



Module score: 0.898862

GO:0006501  
 p (fisher): 0.00038365  
 C-terminal protein lipidation

GO:0006438  
 p (fisher): 0.00234855  
 valyl-tRNA aminoacylation

GO:0034645  
 p (fisher): 0.00351254  
 cellular macromolecule  
 biosynthetic process

brown: multiple overrepresented terms