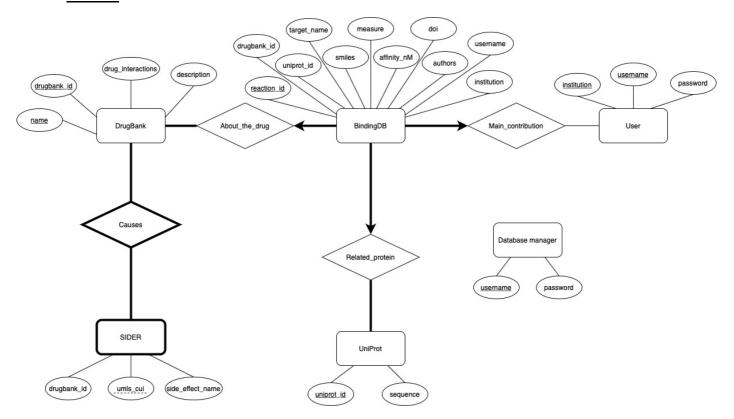
2021 Spring - CMPE321 Project 2 Report

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Part 1



Part 2

Relational schemas and tables are as follows:

- User(username: string, institution: string, password: string)
- Database manager(username: string, password: string)
- DrugBank(drugbank_id: string, name: string, drug_interactions: list<string>, description: string)
- **SIDER**(ulms_cui: **string**, drugbank_id: **string**, side_effect_name: **string**)

• BindingDB(reaction_id: integer, drugbank_id: string, uniprot_id: string, target_name: string, smiles: string, measure: string, affinity_nM: real, doi: string, authors: string, username: string, institution: string)

UniProt(uniprot_id: string, sequence: string)

User(username: string, institution: string, password: string)						Database manager(username: string, password: string)			DrugBank(drugbank_id: string, name: string, drug_interactions: list <string>, description: string)</string>)
username institution		password		us	ername	password	d	rugbank_id		name		drug_intera	actions	descrip	otion	
Abramovitz Merck Frosst Centre for Therapeutic Research		abrar	movitz3232	sele	en.parlar	selen.parlar	1	DB00459		Acitretin		['DB00304', 'DB00367', 'DB00162']		An oral retinoid effective		
Afzelius AstraZeneca R&D		afze	elius4343	riza	riza.ozcelik riza.ozcelik0			DB00523		Alitretinoin		DB00304', 'D DB001'		An important re	egulator of	
Ahlin	Ahlin Uppsala University		ah	nlin5151	arzuo	can_ozgur	arzucan_135		DB00964 Apraclonidine		['DB06237']		Apraclonidine, as .			
SIDER(ulms_cui: string, drugbank_id: string, side_effect_name: string)					BindingDB(reaction_id: integer, drugbank_id: string, uniprot_id: string, target_name: string, smiles: string, measure: string, affinity_nkt.real, doi: string, authors: string, username: string, institution: string) UniProt(unit string, smiles: string, seque: string, institution: string)											
ulms_cui	drugbank_id	side_effect_i	name	reaction_id	drugbank_id	uniprot_id	target_name	smiles	measure	affinity_n	/ doi	authors	usemame	institution	uniprot_id	sequence
C0000737	DB00459	Abdominal	pain	50876947	DB00925	O15245	Adrenergic Alpha	CC(CO	IC50	15100	10.1021/	Ahlin, G;	Ahlin	Uppsala University	P04278	MESRG
C0702166	DB00459	Acne		50876963	DB01162	O15245	Adrenergic Alpha	COc1c	IC50	23700	10.1021/	Ahlin, G;	Ahlin	Uppsala University	Q99808	мттѕн
C0155626	C0155626 DB00459 Acute my			50739658	DB01132	095342	Bile salt export pump	CCc1c	IC50	400	10.1002/	Aleo, MD;	Aleo	Pfizer Inc	P51574	MGMSK

Part 3

Now we will go through all the relations and check their suitability of BCNF.

1- User(*username(U):* **string**, *institution(I):* **string**, *password(P):* **string**):

 $K = \{U, I\}$ is the primary key.

Non-trivial functional dependencies:

 $K \rightarrow P$

Since k is a key for the entity, the requirements for BCNF are met.

2- Database manager(*username*(*U*): **string**, *password*(*P*): **string**):

U is the primary key.

Non-trivial functional dependencies:

 $U \rightarrow P$

Since k is a key for the entity, the requirements for BCNF are met.

3- DrugBank(drugbank_id(Did): string, drug_name(Dn): string, drug_interactions(Di): list<string>, drug_description(Des): string): Did and Dn are both keys since any one of them uniquely determines the entity.

Non-trivial functional dependencies:

Did → DnDiDes

Dn → DidDiDes

Since Did and Dn are both keys for the entity, the requirements for BCNF are met.

4- SIDER(*ulms_cui(U):* **string**, *drugbank_id(D):* **string**, *side_effect_name(S):* **string**) :

K1 = {U, D} and K2 = {S, D} are both keys since they uniquely determine SIDER entities.

Non-trivial functional dependencies:

 $K1 \rightarrow S$

 $K2 \rightarrow U$

Since K1 and K2 are both keys for the entity, the requirements for BCNF are met.

5- BindingDB(reaction_id(R): integer, drugbank_id(Did): string, uniprot_id(Uid): string, target_name(T): string, smiles(S): string, measure(M): string, affinity_nM(An): real, doi(D): string, authors(A): string, username(U): string, institution(I): string):

R is the primary key.

Non-trivial functional dependencies:

R → DidUidTSMAnDAUI

 $Did \rightarrow S$

 $S \rightarrow Did$

 $D \rightarrow A$

Since Did, S and D are not superkeys, the requirements of BCNF are not met. Likewise, since D is not a superkey and A is not part of some key for the relation, the requirements of 3NF are also not met. We can decompose the relation into BCNF when the following functional dependencies hold:

 $R \rightarrow DidUidTMAnDUI$

 $Did \rightarrow S$

 $S \rightarrow Did$

 $D \rightarrow A$

Where the new and updated attributes and entities are (the parts that are newly added are written in green):

BindingDB(reaction_id(R): integer, drugbank_id(Did): string, uniprot_id(Uid): string, target_name(T): string, measure(M): string,affinity_nM(An): real, doi(D): string, username(U): string, institution(I): string)

Non-trivial functional dependencies:

 $R \rightarrow DidUidTMAnDUI$

Since R is a key, the requirements of BCNF are met.

Links(doi(D): string, authors(A): string)

D is a key for Links since it completely determines the entity. Non-trivial functional dependencies:

 $D \rightarrow A$

Since D is a key, the requirements of BCNF are met.

DrugBank(drugbank_id(Did): string, drug_name(Dn): string,
drug_interactions(Di): list<string>, drug_description(Des): string,
smiles(S): string)

Did, Dn and S are all keys since any one of them uniquely determines the entity.

Non-trivial functional dependencies:

Did → DnDiDesS

 $Dn \rightarrow DidDiDesS$

 $S \rightarrow DidDnDiDes$

Since Did, Dn and S are all keys for the entity, the requirements for BCNF are met.

Dependency preserving:

As seen above, the functional dependencies are preserved and distributed among entities. Hence the decomposition is dependency preserving.

• Lossless-join:

-The attributes are preserved, i.e. neither new attributes are introduced nor existing attributes are discarded. Instead, some attributes are shifted to other entities which preserves the union of attributes after decomposition.

-After the decomposition, the intersection of the remaining attributes in **BindingDB** with the removed attributes contains keys for entities **Links** (which is D) and **DrugBank** (which is S).

Because of the conditions above that are shown to hold, the decomposition is lossless-join.

6- UniProt(*uniprot_id*(*Uid*): **string**, *sequence*(*S*): **string**):

Uid is the primary key.

Non-trivial functional dependencies:

 $Uid \rightarrow S$

Since Uid is a key for the entity, the requirements for BCNF are met.

Updated tables for updated and new entities are below:

	BindingDB(reaction_id: integer, drugbank_id: string, uniprot_id: string, target_name: string, measure: string, affinity_nM: real, doi: string, username: string, institution: string)									
reaction_id	drugbank_id	uniprot_id	target_name	measure	affinity_nM	doi	username	institution		
50876947	DB00925	O15245	Adrenergic Alpha	IC50	10.1021/	Ahlin	Ahlin	Uppsala University		
50876963	DB01162	O15245	Adrenergic Alpha	IC50	10.1021/	Ahlin	Ahlin	Uppsala University		
50739658	DB01132	O95342	Bile salt export pump	IC50	10.1002/	Aleo	Aleo	Pfizer Inc		

Links(doi: string, authors: string)							
doi	authors						
10.1021/	Ahlin, G;						
10.1021/	Ahlin, G;						
10.1002/	Aleo, MD;						

DrugBank(drugbank_id: string, drug_name: string, drug_interactions: list <string>, drug_description: string, smiles: string)</string>									
drugbank_id	drug_name	drug_interactions	drug_description	smiles					
DB00459	Acitretin	['DB00304', 'DB00367', 'DB00162']	An oral retinoid effective	CC(CO					
DB00523	Alitretinoin	['DB00304', 'DB00367', 'DB00162']	An important regulator of	COc1c					
DB00964	Apraclonidine	['DB06237']	Apraclonidine, also known as	CCc1c					