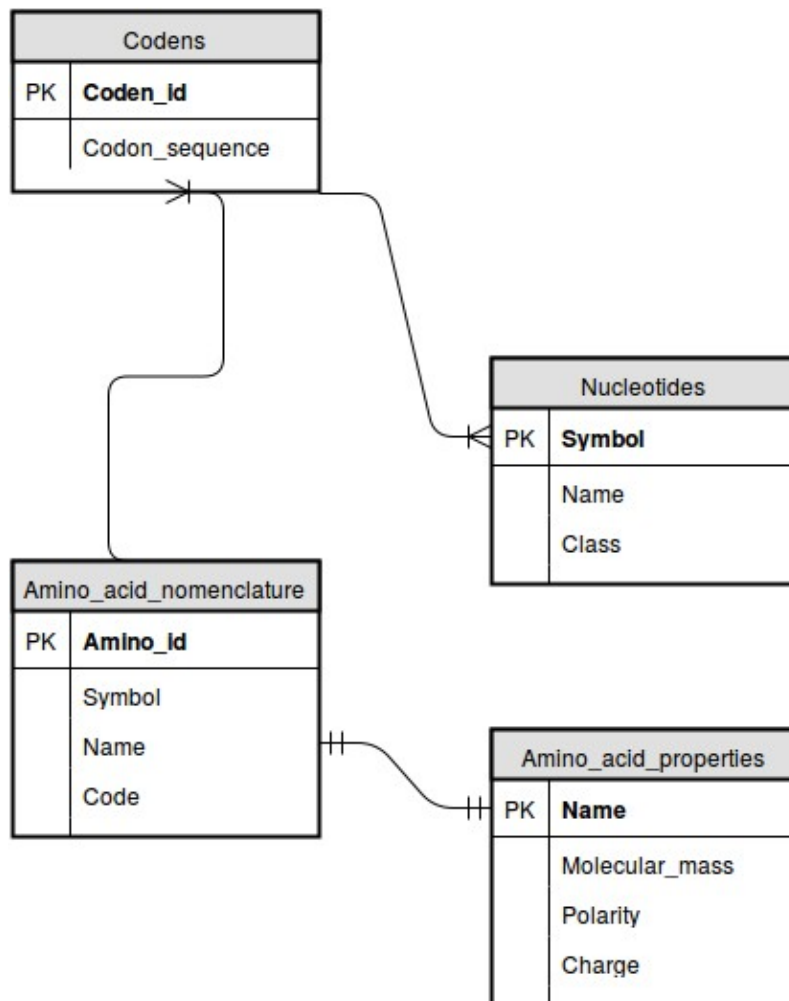


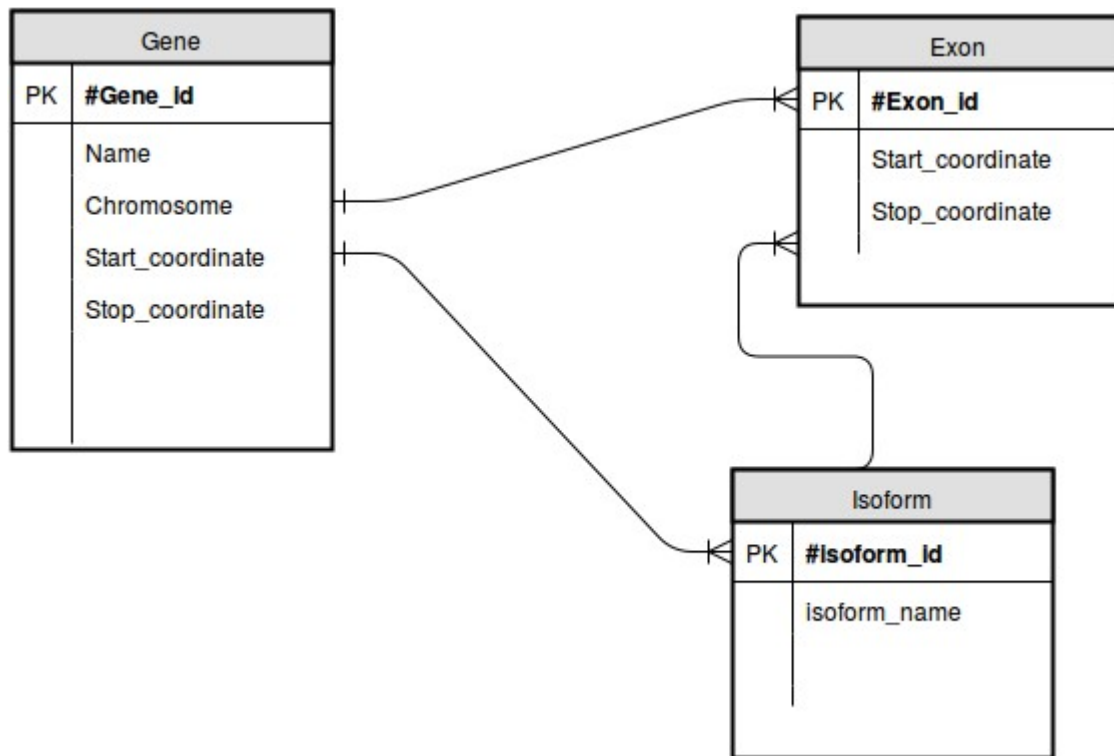
Inf115 compulsory 2

Morten Lohne

1:



2a: Gene, Exon, Isoform



2c:

Genes (#Gene_id, Name, Chromosome*, Start_coordinate, Stop_coordinate)

Exons (#Exon_id, Start_coordinate, Stop_coordinate, Gene_id*)

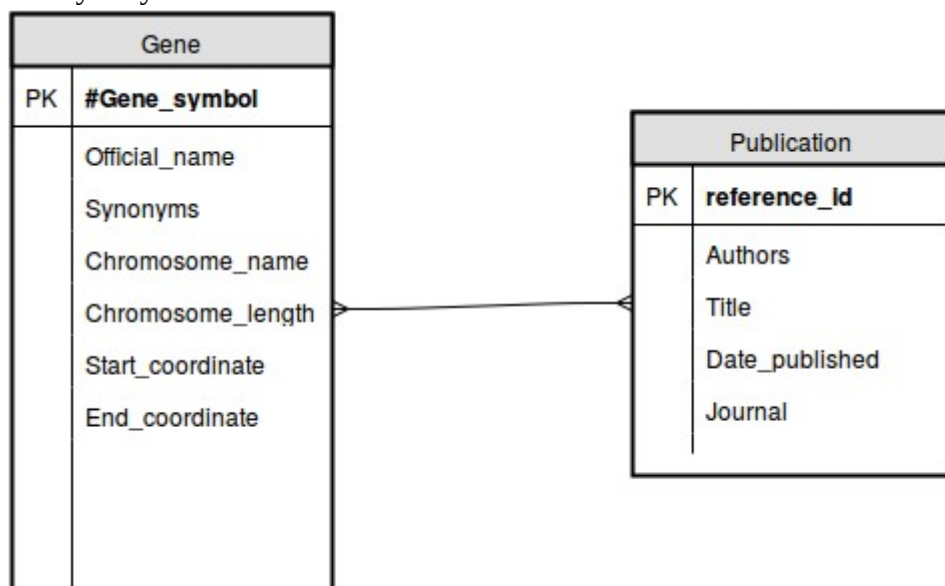
Chromosome(#Id, Chromosome_name)

Isoforms (#Isoform_id, Isoform_Name, Gene_id*)

Exon_Isoform_Bridge(#Exon_id*, #Isoform*)

3a: Gene, Publication

3b: Not on 1NF: Synonyms and authors are not atomic.



3c:

Gene(#Gene_symbol, #Official_name, Chromosome_name, Chromosome_length, Start_coordinate, End_coordinate)

Publication(#Reference_id, Title, Journal, Year_published, Publication_authors*)

Publication_authors(#Reference_id, #Author)
 Gene_synonyms(#Synonym_name, #Gene_symbol)

3d:

Gene(#Gene_symbol, Chromosome_id*, Start_coordinate, End_coordinate)
 Chromosome(#Chromosome_id, Chromosome_name, Chromosome_length)
 Gene_synonyms(#Synonym_name, #Gene_symbol)
 Publication(#Reference_id, Title, Journal, Year_published,
 Publication_authors*)Publication_authors(#Reference_id, #Author)

4:

