

Analysis results by CePa

Method to adjust raw p-values: none

Cutoff: 0.01

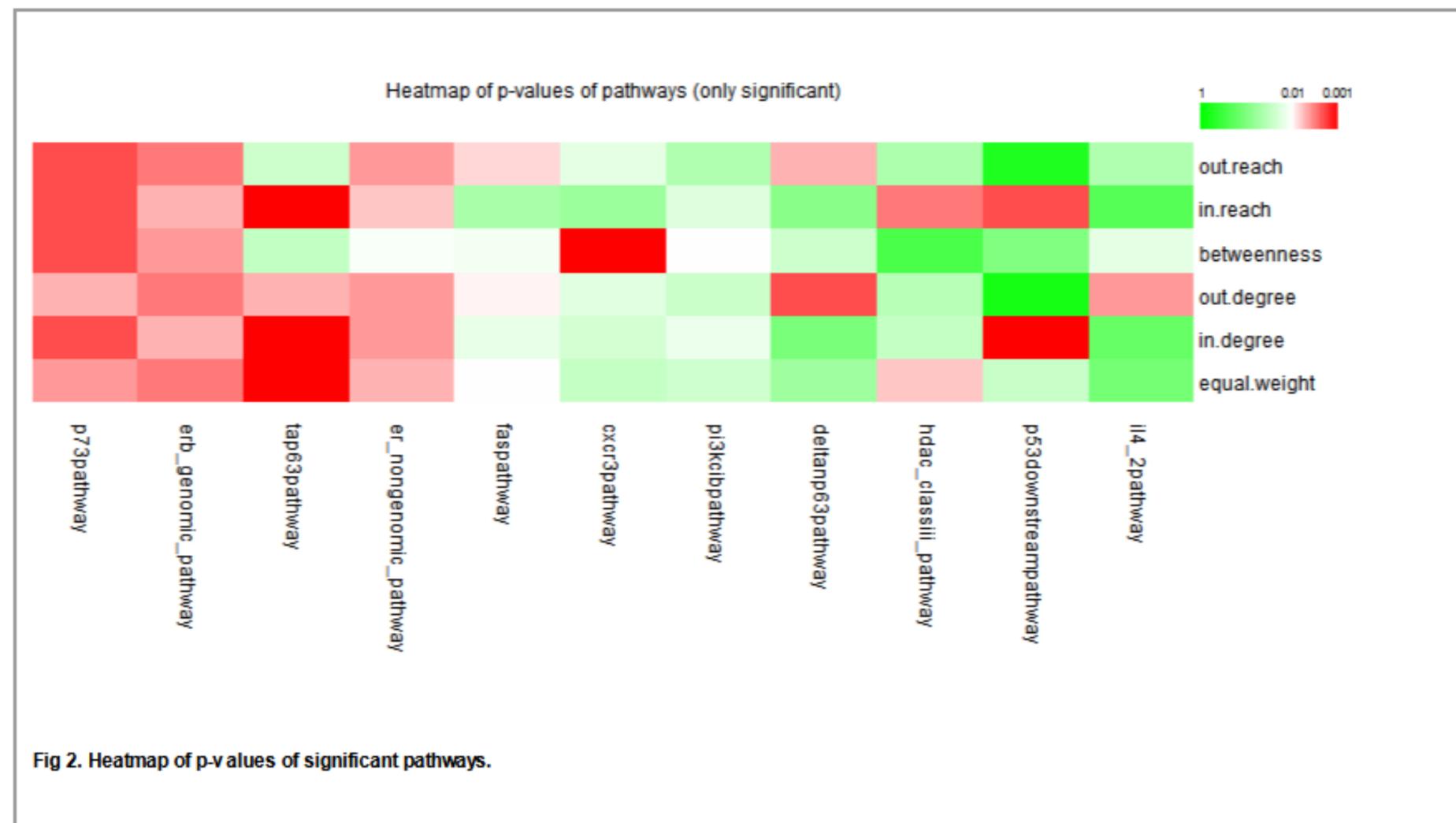
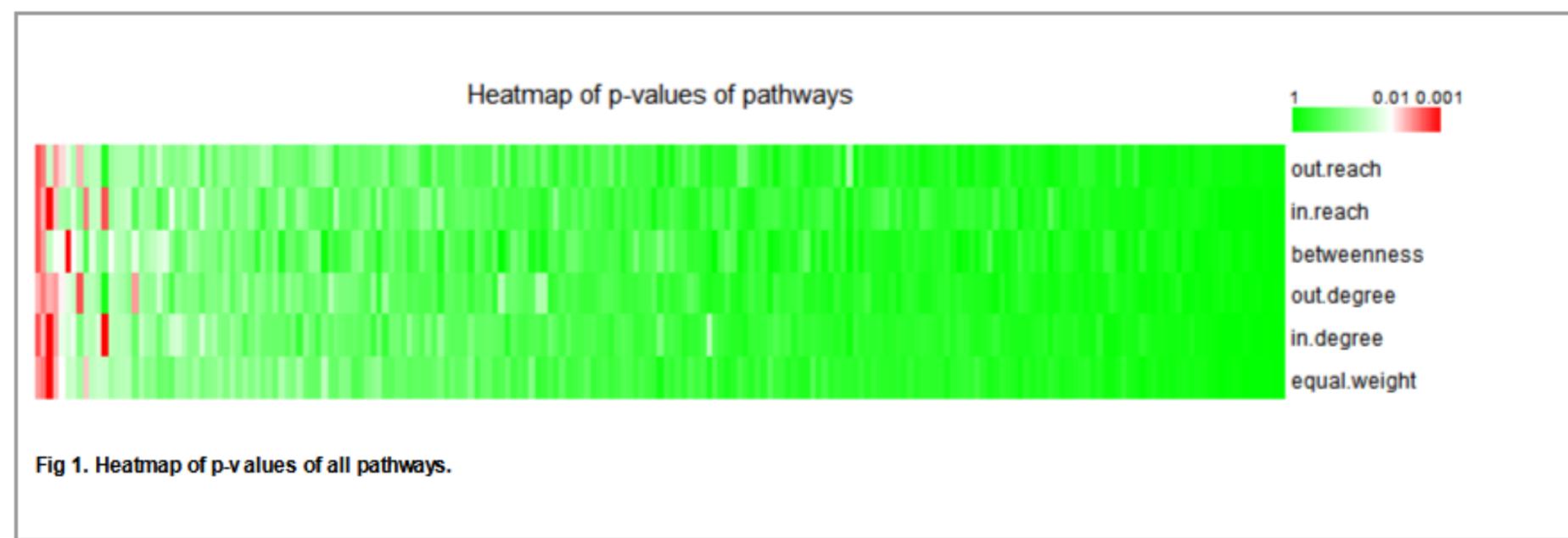


Table 1. Complete list of p-values of all pathways under all centrality measurements.

Table of significance in txt format

Significant pathways

Significant p-v alues

Pathway	equal.weight	in.degree	out.degree	betweenness	in.reach	out.reach
p73pathway	0.004	0.002	0.005	0.002	0.002	0.002
erb_genomic_pathway	0.003	0.005	0.003	0.004	0.005	0.003
tap63pathway	0.001	0.001	0.005	0.03	0.001	0.025
er_nongenomic_pathway	0.005	0.004	0.004	0.011	0.006	0.004
faspathway	0.01	0.015	0.009	0.012	0.047	0.007
cxcr3pathway	0.029	0.022	0.017	0.001	0.0599	0.016
pi3kcbpathway	0.024	0.014	0.026	0.01	0.018	0.041
deltanp63pathway	0.0559	0.1119	0.002	0.025	0.0839	0.005
hdac_classiii_pathway	0.006	0.029	0.036	0.2617	0.003	0.044
p53downstreampathway	0.027	0.001	0.6823	0.0959	0.002	0.5654
il4_2pathway	0.1219	0.1578	0.004	0.016	0.2098	0.041
a6b1_a6b4_integrin_pathway	0.032	0.039	0.036	0.017	0.031	0.034
il27pathway	0.025	0.018	0.0569	0.0649	0.026	0.032
circadianpathway	0.0549	0.033	0.03	0.011	0.0649	0.0599