

# BMI 206 Project Data Summary for Paper "Uncovering disease-disease relationships through the incomplete interactome"

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To obtain the relevant data for this paper, we downloaded the data archive from the online supplement on the Science website

(<https://www.sciencemag.org/content/347/6224/1257601/suppl/DC1>), which

contains datasets in the format of tables with tab-separated columns (.tsv files):

- 1) Human interactome dataset listing pairwise interactions of genes and the types of interactions,

```
> head(DataS1_interactome,5)
  gene_ID_1 gene_ID_2 data_source.s.
1      1394      2778      literature
2 100290337      4214      literature
3     122704     54460      complexes
4      4790     79155 binary;literature;signaling
5      2597        70      signaling
```

containing 141,295

pairs of genes.

```
> nrow(DataS1_interactome)
[1] 141295
```

Here is a summary quantifying the number of instance of each types of interactions and demonstrating how one or more interactions occurs for each gene-gene pair:

```

> table(col_3_split)
col_3_split
      binary complexes kinase literature metabolic regulatory signaling
795362  28653   31276   6066   88349   5325   1335   32706
> head(col_3_split,5)
  [,1] [,2] [,3] [,4] [,5] [,6] [,7]
[1,] "literature" "" "" "" "" "" ""
[2,] "literature" "" "" "" "" "" ""
[3,] "complexes" "" "" "" "" "" ""
[4,] "binary" "literature" "signaling" "" "" "" ""
[5,] "signaling" "" "" "" "" "" ""

```

- 2) Disease genes dataset.
- 3) Network properties dataset for the diseases considered in the study. Here are summary statistics describing number of genes ascribed to each disease and other parts of the dataset:

```

> dataSummary3
      disease number_of_all_genes LCC_size LCC_z_score
abnormalities, multiple : 1 Min. : 20.00 Min. : 1.00 Min. : -0.500
actinomycetales infections: 1 1st Qu.: 29.00 1st Qu.: 3.00 1st Qu.: 2.000
adenocarcinoma : 1 Median : 48.00 Median : 6.00 Median : 4.900
adnexal diseases : 1 Mean : 99.58 Mean : 25.89 Mean : 6.209
adrenal gland diseases : 1 3rd Qu.: 99.00 3rd Qu.: 15.50 3rd Qu.: 8.350
albuminuria : 1 Max. : 1168.00 Max. : 730.00 Max. : 31.100
(Other) :293
      d_s d_s_Glass_delta d_s_p_value
Min. :1.050 Min. : -2.2939 Min. : 0.0000000
1st Qu.:1.566 1st Qu.: -1.0891 1st Qu.: 0.0000000
Median :1.788 Median : -0.7829 Median : 0.0000003
Mean :1.823 Mean : -0.8020 Mean : 0.0251731
3rd Qu.:2.000 3rd Qu.: -0.5424 3rd Qu.: 0.0014005
Max. :3.143 Max. : 0.8095 Max. : 0.4959045

```

- 4) Network relationships for all the pairs of diseases considered in the study, summarized here:

```

> summary(DataS4_disease_pairs)
      disease_A disease_B s_AB..observed. d_AB..observed. z..full.rand.
abnormalities, multiple : 298 vision disorders : 298 Min. : -3.1429 Min. : 0.000 Min. : -50.761
actinomycetales infections: 297 virus diseases : 297 1st Qu.: 0.2037 1st Qu.: 2.036 1st Qu.: 1.700
adenocarcinoma : 296 vasculitis : 296 Median : 0.3646 Median : 2.209 Median : 3.468
adnexal diseases : 295 vascular diseases: 295 Mean : 0.3672 Mean : 2.190 Mean : 3.520
adrenal gland diseases : 294 varicose veins : 294 3rd Qu.: 0.5437 3rd Qu.: 2.386 3rd Qu.: 5.489
albuminuria : 293 uveitis, anterior: 293 Max. : 1.8000 Max. : 3.303 Max. : 18.072
(Other) :42778 (Other) :42778
p..full.rand. q..full.rand. z..MeSH.rand. p..MeSH.rand. q..MeSH.rand.
Min. : 0.00000 Min. : 0.0000000 Min. : -6.247 Min. : 0.00000 Min. : 0.0000000
1st Qu.: 0.00000 1st Qu.: 0.0000009 1st Qu.: 2.146 1st Qu.: 0.00000 1st Qu.: 0.0000004
Median : 0.00000 Median : 0.0004255 Median : 3.829 Median : 0.00000 Median : 0.0002416
Mean : 0.06091 Mean : 0.0859514 Mean : 4.252 Mean : 0.07358 Mean : 0.0745713
3rd Qu.: 0.02682 3rd Qu.: 0.0516171 3rd Qu.: 5.936 3rd Qu.: 0.02000 3rd Qu.: 0.0311266
Max. : 1.00000 Max. : 0.7482555 Max. : 18.903 Max. : 1.00000 Max. : 0.7458767

```