

## Diseases Genes

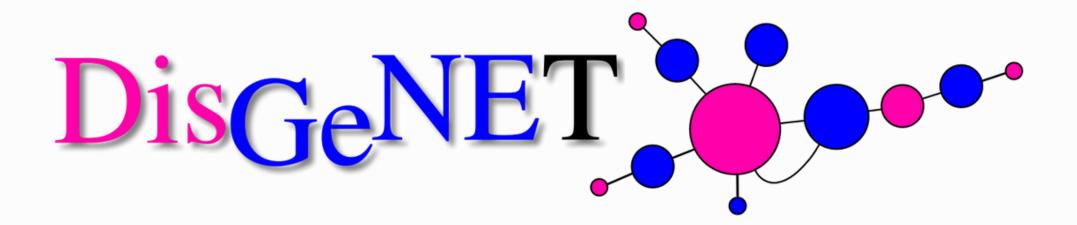
The association between diseases and genes



# How are diseases related?

# What diseases are you likely to get?

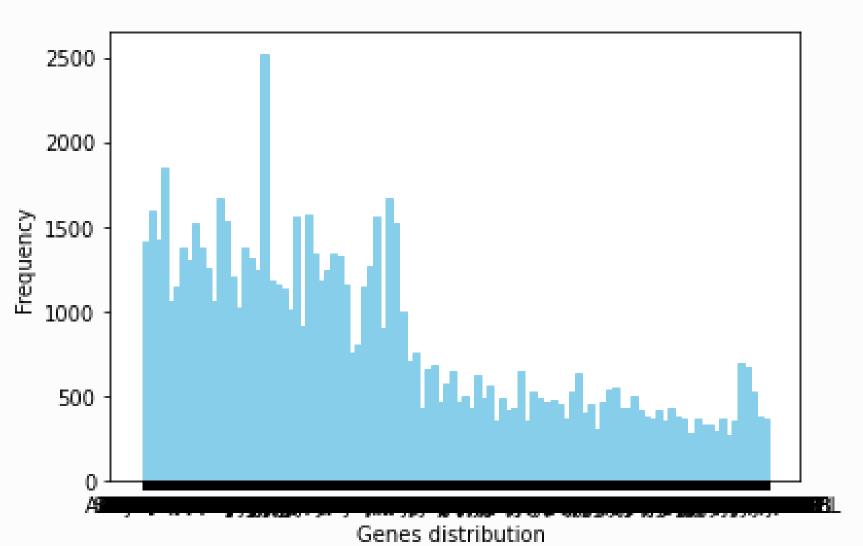
Based on Genes in common



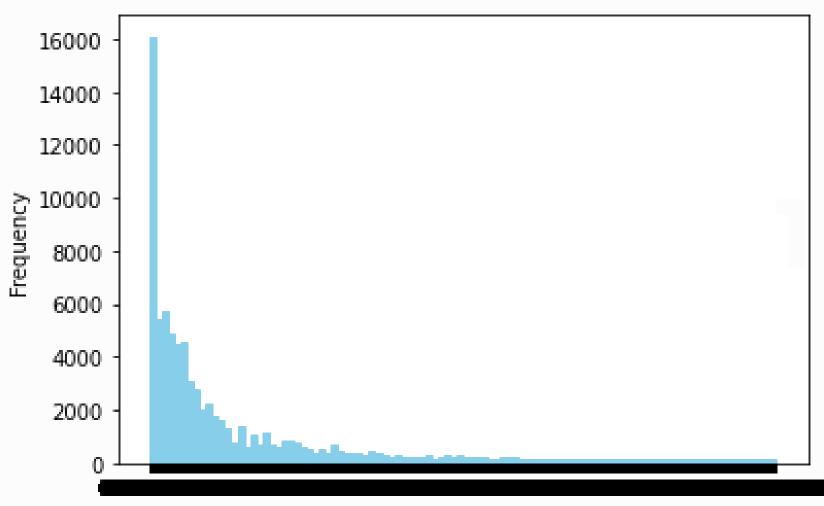
9411

10370

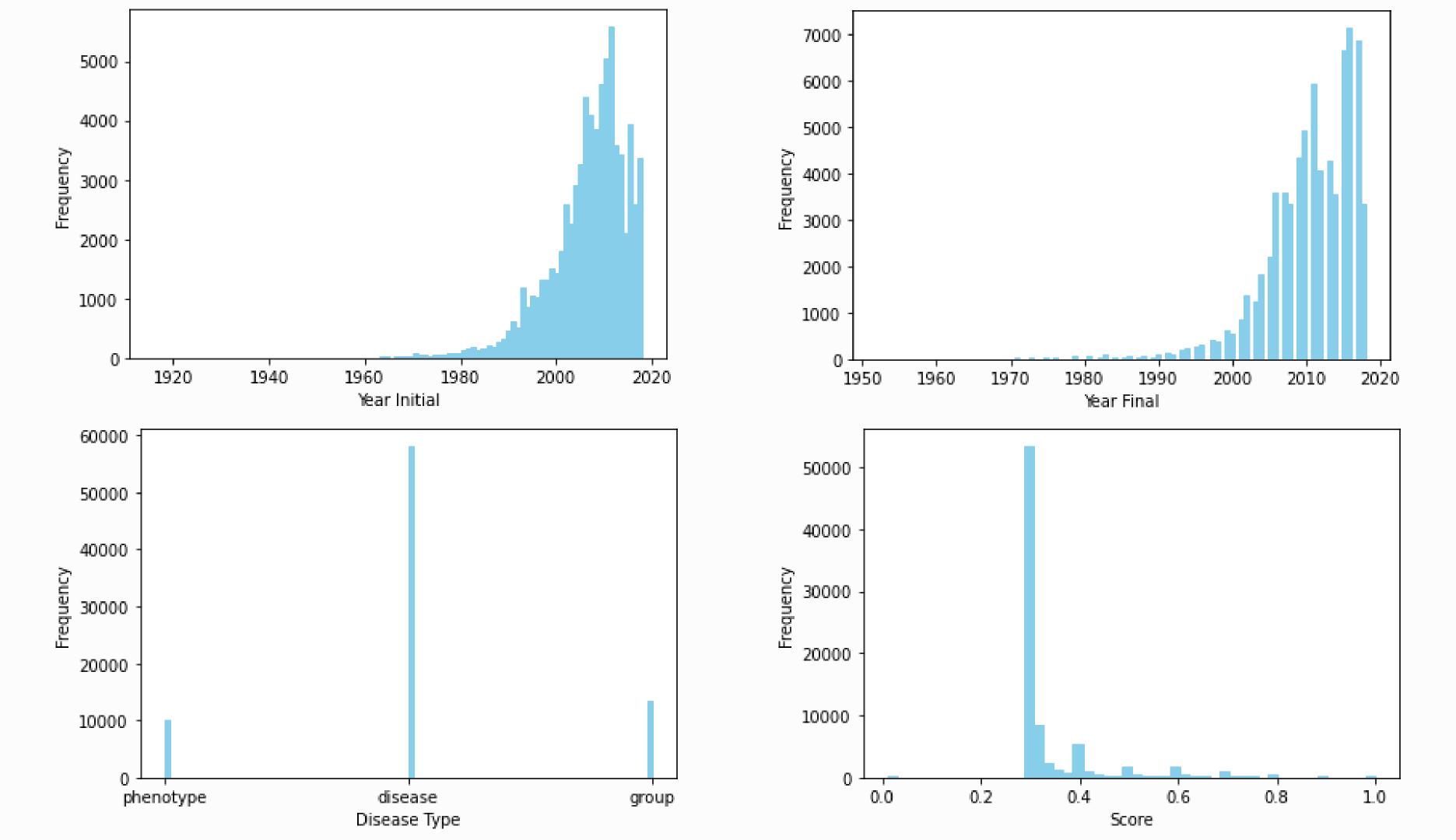




#### Diseases



Disease distribution



#### Process

How data is selected?

Score > 0.5
Recent years
Early years



Hypergeomitrec Filter

Filter the low probabilitis



Work with Gephi

Filter and draw

**A Lucrative Industry** 

# Diseases Network

Edges: 311

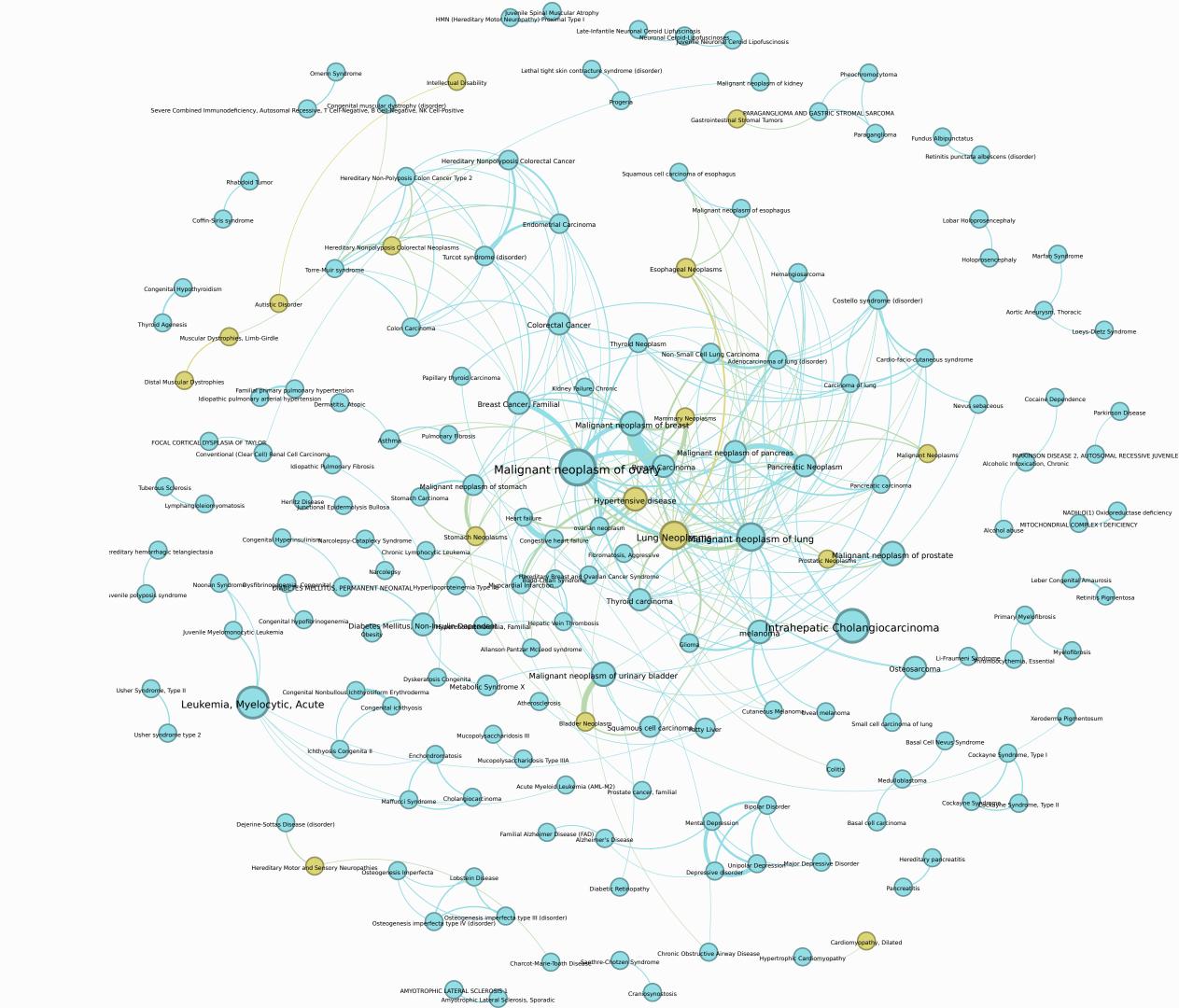
#### Filter:

Years>2014

Score > 0.5

Wieght

Degree



Edges: 311

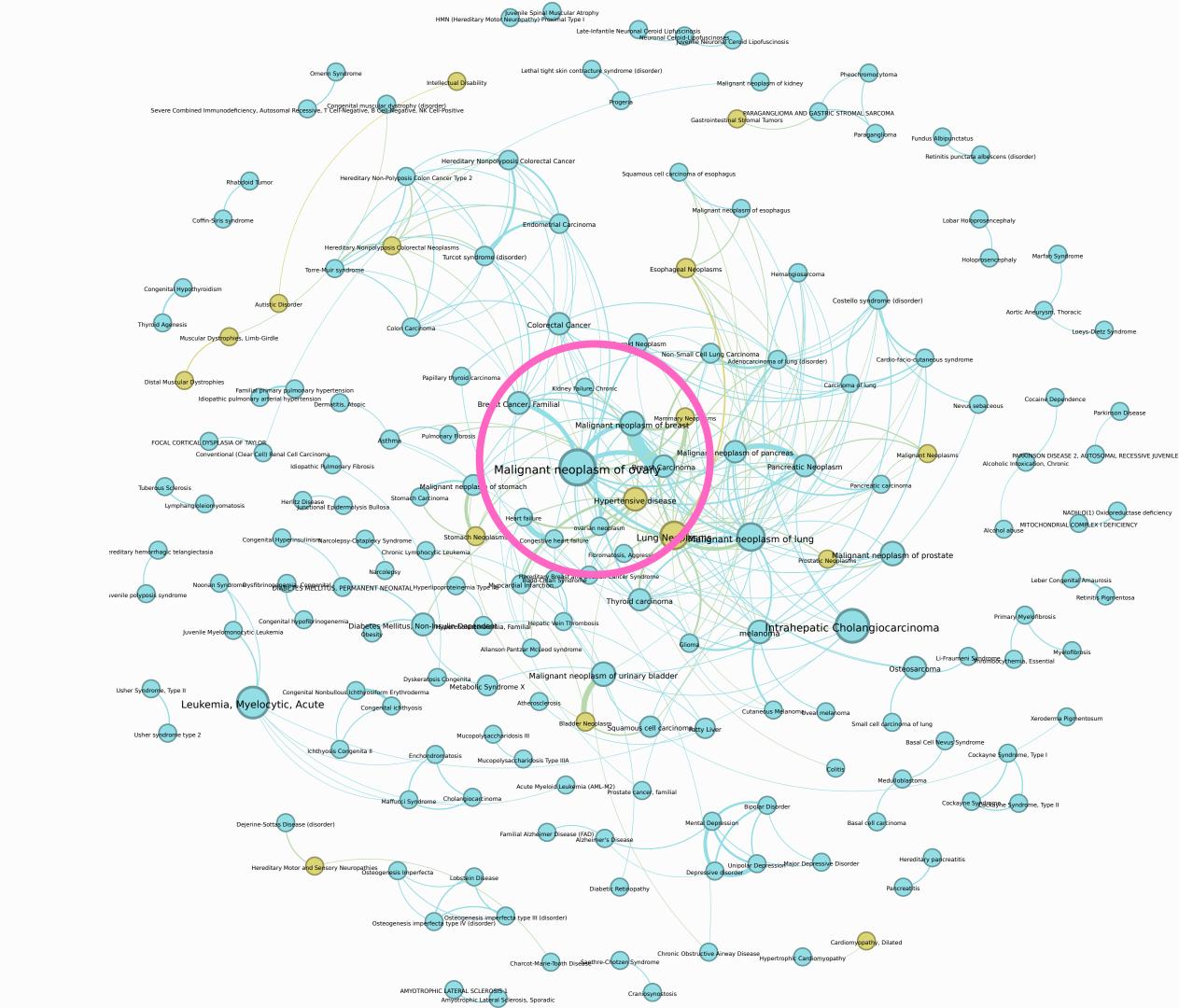
#### Filter:

Years>2014

Score > 0.5

Wieght

Degree



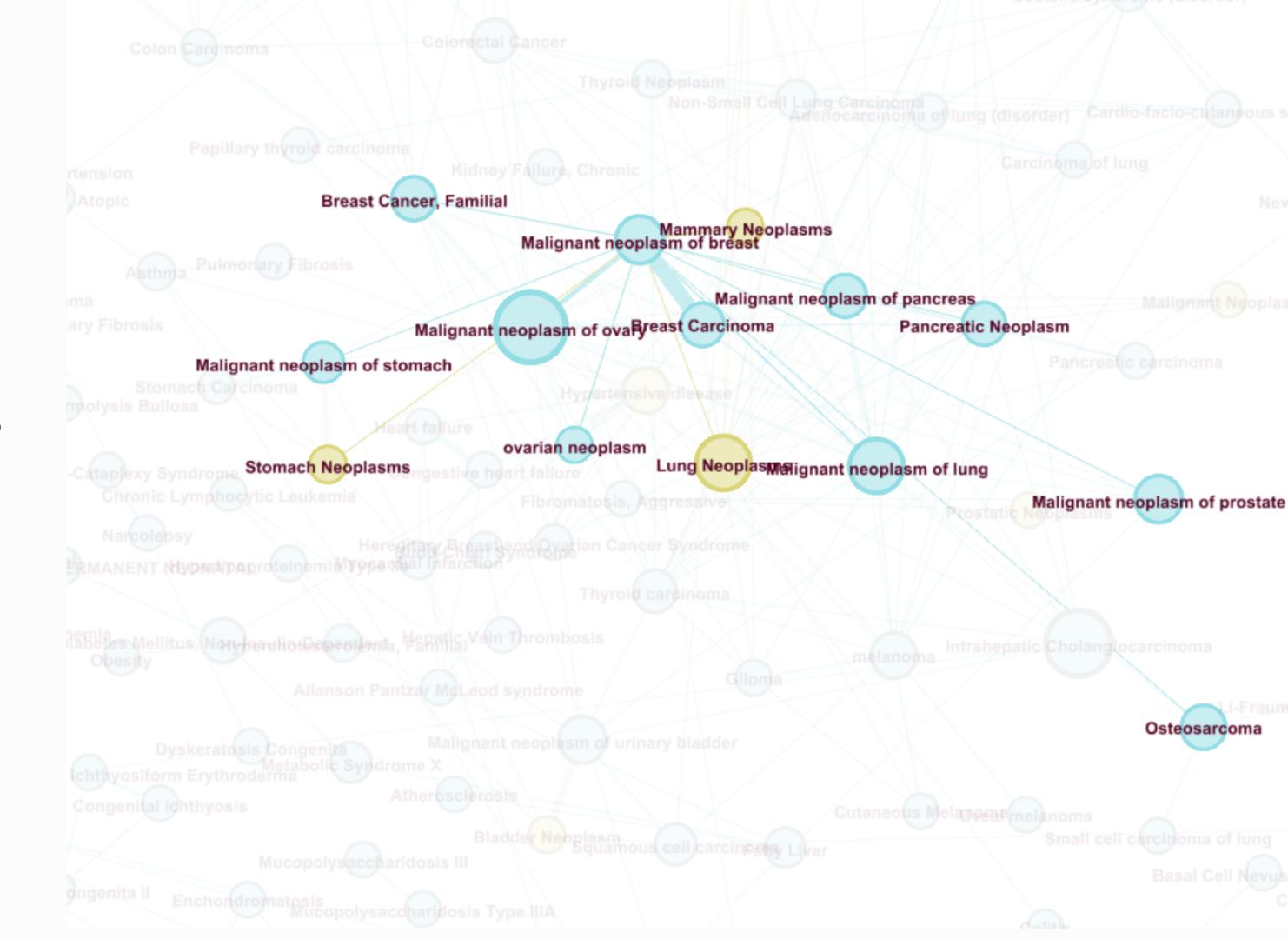
# Malignant neoplasm of breast:

1983-2018

1025 Gene

2008-2018 **26 Gene** 

1998-2008 **16 Gene** 



Edges: 311

Node Size: betweenness centrality

Node Color: Degree

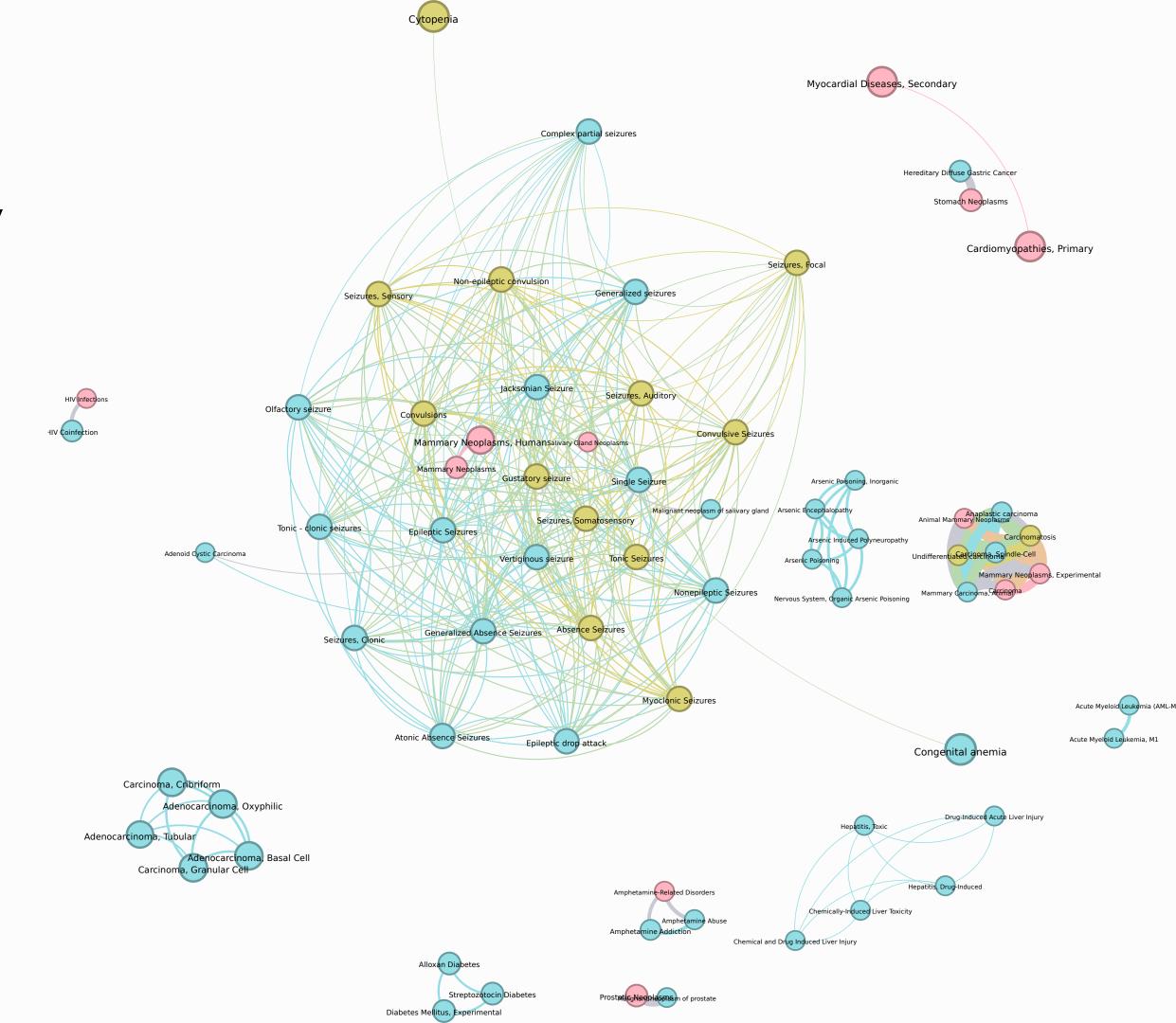
#### Filter:

Years: 1998-2008

Score > 0.5

Edge Wieght

Node Degree



Edges: 311

Node Size: betweenness centrality

Node Color: Degree

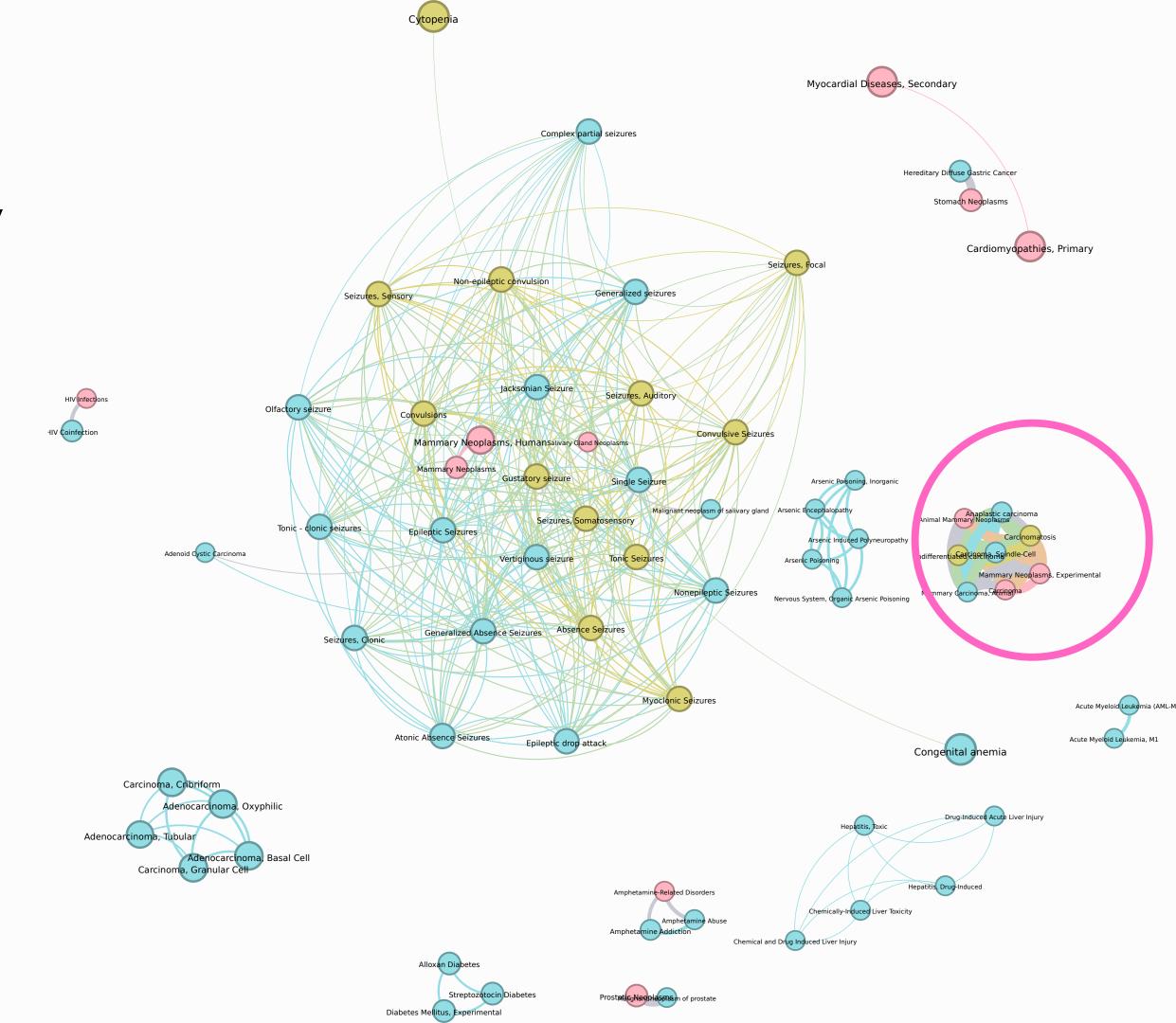
#### Filter:

Years: 1998-2008

Score > 0.5

Edge Wieght

Node Degree

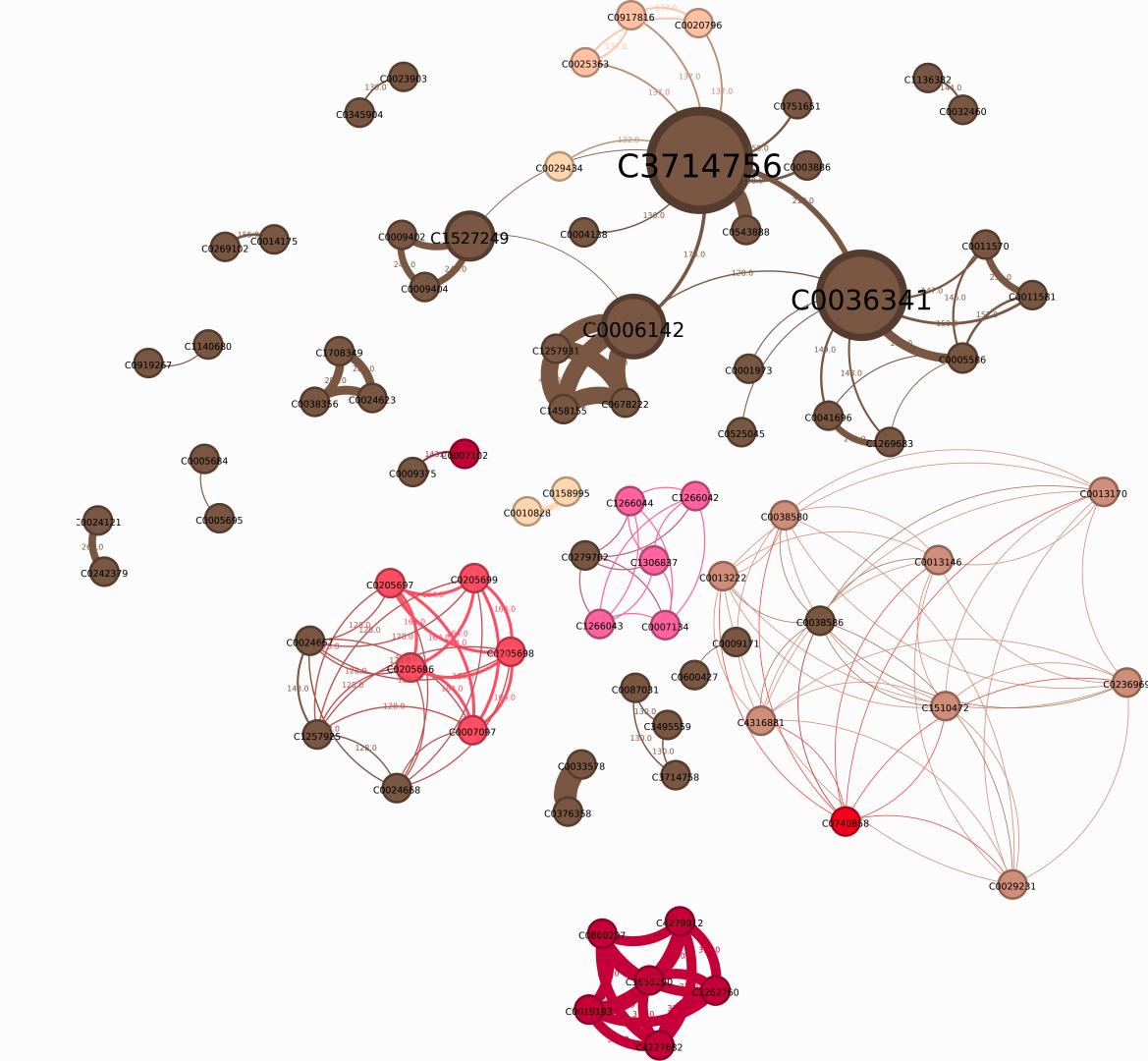


#### Filter:

Edge Wieght Node Degree

Node Size: betweenness centrality

Node Color: Degree



### Github Repository

https://github.com/halaalbahloul/DisgenetDataAnalysis

Thank you for your attension!

## How is the score for a gene-disease pair, or variant-disease pair computed?

The DisGeNET score for GDAs takes into account the number and type of sources (level of curation, organisms), and the number of publications supporting the association, while the score for the VDAs takes into account sources, and number of papers. The scores range from 0 to 1.