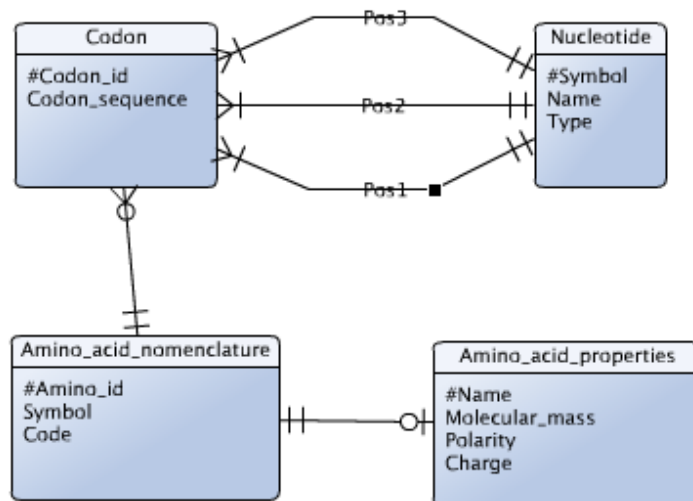
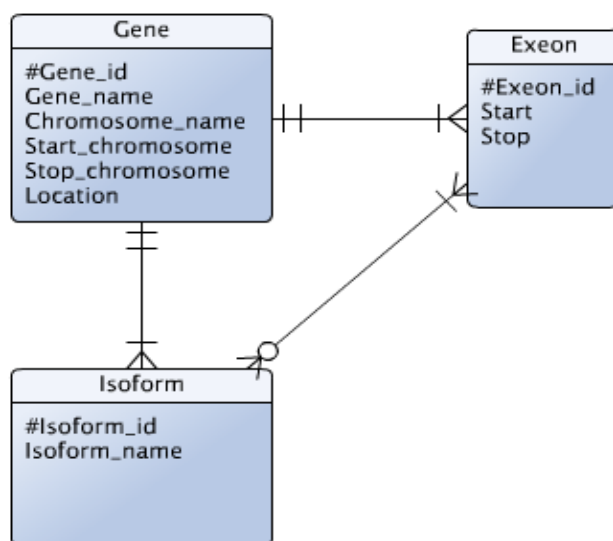


**Task 1****Task 2i**

The entities are Gene, Exon and Isoform. One can eventually add Chromosome as an entity, as we might be handling several Chromosomes.

**Task 2ii**

E/R diagram



## Task 2iii

Convert to 3NF

Gene(#Gene\_id, Gene\_name, Chromosome\_name, Start\_chrom, Stop\_chrom, Location)  
Isoform(#Isoform\_id, Isoform\_name, Gene\_id\*)  
Exeon( #Exon\_id, Start, Stop, Gene\_id\*)  
IsoExo(#Exon\_id, #Isoform\_id)

Dependencies:

Gene:            Gene\_name → Chromosome\_name

Rewriting to 3NF:

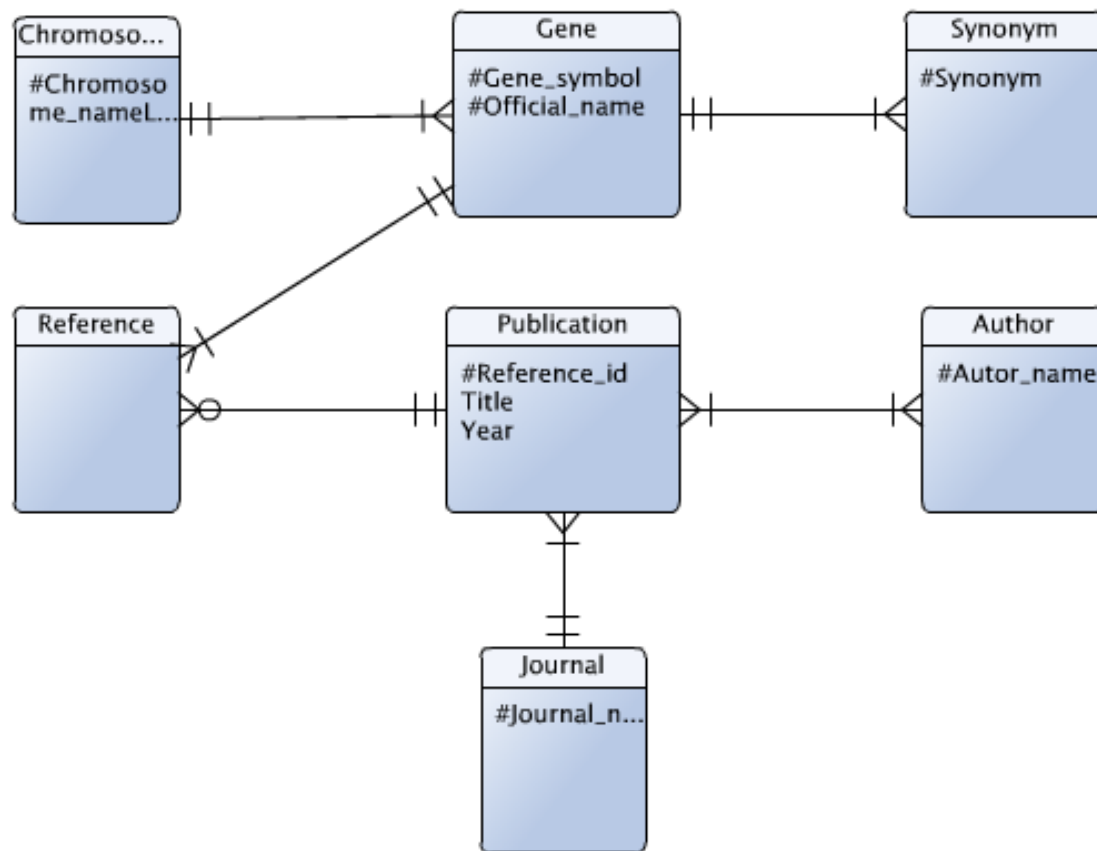
Chromosome (#Gene\_name, Chromosome\_name)  
Gene(#Gene\_id, Gene\_name\*, Start\_chrom, Stop\_chrom, Location)  
Isoform(#Isoform\_id, Isoform\_name, Gene\_id\*)  
Exeon( #Exeon\_id, Start, Stop, Gene\_id\*)  
IsoExe(#Exeon\_id, #Isoform\_id)

## Task 3i

Chromosome  
Gene  
Synonym  
Publication  
Author  
Journal

I have interpreted Journal as a collection of publications.

### Task 3ii



### Task 3iii

In order to convert this to 1NF, one can put publication year in the references entity, (marked in red):

Chromosome( #Chromosome\_name, Length)

Gene( #Gene\_symbol, #Official\_name, Chromosome\_name\*)

Synonym(#Syn\_name, Official\_name\*, Gene\_symbol\*)

Reference(#Official\_name, #Gene\_symbol, #Reference\_id, **Year**)

Publication(#Reference\_id, Title, Journal\_name\*)

Author(#Reference\_id\*, #Author\_name)

Journal(#Journal\_name)

Reference now contains the dependency  $\text{Reference\_id} \rightarrow \text{Year}$ , where  $\text{Reference\_id}$  is a true subset of the primary key “ $\text{Official\_name} + \text{Gene\_symbol} + \text{Reference\_id}$ ”. This is a violation of the 2NF and is therefore of 1NF.

Here, assumed that neither  $\text{Gene\_symbol}$  nor  $\text{Official\_name}$  is necessarily unique.

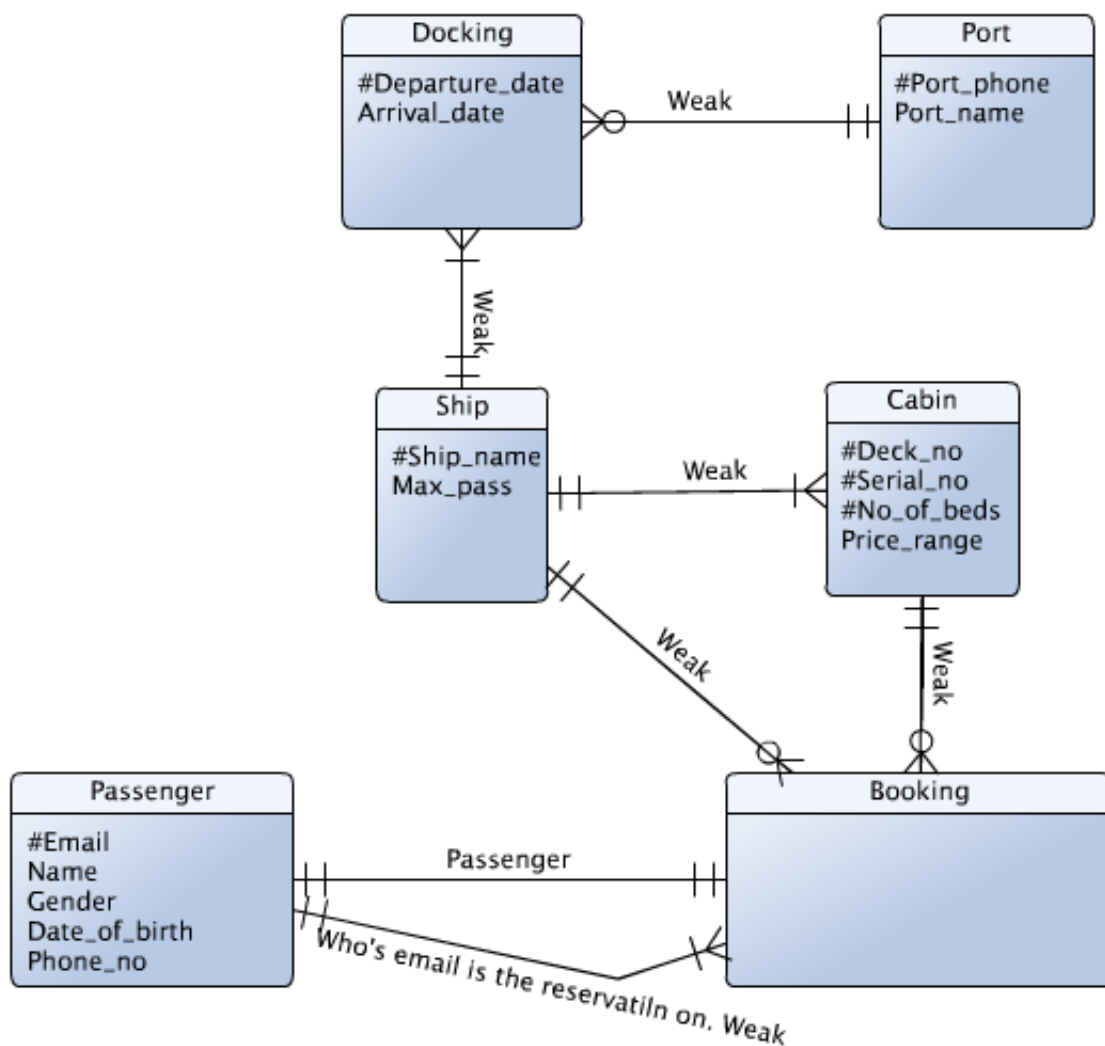
Please note that References is a weak entity.

### Task 3iv

Converting to BCNF

Chromosome( #Chromosome\_name, Length)  
Gene( #Gene\_symbol, #Official\_name, Chromosome\_name\*)  
Synonym(#Syn\_name, Official\_name\*, Gene\_symbol\*)  
Reference(#Official\_name, #Gene\_symbol, #Reference\_id)  
Publication(#Reference\_id, Title, Journal\_name\*)  
Author(#Author\_name, #Reference\_id\*)  
Journal(#Journal\_name, year)

### Task 4



## Task 5i

Truck( #Registration\_number, Registration\_year, Model, Maximum\_weight, Assignment\_number\*)

Truck should be used to keep track of which truck is on which assignment, and not to store permanent information about the truck. We can say that the registration number decides registration year, model and maximum weight. If we put this information in another table, we won't have to repeat all of those attributes every time we mention a specific truck.

## Task 5ii

Registration\_number → Registration\_year  
Registration\_number → Model  
Registration\_number → Maximum\_weight  
Registration\_number → Assignment\_number  
Model+Registration\_year → Maximum\_weight

## Task 5iii

The candidate key for Truck is Registration\_number, as Registration\_number decides all the other attributes and is a minimal super key.

## Task 5iv

Container\_type (#Type\_id, Type\_name, Max\_weight, Cubic\_quantity, Nightly\_rate)

Container (#Container\_number, Type\_id\*)

Customer (#Telephone\_number, Address)

Assignment (#Assignment\_number, Telephone\_number\*, Container\_number\*, Start\_date, End\_date)

Truck\_assignment(#Registration\_number\*, Assignment\_number\*)

Truck(#Registration\_number, Registration\_year\*, Model\*)

Truck\_load(#Registration\_year, #Model, Maximum\_weight)