

**Curation and Analysis of Pharmacogenomics data of  
Dermatological Disorders**

A DISSERTATION  
SUBMITTED IN PARTIAL FULFILLMENT OF THE REQUIREMENTS FOR THE AWARD  
OF THE DEGREE  
OF  
**MASTER OF TECHNOLOGY**  
IN  
**BIOINFORMATICS**

SUBMITTED BY  
**RUCHI SHARMA**  
(2K16/BIO/06)

UNDER THE SUPERVISION  
OF  
**DR. YASHA HASIJA**



**DEPARTMENT OF BIOTECHNOLOGY  
DELHI TECHNOLOGICAL UNIVERSITY**

(Formerly Delhi College of Engineering)  
Shahbad Daultpur, Main Bawana Road  
Delhi-110042, India  
JUNE 2018

# **DELHI TECHNOLOGICAL UNIVERSITY**

(Formerly Delhi College of Engineering)

Bawana Road, Delhi-110042

## **CANDIDATE'S DECLARATION**

I, **Ruchi Sharma**, Roll no. 2K16/BIO/06, student of **M.Tech (Bioinformatics)**, hereby declare that the project Dissertation titled “**Curation and Analysis of Pharmacogenomics data of dermatological Disorders**” which is submitted by me to the Department of Biotechnology, Delhi Technological University, Delhi in partial fulfilment of the requirement for the award of the degree of Master of Technology, is original and not copied from any source without proper citation. This work has not previously formed the basis for the award of any Degree, Diploma Associate ship, Fellowship or other similar title or recognition.

Place: Delhi

Date:

**Ruchi Sharma**

M.Tech. (Bioinformatics)

2K16/BIO/06

# **DEPARTMENT OF BIOTECHNOLOGY**

**DELHI TECHNOLOGICAL UNIVERSITY**

(Formerly Delhi College of Engineering)

Bawana Road, Delhi-110042

## **CERTIFICATE**

I, hereby certify that the Project Dissertation titled “**Curation and Analysis of Pharmacogenomics data of Dermatological Disorders**” which is submitted by **Ruchi Sharma**. 2K16/BIO/06, Department of Biotechnology, Delhi Technological University, Delhi in partial fulfilment of the requirement for the award of the degree of Master of Technology, is a record of the project work carried out by the student under my supervision. To the best of my knowledge this work has not been submitted in part or full for any Degree or Diploma to this University or elsewhere.

Place: Delhi

Date:

**(Dr. Yasha Hasija)**

**Supervisor**

**Assistant Professor**

**Department of Biotechnology**

**Delhi Technological University**

**(Dr. Jaigopal Sharma)**

**Head of Department**

**Professor**

**Department of Biotechnology**

**Delhi Technological University**

## **ABSTRACT**

A significant proportion of patients with skin disease do not respond to treatment and adverse drug reactions are a common problem. Genetic factors are important determinants for both drug efficacy and toxicity. The fields of pharmacogenetics and pharmacogenomics examine inter-individual variations in the DNA sequences that are related to drug efficacy and toxicity. The notion of treating the patient, and not the particular disease, has been emphasized by physicians for some time. In the past decade, this idea advanced with the human genome project, and has been taken further with the advent of personalized dermatology, or using genetics to drive pharmacological treatment. Although some dermatological conditions such as melanoma are being targeted with gene-specific therapy, the idea of choosing a drug based on the genetic makeup to treat other dermatologic conditions might be relevant, since it may increase drug efficacy or decrease adverse drug events. This concept of pharmacogenomics could be applied throughout the field of dermatology. Online libraries have been developed to guide drug efficacy, dose prediction and adverse events. We provide a list of current systemic dermatologic drugs in which the pharmacokinetics and pharmacodynamics have been studied. It would be beneficial to guide patient treatment with these drugs, if we can better understand their pharmacogenomics.

# ACKNOWLEDGEMENT

*I wish to express my deep sense of gratitude and indebtedness to **Dr. Yasha Hasija**, Assistant Professor, Department of Biotechnology, DTU, Delhi; for introducing the present topic and for her inspiring guidance, constructive and valuable suggestion throughout this work.*

*I am heartily thankful for her guidance and support during my project. Her able knowledge and expert supervision with unswerving patience fathered my work at every stage, for without her warm affection and encouragement, the fulfilment of the task would have been very difficult.*

*I would also like to extend my heartfelt gratitude towards all the members of computational biology lab for gratuitously helping me in the successful completion of the project.*

*I am genuinely appreciative my parents and all of my friends for their suggestions and moral support during my work.*

# TABLE OF CONTENTS

CANDIDATE'S DECLARATION.....	iii
CERTIFICATE.....	iv
ABSTRACT.....	v
ACKNOWLEDGEMENT.....	vi
LIST OF FIGURES.....	vii
LIST OF TABLES.....	viii
LIST OF ABBREVIATIONS.....	ix
Chapter 1 Introduction.....	1
Chapter 2 Review of Literature.....	3
2.1 Pharmacogenetics and Pharmacogenomics.....	3
2.2 Genome Wide association Studies (GWAS).....	5
2.3 Other Resources .....	5
2.3.1 PharmGKB.....	5
2.3.2 Drugbank.....	6
2.3.3 CPIC.....	8
Chapter 3 Material and Methods.....	9
3.1 Data Curation.....	9
3.1.1 PharmGKB.....	9
3.1.2 DrugBank.....	13
3.1.3 OMIM.....	15
3.1.4 Orphanet.....	17
3.2 Data Cleaning.....	18
3.3 Data Analysis.....	18

Chapter 4 Result and Discussion.....	20
Chapter 5 Conclusion.....	29
Chapter 6 References.....	30
Chapter 7 Websites.....	32

## LIST OF FIGURES

<b>Figure 1</b>	Pharmacogenomics-genetic markers to identify patients who will benefit
<b>Figure 2</b>	Factors that may affect drug efficacy and drug toxicities
<b>Figure 3</b>	Pharmacogenetics-pharmacogenomics
<b>Figure 4</b>	PharmGKB pyramid
<b>Figure 5</b>	Screenshot of Drugbank homepage
<b>Figure 6</b>	CPIC levels for drugs
<b>Figure 7</b>	PharmGKB homepage
<b>Figure 8</b>	Vemurafenib pathway for EGFR gene
<b>Figure 9</b>	The EGFR gene overview page that shows links to different attributes
<b>Figure 10</b>	Screenshot showing clinical annotation for EGFR gene
<b>Figure 11</b>	Screenshot showing variant annotation for EGFR gene
<b>Figure 12</b>	Detailed page of Drugbank homepage showing browsing details
<b>Figure 13</b>	Detailed page of Drugbank homepage showing searching details
<b>Figure 14</b>	User-interface of OMIM
<b>Figure 15</b>	Number of gene and phenotype entries in OMIM
<b>Figure 16</b>	Orphanet homepage
<b>Figure 17</b>	Flow diagram of methodology



## LIST OF TABLES

<b>Table 1</b>	Drug labels of dermatological disorders by different associations
<b>Table 2</b>	Top prediction of VIPs for dermatological disorders
<b>Table 3</b>	Pharma drugs with their targets in some dermatological disorders

## **LIST OF ABBREVIATIONS**

<b>BioPAX</b>	Biological Pathway Exchange
<b>CPIC</b>	Clinical Pharmacogenetics Implementation Consortium
<b>DNA</b>	Deoxyribonucleic Acid
<b>EGFR</b>	Epidermal Growth Factor receptor
<b>EMA</b>	European Medicines Agency
<b>FDA</b>	Food and Drug Administration
<b>GPML</b>	Gaussian Processes for Machine Learning
<b>GWAS</b>	Genome-wide Association Studies
<b>HCSC</b>	Health care Service Corporation
<b>HGNC</b>	HUGO gene Nomenclature Committee
<b>MS</b>	Mass Spectrometry
<b>NCBI</b>	National Center for Biotechnology
<b>NMR</b>	Nuclear magnetic Resonance
<b>OMIM</b>	Online Mendelian Inheritance of Man
<b>PharmGKB</b>	Pharmacogenomics Knowledge base
<b>PD</b>	Pharmacodynamics
<b>PK</b>	Pharmacokinetics
<b>PGx</b>	Pharmacogenomics
<b>PMDA</b>	Pharmaceuticals and Medical Devices agency
<b>TSV</b>	Tab-separated Value
<b>USA</b>	United States of America
<b>VIPs</b>	Very important pharmacogenes

## **Chapter-1 INTRODUCTION**

Every person has different effects on medication because of their variation in genomic structure. How the drugs respond at germ line and somatic level can be predicted through a study known as pharmacogenetics. Development of pharmacogenetics is based on how the genes identified and their allelic variants affects drug response. While PGx is a new field that explains the identification of human genes, their products, and their variation express and how they function. Pharmacogenomics used these data to predict right treatment for patients and helps in development of new drugs. Main difference between pharmacogenetics and pharmacogenomics is that pharmacogenetics used to study single genes and their effect on individual while pharmacogenomics used in broader context i.e. for study the whole genome in respect to all genes function and their interactions and it is caused by pharmacokinetics and pharmacodynamics or both. Study that helps in knowing individual's genetic behavior in response to drugs is known as pharmacogenomics [Berlin DS et al. 2010]. It is a very new field in science that combines two fields i.e. Pharmacology and Genomics. Pharmacology is defined as the study of drugs while genomics is used for study of genes and their functions. It means pharmacogenomics deals with both study of genes and drugs and how they work. Hence, it is widely used to make new drugs that are more effective and safe according to patient's genetic makeup. Drugs that are available in market follow "one size fits all" theorem i.e. one medication is used for several diseases but they are not fit for all individual who are taking it. This becomes very difficult task to prove which medication is perfect for which individual and do not cause any side effects. These effects are known as adverse drug reactions because of these reactions there are more chances for hospitalization and death of an individual [Nelson MR et al. 2009]. After the discovery of Human Genome Project, we can learn genetic differences are affected any patient's response to any medication. These differences help to predict right medication according to patient's genetic makeup [Whirl-Carrillo M, et al. 2012]. As we know, genes are responsible for making of human proteins, enzymes, their receptors and other molecules that are involved in drug and pathways of diseases. Pharmacogenomics can differentiate patients and diseases with the help of variants present in these genes on the basis of their genetic tests and

prescribe a drug that is effective for whom and in what dose it should be given [Evans WE et al. 2003]. We can say that pharmacogenomics helps to design new drugs and in identification of drugs effectiveness before symptoms are apparent. Pharmacogenomics helps companies to bring their drugs that are FDA approved more quickly in the market because it reduces the risk of clinical trials failures by giving exact knowledge of people's genetic behavior and drugs that are effective for them at genetic level. Pharmacogenomics also reduces time and number of trials to prove drugs efficacy and safety with respect to patient's genetic makeup.

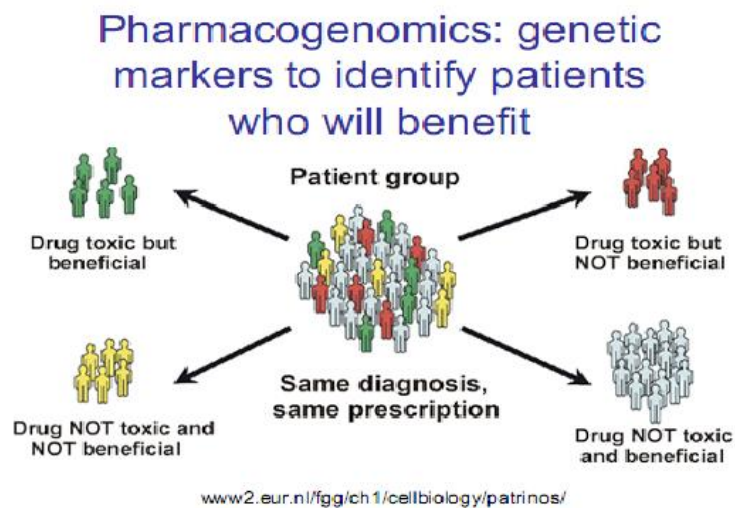


Figure1: Pharmacogenomics – Genetics markers to identify patients who will benefit.

## **Chapter-2 REVIEW OF LITERATURE**

### **2.1 Pharmacogenetics and Pharmacogenomics**

In human genomic sequence, the first major clinical application in advances is pharmacogenetics. There are so many responsibilities to these advances in pharmacology which include maximizing drug efficacy, minimizing toxicity and providing selective medication to patient according to their genetic makeup [Whirl-Carrillo M et al. 2012]. Advancement in pharmacogenetics results into pharmacogenomics. Pharmacogenomics include studies of monogenic to polygenic traits and make genomic science to genome-wide studies. Their effects are classified as those which alter factors that influence the drug concentration reaches to its target, hence called pharmacokinetics factors. Those effects that involve the target itself, known as pharmacodynamics factors. Properties of drug that influence pharmacokinetics include their absorption, distribution, metabolism and excretion. Recently, the main focus in pharmacogenomics is increasing on pathways that include both pharmacokinetic and pharmacodynamics [Giacomini KM et al. 2007].

Figure 1. Factors That May Affect Drugs' Efficacy and Toxicities

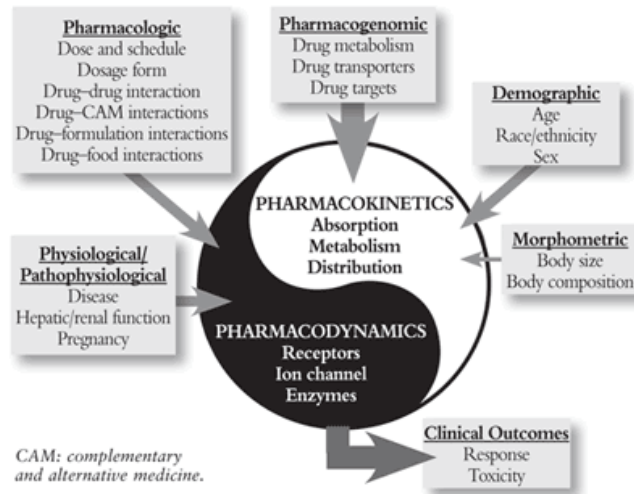


Figure2: factors that may affect drug efficacy and drug toxicities

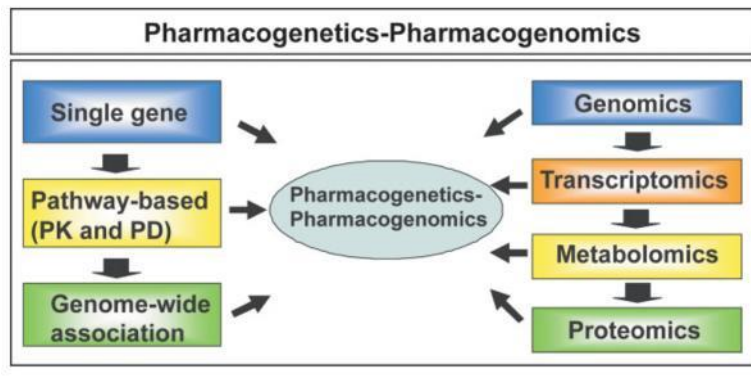


Figure3: pharmacogenetics-pharmacogenomics

## **2.2 Genome-wide association studies (GWAS)**

It is the studies that measures and analyze DNA sequence variation in whole genome so as to identify genetic risk factors that are most common in population [Crowley JJ et al. 2009]. The main goal of GWAS is to use genetic risk factors to predict the risk and identify disease susceptibility for developing new prevention and treatment analyses [Yee SW, et al. 2016]. The success of pharmacogenomics GWAS depends on allelic frequency of genetic variants, their effect size that influence traits. There are so many advantages of GWAS in pharmacogenomics that includes:

1. GWAS provides that information that is not available to genetic contribution of pharmacogenomics traits.
2. Pharmacogenomics GWAS directly investigate the role of genetic variation on clinical trials results.

## **2.3 Other resources**

### **2.3.1 PharmGKB**

Previously there was no standard format to store and describe genotypic and phenotypic data that was collected from pharmacogenetics studies. In 2000, a database as started as one of the first ‘post-genomic’ databases known as PharmGKB [Klein TE et al. 2001]. The main challenge of PharmGKB is to maintain the data quality without compromising privacy of subject [Klein TE et al. 2001]. It presents the data in a new way with increasing volumes of data. PharmGKB built new relationships with other known resources like drugbank, Biopax, university of California Santa Cruz (CA, USA) genome browser [Altman RB et al. 2013]. The main motive to make new relations is to enhance the knowledge to pharmacogenomics. At present, PharmGKB collects and annotate pharmacogenomics data from different sources and data can access from related gene, drug and disease tabs [Owen RP et al. 2008].

The PharmGKB Knowledge Pyramid provides users with an overview of the different types of information found in our knowledgebase and shows how this information is acquired and integrated together. PGx knowledge is accumulated at the bottom of the pyramid, to the implementation of PGx in the clinic at the top.

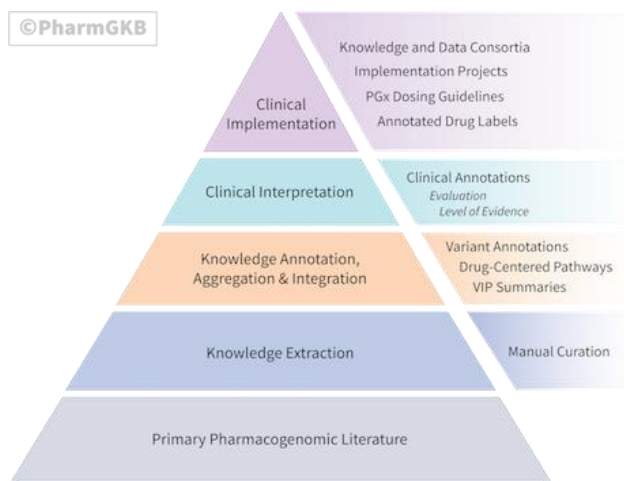


Figure4: PharmGKB pyramid

### **2.3.2 Drugbank**

A unique bioinformatics/cheminformatics resource that provides complete data about drug and their targets information is called DrugBank. It contains approximately >4100 drug entries that includes >800 FDA approved drug, >14000 protein or drug targets sequences [Knox C et al. 2011]. Its main focus is on quantitative, analytic or molecular-scale information about drugs and their targets. It is a fully searchable web-based resource that contains many built-in tools and features that allow us to view, sort and extract drug or drug targets data [Wishart DS et al. 2016]. The aim of Drugbank is to provide comprehensive resources on drugs that provide their pharmacological actions, biochemical information, their mechanism and targets. It is a database of drugs that facilitate in-silico design of drug, drug targets discovery, prediction of drug metabolism, their interaction etc.



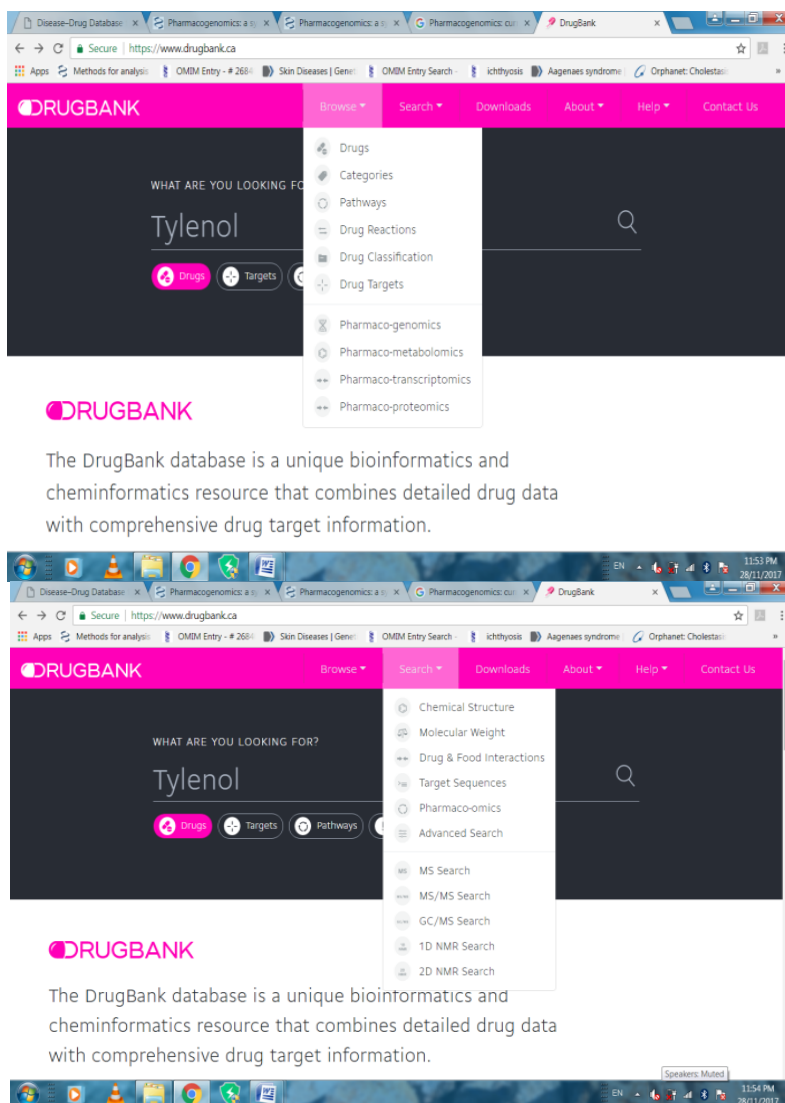


Figure5: Screenshot of DrugBank homepage.

### 2.3.3 CPIC

A shared project was started in late 2009 between Pharmacogenomics knowledge base and pharmacogenomics research network (PGRN) is named as Clinical Pharmacogenetics Implementation Consortium (CPIC). It is an international consortium in which an individual person volunteers and staff containing very small number of peoples who are dedicated to their work are come [Relling MV et. Al. 2011]. These staff members are facilitated to use pharmacogenetics tests for patient care. Its main targets is to translate genetic laboratory test results into taking decisions for drug prescription by provide all the information about gene/drug clinical practice guidelines and make them freely available and updatable data.

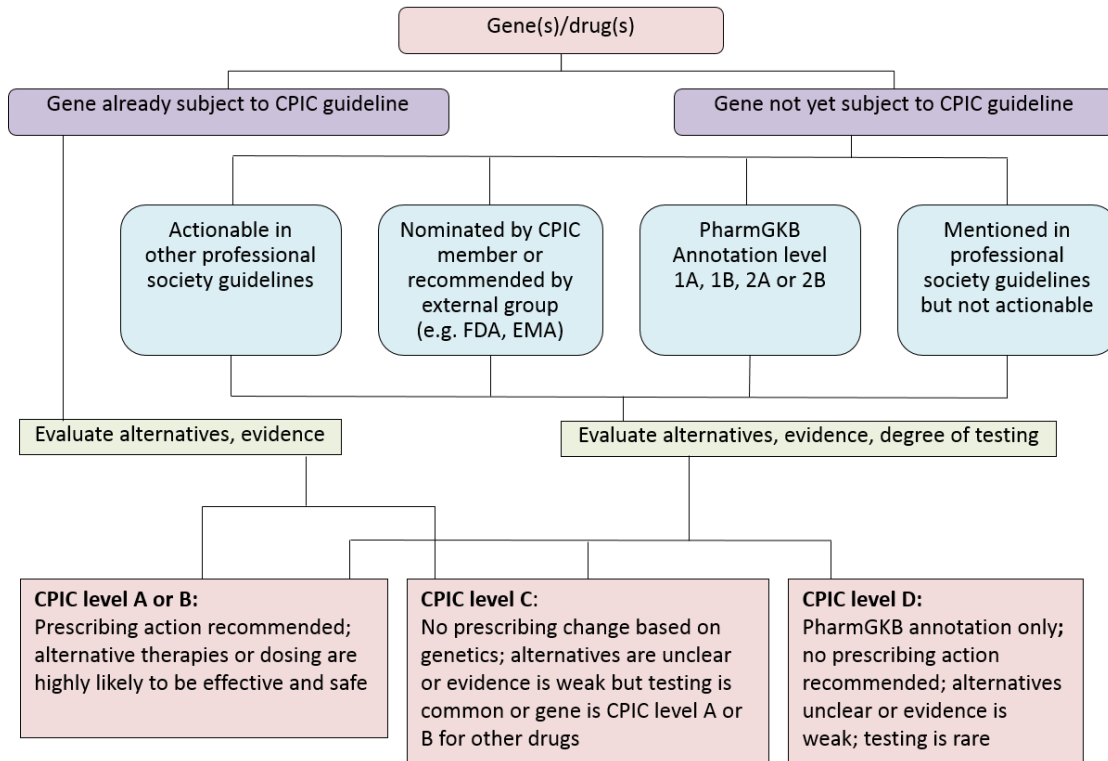


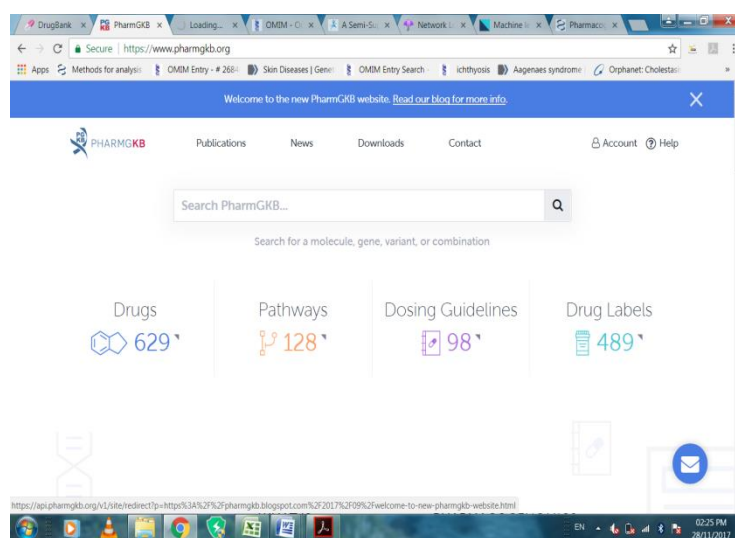
Figure6: CPIC levels for drugs

# **Chapter-3 MATERIAL AND METHOD**

## **3.1 Data curation**

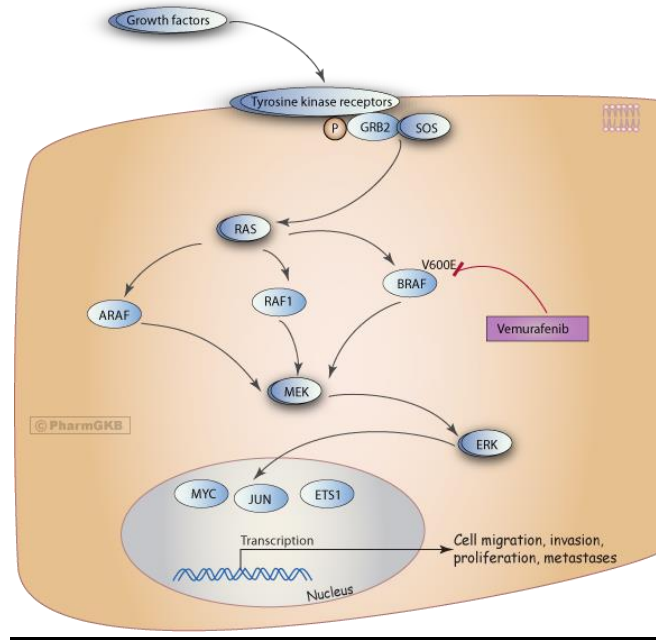
### **3.1.1 PharmGKB**

PharmGKB is a tool that provides access to all the pharmacogenetics and pharmacogenomics data. It contains information in a fully structured format [Sanguhl K et al. 2008]. In the homepage of PharmGKB, we can search by using disease name, drug, variant or combination. Some tabs are also present where we can search for clinical pharmacogenomics, PGx research, overview, VIPs, haplotypes, pathways etc. as shown in figure given below [Thorn CF et al. 2009].



**Figure7:** Homepage of PharmGKB

- From PharmGKB tool, we opt for pathways, clinical annotation, variant annotation, gene location. Pathways that are present in PharmGKB are evidence-based diagrams that depict the PK and PD of a drug [Hodge AE et al. 2007]. Drugs that are used in PharmGKB pathways are taken from various other sources. It also contains other important PGx related information. We can easily download all the information present in pathways in the format of TSV, BioPAX and GPML.
- Clinical annotation provides information about variant-drug relations. In clinical annotation, phenotype for known genotype can be related to other genotypes that are associated. The figure below contains different level of clinical annotations. For viewing clinical annotation, one can use search box at the top of the page and search for their interested gene or drug. A new page is open where you can click on “clinical annotation” tab in left-hand menu. Each row represents single clinical annotation. From clinical annotation we choose here levels, variants, molecules and their types.
- Variant annotation provides information about the association between variant and phenotype information that are present here, are directly taken from publication. To search for variant annotation, click on “variant annotation” tab in left-hand menu on gene page.



**Figure8:** the Vemurafenib Pathway for EGFR gene

The screenshot shows the PharmGKB website for the EGFR gene overview. The page includes navigation tabs for Overview, Prescribing Info, Clinical Annotations, Haplotypes, and Pathways. It displays a 'Very Important PGx Gene Summary' and a table for the gene's location on chromosome 7.

Location	
Strand	chr7 : Plus
Cytogenetic	chr7 : p11.2 - p11.2
GRCh37.p10	chr7 : 55086678 - 55279262
GRCh38.p7	chr7 : 55019032 - 55207338

**Figure9:** the EGFR gene overview page that shows links to different attributes.

LEVEL	VARIANT	GENE	MOLECULE	TYPE	PHENOTYPE
Level 1B	rs121434568	EGFR	gefitinib	efficacy	
Level 1B	rs121434568	EGFR	erlotinib	efficacy	
Level 2A	rs121434568	EGFR	carboplatin, gefitinib, paclitaxel	efficacy	
Level 2A	rs121434569	EGFR	erlotinib, gefitinib	efficacy	
Level 2A	rs121434568	EGFR	carboplatin, docetaxel, erlotinib, gemcitabine, paclitaxel	efficacy	
Level 2B	rs11568315	EGFR	gefitinib	efficacy	
Level 3	rs2227983	EGFR	egfr inhibitors	toxicity	
Level 3	rs2293347	EGFR	gefitinib	efficacy	
Level 3	rs712830	EGFR	cetuximab, irinotecan, leucovorin, teniposide	toxicity	

Figure10: screenshot showing clinical annotation for EGFR gene

VARIANT	PMID	MOLECULES	ASSOCIATION	SIGNIFICANT	PVALUE	# OF
rs2227983	16788380	cetuximab	Sign in to view	no		39
rs712829	18652519	erlotinib	Sign in to view	not stated		
rs712829	18652519	geldanamycin	Sign in to view	not stated		
rs712829	18652519	topoisomerase Inhibitors	Sign in to view	not stated		
rs712829	18652519	Alkylating Agents	Sign in to view	not stated		
rs712829	17375033	gefitinib	Sign in to view	yes	0.005	92
rs712829	18006781		Sign in to view	yes	0.005	
rs712829	18006781	egfr inhibitors	Sign in to view	yes		

Figure11: screenshot showing variant annotation of EGFR gene

### 3.1.2 DrugBank

Drugbank is a resource which is used to search information about drug structure and their targets, their metabolism, actions, side-effects, cost and we can also identify drug by their MS or NMR spectra. It is also used for repurpose existing drugs and to design new drugs based on the information that it provides [ Wishart DS et al. 2017]. On the DrugBank homepage, we have different tabs for browse, search, downloads, about, help and contact us. From browse we can search any drug on the basis of their category, pathways etc. from the data available on Drugbank we use here drug, drug group, their targets and enzymes.

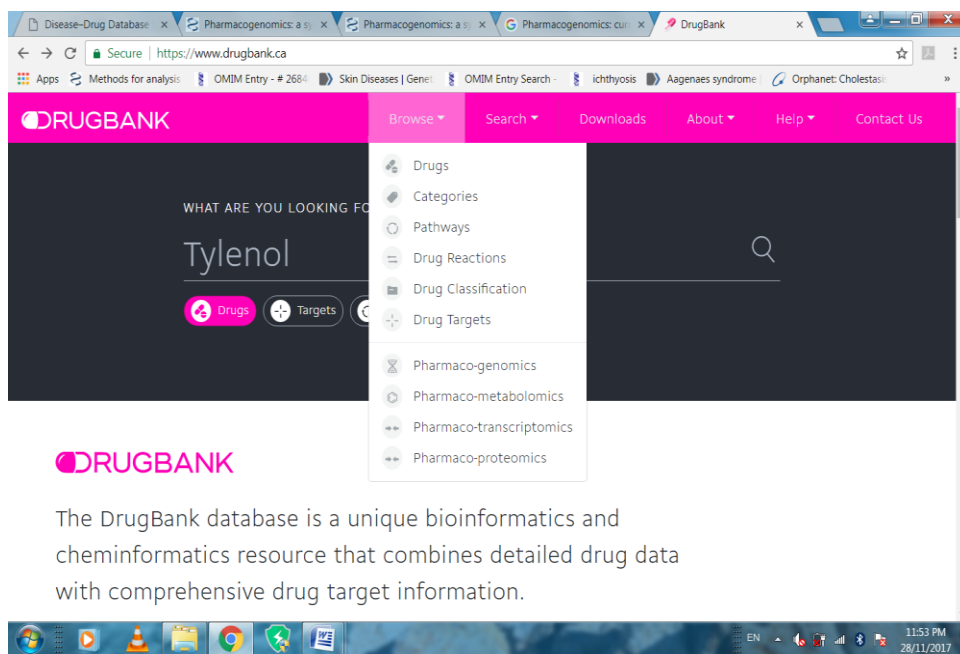


Figure12: detailed page of DrugBank homepage showing browsing details.

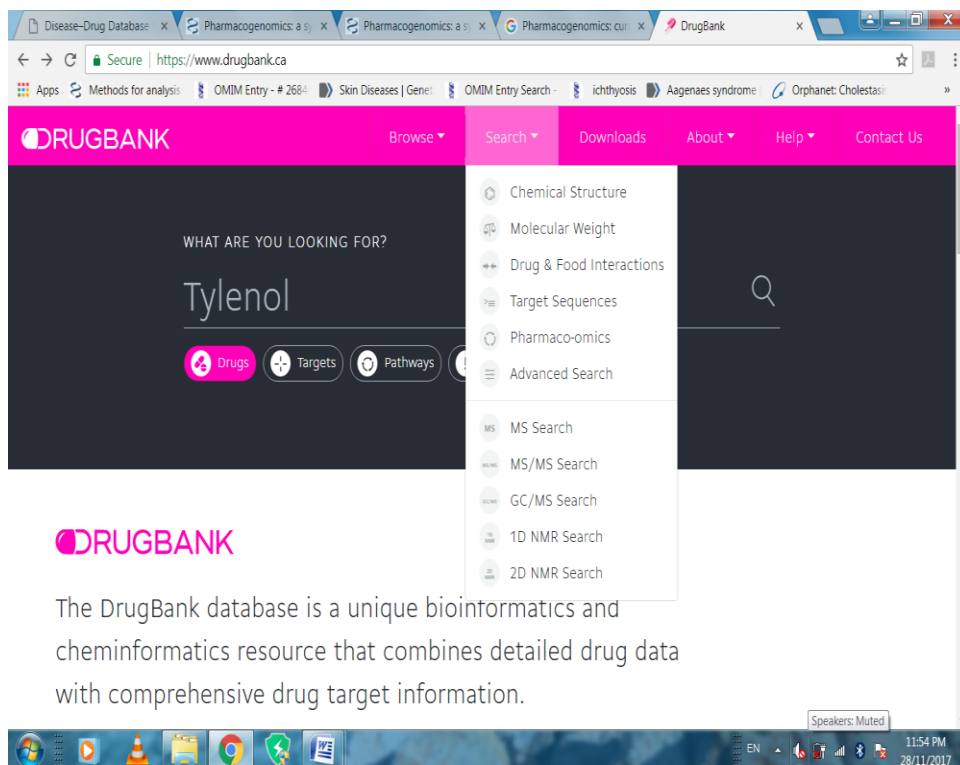


Figure13: detailed page of Drugbank homepage showing searching details.



### **3.1.3 OMIM**

Online Mendelian Inheritance in Man (OMIM) is a complete, reliable and punctual knowledgebase of human genes and genetic disorders that are assemble to support human genetics research and education as well as clinical genetics practices. Each OMIM entry contains full abstract of a genetically determined phenotype and/or gene and it contains number of links to other genetic databases such as DNA and protein sequence [Amberger J et al. 2009]. OMIM is an easy, simple and straight portal to the expand information in human genetics. We can search OMIM from its homepage or from any other page in the NCBI Entrez suite of databases.

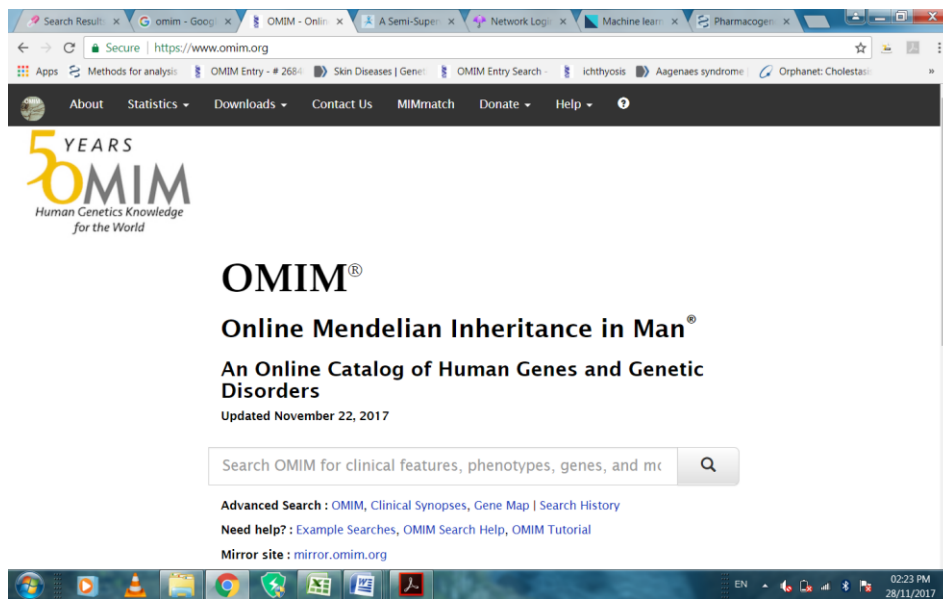


Figure14: User interface of OMIM.

**Table 1.**  
Number of gene and phenotype entries in OMIM as on September 13, 2004

	Autosomal	X-linked	Y-linked	Mitochondrial	Total
* Gene with known sequence	9318	414	47	37	9816
+ Gene with known sequence and phenotype	356	36	0	0	392
# Phenotype description, molecular basis known	1462	134	1	25	1622
% Mendelian phenotype or locus, molecular basis unknown	1288	130	4	0	1422
Other, mainly phenotypes with suspected mendelian basis	2185	154	2	0	2341
Total	14 609	868	54	62	15 593

**Figure15:** number of gene and phenotype entries in OMIM

### 3.1.4 Orphanet

The nomenclature of rare disorders used by Orphanet, the reference resource for information on rare diseases and ORPHAN drugs. Since 1997 Orphanet maintains an inventory of rare diseases. Orphanet indexes any disease described at least two patients and in less than 50 per 100,000 persons in the General European Population. To search for a particular disease or several diseases and to identify their corresponding ORPHA numbers several options are available. The web-based Orphanet information portal allows you to search for one disease at a time. Now on entering the gene name that we wish to consult click on search represented with the list of results that could interest, we select the gene you wish to consult then we will arrive on the page concerning the gene. We will find the name symbol of the gene, at the top of the page we will then find the gene identity card with the synonyms previous symbols and names of the gene type of gene and chromosome location. We also provide cross reference with other nomenclature and resources such as OMIM, HGNC, uniprot, ensembl, etc.

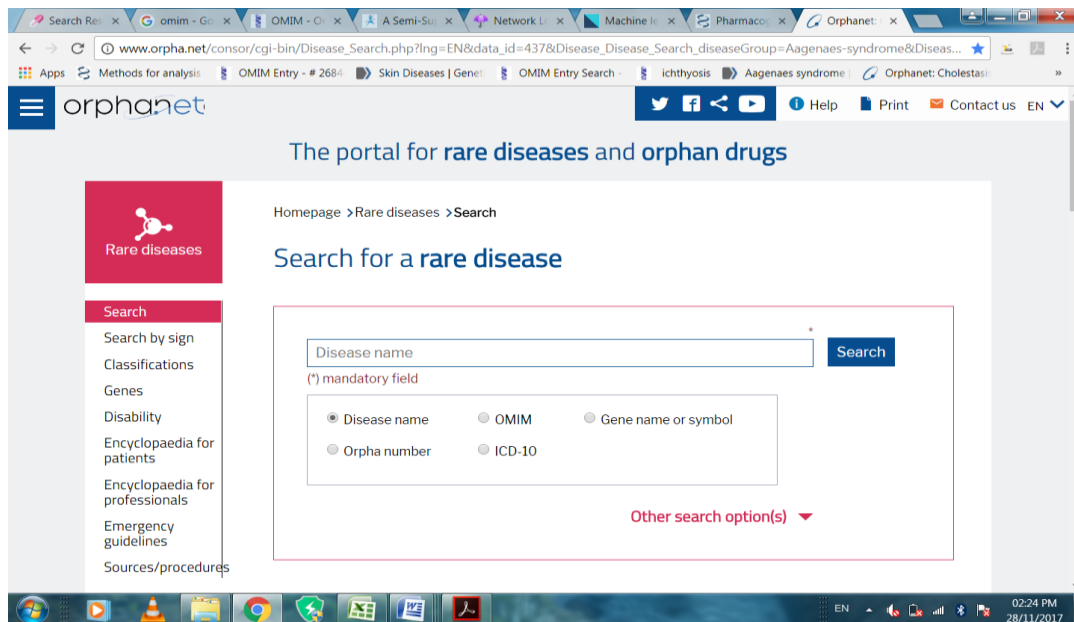


Figure16: Orphanet homepage

## **3.2 Cleaning data**

Now we are going to clean-up the raw data that isn't exactly perfect for analyzing. Some of the data are not present for all genes so we exclude that data. Genes whose data is not present we remove that gene from our citation process.

## **3.3 Data analysis**

We have around 1532 genes of approximately 214 diseases that are taken from mesh browser. Through OMIM we found attributes of 176 diseases out of 214 diseases and we curate them. From all the data present in PharmGKB, we analysis that total of 65 VIPs gene's data are present in PharmGKB out of which only 29 VIPs gene are for our dermatological disorders which we are taken here.

There are 501 drug labels entries in PharmGKB, from which only 49 drug labels are for our dermatological disorders present. From these drug labels we found 24 drugs whose targets we had taken from drugbank database.

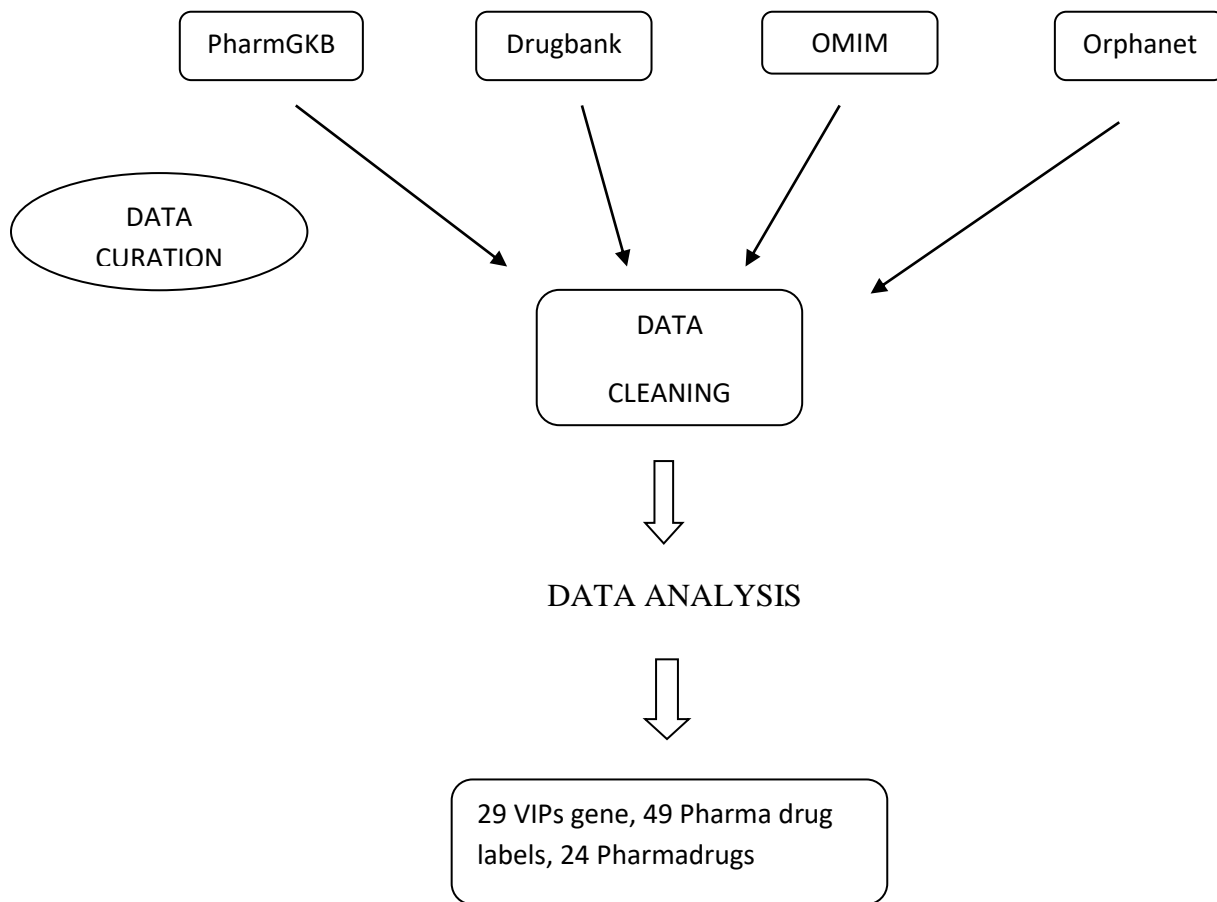


Figure17: flow chart of methodology

## **Chapter- 4 RESULT AND DISCUSSION**

- From the above methods that are mentioned, we found different drugs that contain PGx information approved by FDA, EMA, PMDA and HCSC. There are around 501 drug labels on PharmGKB, from which around 49 drug labels are for dermatological disorders. Table containing all the drug labels of dermatological disorders.

**Table1:** Drug labels of dermatological disorders approved by different associations.

<b><u>DISEASES</u></b>	<b><u>DRUGS</u></b>	<b><u>FDA</u></b>	<b><u>EMA</u></b>	<b><u>HCSC</u></b>	<b><u>PMDA</u></b>
Bullous Congenital Ichthyosiform Erythroderma	Vandetanib	-	Testing recommended	-	-
	Sulfadiazine	Actionable PGx	-	Actionable PGx	Actionable PGx
Harlequin Fetus	peginterferon alfa-2b	Actionable PGx	-	-	-
Ichthyosis Hystrix	Tretinoin	Testing required	-	Testing required	Actionable PGx
Ichthyosis Vulgaris	Tretinoin	Testing required	-	Testing required	Actionable PGx
	Nalidixic Acid	Actionable PGx	-		Actionable PGx
Hereditary Palmoplantar Keratoderma Epidermolytic	Tretinoin	Testing required	-	Testing required	Actionable PGx
Hyperkeratosis	Dabrafenib	Testing required, actionable PGx, informative PGx	Testing required	Testing required	-
	Tretinoin	Testing required	-	Testing required	Actionable PGx
Keratosis Follicularis Spinulosa Decalvans	Tretinoin	Testing required	-	Testing required	Actionable PGx

Pachyonychia Congenita	Quinine	Actionable PGx	-	Actionable PGx	-
	Tretinoin	Testing required	-	Testing required	Actionable PGx
Dyskeratosis Congenita	Quinine	Actionable PGx	-	Actionable PGx	-
Ataxia Telangiectasia	Carisoprodol	Actionable PGx	-	-	-
	Succimer	Informative PGx	-	-	-
	Cerliponase alfa	Testing required	-	-	-
	Fosphenytoin	Actionable PGx	-	Testing recommended	-
	Pantoprazole	Actionable PGx	-	-	-
	Carbamazepine	Testing required, actionable PGx	-	Testing recommended	Actionable PGx
	Diazepam	Actionable PGx	-	-	-
	Imipramine	Actionable PGx	-	-	-
Cutis Marmorata Telangiectatica Congenita	<u>Quinine</u>	Actionable PGx	-	Actionable PGx	-
	<u>Tretinoin</u>	Testing required	-	Testing required	Actionable PGx
Hereditary Hemorrhagic	peginterferon alfa-2b	Actionable PGx	-	-	-
Aplasia Cutis Congenita	<u>Enasidenib</u>	Testing required	-	-	-
	<u>Quinine</u>	Actionable PGx	-	Actionable PGx	-
	Tretinoin	Testing required	-	Testing required	Actionable PGx
Focal Dermal Hypoplasia	<u>Azathioprine</u>	Testing recommended	-	Actionable PGx	Actionable PGx
	<u>Carbamazepine</u>	Testing required, actionable PGx	-	Testing recommended	Actionable PGx
Albright Hereditary Osteodystrophy	Anastrozole	Testing required	-	Testing required	-

Familial Dysautonomia	<u>Evolocumab</u>	Informative PGx	-	-	-
	<u>Lomitapide</u>	Informative PGx	Testing required	Informative PGx	-
	<u>Celecoxib</u>	Actionable PGx has dosing info	-	Actionable PGx	Actionable PGx
	<u>Alirocumab</u>	Informative PGx	-	-	-
	<u>Mipomersen</u>	Informative PGx	-	-	-
	<u>Rosuvastatin</u>	Actionable PGx	-	Actionable PGx	-
Familial Dysautonomia	<u>Simvastatin</u>	Informative PGx	-	-	-
	<u>Atorvastatin</u>	Informative PGx	-	Informative PGx	Informative PGx
Hereditary Lymphedema	<u>Trametinib</u>	Testing required, Actionable PGx, informative PGx	Testing required	Testing required	-
NOMID/CINCA	<u>Donepezil</u>	Actionable PGx	-	-	-
Urticaria Pigmentosa	<u>Desloratadine</u>		Informative PGx	-	-
	<u>Doxepin</u>	Actionable PGx	-	-	-
Epidermal Nevus	<u>Cetuximab</u>	Testing required	Testing required	Testing required	Testing required
	Panitumumab	Testing required	Testing required	Testing required	Testing required
	Lapatinib	Testing required, actionable PGx	Testing required	Testing required	Testing required
	<u>Gefitinib</u>	Testing required, actionable PGx	Testing required	Testing required	Testing required
	<u>Neratinib</u>	Testing required, informative PGx	-	-	-
	<u>Osimertinib</u>	Testing required	-	-	-
	<u>Erlotinib</u>	Testing required	Testing required	Testing required	Testing required
	<u>Pertuzumab</u>	Testing	Testing	Testing	Testing



		required, actionable PGx	required	required	required
	<u>Palbociclib</u>	Testing required	-	-	-
	<u>Vandetanib</u>		Testing recommended	-	-
	<u>Afatinib</u>	Testing required	Testing required	Testing required	-
	<u>Abemaciclib</u>	Testing required	-	-	-
	<u>Brigatinib</u>	Testing required	-	-	-
	<u>Flurbiprofen</u>	Actionable PGx	-	-	-
	Telaprevir	Actionable PGx	Actionable PGx	-	-
Congenital Erythropoietic Porphyria	<u>Vandetanib</u>		Testing recommended	-	-
	<u>Sulfadiazine</u>	Actionable PGx	-	Actionable PGx	Actionable PGx
Familial Cutaneous Amyloidosis	<u>Evolocumab</u>	Informative PGx	-	-	-
	<u>Tauroursodeoxycholic acid</u>	Testing required	-	-	-
	<u>Lomitapide</u>	Informative PGx	Testing required	Informative PGx	-
	<u>Denileukin diftitox</u>	Testing required	-	-	-
	<u>Celecoxib</u>	Actionable PGx	-	Actionable PGx	Actionable PGx
	<u>Alirocumab</u>	Informative PGx	-	-	-
	<u>Mipomersen</u>	Informative PGx	-	-	-
	<u>Rosuvastatin</u>	Actionable PGx	-	Actionable PGx	-
	<u>Simvastatin</u>	Informative PGx	-	-	-
	<u>Vemurafenib</u>	Testing required, actionable PGx	Testing required	Testing required	Testing required
	<u>Atorvastatin</u>	Informative PGx	-	Informative PGx	Informative PGx
	<u>Evolocumab</u>	Informative PGx	-	-	-
	<u>Lomitapide</u>	Informative PGx	Testing required	Informative PGx	-

Familial Mucocutaneous Candidiasis	<u>Celecoxib</u>	Actionable PGx	-	Actionable PGx	Actionable PGx
	Posaconazole	-	Informative PGx	-	-
	<u>Voriconazole</u>	Actionable PGx	Informative PGx	Actionable PGx	Actionable PGx
	<u>Alirocumab</u>	Informative PGx	-	-	-
	<u>Mipomersen</u>	Informative PGx	-	-	-
	<u>Rosuvastatin</u>	Actionable PGx	-	Actionable PGx	-
	Tretinoin	Testing required	-	Testing required	Actionable PGx
	<u>Simvastatin</u>	Informative PGx	-		
	<u>Atorvastatin</u>	Informative PGx	-	Informative PGx	Informative PGx

- VIPs provide an overview of significant gene that is involved in drug metabolism. It includes all the information on the gene that includes any disease association in-depth information on the gene's pharmacogenetics. From this data we curate around 29 VIPs of dermatological disorders from total 65 VIPs whose information is present in PharmGKB.

**Table2:** Top predictive VIPs for dermatological disorders

<b><u>S.NO.</u></b>	<b><u>VIPs GENES IN PharmGKB</u></b>	<b><u>DISEASES</u></b>
1	ABCG2	Breast Neoplasms
2	ABCB1	Breast Neoplasms
3	AHR	Breast Neoplasms, Dermatitis_ Contact, Hyper pigmentation
4	ALOX5	Drug Eruptions, Urticaria
5	BRAF	LEOPARD SYNDROME 3, Cardiofaciocutaneous syndrome
6	BRCA1	Breast Neoplasms, Breast-Ovarian Cancer familial susceptibility to 1, Hereditary Breast and Ovarian Cancer Syndrome
7	COMT	Breast Neoplasms
8	CYP1A2	Porphyria Cutanea Tarda
9	CYP2E1	Drug Eruptions
10	CYP3A4	Breast Neoplasms
11	DPYD	Breast Neoplasms
12	EGFR	Chloracne, Breast Neoplasms
13	ERBB2	Breast Neoplasms
14	F5	Scleroderma_ Systemic
15	G6PD	Dermatitis_ Contact
16	GSTP1	Breast Neoplasms, Dermatitis_ Contact
17	GSTT1	Skin Neoplasms
18	HLA-B	Dermatitis, Drug Eruptions, Stevens-Johnson Syndrome, Dermatomyositis, Exanthema, Behcet Syndrome
19	KIT	Breast Neoplasms, Mastocytosis_ Systemic, Piebaldism
20	KRAS	Breast Neoplasms, Cardiofaciocutaneous syndrome, Nevus_ Sebaceous of Jadassohn
21	MTHFR	Breast Neoplasms, Drug Eruptions, Alopecia
22	NAT2	Breast Neoplasms, Dermatitis_ Occupational
23	NQO1	Dermatitis_ Contact
24	PTGIS	Stevens-Johnson Syndrome
25	SLC22A1	Breast Neoplasms
26	SLCO1B1	Breast Neoplasms
27	TYMS	Breast Neoplasms
28	UGT1A1	Dermatitis_ Contact
29	VDR	Breast Neoplasms, Alopecia

- DrugBank data releases its latest version that contains 11,678 drug entries which includes 2,606 approved small molecules drugs, 1,079 approved biotech drugs, 128 nutraceuticals and over 5,504 experimental drugs. From drugbank database we curate some Pharmadrugs and their targets [Imming P et al. 2006].

**Table3-** Pharma drugs with their targets in some dermatological disorders

<b><u>S.NO.</u></b>	<b><u>DRUGS</u></b>	<b><u>TARGETS</u></b>
1	Vandetanib	Vascular endothelial growth factor A, epidermal growth factor receptor, protein-tyrosine kinase 6, angiotensin-1 receptor, proto-oncogene trypsin-protein kinase receptor ret
2	Sulfadiazine	Dihydropteroate synthetase
3	Peg interferon alfa-2b	Interferon alpha/beta receptor 1, Interferon alpha/beta receptor 2
4	Tretinoin	Retinoic acid receptor (RXR-alpha, RXR-beta, RXR-gamma, alpha, beta, gamma), Retinal dehydrogenase (1, 2), Retinoic acid-induced protein 3, Nuclear receptor subfamily 0 group B member 1, Retinoic acid receptor responder protein 1, Lipocalin-1, Odorant-binding protein 2a, Retinol-binding protein 4, [pyruvate dehydrogenase [lipoamide]] kinase isozyme4, cytochrome P450 (26A1, 26B1, 26C1), Hematopoietic prostaglandin D synthase
5	Nalidixic acid	DNA
6	Dabrafenib	Serine/threonine-protein kinase (B-raf, Nek11, SIK1), RAF proto-oncogene serine/threonine-protein kinase, LIM domain kinase 1
7	Quinine	Fe(II)-protoporphyrin IX, Platelet glycoprotein IX, Intermediate conductance calcium-activated potassium channel protein 4
8	Carisoprodol	Gamma-amino butyric acid receptor subunit (alpha-1,beta-2, gamma-2)
9	Succimer	Lead, Mercury, Cadmium, Arsenic
10	Cerliponase alfa	Cation-independent mannose-6-phosphate receptor
11	Fosphenytoin	Sodium channel protein type 5 subunit alpha
12	Pantoprazole	Potassium-transporting ATPase alpha chain 1
13	Carbamazepine	Sodium channel protein type 5 subunit alpha, Neuronal acetylcholine receptor subunit (alpha-4,beta-2), Neuronal receptor subfamily 1 group I member 2
14	Diazepam	Gamma-amino butyric acid receptor subunit alpha-(1/ 2/ 3/ 5), gamma-amino butyric acid receptor subunit beta-(1/ 2/ 3), gamma-alpha butyric acid receptor subunit gamma-(1/

		2/ 3), gamma-amino butyric acid receptor subunit (delta/ epsilon/ pi/ rho-1/ rho-2/ rho-3/ theta), translocator protein, GABA-A receptor (anion channel) (protein group)
15	Imipramine	Sodium-dependent (noradrenaline/serotonin) transporter, sodium dependent serotonin transporter, 5-hydroxytryptamine receptor-(1A/ 2A/6), histamine H1 receptor, alpha-1a adrenergic receptor, alpha-1d adrenergic receptor, muscarinic acetylcholine receptor (M1/M2/M3/M4/M5), potassium voltage-gated channel subfamily D member (2, 3), 5-hydroxytryptamine receptor-(2C/7), alpha-1B adrenergic receptor, [D(1)/ D(2)] dopamine receptor (protein group), Potassium voltage-gated channel subfamily H member (1,2), sodium dependent dopamine transporter, alpha-1-acid glycoprotein 2
16	Enasidenib	Isocitrate dehydrogenase [NADP], mitochondrial
17	Azathioprine	Hypoxanthine-guanine phosphoribosyltransferase, Ras-related C3 botulinum toxin substrate 1
18	Anastrozole	Cytochrome P450 19A1
19	Evolucumab	-
20	Lomitapide	Microsomal triglyceride transfer protein large subunit
21	Celecoxib	Prostaglandin G/H synthase2, 3-phosphoinositide-dependent protein kinase 1, carbonic anhydrase (2/3), ATP-binding cassette sub-family B member 5, ATP-binding cassette sub-family G member 2, Multidrug resistance protein 1
22	Alirocumab	Proprotein convertase subtilisin/kexin type 9
23	Mipomersen	mRNA of ApoB-100
24	Rosuvastatin	3-hydroxy-3-methylglutaryl-coenzyme A reductase
25	Simvastatin	3-hydroxy-3-methylglutaryl-coenzyme A reductase, Integrin beta-2, Integrin alpha-1
26	Atorvastatin	3-hydroxy-3-methylglutaryl-coenzyme A reductase, Dipeptidyl peptidase 4, Aryl hydrocarbon receptor
27	Tremetinib	-
28	Donepezil	Acetylcholinesterase, 5-hydroxytryptamine receptor 2A
29	Desloratadine	Histamine H1 receptor
30	Doxepin	Histamine (H1/H2/H4) receptor, Sodium-dependent noradrenaline transporter, sodium-dependent serotonin transporter, 5-hydroxytryptamine receptor(2A/ 2B/2C), Muscarinic acetylcholine receptor (M1/M2/M3/M4/M5), (Alpha-1A/1B/1D/2A/2B/2C) adrenergic receptor, 5-hydroxytryptamine receptor (1A/6), D(2) dopamine receptor, Potassium voltage-gated channel subfamily H member2
31	Cetuximab	Epidermal growth factor receptor, low affinity immunoglobulin gamma Fc region receptor (II-a/II-b/II-c/III-A/III-B), complement C1r subcomponent, complement C1q subcomponent subunit (A/B/C), complement C1s subcomponent, high affinity immunoglobulin gamma Fc receptor I

32	Panitumumab	Epidermal growth factor receptor
33	Lapatinib	Epidermal growth factor receptor, Receptor tyrosine-protein kinase erbB-2
34	Gefitinib	Epidermal growth factor receptor
35	Neratinib	Epidermal growth factor receptor
36	Osimertinib	Epidermal growth factor receptor
37	Pertuzumab	Receptor tyrosine-protein kinase erbB-2
38	Palbociclib	Cyclin-dependent kinase 4, Cyclin-dependent kinase 6
39	Afatinib	Epidermal growth factor receptor, Receptor tyrosine-protein kinase erbB-2, Receptor tyrosine-protein kinase erbB-4
40	Abemaciclib	Cyclin-dependent kinase 4, Cyclin-dependent kinase 6
41	Brigatinib	ALK tyrosine kinase receptor, Epidermal growth factor receptor, Tyrosine-protein kinase ABL-1, Insulin like growth factor 1 receptor, Receptor-type tyrosine-protein kinase FLT3, Insulin receptor, Hepatocyte growth factor receptor, receptor tyrosine-protein kinase erbB-4, Receptor tyrosine-protein kinase erbB-2
42	Flurbiprofen	Prostaglandin G/H synthase 1, Prostaglandin G/H synthase 2
43	Telaprevir	NS3/4A protein, Solute carrier organic anion transporter family member 1B1, Solute carrier organic anion transporter family member 2B1
44	Tauroursodeoxy cholic acid	-
45	Denileukin diflitox	Interleukin-2 receptor subunit alpha, Interleukin-2 receptor subunit beta, cytokine receptor common subunit gamma
46	Vemurafenib	Serine/threonine-protein kinase B-raf
47	Posaconazole	Lanosterol 14-alpha demethylase
48	Voriconazole	Lanosterol 14-alpha demethylase
49	Erlotinib	Epidermal growth factor receptor, nuclear receptor subfamily 1 group I member 2

## **Chapter- 5 CONCLUSION**

We can conclude that advances in pharmacogenomics will lead to greater significant for clinical improvement in terms of safety and efficiency of medication that increase hundreds of times. Medication which is given to a patient depends on its pharmacokinetics, pharmacodynamics, and pharmacogenetics in combination. Currently, applications of pharmacogenomics in clinical practices are limited but this promise to expand more in future. Pharmacogenomics is used to identify new pharmacological targets and new medications by using biotechnologies such as genomic high-throughput. As the advancement in knowledge has been increasing day by day, this shows that both pharmacogenetics and pharmacogenomics has a great impact on drug research and development, their clinical trials and practices. Current drug databases provide may tool to predict patient's drug effects and also give PK and PD information about so many different drugs. By using this information one can easily increase drug efficacy and using an individual genetic makeup we can provide correct pharmacologic treatments. Although it is not available for all dermatological drugs. It gives an opportunity for doctors to correlate clinical trials with the above-mentioned database and can change the patient treatment in future.

## **Chapter-6 REFERENCES**

1. Altman RB, Flockhart DA, Sherry ST, Oliver DE, Rubin DL, Klein TE. Indexing pharmacogenetics knowledge on the World Wide Web. *Pharmacogenetics*. 2003; 13(1):3–5.
2. Amberger J, Bocchini CA, Scott AF, Hamosh A. *McKusick's Online Mendelian Inheritance in Man (OMIM)*. *Nucl Acids Res* 2009; **37**: D793–D796.
3. Berlin DS, Person MG, Mittal A, et al. DNATwist: a web-based tool for teaching middle and high school students about pharmacogenomics. *Clin Pharmacol Ther*. 2010 In Press.
4. Crowley JJ, Sullivan PF, McLeod HL. Pharmacogenomic genome-wide association studies: lessons learned thus far. *Pharmacogenomics*. 2009; 10(2):161–163
5. Evans WE, McLeod HL. Pharmacogenomics: drug disposition, drug targets, and side effects. *N Engl J Med* 2003; 348: 538–549.
6. Giacomini KM, Brett CM, Altman RB, et al. The pharmacogenetics research network: from SNP discovery to clinical drug response. *Clin Pharmacol Ther*. 2007;81(3):328–345.
7. Hodge AE, Altman RB, Klein TE. The PharmGKB: integration, aggregation, and annotation of pharmacogenomic data and knowledge. *Clin. Pharmacol. Ther*. 2007; 81:21–24.
8. Imming P, Sinning C, Meyer A. Drugs, their targets and the nature and number of drug targets. *Nat. Rev. Drug Discov*. 2006; 5:821–834.
9. Klein TE, Chang JT, Cho MK, et al. Integrating genotype and phenotype information: an overview of the PharmGKB project. *Pharmacogenetics Research Network and Knowledge Base. Pharmacogenomics J*. 2001; 1(3):167–170.
10. Knox C, Law V, Jewison T, et al. DrugBank 3.0.: A comprehensive resource for ‘omics’ research on drugs. *Nucleic Acids Res*. 2011; 39(Database issue):D1035–D1041.
11. Nelson MR, Bacanu SA, Mosteller M, et al. Genome-wide approaches to identify pharmacogenetics contributions to adverse drug reactions. *Pharmacogenomics J*. 2009; 9(1):23–33.



12. Owen RP, Altman RB, Klein TE. PharmGKB and the International Warfarin Pharmacogenetics Consortium: the changing role for pharmacogenomic databases and single-drug pharmacogenetics. *Hum Mutat.* 2008; 29(4):456–460.
13. Relling MV, Klein TE. CPIC: Clinical Pharmacogenetics Implementation Consortium of the Pharmacogenomics Research Network. *Clin Pharmacol Ther.* 2011;89:464–467
14. Sangkuhl K, Berlin DS, Altman RB, Klein TE. PharmGKB: understanding the effects of individual genetic variants. *Drug Metab Rev.* 2008; 40(4):539–551.
15. Thorn CF, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for angiotensin-converting enzyme. *Pharmacogenet Genomics.* 2009; 20(2):143–146.
16. Thorn CF, Grosser T, Klein TE, Altman RB. PharmGKB summary: very important pharmacogene information for PTGS2. *Pharmacogenetic Genomics.* 2010; 21(9):607–613. Example of a Pharmacogenomics Knowledge Base Very Important Pharmacogene (VIP) summary.
17. Whirl-Carrillo M, McDonagh EM, Hebert JM, Gong L, Sangkuhl K, Thorn CF *et al.* *Pharmacogenomics knowledge for personalized medicine. Clin Pharmacol Ther* 2012; **92**: 414–417.
18. Wishart DS, Knox C, Guo AC, et al. DrugBank: a comprehensive resource for *in silico* drug discovery and exploration. *Nucleic Acids Res.* 2006; 34(Database issue):D668–D672.
19. Wishart DS, Feunang YD, Guo AC, Lo EJ, Marco A, Grant JR, Sajed T, Johnson D, Li C, Sayeeda Z, Assempur N, Iynkkaran I, Liu Y, Maciejewski A, Gale N, Wilson A, Chin L, Cummings R, Le D, Pon A, Knox C, Wilson M. DrugBank 5.0: a major update to the drugbank database for 2018. *Nucleic Acids Res.* 2017 nov 8. Doi: 10. 1093/nar/gkx 1037.
20. Yee SW, et al. Genomewide Association Studies in Pharmacogenomics: Meeting Report of the NIH Pharmacogenomics Research Network-RIKEN (PGRN-RIKEN) Collaboration. *Clin Pharmacol Ther.* 2016; 100:423–426.

## **Chapter-7 WEBSITES**

- PharmGKB. [www.pharmgkb.org](http://www.pharmgkb.org).
- National Library of Medicine's Medical Subject Headings (MeSH browser) [www.nlm.nih.gov/mesh/2011/mesh\\_browser/MBrowser.html](http://www.nlm.nih.gov/mesh/2011/mesh_browser/MBrowser.html).
- DrugBank. [www.drugbank.ca](http://www.drugbank.ca).
- PubChem<sup>®</sup> <http://pubchem.ncbi.nlm.nih.gov/>
- OMIM<sup>®</sup>, Online Mendelian Inheritance in Man<sup>®</sup> [www.ncbi.nlm.nih.gov/omim](http://www.ncbi.nlm.nih.gov/omim).
- US FDA. Table of pharmacogenomics biomarkers in drug labels. [www.fda.gov/Drugs/ScienceResearch/ResearchAreas/Pharmacogenetics/ucm083378.htm](http://www.fda.gov/Drugs/ScienceResearch/ResearchAreas/Pharmacogenetics/ucm083378.htm).
- Orphanet. <https://www.orpha.net/consor/cgi-bin/index.php>