
CONNECTING THE DOTS: A NETWORK ANALYSIS OF CKD AND AKI GENES

Student: Katelyn Hur, Red River High School, Grand Forks, ND

Mentor: Junguk Hur, PhD, University of North Dakota, Grand Forks, ND

BACKGROUND

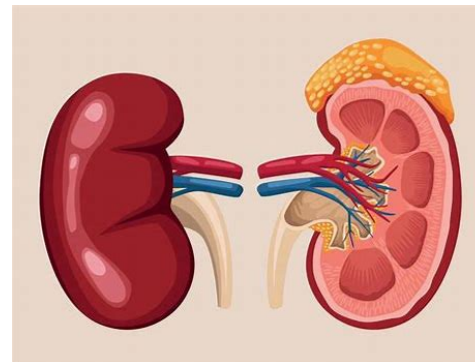
- Was intrigued to learn about the severity of kidney diseases and the fact that we are still trying to find effective cures.
- Kidney diseases are primarily classified into two main types: chronic kidney disease (CKD) and acute kidney injury (AKI).

What is Chronic Kidney Disease (CKD)?

Chronic Kidney Disease is a gradual loss of function in your kidneys often because of diabetes and high blood pressure (long-term conditions), it is incurable.

What is Acute Kidney Injury (AKI)?

Acute Kidney Injury is a sudden loss of function in your kidneys often because of dehydration, blood loss, or urinary tract infection, it is curable once the underlying cause is cured.



Kidney

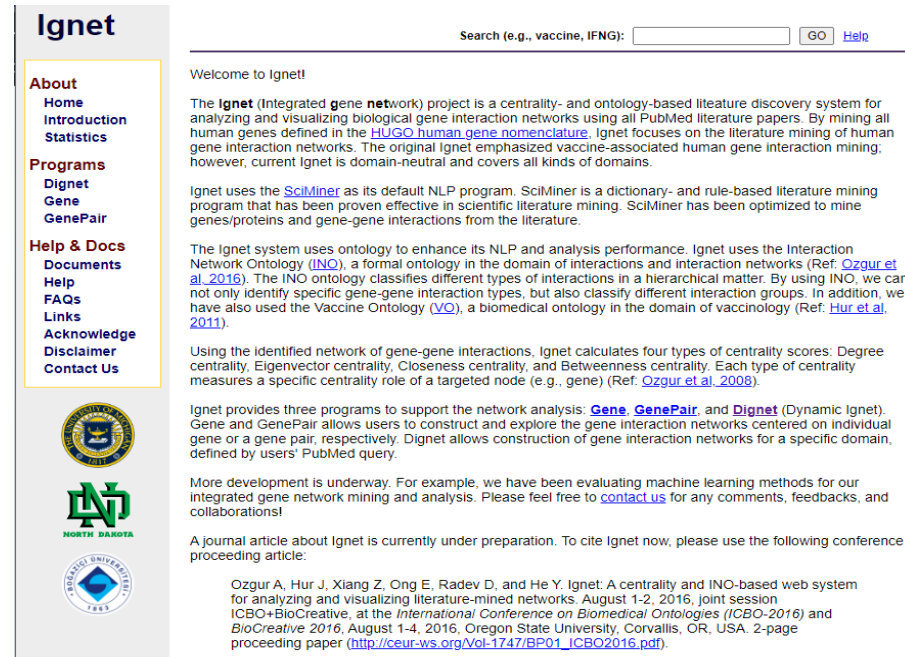
<https://www.vecteezy.com/free-vector/kidney>

WHY ARE CKD & AKI IMPORTANT?

- 800 million people (10% of the world's population) are affected by Chronic Kidney Disease (CKD)
- The annual cost of treating CKD is estimated to be over \$120 billion
- CKD is a leading cause of death globally, with millions of people dying each year due to complications related to CKD
- Acute Kidney Injury (AKI) affects about 13.3 million people globally each year
- The cost of treating AKI in hospitalized patients in the United States is estimated to be around \$10 billion annually
- AKI causes long-term health issues and increased mortality if not treated promptly

How can we better understand the differential gene interactions in CKD and AKI to discover improved treatment options for kidney failure?

METHODOLOGY (DATASETS)

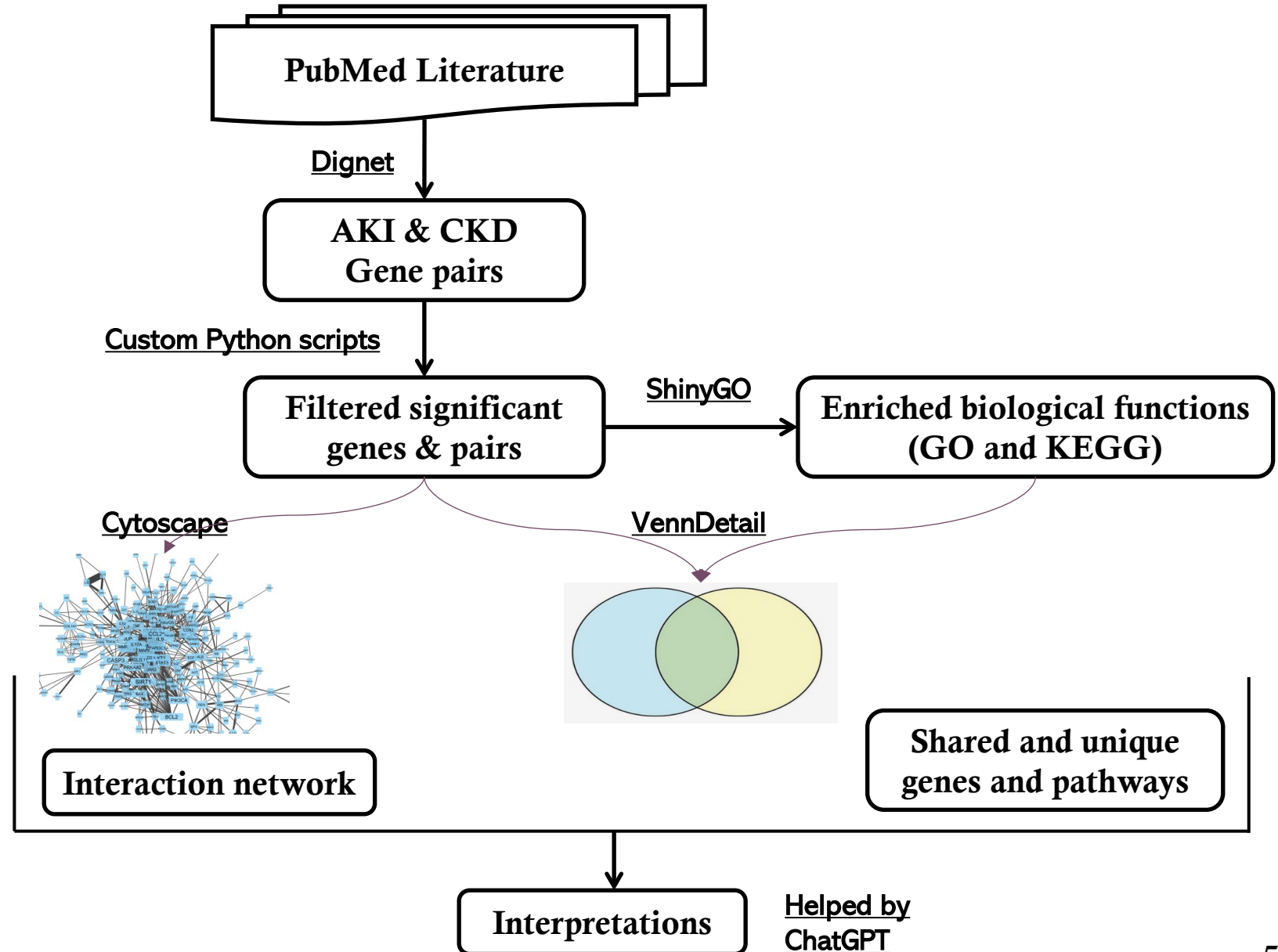


- The abstracts of biomedical literature in **PubMed**
 - “Acute Kidney Injury”: 77,050 papers
 - “Chronic Kidney Disease”: 86,023 papers
- Gene-pairs downloaded from these articles identified by dynamic Ignnet (Dignet)

The **Ignnet (integrated gene network; <https://ignnet.org>)** project is an ontology-based system of discovering literature (from PubMed) analyzing and visualizing biological gene interaction networks.

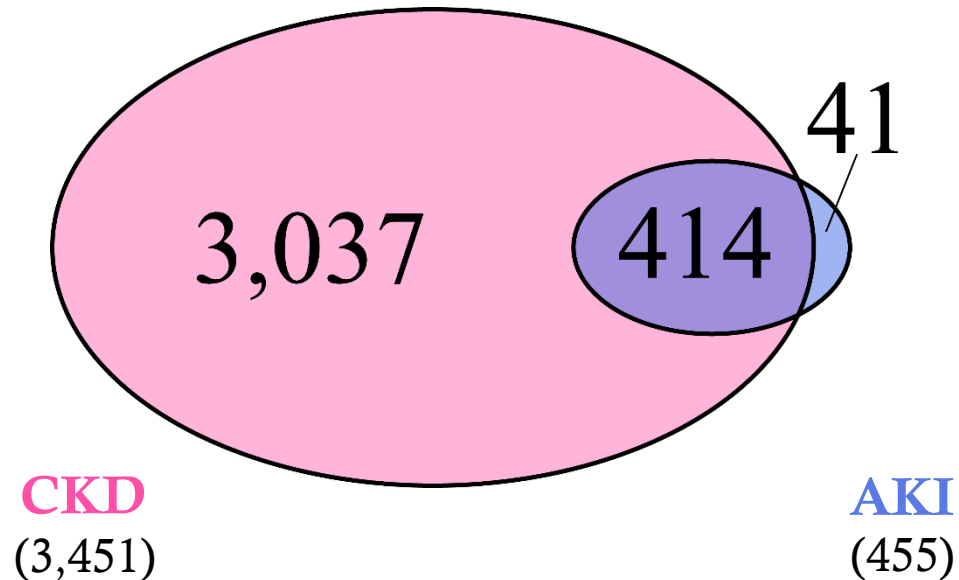
WORKFLOW

- **Dignet: Dynamic Iagnet** with PubMed search
- **Custom Python scripts:** my own scripts for handling gene-pair files, calculating overlap, generating a network file
- **ShinyGO:** web-based gene set analysis system
- **VennDetail:** web-based overlap analysis (Venn diagram)
- **Cytoscape:** Network visualization tool
- **ChatGPT:** Artificial Intelligence (AI) ChatBot

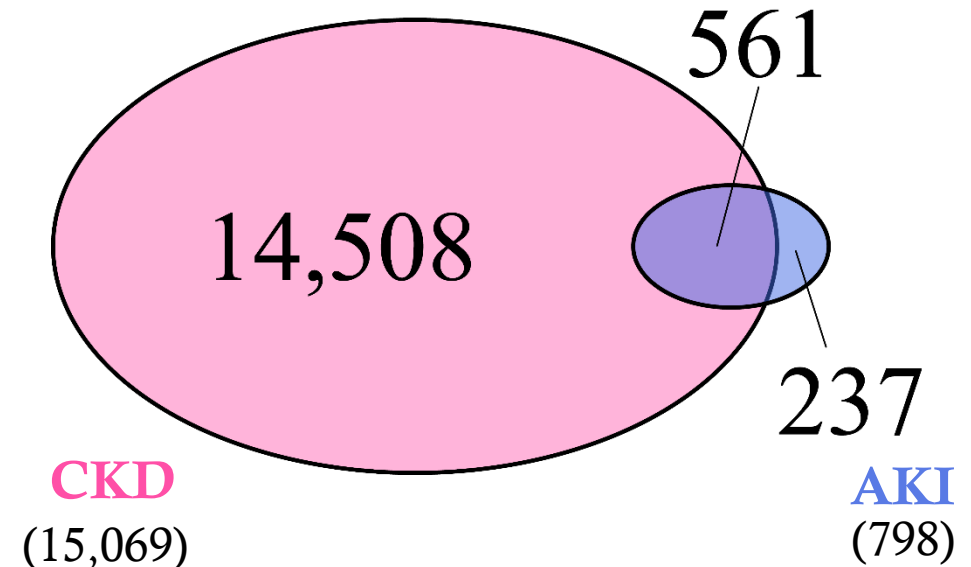


OVERLAP - VENN DIAGRAMS

Genes

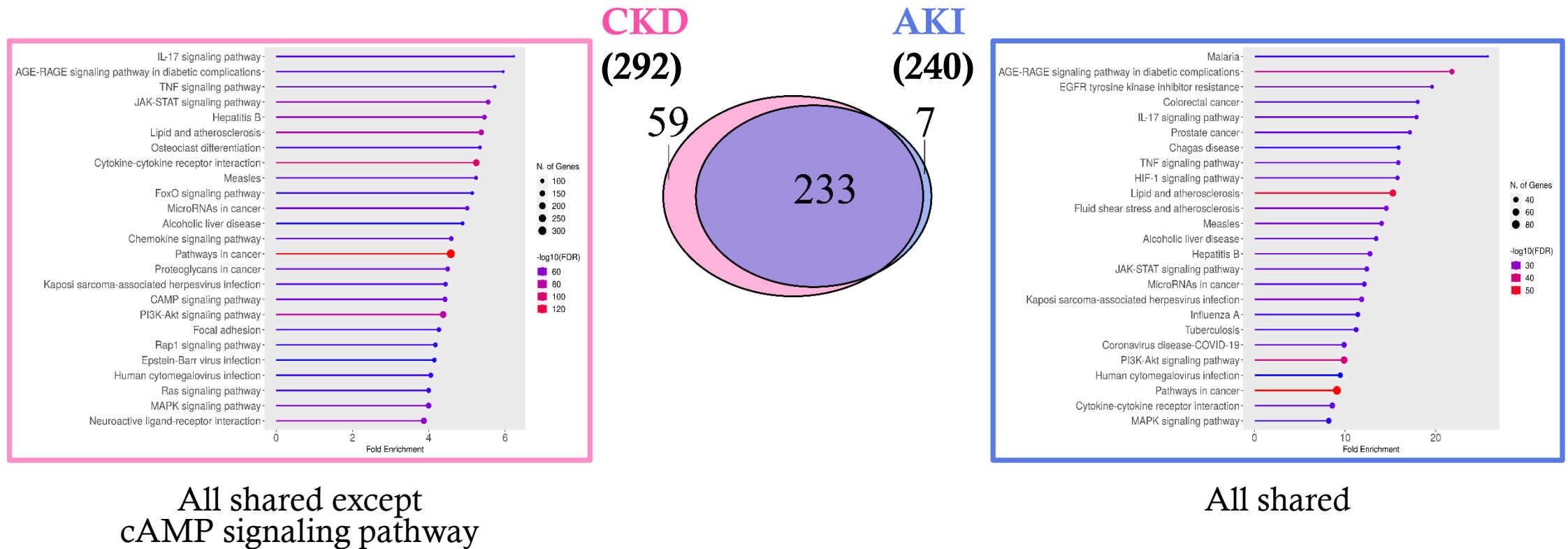


Gene Pairs



A lot more genes and gene-pairs in CKD than AKI

KYOTO ENCYCLOPEDIA OF GENES AND GENOMES (KEGG) PATHWAYS – TOP 25

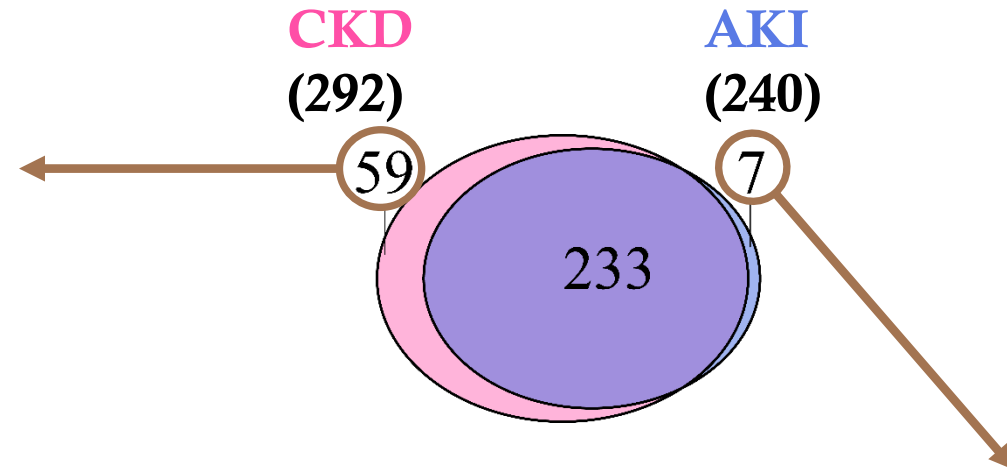


UNIQUE KEGG PATHWAYS

hsa04911 Insulin secretion

hsa00010 Glycolysis/Gluconeogenesis
 hsa00980 Metabolism of xenobiotics by cytochrome P450
 hsa00620 Pyruvate metabolism
 hsa01230 Biosynthesis of amino acids
 hsa04972 Pancreatic secretion
 hsa04971 Gastric acid secretion
 hsa05204 Chemical carcinogenesis-DNA adducts
 hsa04970 Salivary secretion
 hsa00140 Steroid hormone biosynthesis
 hsa00830 Retinol metabolism
 hsa00230 Purine metabolism
 hsa05032 Morphine addiction
 hsa04142 Lysosome
 hsa00983 Drug metabolism-other enzymes
 hsa05033 Nicotine addiction
 hsa04330 Notch signaling pathway
 hsa04340 Hedgehog signaling pathway
 hsa04727 GABAergic synapse
 hsa04721 Synaptic vesicle cycle
 hsa04120 Ubiquitin mediated proteolysis
 hsa01232 Nucleotide metabolism
 hsa00071 Fatty acid degradation
 hsa04978 Mineral absorption
 hsa00591 Linoleic acid metabolism
 hsa00270 Cysteine and methionine metabolism
 hsa00910 Nitrogen metabolism
 hsa01040 Biosynthesis of unsaturated fatty acids
 hsa04961 Endocrine and other factor-regulated calcium reabsorption
 hsa04975 Fat digestion and absorption
 hsa03250 Viral life cycle-HIV-1

hsa01212 Fatty acid metabolism
 hsa00500 Starch and sucrose metabolism
 hsa00600 Sphingolipid metabolism
 hsa03430 Mismatch repair
 hsa02010 ABC transporters
 hsa04392 Hippo signaling pathway-multiple species
 hsa01210 2-Oxocarboxylic acid metabolism
 hsa03420 Nucleotide excision repair
 hsa00603 Glycosphingolipid biosynthesis-globo and isoglobo series
 hsa00100 Steroid biosynthesis
 hsa00430 Taurine and hypotaurine metabolism
 hsa00640 Propanoate metabolism
 hsa00564 Glycerophospholipid metabolism
 hsa04962 Vasopressin-regulated water reabsorption
 hsa04130 SNARE interactions in vesicular transport
 hsa04742 Taste transduction
 hsa03460 Fanconi anemia pathway
 hsa00280 Valine leucine and isoleucine degradation
 hsa00053 Ascorbate and aldarate metabolism
 hsa00565 Ether lipid metabolism
 hsa04136 Autophagy-other
 hsa00240 Pyrimidine metabolism
 hsa00650 Butanoate metabolism
 hsa00232 Caffeine metabolism
 hsa00130 Ubiquinone and other terpenoid-quinone biosynthesis
 hsa00515 Mannose type O-glycan biosynthesis
 hsa00120 Primary bile acid biosynthesis
 hsa04070 Phosphatidylinositol signaling system

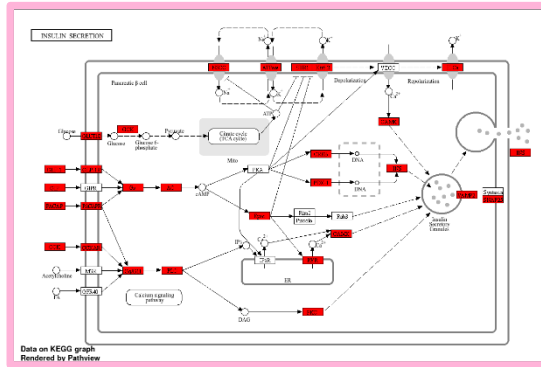


ChatGPT found
 the pathways
 highlighted in
 red to be the
 most relevant.

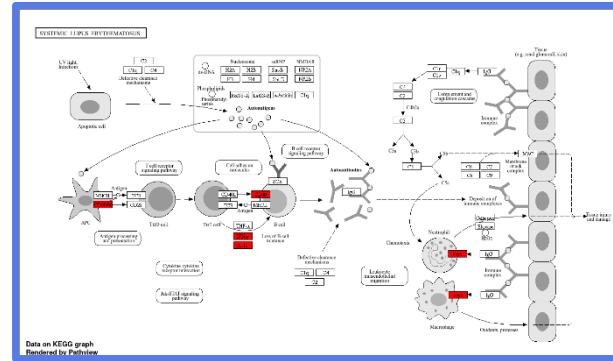
hsa00470 D-Amino acid metabolism
 hsa00730 Thiamine metabolism
 hsa00061 Fatty acid biosynthesis
 hsa00520 Amino sugar and nucleotide sugar metabolism
 hsa00524 Neomycin kanamycin and gentamicin biosynthesis
 hsa05322 Systemic lupus erythematosus
 hsa00750 Vitamin B6 metabolism

KEY KEGG PATHWAYS

hsa04911 Insulin secretion



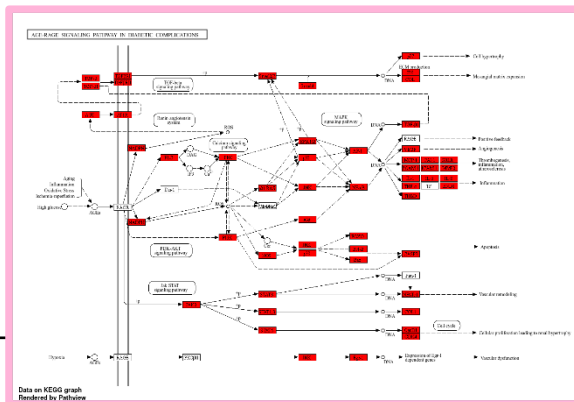
hsa05322 Systemic lupus erythematosus



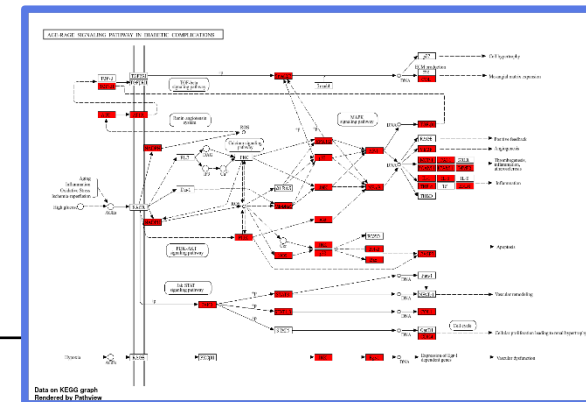
Unique

: genes from CKD (for pink outline) and AKI (for blue outline)

Common Key Pathway: hsa04933 AGE-RAGE signaling pathway in diabetic complications

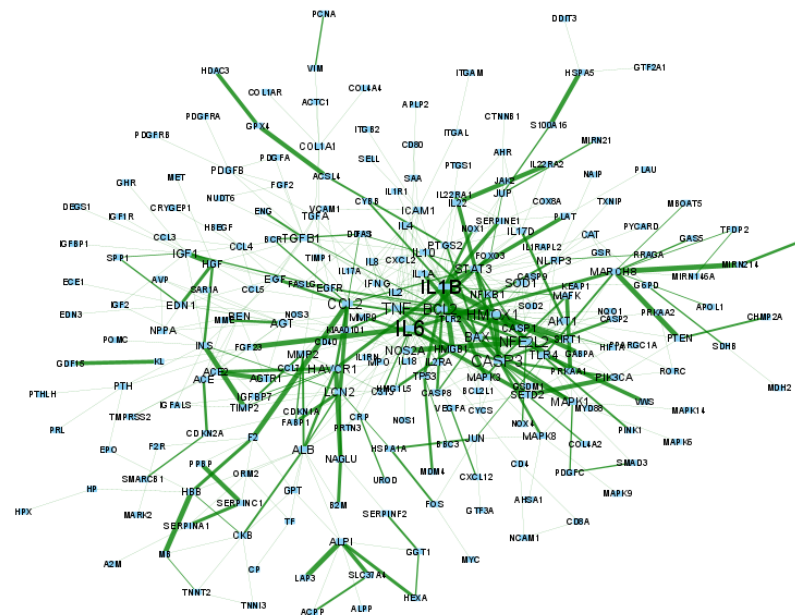


Very high overlap
between CDK and AKI

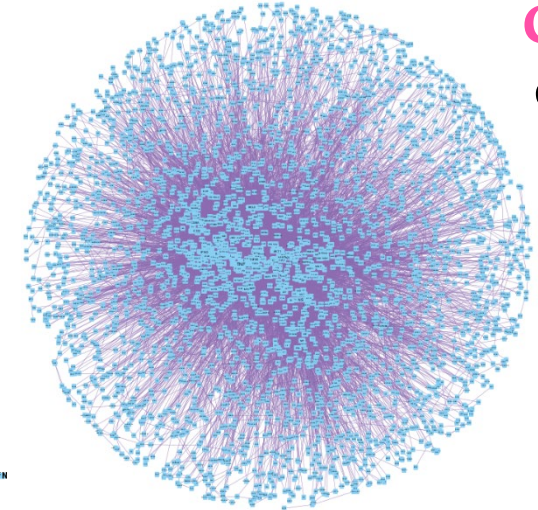


GENE PAIR NETWORK

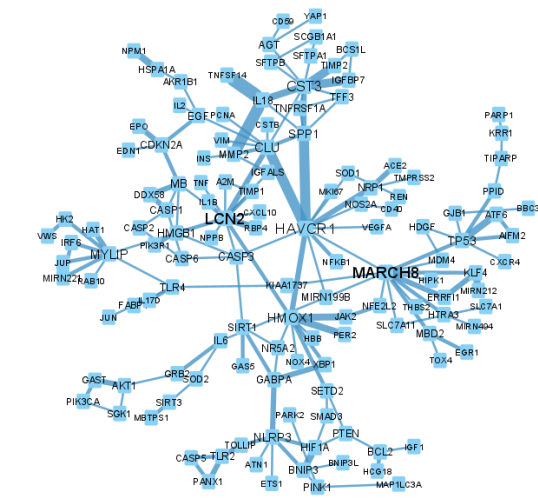
- Using Cytoscape to visualize the gene pair networks
- Three networks
 - Common: centered around IL1B, IL6, ...
 - CKD-unique: too big. Top edges around TNF, AKT1, INS, IFNG, ...
 - AKI-unique: LCN2, MARCH8, ...



Common



CKD
only



AKI
only

CONCLUSIONS & DISCUSSION

- **Major findings**

- Substantially more genes and pairs in CKD because there are much more publications on CKD than AKI
- AGE-RAGE is the top common enriched biological function of CKD and AKI
- Insulin secretion is a unique enriched biological function of CKD
- Systemic lupus erythematosus is a unique biological function of AKI

- **What are the best treatment targets for kidney failures (CKD and AKI)?**

- The three enriched biological functions mentioned above could be important potential targets.

ACKNOWLEDGEMENTS

Mentor

Jungkuk Hur, Ph.D., University of North Dakota

Red River High School

Dr. Darwin Walters
(Associate Principal)

No funding was provided for this research.