

KEGG diagram legend

Node Types

Edge Types

| compound | | | |
|---------------------|---|---------------------------------|--------------|
| hidden compound | | | |
| activation | | gene (protein/enzyme) | |
| inhibition | | | |
| expression | | | |
| repression | | group (complex) | |
| indirect effect | ••••••••••••••••••••••••••••••••••••••• | | |
| state change | | | |
| binding/association | | compound (metabolite/glycan) | |
| dissociation | ••••• | | |
| phosphorylation | | | |
| dephosphorylation | <u>−p</u> | map (pathway) | Pathway name |
| glycosylation | <u>+g</u> | | |
| ubiquitination | <u>+u</u> | | |
| methylation | <u>+m</u> | | |
| others/unknown | ? → | | |