

*Study Data Specifications*

## Revision History

Date	Version	Summary of Changes
2004-07	1.0	Original version
2005-03-18	1.1	Addition of specifications for define.xml and SAS XPORT transport files specifications. Changes in document organization.
2006-03-04	1.2	Update information on annotated ECG waveform data. Delete ecg folder under Specifications for Organizing the Datasets.
2006-11-27	1.3	Addition of specifications for submitting tumor datasets (tumor.xpt) from rodent carcinogenicity studies.
2007-08-01	1.4	Addition of hyperlink to information for 3.1.1 datasets
2009-10-30	1.5	Modified introduction. Additional specifications for submitting data tabulation datasets. Additional specifications for analysis datasets. Revision of maximum file size restrictions. Addition of hyperlink to information for 3.1.2 datasets.
2010-01-04	1.5.1	Modified directory tree structure for organizing datasets and made minor technical corrections.
2011-06-22	1.6	Addition of specifications for general toxicology and carcinogenicity data tabulation datasets

## STUDY DATA SPECIFICATIONS

These specifications are for submitting animal and human study datasets in electronic format. Datasets are views of the study data used by reviewers to conduct specific analyses of the study data. They may include both raw and derived data. Because of the unpredictability of the scientific review process, it is impossible to enumerate *a priori* all datasets needed for review. Prior to the submission, sponsors should discuss with the review division the datasets that should be provided, the data elements that should be included in each dataset, and the organization of the data within the file. Additionally, not all FDA centers have adopted all aspects of these specifications; sponsors are advised to discuss with the reviewing division data needs prior to preparing data for submission.

### SAS XPORT TRANSPORT FILE FORMAT

SAS XPORT transport format, also called Version 5 SAS transport format, is an open format published by the SAS Institute. The description of this SAS transport file format is in the public domain. Data can be translated to and from this SAS transport format to other commonly used formats without the use of programs from SAS Institute or any specific vendor.

#### Version

In SAS, SAS XPORT transport files are created by PROC XCOPY in Version 5 of SAS software and by the XPORT SAS PROC in Version 6 and higher of SAS Software. SAS Transport files processed by the CPORT SAS PROC cannot be processed or archived by the FDA.

Sponsors can find the record layout for SAS XPORT transport files through SAS technical support technical document TS-140. This document and additional information about the SAS Transport file layout can be found on the SAS World Wide Web page at <http://www.sas.com/fda-esub>.

#### Transformation of Datasets

SAS XPORT transport files can be converted to various other formats using commercially available off the shelf software.

#### SAS Transport File Extension

All SAS XPORT transport files should use *.xpt* as the file extension.

## Compression of SAS Transport Files

SAS transport files should not be compressed. There should be one dataset per transport file.

## Maximum Size of User Supplied Elements

Element	Maximum Length in Characters
Variable Name	8
Variable Descriptive Label	40
Dataset Label	40

## CONTENT OF DATASETS AND SIZE OF DATASETS

Each dataset is provided in a single transport file. The maximum size of individual dataset is dependent on many factors. In general, datasets other than SDTM datasets, should be less than 400 MB; SDTM datasets should not be divided. Datasets divided to meet the maximum size restrictions should contain the same variable presentation so they can be easily concatenated.

Datasets which are divided should be clearly named to aid the reviewer in reconstructing the original dataset, e.g., xxx1, xxx2, xxx3, etc. The files that have been divided and need to be concatenated should be noted in the data definition document. This documentation should identify the range of subject numbers (or other criteria used for division) in the label for each of the divided datasets. For further information on file size limitations for files submitted to CDER, contact [eData@fda.hhs.gov](mailto:eData@fda.hhs.gov), for files submitted to CBER, contact CBER.CDISC@fda.hhs.gov

## SPECIFICATIONS FOR SPECIFIC DATASETS AND DOCUMENTATION

Study data are provided using different presentations: Data Tabulation Datasets, Data Listing Datasets, Subject Profiles, and Analysis Datasets.

### Data tabulation datasets

#### *Definition*

Data tabulations are datasets in which each record is a single observation for a subject.

#### *Specifications*

Specifications for the Data Tabulation datasets of human drug product clinical and animal toxicology studies<sup>1</sup> are provided by the SDTM<sup>2</sup>. FDA centers and reviewing

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<sup>1</sup> Here, “drug product” also includes biologic products (submitted as BLAs) that are reviewed in CDER

<sup>2</sup> SDTM refers to the Study Data Tabulation Model (SDTM) developed by the Submission Data Standard working group of the Clinical Data Interchange Standard Consortium (CDISC). The

divisions differ in their use of SDTM data. CDER currently accepts SDTM datasets prepared in accordance with the SDTM IG or SEND IG versions listed in the following table. CBER is currently testing datasets prepared in accordance with the SDTM IG for clinical studies of biologic products. Individual centers or reviewing divisions may specify the version of SDTM needed for review (see below). Follow the corresponding hyperlink to view the appropriate SDTM and implementation guide.

SDTM Model Version	IG Version	Implementation Guide	Support Begins	Support Ends
1.0	SDTM IG 3.1	<a href="http://www.cdisc.org/models/sds/v3.1/index.html">http://www.cdisc.org/models/sds/v3.1/index.html</a>	2004-07-01	2010-03-31
1.1	SDTM IG 3.1.1	<a href="http://www.cdisc.org/content1605">http://www.cdisc.org/content1605</a>	2007-08-01	
1.2	SDTM IG 3.1.2	<a href="http://www.cdisc.org/content1055">http://www.cdisc.org/content1055</a>	2009-10-30	
1.2	SEND IG 3.0	<a href="http://www.cdisc.org/send">http://www.cdisc.org/send</a>	2011-06-22	

While the SDTM provides a valuable representation that may facilitate review, it does not always provide data structured in a way that supports all analyses needed for review. Sponsors should therefore augment SDTM with analysis datasets as described in the *Analysis datasets* section.

## Data Listings

### *Definition*

Data listings are datasets in which each record is a series of observations collected for each subject during a study or for each subject for each visit during the study organized by domain.

### *Specifications*

Each dataset is provided as a SAS Transport (XPORT) file. Currently, there are no further specifications for organizing data listing datasets.

## Subject profiles

### *Definition*

Subject profiles are displays of study data of various modalities collected for an individual subject and organized by time.

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model has two implementation guides: human clinical (SDTM IG) and animal nonclinical studies (SEND IG). In this document, SDTM refers to both models unless a specific IG is stated.

### *Specifications*

Each individual subject's complete subject profile is in a single PDF file. Including the subject ID in the file name will help identify the file. Alternatively, all subject profiles for an entire study may be in one file if the size of each individual subject profile is small and there are not a large number of subject profiles needed for the study. If a sponsor does the latter, the PDF file should be bookmarked using the subject's ID. Including the study number in the file name will help identify the file.

### **Analysis datasets**

#### *Definition*

Analysis datasets are datasets created to support results presented in study reports, the ISS and the ISE and to support other analyses that enable a thorough regulatory review. Analysis datasets contain both raw and derived data.

#### *Specifications*

Each dataset is provided as a SAS Transport (XPORT) file. Prior to submission, sponsors should contact the appropriate center's reviewing division to determine the division's analysis dataset needs. CDISC/ADaM standards for analysis datasets (<http://www.cdisc.org/adam>) may be used if acceptable to the review division.

Any requested programs (scripts) generated by an analysis tool should be provided as ASCII text files and should include sufficient documentation to allow a reviewer to understand the submitted programs. If the programs created by the analysis tool use a file extension for ASCII text files other than .txt, the file name should include the native file extension generated by the analysis tool for ASCII text program files, e.g. myRcode.r, mySAScode.sas, etc. If the analysis tool does not save programs in ASCII format, a PDF rendition of the program file should be provided in addition to the program file.

CDER statisticians perform analyses on the tumor data from each rodent carcinogenicity study, and they need this information provided as an electronic analysis dataset. See Appendix 1 on data elements for the analysis dataset recommended by these statistical reviewers. The dataset containing this information should be named tumor.xpt to aid in identification.

#### *General Considerations for Analysis Datasets*

- For an individual study, all dataset names and dataset labels should be unique across the analysis and raw datasets submitted for this individual study. The internal name for an analysis dataset should be the same as the name shown in the data definition file.

- Each analysis dataset should be described by an internal label which is shown in the data definition file. This label should clearly describe the contents of the dataset. For example, the label for an efficacy dataset might be "TIME TO RELAPSE (EFFICACY)". At least one analysis dataset should be labeled in the data definition file as containing the primary efficacy data.
- The key variables (subject identifier and visit for datasets with multiple records per subject) should appear first in the datasets. Each subject should be identified by a single, unique subject identifier within an entire application (including tabulation, listing and analysis datasets). Subjects enrolled in a primary study and then followed into an extension study should retain their unique identifier from the primary study.
- When a dataset contains multiple records per subject, a variable for relative day of measurement or event and variables for visit should be included. In addition to a protocol-scheduled visit variable, include at least two timing variables; a character variable describing the visit (e.g. WEEK 8) and a corresponding numeric variable (e.g. 8). These two variables are measures of time from randomization.
- For unscheduled visits or measurements, numbers are often assigned values between two protocol-scheduled visits. These numbers should be distinct from other visit numbers but retain the chronological order (e.g. two unscheduled visits between visit 3 and visit 4 might be 3.1 and 3.2). The character form of the visit identifier may be UNSCHEDULED or a similar term.
- Core variables should be listed after the key variables and included on each analysis dataset. Core variables include study/protocol, center/site, country, treatment assignment, sex, age, race, analysis population flags (e.g. ITT, safety) and other important baseline demographic variables.
- Variable names and codes should be consistent across studies and where feasible, the NCI CDISC Vocabulary should be used. For example, if glucose is collected in a number of studies, use the CDISC Submission Value "GLUC" for the laboratory test code in all of the studies. The NCI CDISC terminology is available at <http://evs.nci.nih.gov/ftp1/CDISC/SDTM/>. Additional information is also available at <http://www.cancer.gov/cancertopics/terminologyresources/page6>.
- The format of variables for similar types of data should be consistent within and across studies. For example, all variables that include calendar dates (e.g. birth date, screening visit date, randomization date, date of death) should use the same format for representing the date.
- For textual data that have been mapped to numeric codes, provide two variables, one with text and one with numeric codes.

- Dates should be formatted as numeric in the analysis datasets even if dates are in ISO8601 or another character format in the raw data. This formatting will facilitate calculations, such as duration.

## **SPECIFICATIONS FOR DATASETS DOCUMENTATION**

Dataset documentation includes data definitions and annotated case report forms.

### **Data definition file**

#### *Definition*

The data definition file describes the format and content of the submitted datasets.

#### *Specifications*

The specification for the data definitions for datasets provided using the CDISC SDTM IG or SEND IG is included in the Case Report Tabulation Data Definition Specification (define.xml) developed by the CDISC define.xml Team. The latest release of the define.xml is available from the CDISC web site (<http://www.cdisc.org/models/def/v1.0/index.html>). Include a reference to the style sheet as defined in the specification and place the corresponding style sheet in the same folder as the define.xml file.

For datasets not prepared using CDISC specifications, consult Appendix 2 for information concerning the preparation of a define.pdf data definition file.

### **Annotated case report form**

#### *Definition*

An annotated case report form (CRF) is a blank CRF with annotations that document the location of the data with the corresponding names of the datasets and the names of those variables included in the submitted datasets.

#### *Specifications*

The annotated CRF is a blank CRF that includes treatment assignment forms and maps each item on the CRF to the corresponding variables in the database. The annotated CRF should provide the variable names and coding for each CRF item included in the data tabulation datasets. All of the pages and each item in the CRF should be included. The sponsor should write *not entered in database* in all items where this applies. The annotated CRF should be provided as a PDF file. Name the file *blankcrf.pdf*.

## **SPECIFICATIONS FOR OTHER TYPES OF STUDY DATA**

### **Annotated ECG waveform data**

#### *Definition*

Annotated ECG waveform data are raw voltage-versus-time data comprising the electrocardiogram recording, to which have been attached the identification of various intervals or other features.

#### *Specifications*

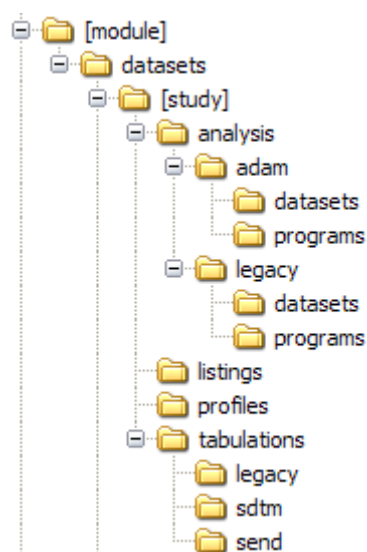
See the HL7 normative standard for creating the annotated ECG waveform data files. This information may be found on the HL7 web site [www.hl7.org](http://www.hl7.org). More information may be found at:

<http://www.fda.gov/Drugs/DevelopmentApprovalProcess/HowDrugsareDevelopedandApproved/ApprovalApplications/AbbreviatedNewDrugApplicationANDAGenerics/ucm154163.htm#ECG>



## SPECIFICATIONS FOR ORGANIZING THE DATASETS

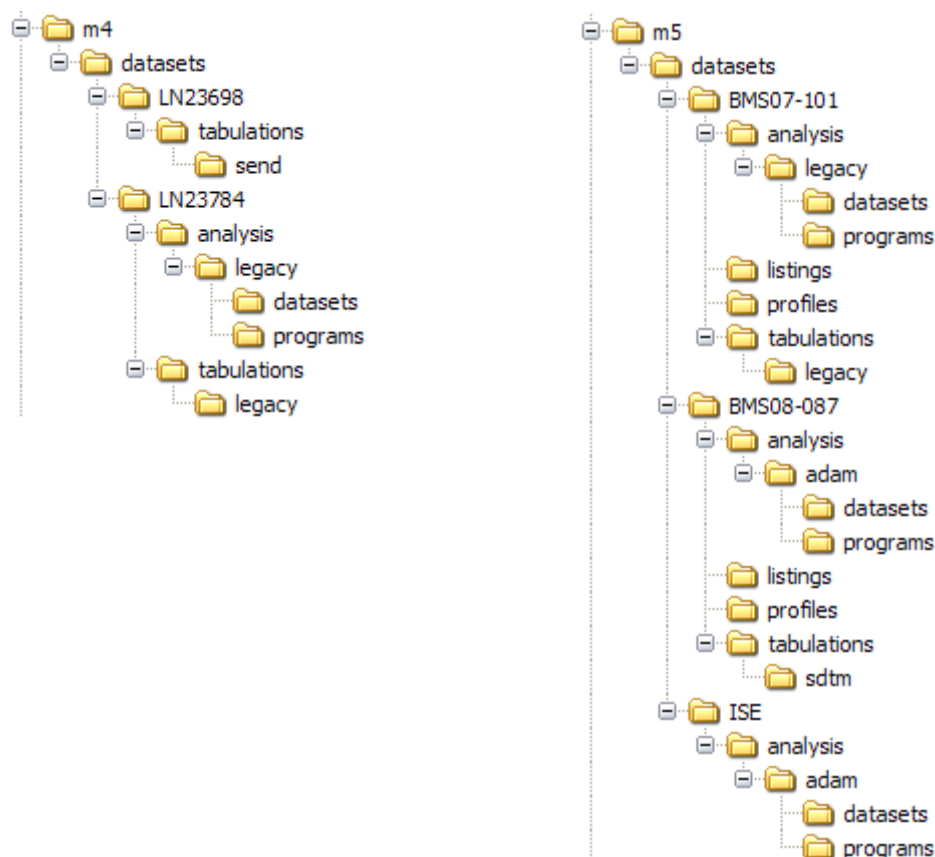
The specifications for organizing study datasets and their associated files in folders are summarized in the following figure and accompanying table. No additional subfolders are needed; unused folders do not need to be supplied.



Folder	Description
[module name]	Name the folder according to the CTD module which the datasets apply to. Use m4 for non-clinical data and m5 for clinical data.
datasets	The folder under which all of the study data being submitted for the module specified is organized.
[study]	Name the folder according to the study identifier or analysis performed for which the data is being supplied, e.g., BMS09-101, ISS, ISE.
analysis	The folder under which analysis datasets and programs will be organized  <b>Note:</b> The analysis datasets and programs are to be placed in specific folders based on their format.
adam	The sub-folder under which ADaM formatted datasets and programs are organized
datasets	The sub-folder in which the analysis datasets are organized
programs	The sub-folder in which the analysis programs are organized
legacy	The sub-folder under which legacy formatted datasets and programs are organized
datasets	The sub-folder in which the analysis datasets are organized
programs	The sub-folder in which the analysis programs are organized
listings	The folder in which miscellaneous datasets that don't qualify as analysis, profile, or tabulation datasets are organized

Folder	Description
profiles	The folder in which patient profiles are organized.
tabulations	The folder under which tabulation datasets will be organized.  <b>Note:</b> The tabulations datasets are to be placed in specific folders based on their format.
legacy	The sub-folder in which tabulations not formatted according to an identified standard format are organized, e.g., non-SDTM datasets.
sdtm	The sub-folder in which tabulations formatted according to the SDTM IG standard are organized. Should only be used in m5 for clinical data
send	The sub-folder in which tabulations formatted according to the SEND IG standard are organized. Should only be used in m4 for animal data

The following example shows the folder structure for a submission containing 2 individual nonclinical studies and 2 clinical studies and an ISE. SEND IG tabulations have been submitted for LN23698. Legacy tabulations and analysis data (such as a tumor.xpt) have been submitted for LN23784. Legacy clinical study data tabulations have been submitted for BMS07-101. SDTM IG and ADaM data has been submitted for BMS08-087.



## APPENDIX 1

Tumor Dataset For Statistical Analysis <sup>1,2</sup> (tumor.xpt)				
Variable	Label	Type	Codes	Comments
STUDYNUM	Study number	char		3
ANIMLNUM	Animal number	char		1,3
SPECIES	Animal species	char	M=mouse R=rat	
SEX	Sex	char	M=male F=female	
DOSEGP	Dose group	num	Use 0, 1, 2, 3,4,... in ascending order from control. Provide the dosing for each group.	
DTHSACTM	Time in days to death or sacrifice	num		
DTHSACST	Death or sacrifice status	num	1 = Natural death or moribund sacrifice 2 = Terminal sacrifice 3 = Planned intermittent sacrifice 4= Accidental death	
ANIMLEXM	Animal microscopic examination code	num	0= No tissues were examined 1 = At least one tissue was examined	
TUMORCOD	Tumor type code	char		3,4
TUMORNAM	Tumor name	char		3,4
ORGANCOD	Organ/tissue code	char		3,5
ORGANNAM	Organ/tissue name	char		3,5
DETECTTM	Time in days of detection of tumor	num		
MALIGNST	Malignancy status	num	1 = Malignant 2= Benign 3 = Undetermined	4
DEATHCAU	Cause of death	num	1 = Tumor caused death 2= Tumor did not cause death 3 = Undetermined	4
ORGANEXM	Organ/Tissue microscopic examination code	num	1 = Organ/Tissue was examined and was usable 2= Organ/Tissue was examined but was not usable (e.g., autolyzed tissue) 3 = Organ/Tissue was not examined	

Each animal in the study should have at least one record even if it does not have a tumor.

1. Additional variables, as appropriate, can be added to the bottom of this dataset.
2. ANIMLNUM limited to no more than 12 characters; ORGANCOD and TUMORCOD limited to no more than 8 characters; ORGANNAM and TUMORNAM limited to no more than 30 characters.
3. A missing value should be given for the variable MALIGNST, DEATHCAU, TUMOR and TUMORCOD when the organ is unusable or not examined.
4. Do not include a record for an organ that was useable and no tumor was found on examination. A record should be included for organs with a tumor, organs found unusable, and organs not examined.

## APPENDIX 2

For datasets not prepared using CDISC specifications, sponsors should include a define.pdf to describe the datasets for each study, specific data analysis (e.g., population PK), and integrated summaries. For the datasets to be useable, the definitions of the variables should be provided. Sponsors should document all of the variables in the datasets in data definition tables. There should be one set of data definition tables for each study, specific data analysis (e.g., population PK) and integrated summary. The first table should include a listing of all datasets provided for the study with a description of the dataset and the location of the dataset file. Provide a hypertext link from the description of the dataset to the appropriate data definition table. Provide a hypertext link from the location listing of the file to the SAS transport file. The reviewer can use the first hypertext link to view the data definition table and the second to open the SAS transport dataset file. For clinical study data, sponsors should also provide a link to the appropriate annotated case report form file (blankcrf.pdf).

In the following table, the dataset for AE is described as adverse events, and the dataset file is located in listings folder for study 1234

<b>Datasets for Study 1234</b>		
<b>Dataset</b>	<b>Description of Dataset</b>	<b>Location</b>
AE	Adverse Events	m5/datasets/study1234/listings/ae.xpt
...	...	...

Subsequent pages should contain a table for each dataset that includes an organized listing of all variable names used in the dataset, a descriptive variable label, data types, codes (and decodes), and comments. The comments field is for further description of the variables. For derived variables, the method for calculating the variable should be included in the comments field. For raw variables in clinical datasets, the location of the variable on the annotated CRF should be provided as well as the CRF field name if different from the variable name in the dataset. Providing a hypertext link from each raw data variable in the data definition table to the appropriate location of the blankcrf.pdf also helps the review process. An example of part of a data definition table for the demographics dataset for study 1234 is provided below.

Study 1234 – Demographics Dataset Variables				
Variable	Label	Type	Codes	Comments <sup>1</sup>
USUBJID	Unique subject ID number	char		Demographics page 3
SEX	Sex of subject	char	f = female m = male	Demographics page 3
BDATE	Birth date	date		Demographics page 3
DUR	Duration of Treatment	num		Derived STOP DATE – START DATE
TRT	Assigned treatment group	num	0= placebo 5= 5mg/day	

<sup>1</sup> Use footnotes for longer comments

The data definition tables should be provided as a single PDF file named *define.pdf* and placed in the appropriate study, specific analysis type or integrated summary folder in the datasets folder. The Title portion of the Document Information field of each data definition file should include the appropriate study report number, specific analysis type or integrated summary name and *data definitions*. For example, the data definition file for study 2001 would be identified as: *study 2001, data definitions*. This file is considered part of the comprehensive table of contents.