

BF768 Homework 0

Problem 1 is due on Tuesday, 1/28/20 and will be discussed in class.
Problems 2 and 3 are due on Thursday 1/30/20.

General Policy on Homework Collaboration:

Except as otherwise noted, all problem sets/homeworks are to represent individual effort, and are to be written up and turned in individually. This does not preclude talking about a problem set with other class members; in fact, working together is encouraged, since it is one of the skills of modern science. However you are not allowed to copy each other's answers. If you work on a problem set with other people, please state that (names included) on your write-up.

Piazza

The website Piazza will be available for class discussion. You're welcome to email questions to the professor or TA, but we will both be monitoring the Piazza website as will other students in the class. I encourage you to post your questions on the class page at <https://piazza.com/class/k5fejzt9pih1bq?cid=7>.

When someone posts a question, if you know the answer, please go ahead and post it. However please **don't provide solutions to homework questions** on Piazza. It's OK to tell people where to look to get answers, or to correct mistakes; just don't provide actual solutions to homeworks.

1. Design and draw an ER diagram that captures the information below.

- Patients are recorded by an id, social security number (SSN), name, address, and date of birth (DOB).
- Doctors are identified by an id, name, specialty, and start date of service.
- Each pharmaceutical company is identified by a name.
- For each drug, the trade name and formula are recorded.
- Each drug is manufactured by a pharmaceutical company and the trade name identifies the drug uniquely from among the products of that company.
- Every pharmacy has a name, address, and phone number.
- Every patient has a primary physician. Every doctor has at least one patient.
- Each pharmacy sells several drugs and has a price for each. A drug could be sold at several pharmacies, and the price could vary from pharmacy to pharmacy.
- Doctors prescribe drugs for patients. Usually, a doctor writes prescriptions for more than one patient and a patient could receive a prescription from more than one doctor. Each prescription has a date and a quantity.
- Pharmaceutical companies have long-term contracts with pharmacies. A pharmaceutical company will contract with more than one pharmacy, and a pharmacy will contract with more than one pharmaceutical company. For each contract, there is a start date, end date, and the text of the contract.

2. Design and draw an ER diagram that captures the information below.

- A gene is described by a chromosome number, an organism, a name, a start position in the chromosome, and an end position.
- A gene is composed of one or more exons and zero or more introns.
- An exon has a start position, a stop position, and a number in the gene sequence (1 means first exon, etc.).
- An intron has a start position, a stop position, and a number in the gene sequence.
- Each intron and each exon belongs to exactly one gene.
- Each gene has zero or more associated transcription promoter elements.
- A promoter element has a name and a type (examples: TATA box, transcription factor binding site).
- Each promoter stored will be associated with at least one gene.
- Experiments are run to test gene activity.
- Each experiment involves one or more genes and is run by one student using one of several protocols. The lab, date, and gene activity are recorded with each experiment.
- Each student has an id number and a name.
- Each protocol has a set of experimental conditions (this could be a text file listing the conditions). Not all protocols are necessarily used and some will be used repeatedly.
- Students can run more than one experiment. Some students will not run any experiments.
- No two experiments are the same on the same day.

3. Design and draw an ER diagram that captures the information below.

- A protein is described by a name, a sequence, and a structure (this could be a file storing PDB structure data).
- Each protein has one or more functions.
- A function has a name, a class, and a subclass (for example, a class could be DNA binding protein, a subclass could be zinc finger, which is a particular protein structure that binds DNA).
- A protein's function is either predicted or experimentally confirmed.
- If confirmed, there is at least one journal reference.
- If predicted, there is a gene annotation program that was used for the prediction.
- Each gene annotation program has a unique journal references and no single reference discusses more than one gene annotation program.
- Some proteins are known to function in one or more cellular pathways. The protein's function can be different in different pathways and a protein can have more than one function in the same pathway.