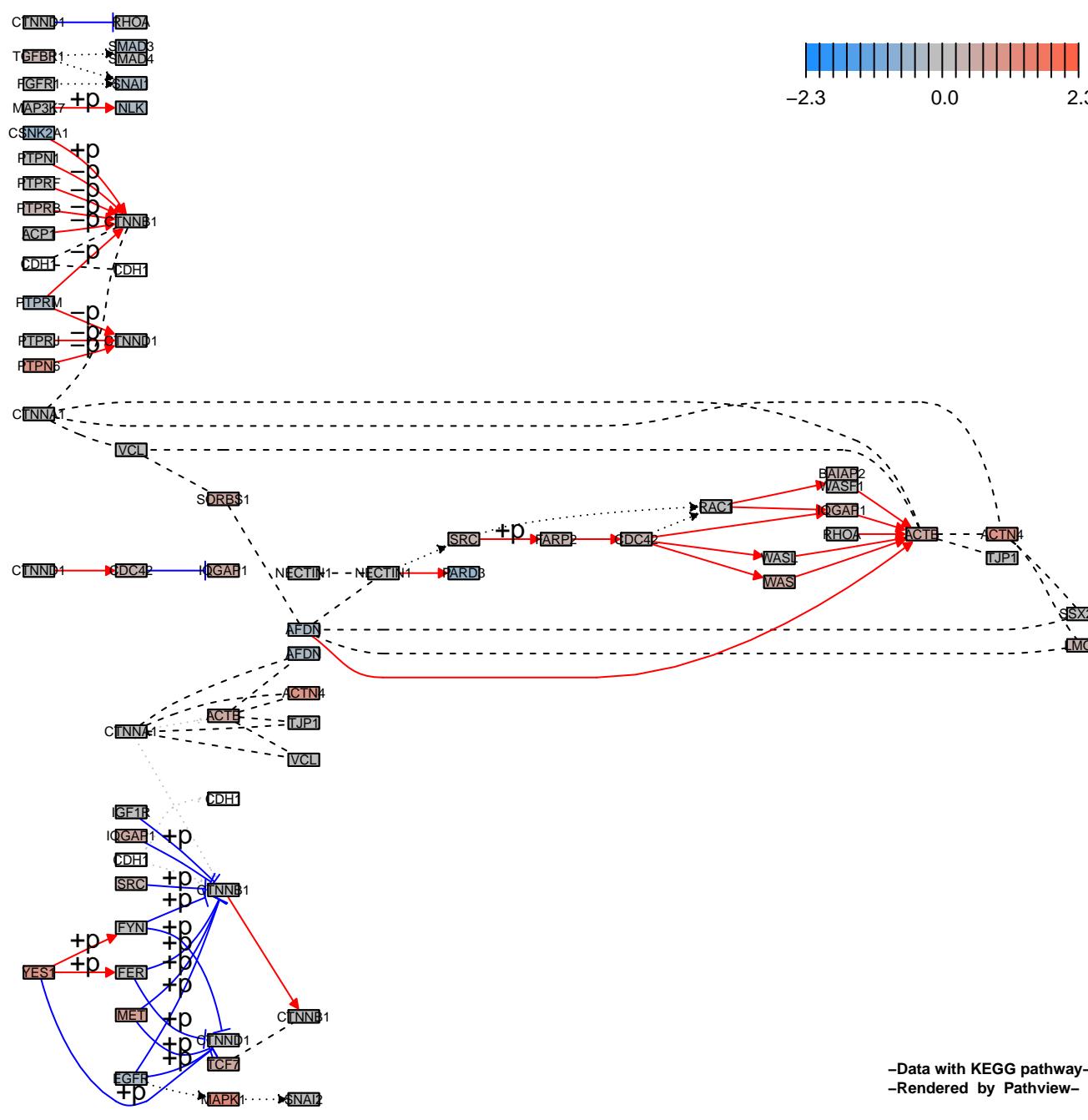
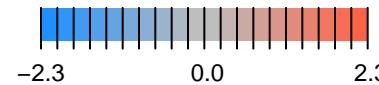


Edge types



-Data with KEGG pathway-
-Rendered by Pathview-

compound



hidden compound



activation



inhibition



expression



repression



indirect effect



state change



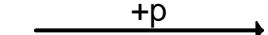
binding/association



dissociation



phosphorylation



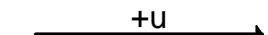
dephosphorylation



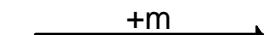
glycosylation



ubiquitination



methylation



others/unknown

