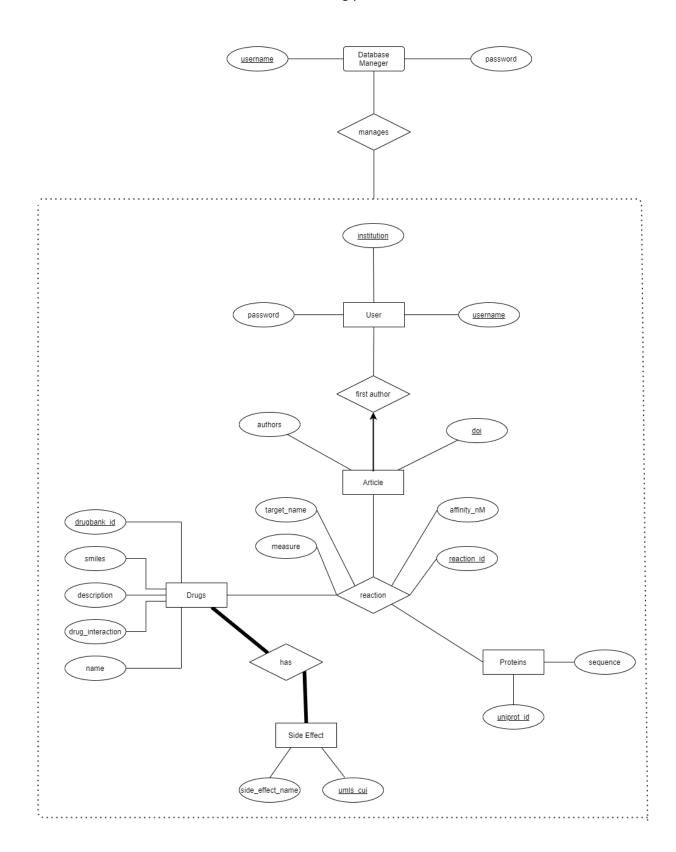
Part 1:

We designed ER according to relationships between entities and attributes. We tried to make it simple as possible. After designing ER diagram, we looked for participation constraints and key constraints. We draw arrows and bold lines accordingly.



### Part 2:

Drug(drugbank\_id:char(7),name:char(50),smiles:char(200),drug\_interaction:char(100),description:char(1000))

User(username:char(50),institution:char(100),password:char(50))

Article(doi:char(100),authors:char(500),institution:char(100),username:char(50))

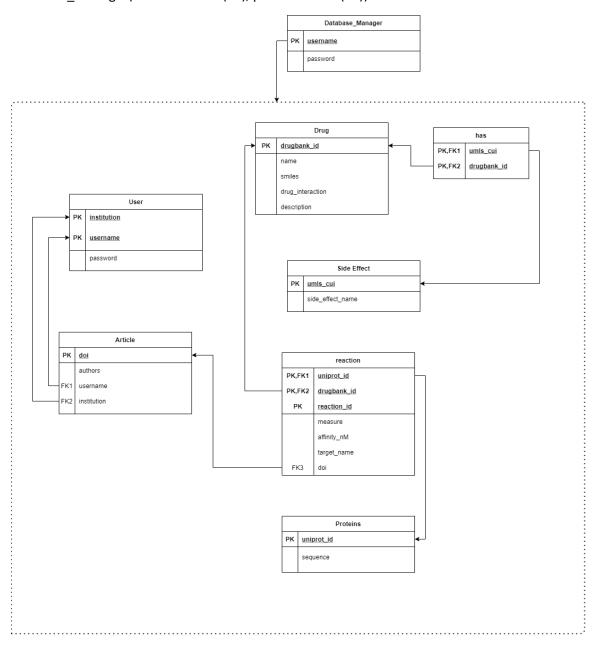
Proteins(uniprot\_id:char(6),sequence:char(10000))

Side\_Effect(umls\_cui:char(8),side\_effect\_name: char(100))

Has(umls\_cui:char(8), drugbank\_id:char(7))

Reaction(uniprot\_id:char(6),drugbank\_id:char(7):reaction\_id:char(8),measure:char(4),affinity\_nM:real,target\_name:char(1000),doi:char(1000)

Database\_Manager(username:char(50), password:char(50))



#### Part 3:

#### **User entity:**

User(username:char(50),institution:char(100),password:char(50))

username: U, institution: I, password: P

UI-> P

since U and I are primary keys it does not violate the BCNF

### **Article entity:**

Article(doi:char(100),authors:char(500))

doi: D, authors: A

D -> A

since D is a primary key it does not violate BCNF

### Drug entity:

Drug(drugbank\_id:char(7),name:char(50),smiles:char(200),drug\_interaction:char(100),description:char(1000))

drugbank\_id: D name: N smiles: S drug\_interaction: I description: R

 $D \rightarrow N$   $D \rightarrow S$   $D \rightarrow I$   $D \rightarrow R$  so  $D \rightarrow NSIR$ 

 $N \rightarrow D$   $N \rightarrow S$   $N \rightarrow I$   $N \rightarrow R$  so  $N \rightarrow DSIR$ 

 $S \rightarrow D$   $S \rightarrow N$   $S \rightarrow I$   $S \rightarrow R$  so  $S \rightarrow DNIR$ 

 $R \rightarrow D R \rightarrow N R \rightarrow I R \rightarrow S$  so  $R \rightarrow DNIS$ 

Then D, S, N and R are candidate keys. We can choose D as primary key.

D -> NSIR So it does not violate BCNF rule.

### **Reaction relationship:**

Reaction(uniprot\_id:char(6),drugbank\_id:char(7):reaction\_id:char(8),measure:char(4),affinity\_nM:re al,target name:char(1000),doi:char(1000)

drugbank\_id: D uniprot\_id: U reaction\_id: R measure: M affinity\_nM: A target\_name: T doi: Do

UDR -> MATDo

Since UDR are primary keys, it does not violate BCNF

## Side\_effect entity:

Side Effect(umls\_cui:char(8),side\_effect\_name:char(100))

umls\_cui : U side\_effect\_name : S

U -> S

since U is primary key, it does not violate BCNF

#### **Proteins entity:**

Proteins(uniprot\_id:char(6),sequence:char(10000))

uniprot\_id: U, sequence: Q

U -> Q

since U is primary key it does not violate BCNF

# Database\_Manager entity:

Database\_Manager(username:char(50), password:char(50))

username: U password: P

U -> P

Since U is primary key it does not violate BCNF

# Has relationship:

Has(umls\_cui:char(8), drugbank\_id:char(7))

There is no functional dependency in Has relation. (Between User and Side effect entities)

In conclusion, our database design is appropriate for BCNF rules. So we did not change anything in our database design.