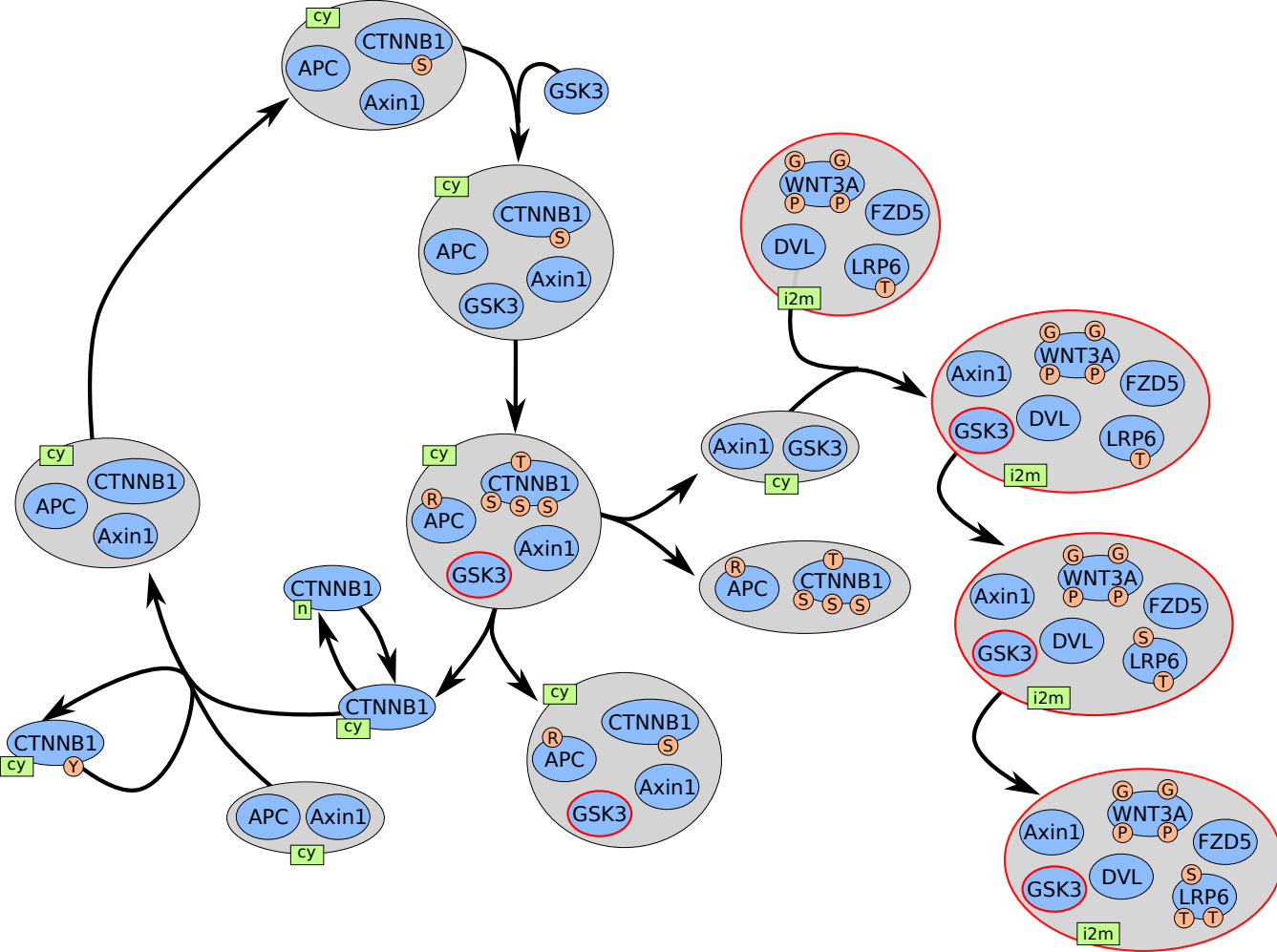




Key

Y: Tyrosine phosphorylation
 T: Threonine phosphorylation
 S: Serine phosphorylation
 R: phosphorylated residue
 Red: activating modification
 G: glycosylation
 P: palmitoylation

n: location=nucleus
 cy: location=cytoplasm
 i2m: location=integral to membrane

beta-cat,cytoplasm: pid_26794
 beta-cat, cytoplasm Y@654: pid_26317
 beta-cat: pid_26214
 beta-cat, S@45: pid_77223
 beta-cat,S@33,S@37,S@45,T@41: pid_77230

APC: pid_3479
 APC, R@NULL: pid_77234
 Axin1: pid_48362
 GSK3: pid_68387
 GSK3, activating modification: pid_77221

WNT3A,G@298,G@87,P@209,P@77: pid_206
 FZD5: pid_207
 DVL: pid_77290
 LRP6, T@1479: pid_88770
 LRP6, S@1490,T@1479: pid_77301
 LRP6, S@1490,T@1479, T@1493: pid_88665