



RAB5A

EEA1

## **KEGG** diagram legend

**Node Types** 

**Edge Types** 

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