Relational Databases, SQL

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Databases and SQL

- Lecture Goals
 - Create a mirna-mrna targeting database from text file(s).
 - Query the database table(s) to find the genes that are predicted to be targeted by a mirna.

What you'll need

- · Firefox, SQLite plugin
- SqliteStudio

Mirdb and Targetscan databases

DB Definitions

- Relational database: collection of tables (also called relations)
- Database Management System (DBMS): a software that is used to create, access, and maintain a database; e.g., sqlite, mysql, postgresql, oracle, sql server.
- Table: Collection of rows (also called tuples or records).
- Each row in a table contains a set of columns (also called fields or attributes).
- Each column has a type:
 - Text: VARCHAR(20)
 - Integer: INTEGER
 - Floating-point: FLOAT, DOUBLE
 - Date/time DATE, TIME, DATETIME
- Primary key: provides a unique identifier for each row (optional).
- Schema: the structure of the database tables
 - The table name
 - The names and types of its columns
 - Various optional additional information (defaults, constraints, etc.)

SQL

- SQL = "Structured Query Language"
 - Non-procedural
 - Set-oriented
 - Relationally complete
 - Functionally incomplete
- Four main types of queries:
 - Insert, Delete, Select, Update

Create/drop table

- Syntax:
 - https://www.sqlite.org/lang_createtable.html
- Create a table for the students:

```
CREATE TABLE students (
id INTEGER PRIMARY KEY,
name VARCHAR(30),
birth DATE,
gpa FLOAT,
grad INTEGER
);
```

Drop table:

DROP TABLE students;

Insert/Delete

Add rows to the students table:

Delete row(s):

DELETE FROM students;

Select Queries

- 3 main elements:
 - What you want
 - Where it is found
 - How you want it filtered
- Show entire contents of a table:

Select

Show just a few columns from a table:

• Filtering: only get a subset of the rows:

Select

• Sorting:

```
SELECT gpa, name, grad FROM students WHERE gpa > 3.0 ORDER BY gpa DESC;
+----+
| gpa | name | grad |
+----+
| 3.9 | Anderson | 2009 |
| 3.2 | Chen | 2011 |
| 3.1 | Hernandez | 2011 |
+----+
```

• Limiting: only get a certain number of rows:

Update / Delete

Update:

```
UPDATE students
   SET gpa = 2.6, grad = 2013
WHERE id = 2;
```

• Delete:

```
DELETE FROM students
WHERE id = 2;
```

Joins

- Join: a query that merges the contents of 2 or more tables, retrieves information from the merged results.
- · Join example: many-to-one relationship
- Students have advisors; add a new table describing faculty.

Join example: many-to-one relationship

 Add new column advisor_id to the students table. This is a foreign key.

Perform the join query: Get the students who are advised by Fujimura

+		+
id na	me	title
+	+-	+
1 Fu	jimura	assocprof
2 Bo	losky	prof
+		+

Join example: many-to-many relationship

- Courses: students take many courses, courses have many students
- Add a new table describing courses:

1	id	İ	number	İ	name	1	quarter	-+ -+
	1 2 3		CS142 ART101 ART101		Web stuff Finger painting Finger painting Mud wrestling	 	Winter 2009 Fall 2008 Winter 2009	
+		+-		+		+		-+

 Create a "join table" courses_students describing which students took which courses.

course_id	++ student_id
1	1
3	1
4	1
1	2
2	2
1	3
2	4
4	4
+	++

Join example: many-to-many relationship

• Find all students who took a particular course ('ART101'):

Additional Database Features

- Indexes: used to speed up searches
- Concurrency: Allow more than one program use the database at the same time.
- Transactions: used to group operations together to provide predictable behavior even when there are concurrent operations on the database.
- · Locks: Used to limit concurrent access.
- · Views: a virtual table for results of a stored query
- Procedures: a set of sql statements stored in the database

Exercise

- GO database
 - http://geneontology.org/page/lead-database-schema
 - http://geneontology.org/sites/default/files/public/diag-godb-er.jpg
 - http://www.berkeleybop.org/goose/
- Retrieve the names of the species that are under the genus 'Drosophila'
- Retrieve the genus and species name of the organisms whose species name has a prefix 'mel' (use LIKE function).
- Retrieve the gene symbols (gene_product.symbol) of all Drosophila melanogaster (species.genus, species.species) genes that are annotated to the 'nucleus'.
 - Join path: term.id -> graph_path.term1_id /term2_id -> association.term_id /gene_product_id -> gene_product.id
 - Join path: gene_product.species_id -> species.id

mirdb exercise

- Create a database from mirdb data.
 - http://mirdb.org/miRDB/download.html
- How many miRNAs are predicted to target XM_539064?
- How many predicted targets of cfa-let-7a have a prediction score of at least 80?
- List the miRNAs and the number of their targets. (Each row of the result should contain a distinct miRNA). (Use count() and GROUP BY). Show only top 10 rows of the result.