MCODE cluster 5
Nodes: 8; edges: 26; score ~7.4
KEGG: Glycerophospholipid metabolism (8/8); phosphatidylinositol signaling (3/8)

MCODE cluster 6
Nodes: 77; edges: 268; score ~7.1
KEGG: RNA transport (15/77); ubiquitin mediated proteolysis (10/77); purine metabolism (10/77); glycolysis (7/77)

MCODE cluster 7
Nodes: 80; edges: 258; score ~6.5
KEGG: Endocytosis (7/80); DNA replication (5/80)

MCODE cluster 8
Nodes: 11; edges: 32; score ~6.4
KEGG: Spliceosome (6/8)

MCODE cluster 9
Nodes: 73; edges: 204; score ~5.7
KEGG: mRNA surveillance (9/73); inositol phosphate metabolism (6/73); aminoacyl-tRNA biosynthesis (6/73)

MCODE cluster 10
Nodes: 6; edges: 14; score ~5.6
KEGG: SNARE interactions in vesicular transport (3/6)