype results when no gene criteria**

## 20.7 Browse Sample Storage Locations

Use this form to display a listing of all samples in a particular location.

**Figure 20-15 Browse Sample Storage Locations Form**

Check "Recurse" to see all samples in nodes below the selected one.

## 20.8 Print Sample Labels

A “Print Sample Labels” button found on the Main Menu reports tab, Add Sample, Add Embryo Litter, and Edit Sample forms will open the Print Sample Labels form, shown below in Figure 20-16. Use this form to print samples selected by searching on the Sample ID value. First configure the label parameters by indicating the height of the label, font size, and whether you want text or bar coded labels. After that, select the samples to print. Similar to the mechanism for selecting samples to edit, enter a full or partial sample ID value, click “Search,” and then select samples for printing based on the search results. To select a sample for printing, highlight it in the search results list and click the right arrow (“>”) button. Multiple values may be selected by using the shift (consecutive select) and control (non-consecutive select) keys. Samples may be removed from the samples to print list by highlighting them and clicking the remove (“X”) button, or clear the entire list by clicking the “Clear List” button. After setting the label parameters and selecting samples to print, click the “Preview” button. This will bring up the print preview window, and if everything looks correct, print the labels on the printer from there.

See Section 2.3.11 for instructions on obtaining a bar code font in order to use bar code labels.

The screenshot shows the 'Print Sample Labels' dialog box. At the top, it displays 'Wean Work Report (mtsadmin) {Secretaries and Owners}' in the title bar. The main area is divided into two sections: 'Select label format' and 'Select samples for printing'. In 'Select label format', the 'Label height' is set to '0.5 inche(s)' and the 'Font size' is set to '10'. There are two radio buttons: 'Print text labels' (selected) and 'Print bar code labels'. In 'Select samples for printing', there is a 'Sample ID like:' input field with a placeholder '(empty)', a 'Search' button, and a 'Resulting Rows Found : 5' label. Below this is a 'Search results:' list box containing five items, with a right arrow ('>') button to move selected items to the 'Samples to print:' list box. The 'Samples to print:' list box contains one item, with a red 'X' button to remove it. A 'Clear List' button is also present. At the bottom, there is a 'Preview' button and a navigation bar with 'Record: < < 1 of 1 > >> No Filter Search'.

Figure 20-16 Print Sample Labels

## **21 FAQ's (Frequently Asked Questions)**

### **21.1.1 JCMS starts up and shows the database, but no welcome window appears.**

Find the answer to this question in the Configuration Issues section 2.3.13.7.

### **21.1.2 Every time I add, edit, or delete a record in JCMS I'm prompted with a dialog box.**

MS Access has an option that allows this feature to be turned on or off. Section 2.3.5 describes how to configure these database options.

### **21.1.3 What to do about an “end/debug” error message**

Occasionally an error dialog box will be displayed with two buttons, *end* and *debug*. This dialog box is displayed only when the program encounters a serious bug. If you see this dialog box you should write down a description about what you were doing at the time and the exact text of the error message. Then contact your Administrator as soon as possible. If your Administrator is not available, select the “end” button and close any program code windows that are left open. You can check your data entry via the various JCMS query forms. If the data looks okay, then continue to work, but be sure to report the bug so it can be fixed in the next release.

Oftentimes, the bug is a minor thing that can be worked around until the next release. If you detect a data problem, stop all data entry and report the problem immediately.

### **21.1.4 Error messages when editing date fields**

Why are date fields giving error messages when edited?

Problem: If the format the dates are in is MM/DD/YYYY (a 4 digit year), when this is edited, JCMS expects the date to be in the format MM/DD/YY (a 2 digit year) and gives an error. Most of the dates have now been changed to use a pick control, eliminating this problem.

Solution: Remember to highlight the whole field when editing and enter the date with a two digit year. A second, more permanent solution is to change the computer to use two digit dates. From the Windows Start button, choose Settings – Control Panel. Double click on the Regional Settings Icon. Click on the date tab. Change the “Short date style” to mm/dd/yy. This will change how all dates on this computer are displayed.

### **21.1.5 Error message: “user Admin does not have permission to use this form”**

JCMS does not recognize user *Admin* as a JCMS Administrator. In fact, JCMS prevents user *Admin* from working in JCMS at all. In order to use JCMS you need to be logged on as *mtsadmin*, an owner, or a secretary of an owner. NOTE: Microsoft Access will not prompt for a logon unless user Admin has a password. So assign a password to the Admin user before using JCMS. See the Installation section 2.2.3 on initializing passwords for more information about user accounts.

### **21.1.6 Error message about “could not find file”**



**Figure 21-1 Could Not Find File Error**

The most common reason for this error message is that the file server is not available over the network. JCMS is looking for the linked data tables located on the file server. This message also occurs if the data tables have been moved and the JCMS access interface has not been updated to re-link the tables.

### **21.1.7 A note about session boxes**

Session boxes provide a history list for the data entry person. MS Access limits the size of this list to 2048 characters long (including some format characters that do not display in the box). If the session box string grows to long, JCMS will warn that the string will be reset soon. You can keep working and the session box will reset before the MS Access limit is reached. The only loss to the user is the history list starts over again. NO DATA IS LOST when you see the warning message or when the session box is reset.

### **21.1.8 List boxes of mice information are scrambled!**

On several forms, there are list boxes with rows of mouse information or session boxes. The rows of data in these list boxes sometimes get messed up if there are semicolons or commas in the data. The easy fix: remove any commas or semicolons from the terms in the controlled vocabulary tables. If this doesn't solve the problem, report the bug to the Jackson Laboratory.

### **21.1.9 Warning message about not saving a record.**



**Figure 21-2 Error: You can't save this record at this time**

This message appears when there was an error on the form that was not corrected. This type of error may be something such as an invalid date or entering something that is not in a pick list. Answer yes if the information on the form was not supposed to be submitted / entered into

the database and the form should just be closed. Answer no if the information needs to be entered and correct the information on the form, then resubmit it.

### **21.1.10 The database keeps telling me it is “Read only”**

Logged in with administrator permissions, navigate to the folder containing your JCMS files. For example c:\Program Files\The Jackson Laboratory\JAX-CMS\. Right click on the folder containing the files (For example: JAX-CMS). Select ‘properties’. Open the ‘Securities’ tab. Give ‘Users’ Read/Write privileges.

### **21.1.11 The Connector/ODBC Window is Popping Up**

If this occurs when using the Experiment Work Report, edit the MySQL setup variables to reflect the current status. See Section 19.16.1 for more details.

## 21.1.12 Other FAQs

Also check the Configuration issues in section 2.3.13, the online [JCMS Discussion Forum](#), and the online colony management website [FAQs](#).

# 22 Technical Guide

## 22.1 Security

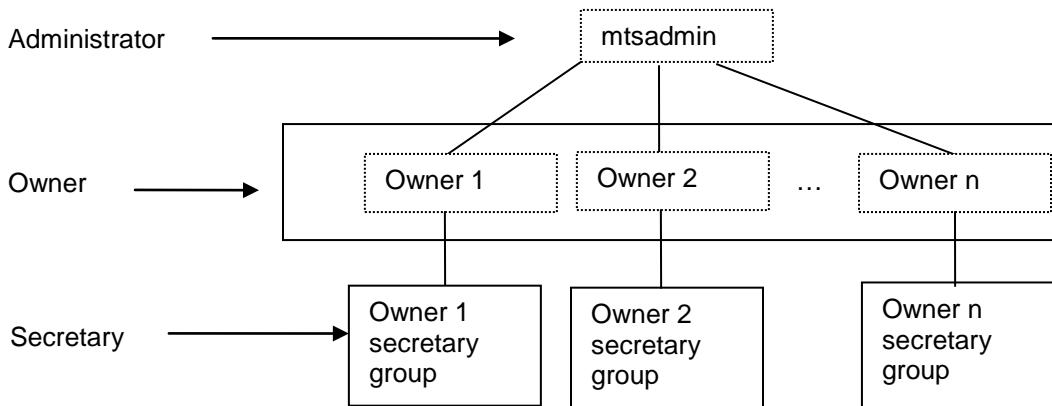
The JCMS security implementation is based on the premise that people who have access to JCMS are not malicious. JCMS security is therefore limited to trying to prevent accidental changes to the database.

JCMS has two components, an interface component and a database component. The interface component is installed on all workstations that require access to the database. Each installation of the interface component has its own set of user accounts.

Data security is implemented as a hierarchy with two parameters: access level (horizontal) and ownership (vertical). Higher access levels inherit all access permissions from lower access levels plus additional permissions associated with the higher level.

The vertical access parameter (ownership) comes into play only when a user attempts to submit a record-update or a record-addition to an ownership-protected table in the database.

We define three security access levels: Administrator, Owner, and Secretary. In the diagram below, groups are illustrated in solid outlined boxes. An individual logon account can belong to only one access level. In practice this means that secretaries can belong to more than one secretary group (secretary level is the only level that has more than one group associated with it). Also note there is exactly one mtsadmin.



**Figure 22-1 Administrator - Owner - Secretary Security Relationship**

Individual secretaries can belong to many secretary groups. Thus, it is possible to for an owner to share the services of a single secretary; however, you need to be careful. Any secretary that belongs to more than one owner-secretary group will have the ability to edit records owned by all owners associated with the secretary in the same edit session. It may be a better administrative policy to give one person more than one access account (log on name) if needed.

Several tables in the database have an owner field: Mouse, Mating, ExpPlan, ExpData, and Sample. Litters inherit ownership from their mating. When editing involves Mouse, Mating, Litter, Experimental Plan, Experimental Data, or Sample, access to the records is restricted by owner.

## **22.2 Changing Security Access to Forms**

Access to the database is restricted by the form interface. That is, each form is assigned an access security level. Users *cannot* use forms that have a higher security level than their assigned level. The opposite is also true; users can use all forms with security level less than or equal to their assigned security level. The mtsadmin has the highest access level and therefore has complete access to the database.

The DbFormPrivileges table controls the access security level (or privilege level) for certain forms. The administrator may change the level between secretary, owner, and administrator for these forms by using the Set Form Access Privileges button on the Administrator button bar. The forms not listed in this table are required to stay at a certain security level. Any changes will take effect the next time JCMS is started by a particular user.

## **22.3 Data Integrity**

Data integrity is only loosely enforced. If all data are entered through the user interface forms, then data integrity will not be a problem. There is one important exception. It is possible for users to conflict in their usage of the database. If multiple users access the database simultaneously and change records there is the possibility of conflicting edits.

## **22.4 Back up the Database**

Having a backup strategy that is followed regularly is very important. Routinely make full copies of the JCMS installation folder and all of its contents. Be sure when backing up to copy the interface (JCMS.mdb) file from wherever you have installed it.

MySQL backups should be conducted by your MySQL system administrator according to published industry recommendations.

## **22.5 The dbInfo Table**

The dbInfo table has release information in it as well as counters for maximum pen ID and maximum auto mouse ID number and maximum auto litter ID number.

## **22.6 Access to the Primary Data Tables**

JCMS allows access to the primary tables through a set of data-sheet forms. Use this level of access to the data only sparingly to correct mistakes in data entry. Entering or changing data this way can cause the data to have multiple problems because the normal forms interface enforces business rules which will not be enforced here.

From the Administrator button bar (click the Administrator button on the Administrator tab), click the button labeled *Access to primary data tables*.

Each table can be opened in "datasheet view," which appears similar to a spreadsheet. Use the scroll bars to navigate. Click on the column headers and drag to resize them if some of the information in the field is not visible.

Note: the Pen Group table is maintained for historical purposes for those users who upgraded from releases prior to 4.0.0. Changes to this table will have no effect on current pens and rooms.

### **22.6.1 Editing Records in Datasheet View**

Any field that is typed in will be changed by the database as soon as the cursor moves into another field. Some changes will not be allowed and will generate an error message.

Add records by scrolling to the bottom of the list and entering the new record into the bottom row. It is not allowed to add the record unless values are entered into all of the required fields. Adding will occur when the mouse is clicked on a different row. A **key** must be entered into any field with a name that begins `_xxx_key`. The key should be a number one larger than in the previous row. This is an internal number that normally is not visible. The database is using this key field to keep track of the records. When the forms are used, the database is generating this number automatically. When the tables are used in datasheet view, this number must be added manually.

To delete records from the table, click on the box at the left side of the row in the table. The whole row will be highlighted. Press the delete key. A dialog box will ask confirmation that a record should be deleted.

## **22.7 Temporary Tables**

JCMS creates temporary tables for storing query results that are generated by the user query forms (e.g. query matings) and some of the experimental data forms. Temporary tables are bound to the query output form (i.e., the results of the query) or other screen forms. When the query results form is deleted or other forms are closed, the temporary tables can be removed.

When JCMS starts up it attempts to delete all temporary tables. If a table is still bound to a form then it may not get deleted. An Administrator can delete temporary tables, but be aware that if one of the client installations currently has a query results form bound to the table, the temporary table could be locked (undeletable).

MTSTemp – These temporary tables are named MTSTempX. X in the name MTSTempX is an integer that makes the current temporary table name unique in the tables collection. Temporary tables that are not bound to any form are deleted on startup.

## **22.8 Temporary Queries**

The cage use summary report creates temporary queries named `__BillingReportQueryxxxx` and the Query Samples form creates temporary queries named `__sampleQueryxxxx`. These are deleted when JCMS starts up.

## **22.9 Screen Resolution**

The forms used by this system are rather large and take up quite a bit of screen space. Every effort has been made to make them fit on a normal screen. Many just fit with a screen resolution of 800x600 on a 17" screen. For a smaller screen, try using a higher resolution such as 1024x768. Otherwise, some forms will have to be scrolled to see the entire contents.

## **22.10 Printer Notes**

The cage card reports use the default printer and default paper location. Many modern printers will use the sheet feeder as the default whenever there is paper in the sheet feeder. To print cage cards, open the sheet feeder and load the cards into the envelope feeder part of it. Also set the printer to use as straight a paper path as possible. Many printers have an option for sending sheets out the back if it is open or have a toggle switch to change the path.

The cage cards are designed to print on either the upper left side or center of the paper depending on the card format used. If the envelope feeder places the cards in another location, a programmer will have to change the margin settings to match the location of the envelope feeder.

It is possible to have a programmer customize what printer and paper location is used by changing the File-Page Settings information in the Design View for an individual report.



## 23 Appendix 1: Automated TGS Genotype Loader Submissions

JCMS has an automated interface into The Jackson Laboratory's Transgenic Genotyping Service (TGS). This interface is only visible to users whose setup variable JCMS\_JAXLAB\_INSTALLATION is set to true.

### 23.1 Adding a TGS Genotype Request

Click on the "Add Genotype Request" button from the 'Manage Mice' tab.

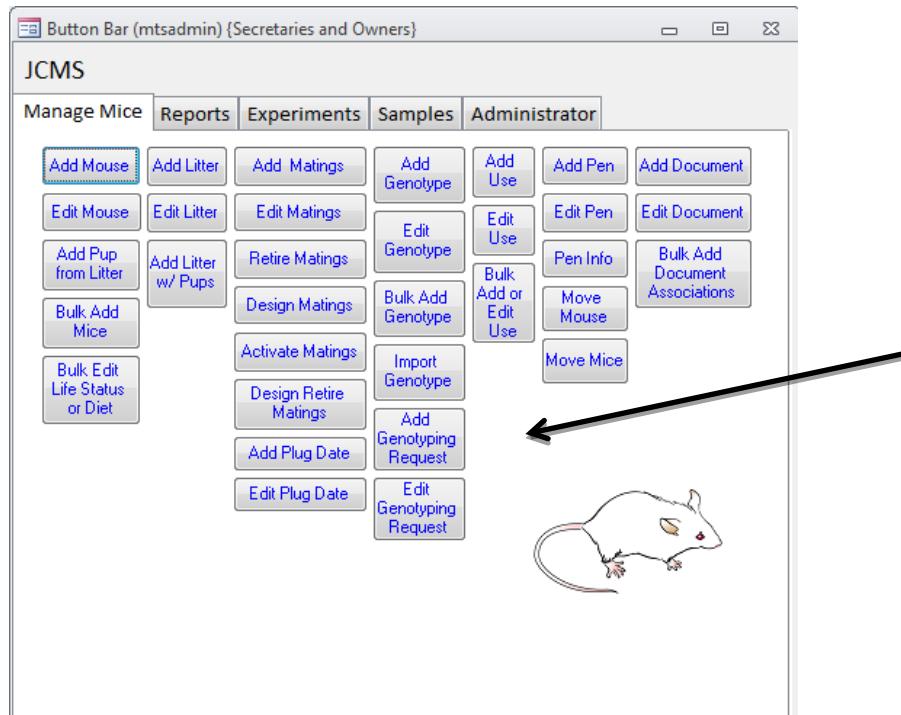


Figure 23-1 The Add Genotyping Request button

Add samples (mice) by

- Selecting the mouse ID
- Selecting a pattern of mouse IDs
- Select a pen
- Filtering by strains
- Filtering by life status

Below is an example of select all mice that begin with the letters "A1".

Mouse ID:  Mouse ID like: A1| Pen ID:

Stock #  Any  Selected from list

07-60	0
07-61	0
07-62	2921
07-63	6777
00-00	0
05-02	0
05-12	0
06-02	0
06-15	5810
06-16	n

Life Status  Any  Selected from list

A	Alive
D	Dead
K	Killed
M	Missing
S	Shipped
E	Euthanized
FD	Found Dead

**Run Query** **Run Query and Add iff One** **Clear Filters**

**Query Results (Mice):** 74 matches

Mouse ID	Pen ID	Date born	Age in Day	Sex	Owner	Status	Strain	JRNum
A1	46	12/11/2003	0	M	PGN	K	Rnl5	0
A10	177	8/23/2006	0	M	PGN	K	Rnl5	0
A11	178	6/20/2007	0	F	PGN	K	Rnl5	0
A12	178	6/20/2007	0	F	PGN	K	Rnl5	0
A13	178	6/20/2007	0	F	PGN	K	Rnl5	0
A14	179	6/20/2007	0	M	PGN	K	Rnl5	0
A15	179	6/20/2007	0	M	PGN	K	Rnl5	0
A16	179	6/20/2007	0	M	PGN	K	Rnl5	0
A17	179	6/20/2007	0	M	PGN	K	Rnl5	0
A18	180	8/21/2006	0	F	PGN	K	Rnl5	0
A19	180	8/21/2006	0	F	PGN	K	Rnl5	0
A103	6749	8/12/2009	0	F	PGN	K	08-43	0
A104	6749	8/12/2009	0	F	PGN	K	08-43	0
A105	6749	8/12/2009	0	F	PGN	K	08-43	0
A106	6749	8/12/2009	0	F	PGN	K	08-43	0
A107	6964	8/4/2009	0	F	pgn	D	08-47	0
A108	6964	8/4/2009	0	F	pgn	D	08-47	0
A109	6964	8/4/2009	0	F	par	D	08-47	0

**Add Selected**

**Figure 23-2 Select mice whose IDs begin with “A1”**

To place mice on the well plate, select them by single clicking on them. Then click “Add Selected”. If this is the first time a dialog will appear that will ask for an ID for the plate.

SelectGenotypingPlate

The mouse with ID 'A10' cannot be added to any plate currently in the request. Please enter data for a new plate. You may also choose to skip this mouse, or cancel (which stops adding any mice that haven't been processed yet).

New Plate ID:	<input type="text" value="AAAA"/>	Unfillable Region Row Start:	<input type="text" value="5"/>
Number of Columns:	<input type="text" value="12"/>	Unfillable Region Col Start:	<input type="text" value="12"/>
Number of Rows:	<input type="text" value="8"/>	Unfillable Region Row End:	<input type="text" value="8"/>
		Unfillable Region Col End:	<input type="text" value="12"/>

**OK** **Skip** **Cancel**

**Figure 23-3 Setting the plate ID**

Any number of plates can be contained within a request.

Samples are added to the well plate grouped by Stock number (JRNum). A column cannot contain samples from more than a single Stock number. The example below has mice from Stock # 000000 and Stock # 008003.

AAAA												
Plate Status: Open												
	1	2	3	4	5	6	7	8	9	10	11	12
A	A10 000000 (1)	A141 008003 (1)										
B	A19 000000 (2)	A142 008003 (2)										
C	A106 000000 (3)	A143 008003 (3)										
D		A144 008003 (4)										
E											[Unfillable]	
F											[Unfillable]	
G											[Unfillable]	
H											[Unfillable]	

**Figure 23-4 A well plate with samples from seven mice of two strains**

To remove samples (mice) select the cells to be deleted by holding down the shift key and clicking on the cells. (If you are only removing one cell then you do not need to hold down the shift key.) Then click “Remove Selected”.

Before saving a Genotyping Request ID must be entered.

Genotyping Request ID: <input type="text"/>											
Change ID											
7	8	9	10	11	12						

**Figure 23-5 The genotyping request ID field**

Save your work at any time by clicking on the “Save” button. To exit this form, click on the “Cancel” button.

To submit a request, first seal the plate. After a plate is sealed no more mice may be added.  
ie Query Results in the lower left or from the current plate on the right.

Figure 23-6 To seal the plate select “Sealed”

The plate is now ready for submission.

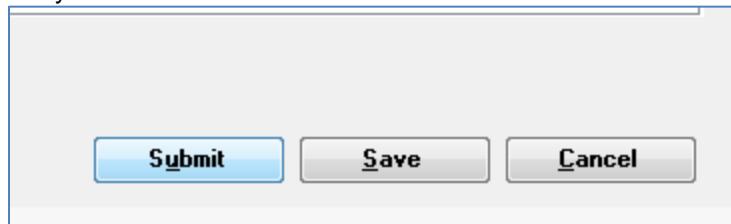


Figure 23-7 The Submit button

Upon submission a file name komp2typ.txt is written into the folder identified by the JCMS setup variable JCMS\_TGS\_REQUEST\_PATH. **Only one submission file at a time is allowed.** The TGS application will move the file when it is processed. If successful you will see some variation of the dialog box below.

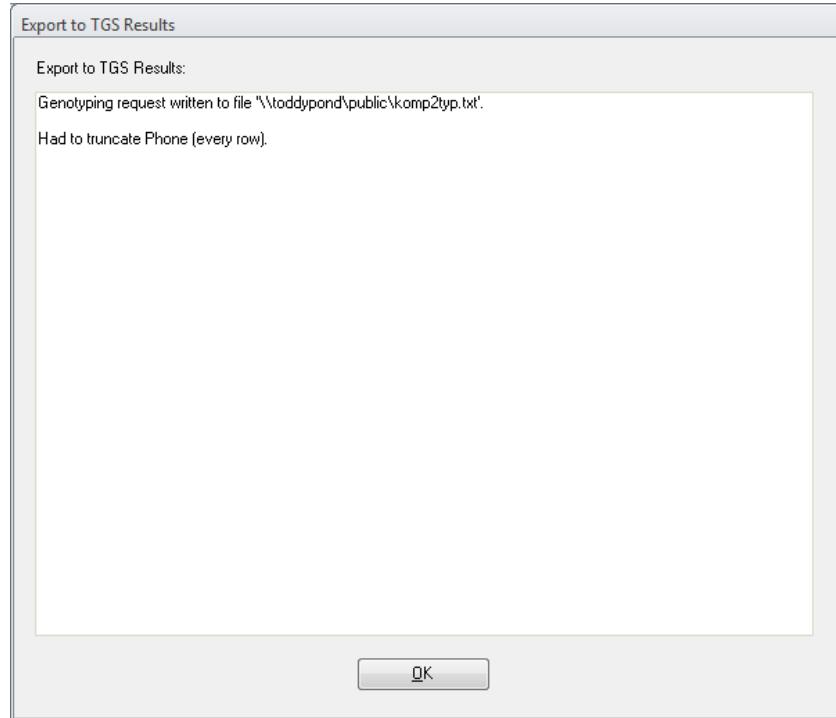


Figure 23-8 Notification of a successful submission.

## 23.2 Editing a TGS Genotype Submission Request

To edit a submitted or un-submitted genotype request click on the “Edit Genotyping Request” button on the ‘Manage Mice’ tab.

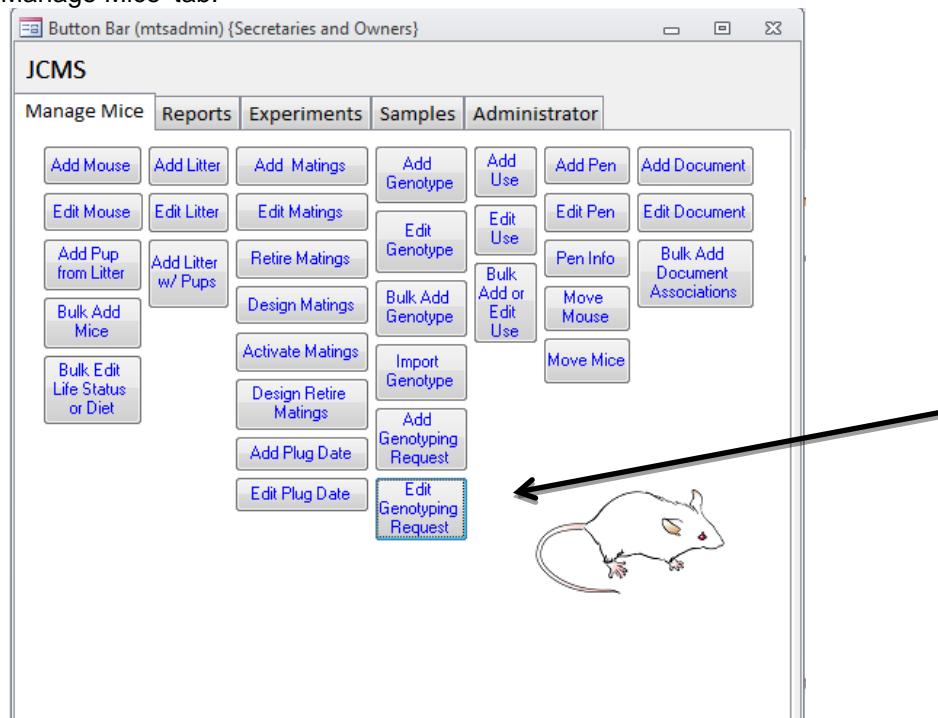


Figure 23-9 To edit an existing genotyping request

A selection dialog appears.

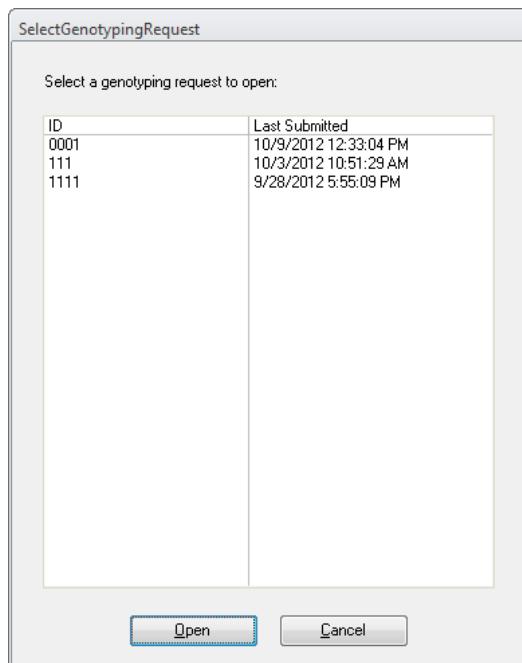


Figure 23-10 Select a request to be edited.

Select a request ID and click “Open”. The ‘Edit Genotype Request’ form will open. It operates exactly like the ‘Add Genotype Request’ form.



When the results are ready the person identified by the setup variable MTS\_PI\_NAME will receive an email.

### 23.3 Importing TGS Genotyping Results

Click on the “Import Genotype” button.

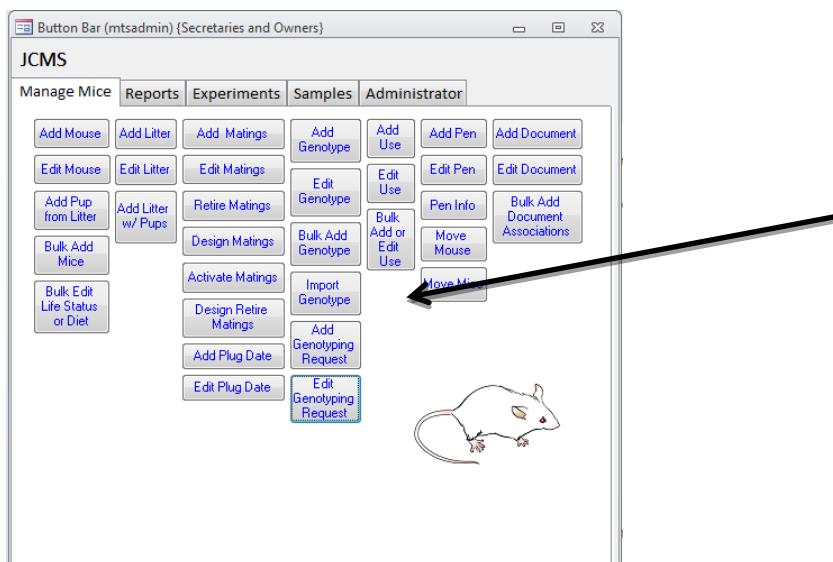


Figure 23-11 Import Genotype data

Select “Import TGS format file”. Click OK.

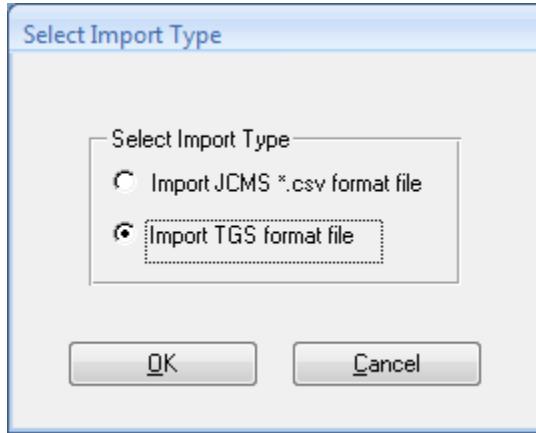
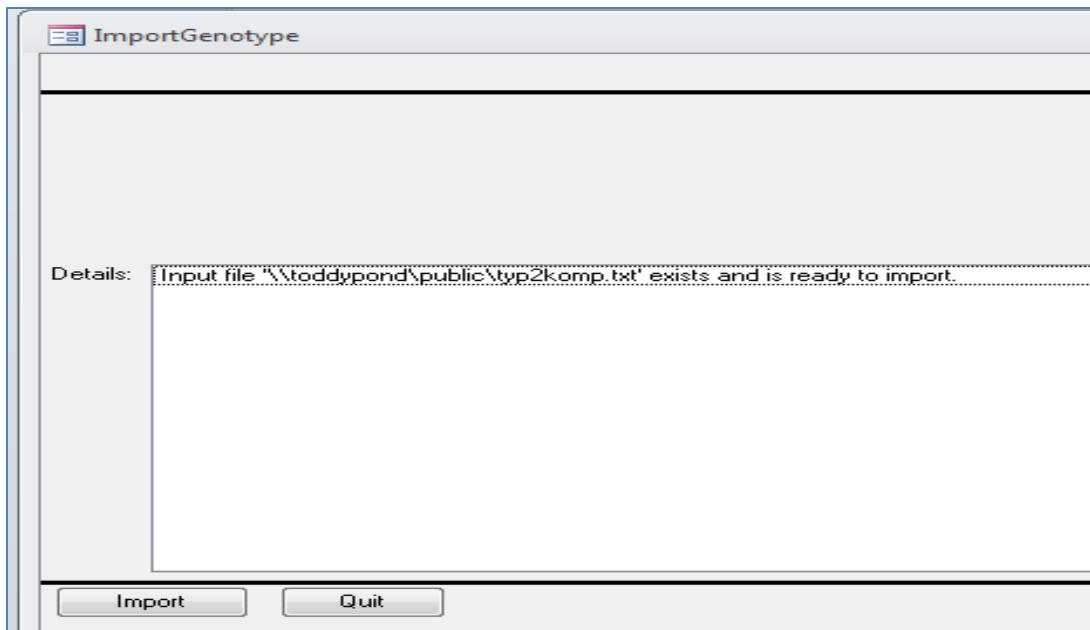


Figure 23-12 Select the format

It will take you to the folder identified by the setup variable JCMS\_TGS\_RESPONSE\_PATH. Select the file typ2komp.txt. When ready click the “Import” button.



**Figure 23-13 To import the results from the selected file.**

The results will appear in the 'Details' dialog box.



**It is up to the JCMS user to rename or delete the input file after a successful import. TGS will not overwrite the file.**