

# Homework #2:

## Entities

- We have 6 entity sets: gene, transcript, proteins, modified proteins, modified protein activity and modified protein localization (make sure there are at least 5 entity sets).
- The attributes used will include at least 3 each (they must at least cover the key attributes)
  - GENE (Gene\_ID, Gene\_name, MaximumCDS)
  - TRANSCRIPT (Transcript\_id, Gene\_Name, Transcript\_Sequence)
  - PROTEIN (Protein\_ID, Transcript\_ID, Protein\_sequence)
  - MOD\_PROTEIN (ModProtein\_ID, Protein\_ID, ModProtein\_Type)
  - MODPROTEIN\_ACTIVITY (ModProtein\_ID, EC\_No, Activity\_Name,)
  - MODPROTEIN\_LOCATION (ModProtein\_ID, Location\_ID, CellCycleStage)

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## Relationship Definitions

- Business rules will include the following
  - Each gene can produce zero to many transcripts
  - Each transcript comes from one and only one gene and produces exactly one protein
  - A protein sequence can be either measured or inferred.
  - A transcript can exist without a gene to which it maps
  - Each protein is produced by one and only one transcript.
  - A protein can exist without a transcript to which it maps.
  - A protein can have zero to many modified amino acids
  - A protein can be modified on more than one amino acid
  - A protein is a type of modified protein (type none) - recursive
  - Amino acids may be modified by more than one type of group.
  - A modified protein is existence-dependent on a protein
  - Proteins can be located at multiple locations in the cell

# Homework #2

- Using MySQL Workbench
  - Create and ER IE model of the given entities
  - The model should include the relationships covering the rules defined above.
  - Due Thursday Feb 4 at 8:00 a.m.
  - Turn in the saved model file.