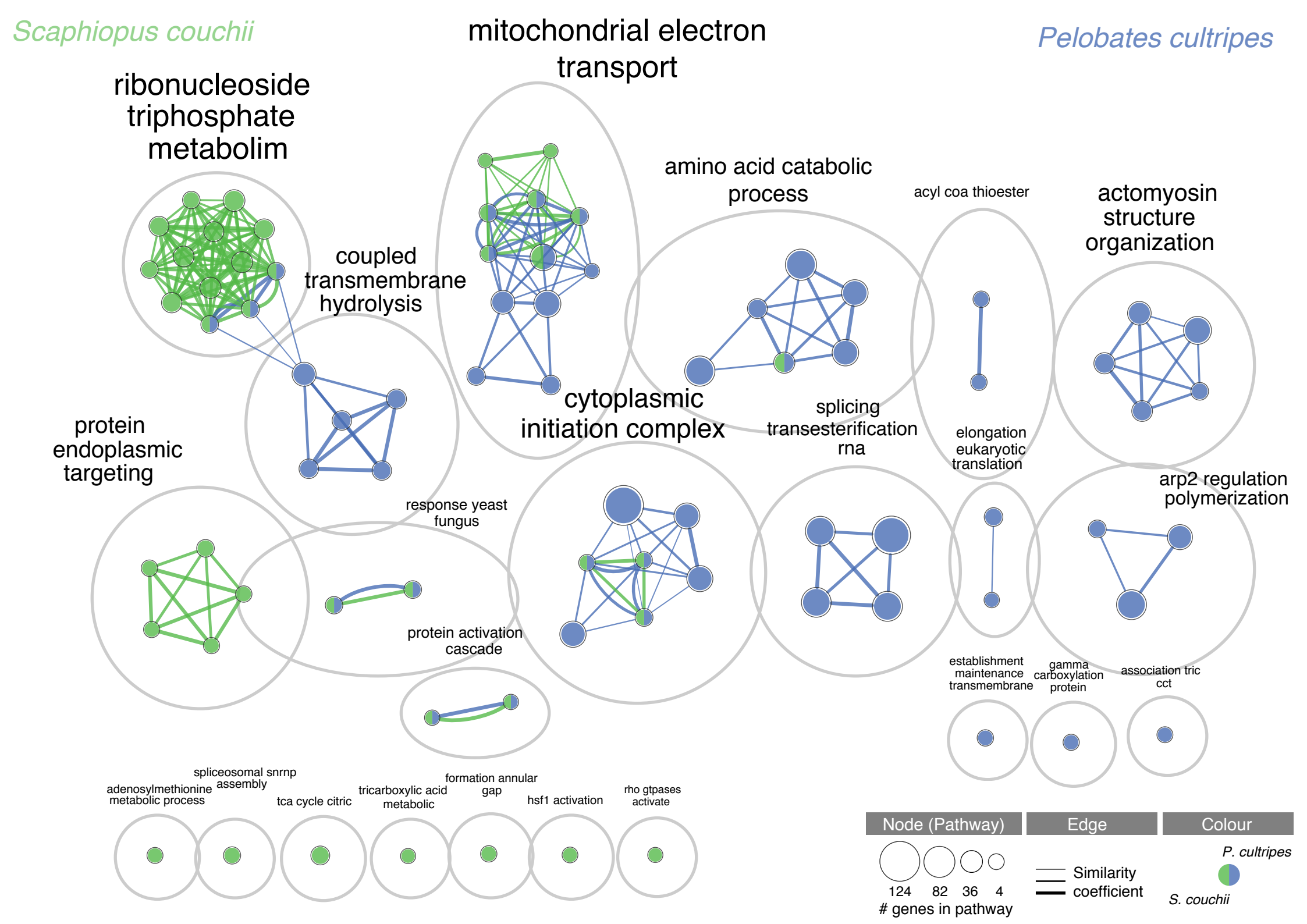
Scaphiopus couchii
Pelobates cultripes
mitochondrial electron transport
ribonucleoside triphosphate metabolim
coupled transmembrane hydrolysis
amino acid catabolic process
acyl coa thioester
actomyosin structure organization
protein endoplasmic targeting
cytoplasmic initiation complex
splicing transesterification rna
elongation eukaryotic translation
arp2 regulation polymerization
response yeast fungus
protein activation cascade
establishment maintenance transmembrane
gamma carboxylation protein
association tric cct
adenosylmethionine metabolic process
spliceosomal snrnp assembly
tca cycle citric
tricarboxylic acid metabolic
formation annular gap
hsf1 activation
rho gtpases activate
Node (Pathway)
Edge
Colour
124 82 36 4
genes in pathway
Similarity coefficient
P. cultripes
S. couchii