



Node size

Transparency

Edges removed for clustering: 0

Grouping

KEGG pathway

Select a pathway to turn the common edges with your dataset in red

 Only available if you open your dataset and choose ENTREZ id's.

Shortest path

Select path end

Select path other end

Displayed node deepness

0 1 2 3 4 5 All

Listed nodes (2000 available)

Ordered by node

1007_s_at
1053_at
117_at
121_at
1255_g_at
1294_at
1316_at
1320_at
1405_l_at
1431_at
1438_at
1487_at

EDGE WEIGHT
THRESHOLD
0.5



Highlight deepness (1+21)

0 1 2 3 4 5

Save selection

Displayed nodes (43)

100132406

Selected node info

10971 [\[List\]](#)

Probe set is used as reference

Other hub members (8) :

[\[+\]](#) *Displayed* (14)
[\[+\]](#) *Hidden* (0)
[\[+\]](#) *Under threshold* (0)

Probe set identifiers :

[\[+\]](#) *Probe set id* (1)

[\[+\]](#) **Gene information :**

Gene identifiers :

[\[+\]](#) *Gene symbol* (1)
[\[+\]](#) *Entrez gene id* (1)
[\[+\]](#) *KEGG id* (1)
[\[+\]](#) *Ensembl gene id* (1)
[\[+\]](#) *Unigene id* (1)

Gene Ontology :

[\[+\]](#) *Biological processes* (3)
[\[+\]](#) *Cellular components* (3)
[\[+\]](#) *Molecular functions* (3)

Protein identifiers :

[\[+\]](#) [Sequence] *Swiss-Prot id* (5)
[\[+\]](#) [Domain] *InterPro id* (0)

Literature identifiers :

[\[+\]](#) *PubMed references* (123)

Disease identifiers :