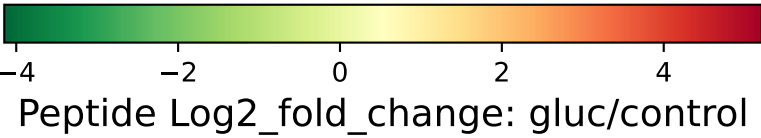
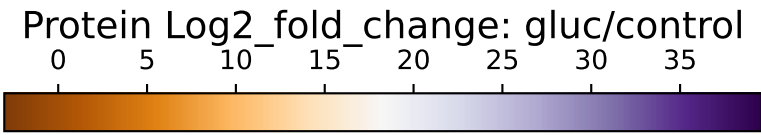
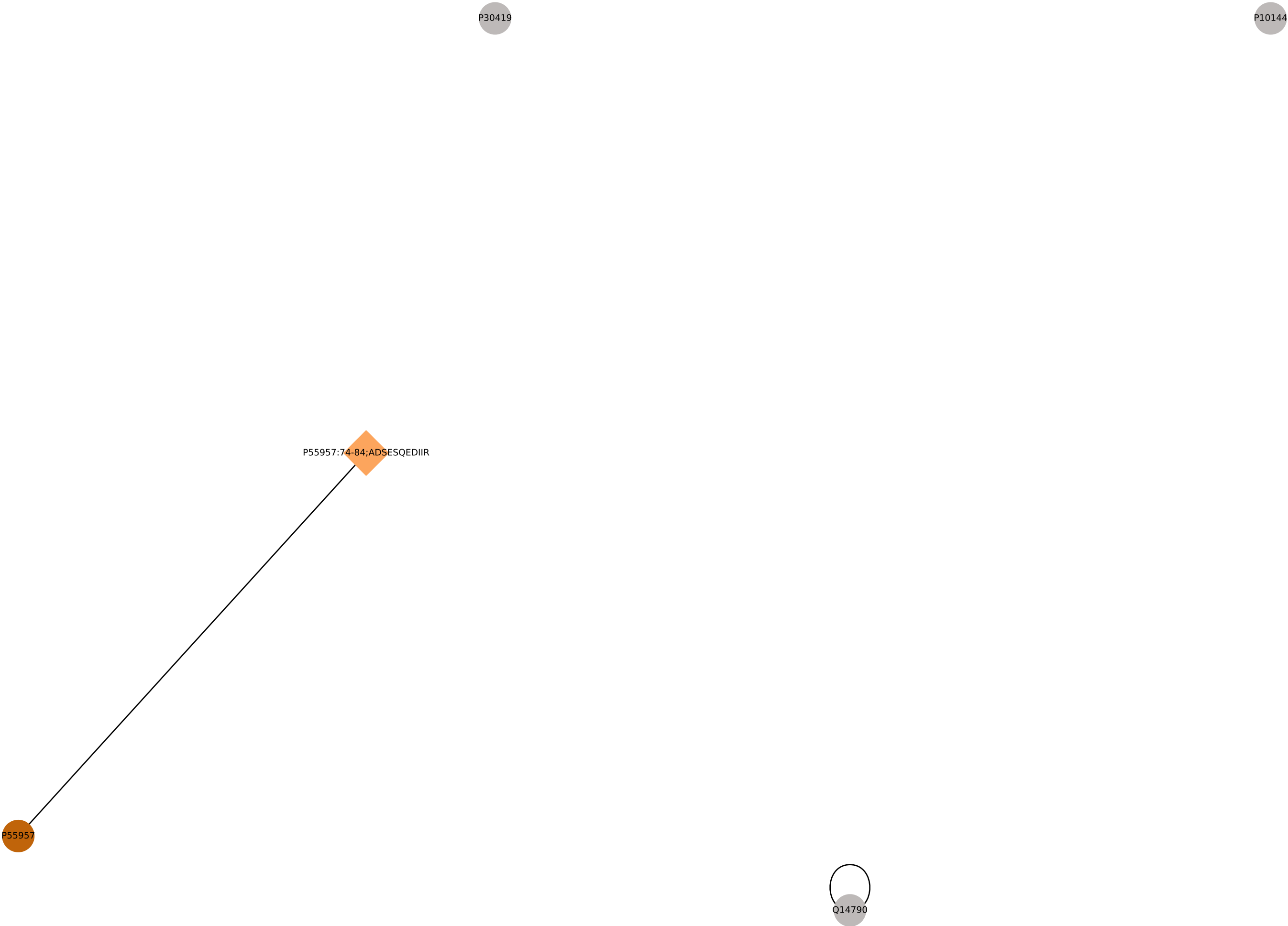
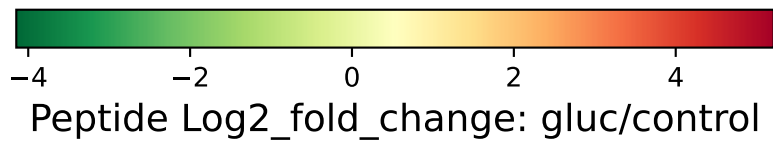
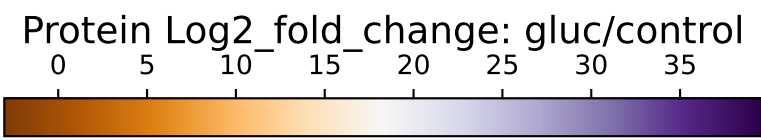
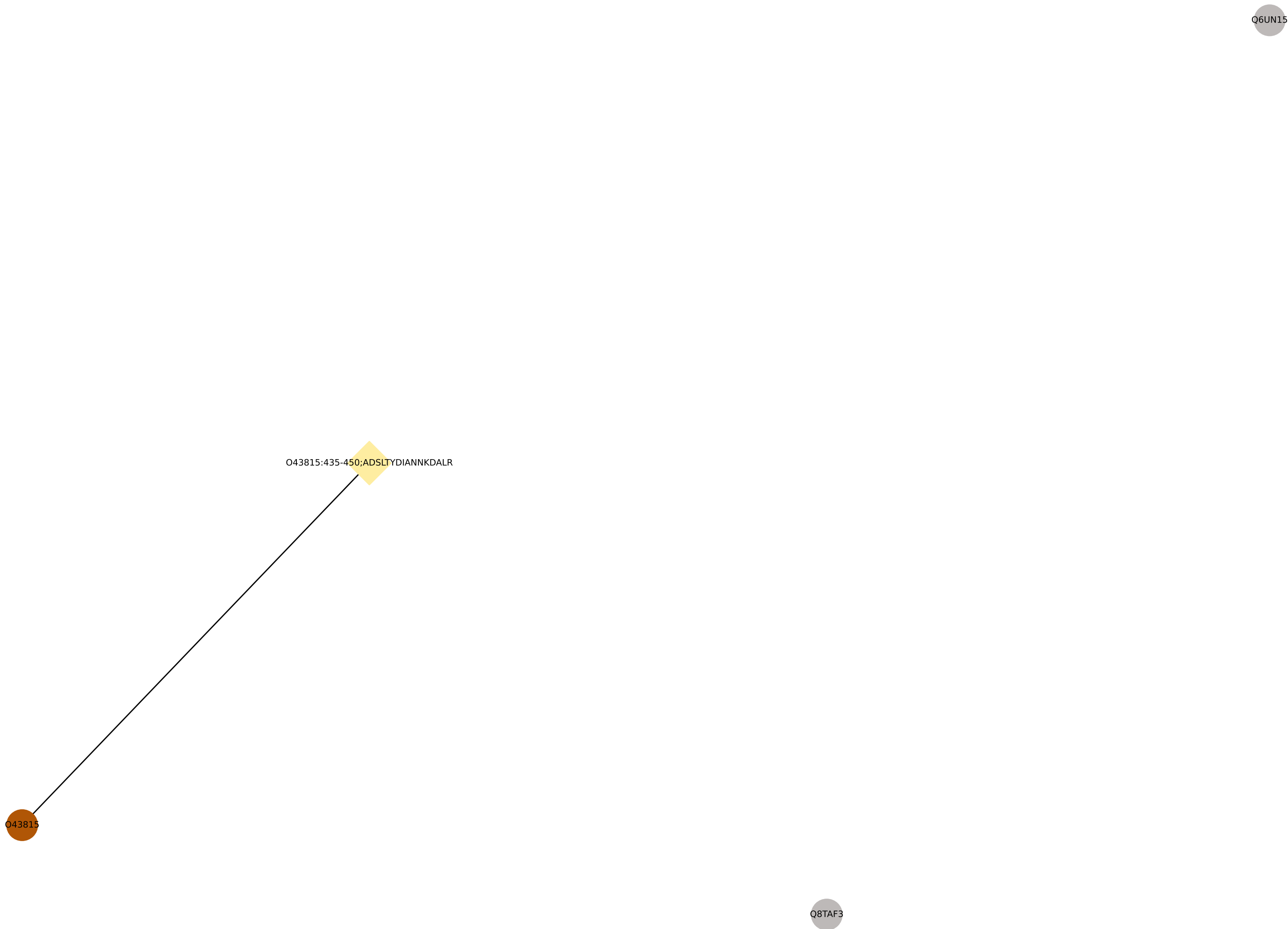


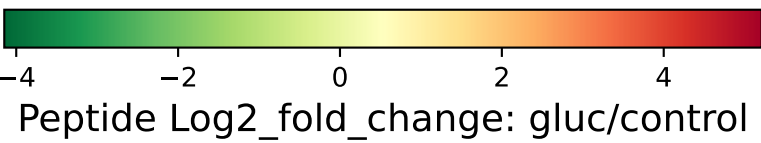
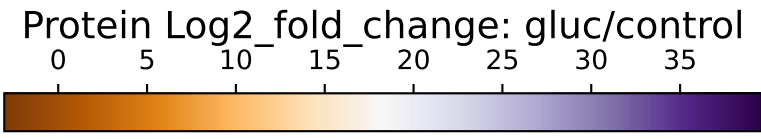
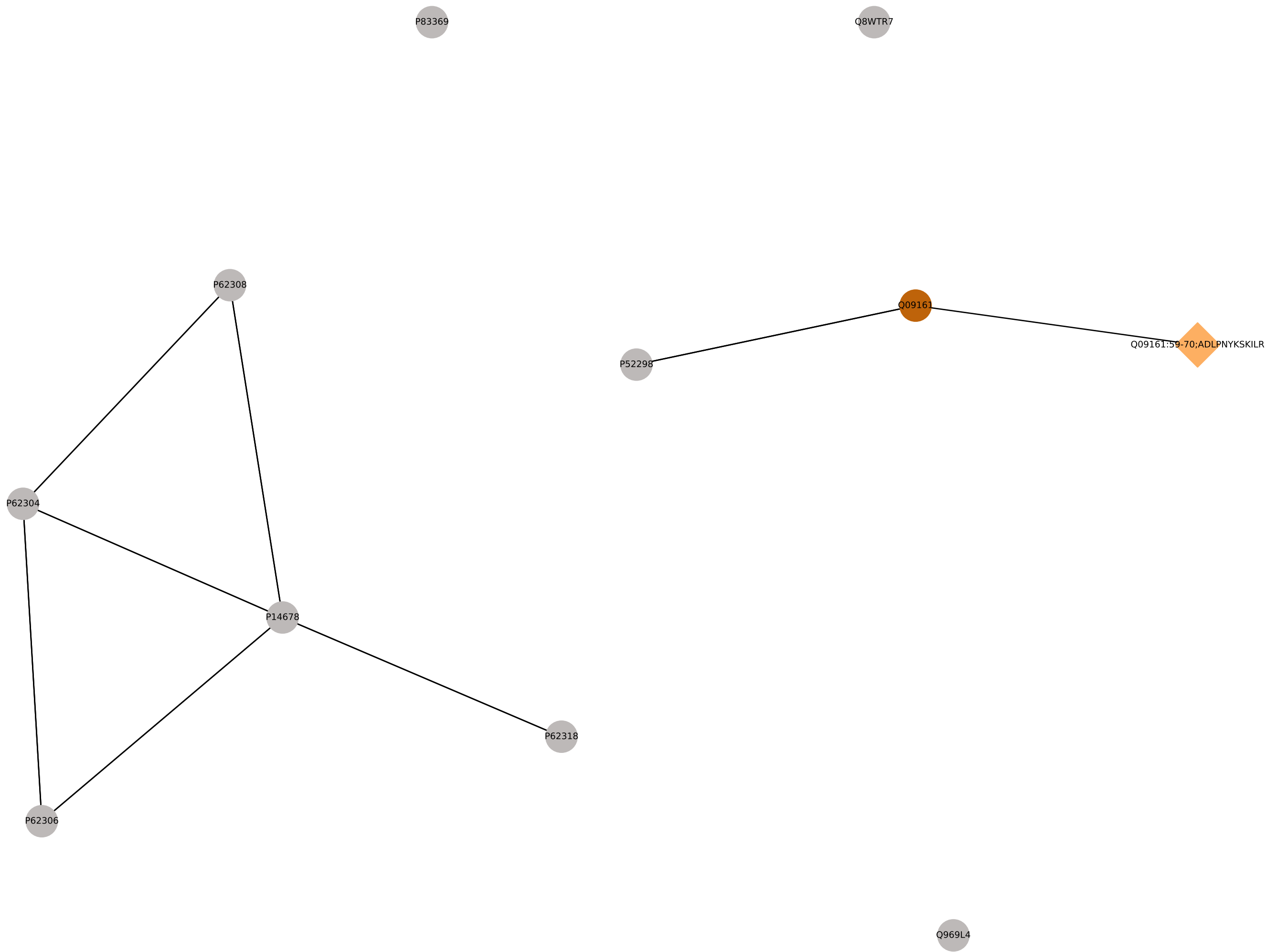
Log2\_fold\_change: gluc/control  
R-HSA-75108: Activation, myristoylation of BID and translocation to mitochondria  
p-value: 0.0151760001661293



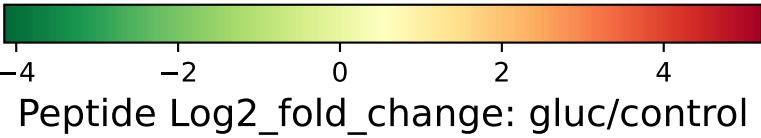
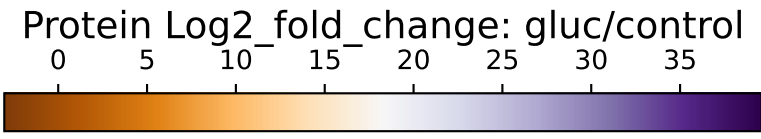
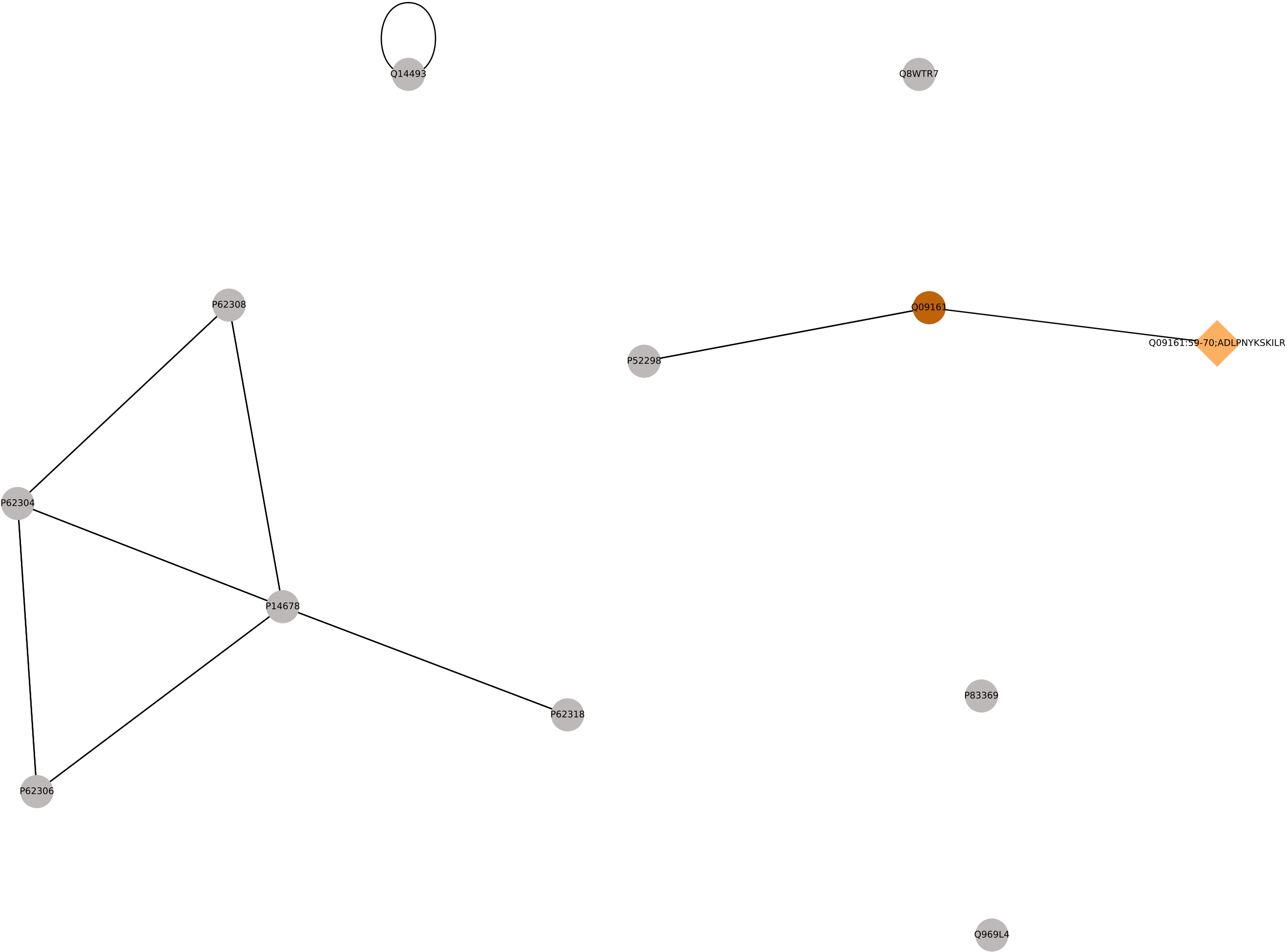
Log2\_fold\_change: gluc/control  
R-HSA-9673766: Signaling by cytosolic PDGFRA and PDGFRB fusion proteins  
p-value: 0.01893449482309495



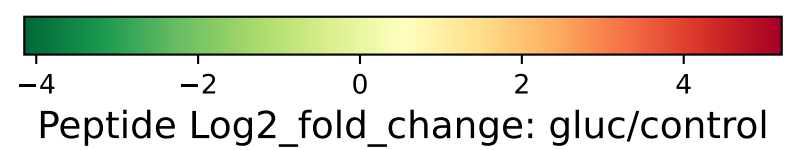
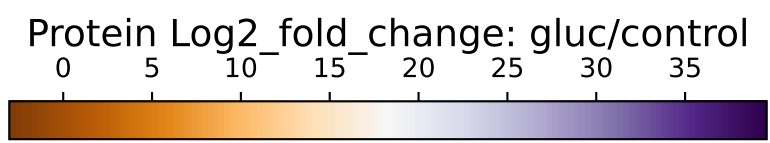
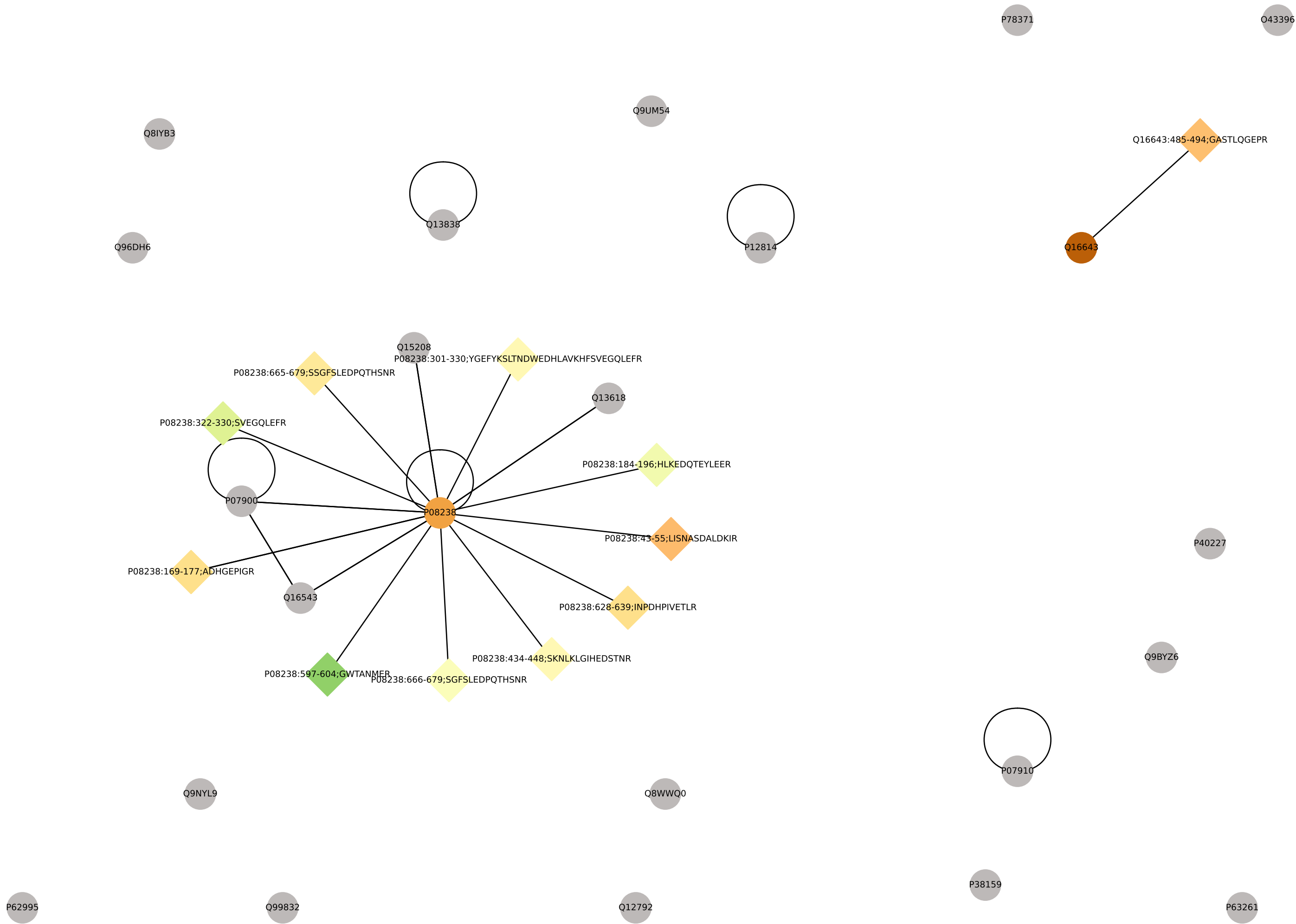
Log2\_fold\_change: gluc/control  
R-HSA-111367: SLBP independent Processing of Histone Pre-mRNAs  
p-value: 0.03751654060258469



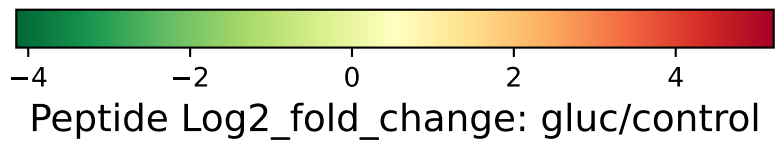
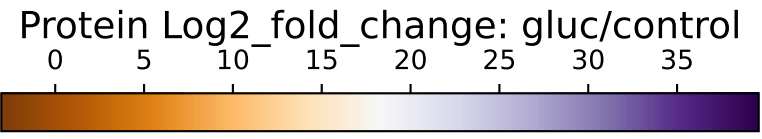
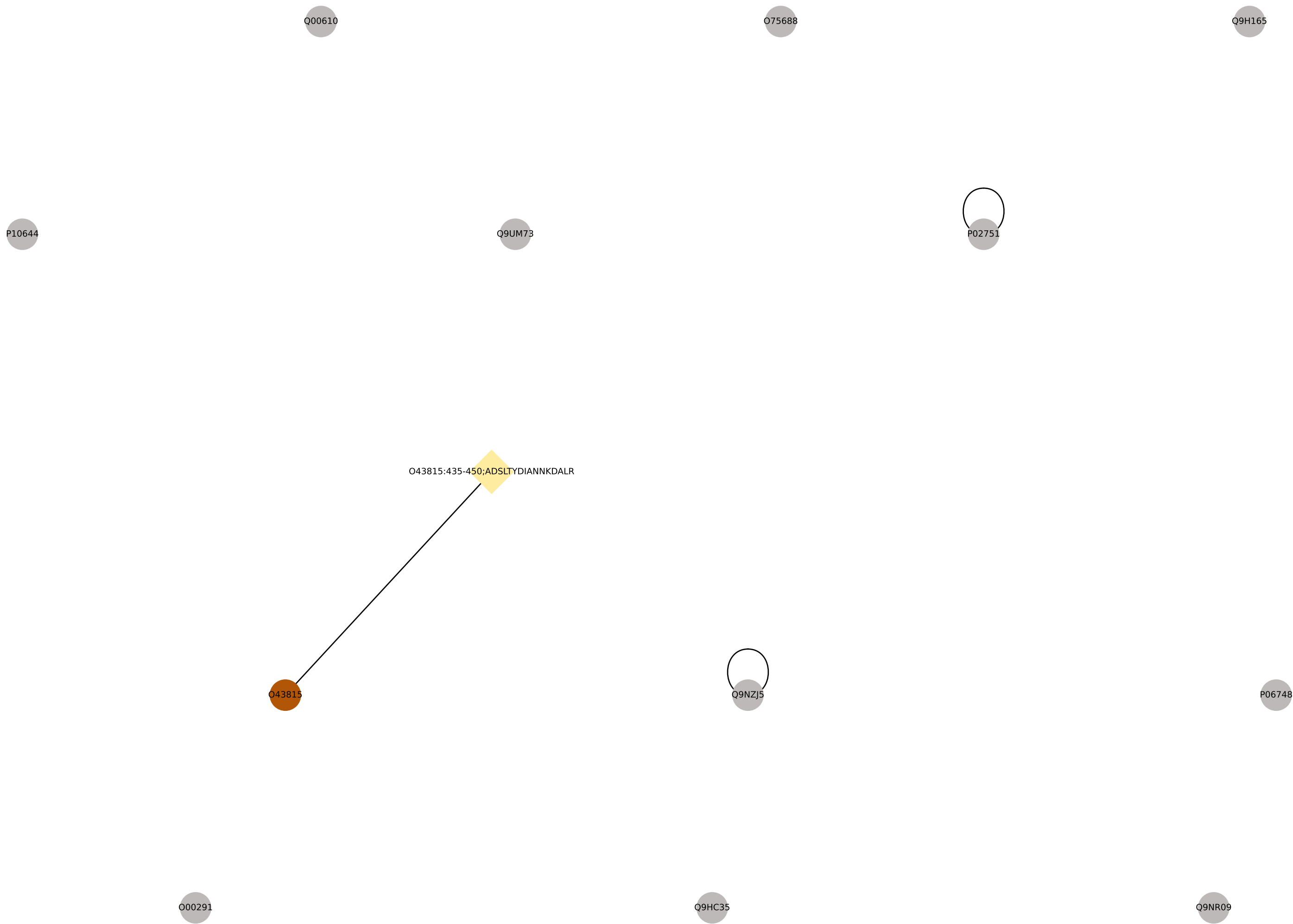
Log2\_fold\_change: gluc/control  
R-HSA-77588: SLBP Dependent Processing of Replication-Dependent Histone Pre-mRNAs  
p-value: 0.04119122530952668



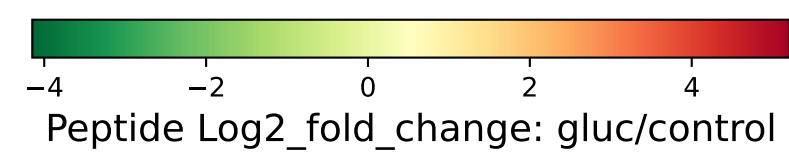
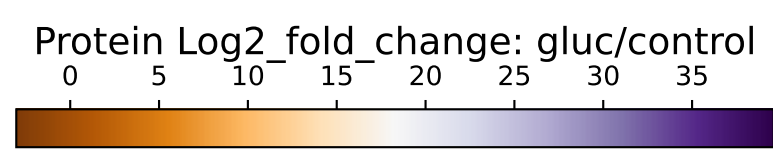
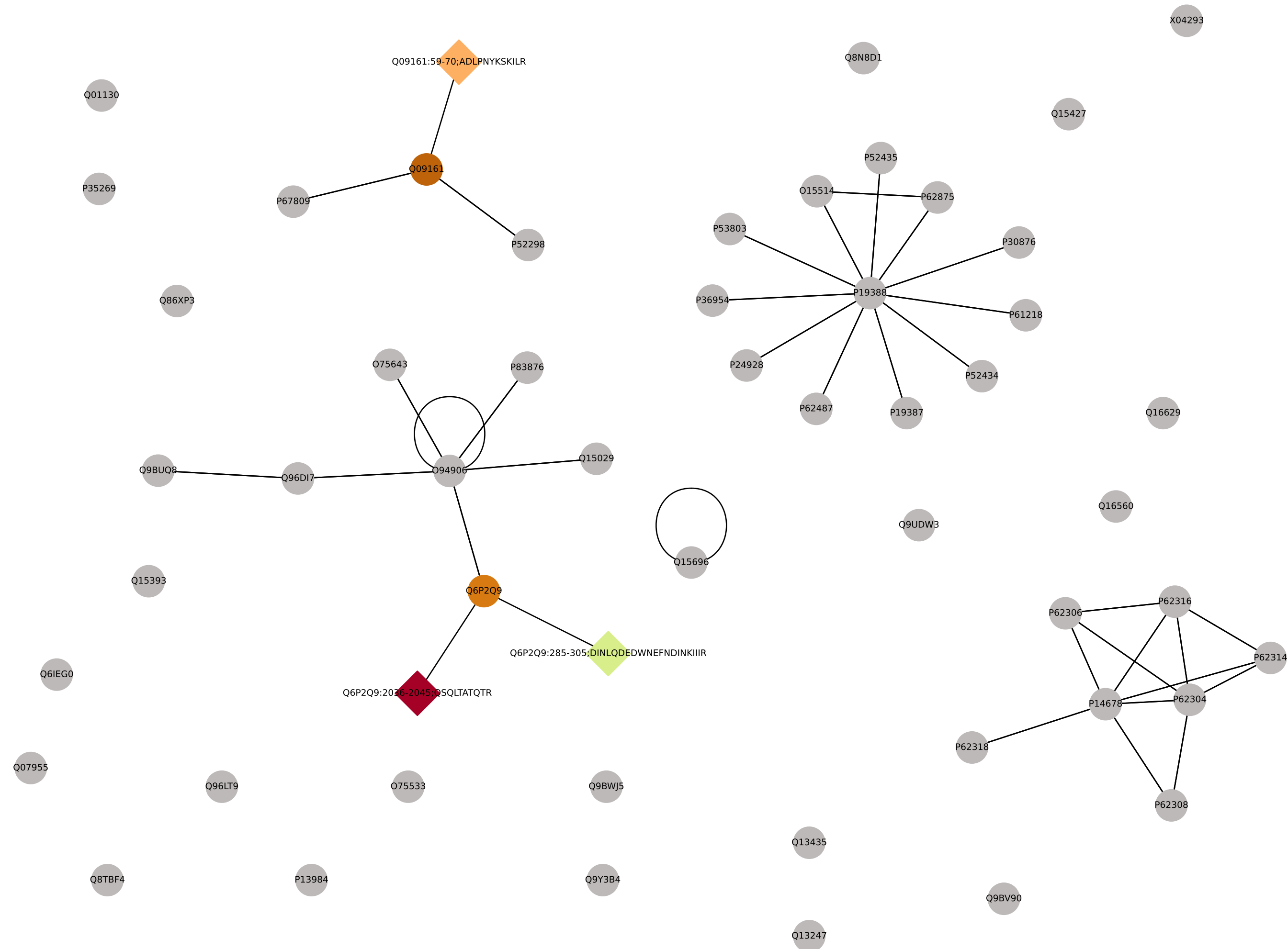
Log2\_fold\_change: gluc/control  
R-HSA-9013418: RHOBTB2 GTPase cycle  
p-value: 0.0038992971180269453



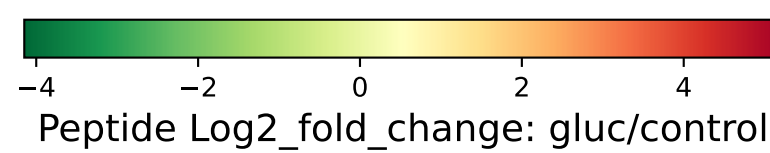
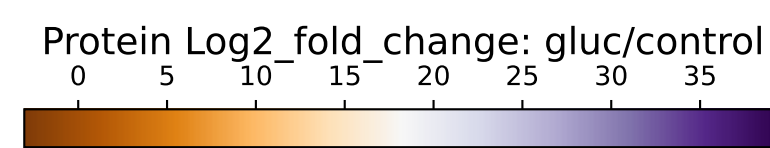
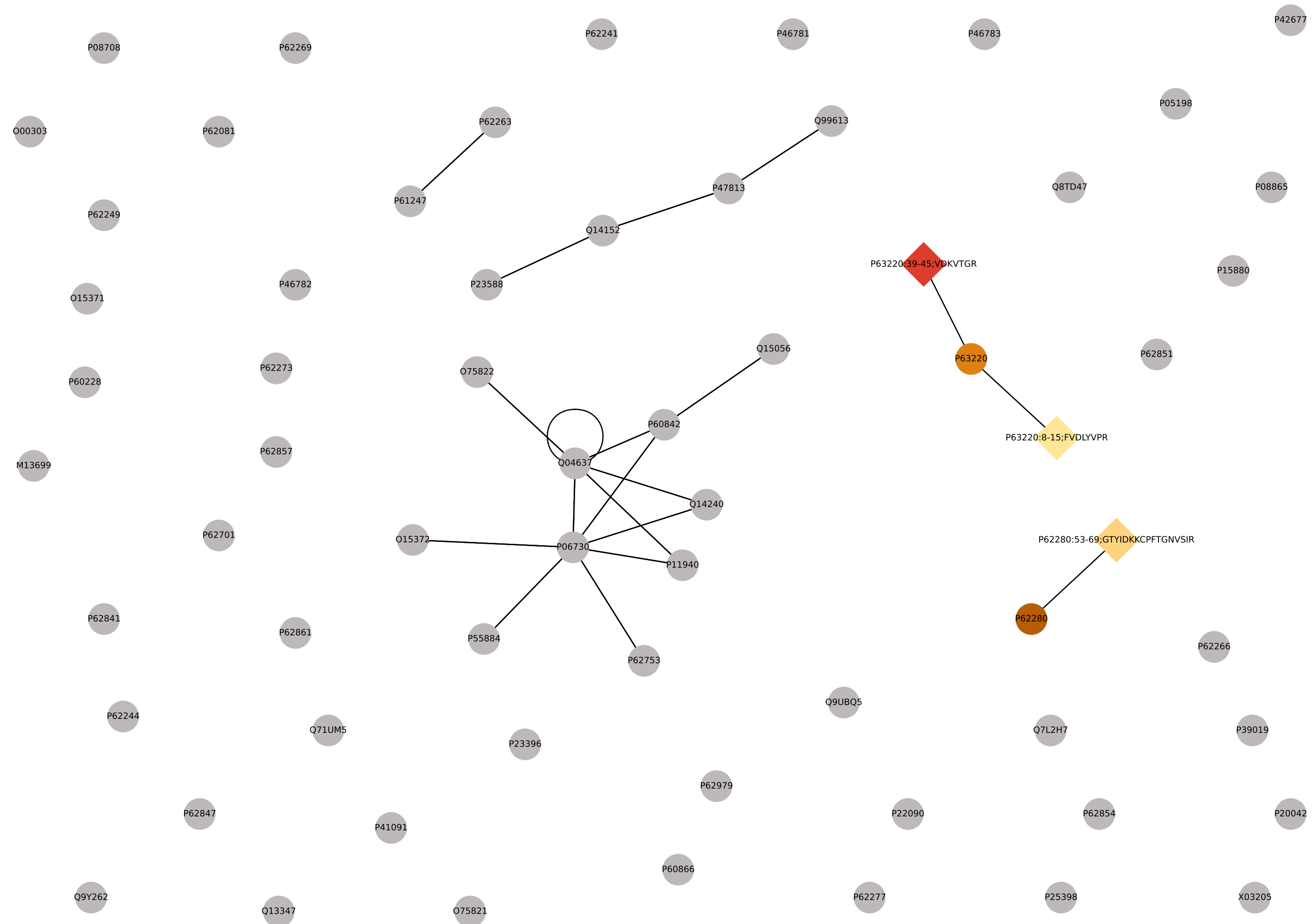
Log2\_fold\_change: gluc/control  
R-HSA-9700645: ALK mutants bind TKIs  
p-value: 0.04485212134737271



Log2\_fold\_change: gluc/control  
R-HSA-72165: mRNA Splicing - Minor Pathway  
p-value: 0.01834940897301718

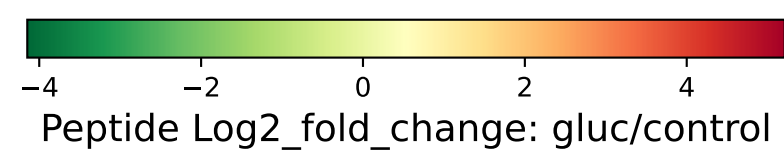
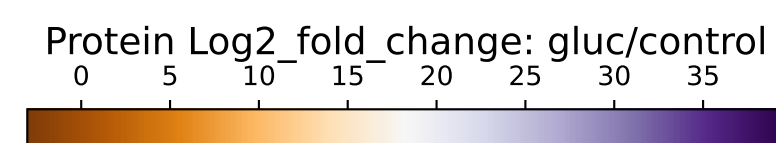
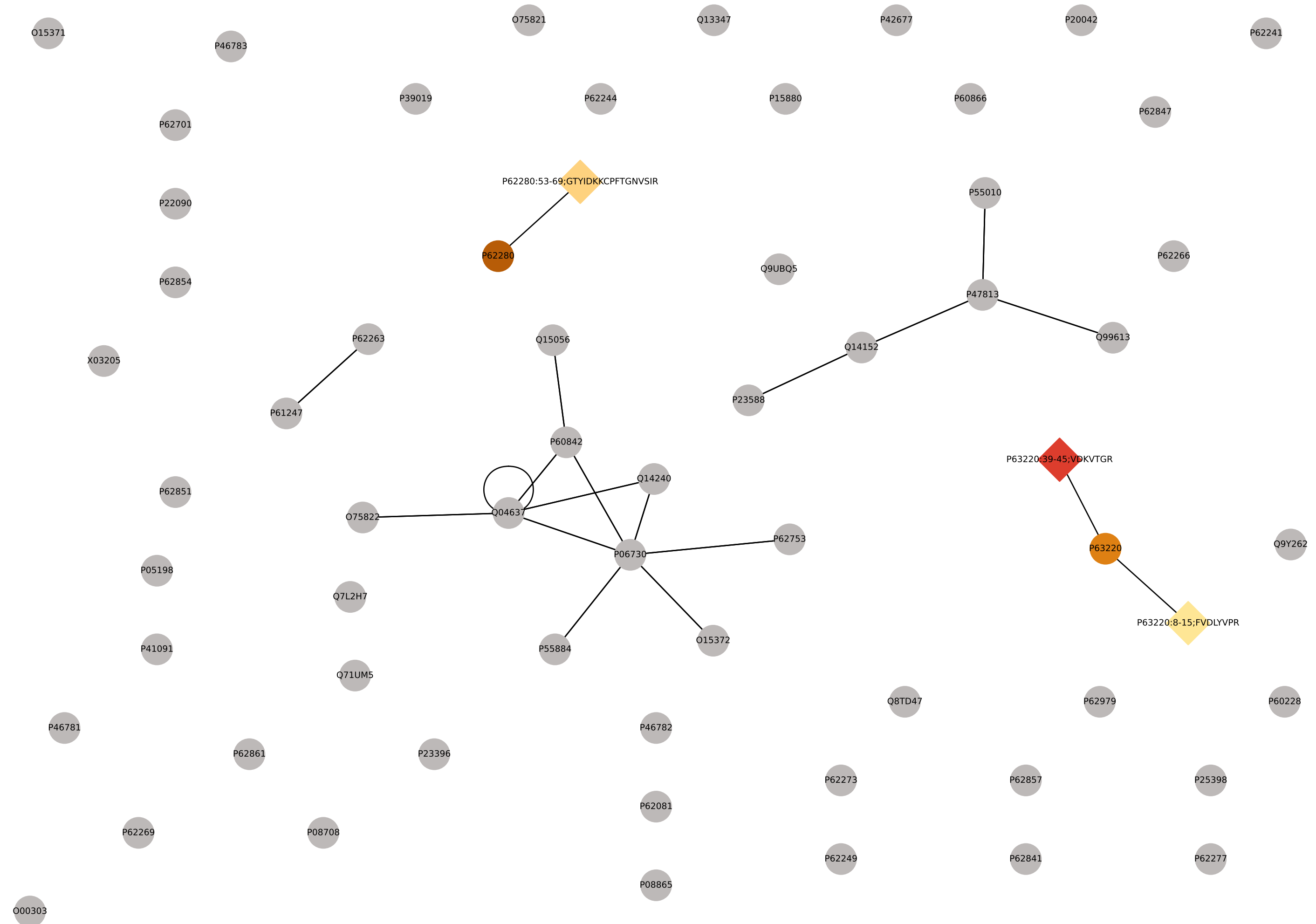


Log2\_fold\_change: gluc/control  
R-HSA-72649: Translation initiation complex formation  
p-value: 0.02372468033749031

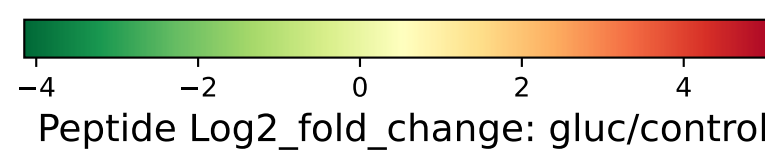
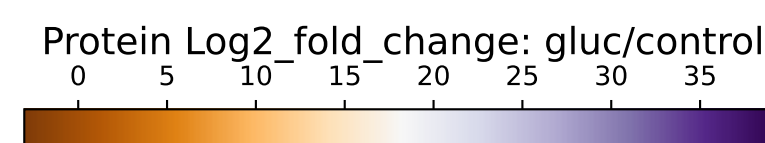
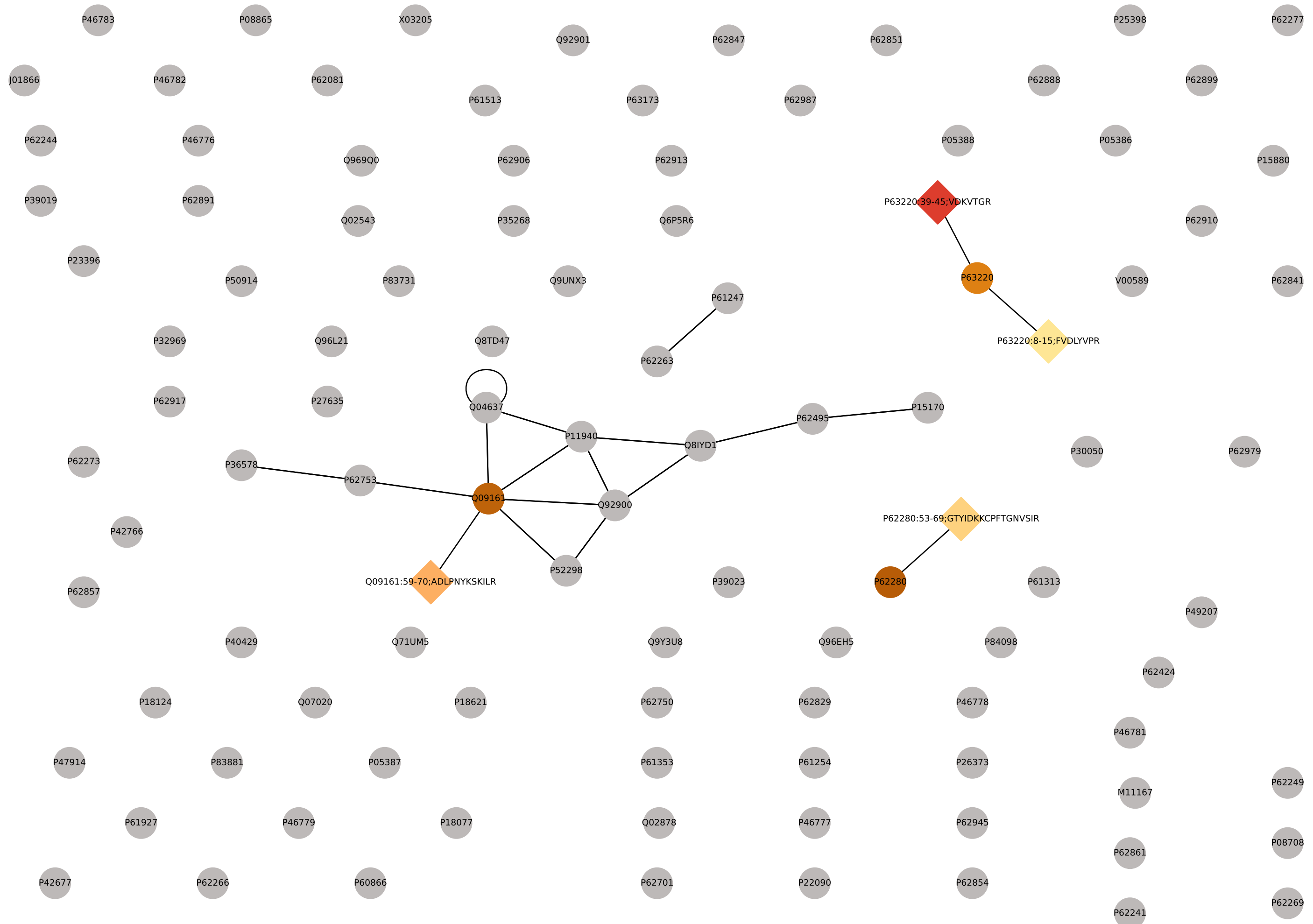




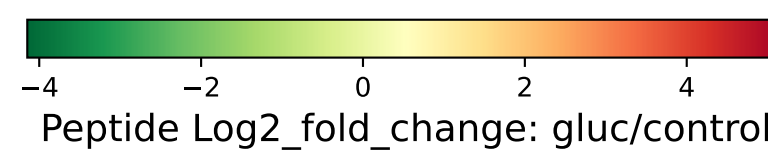
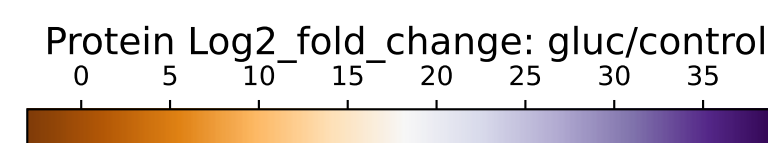
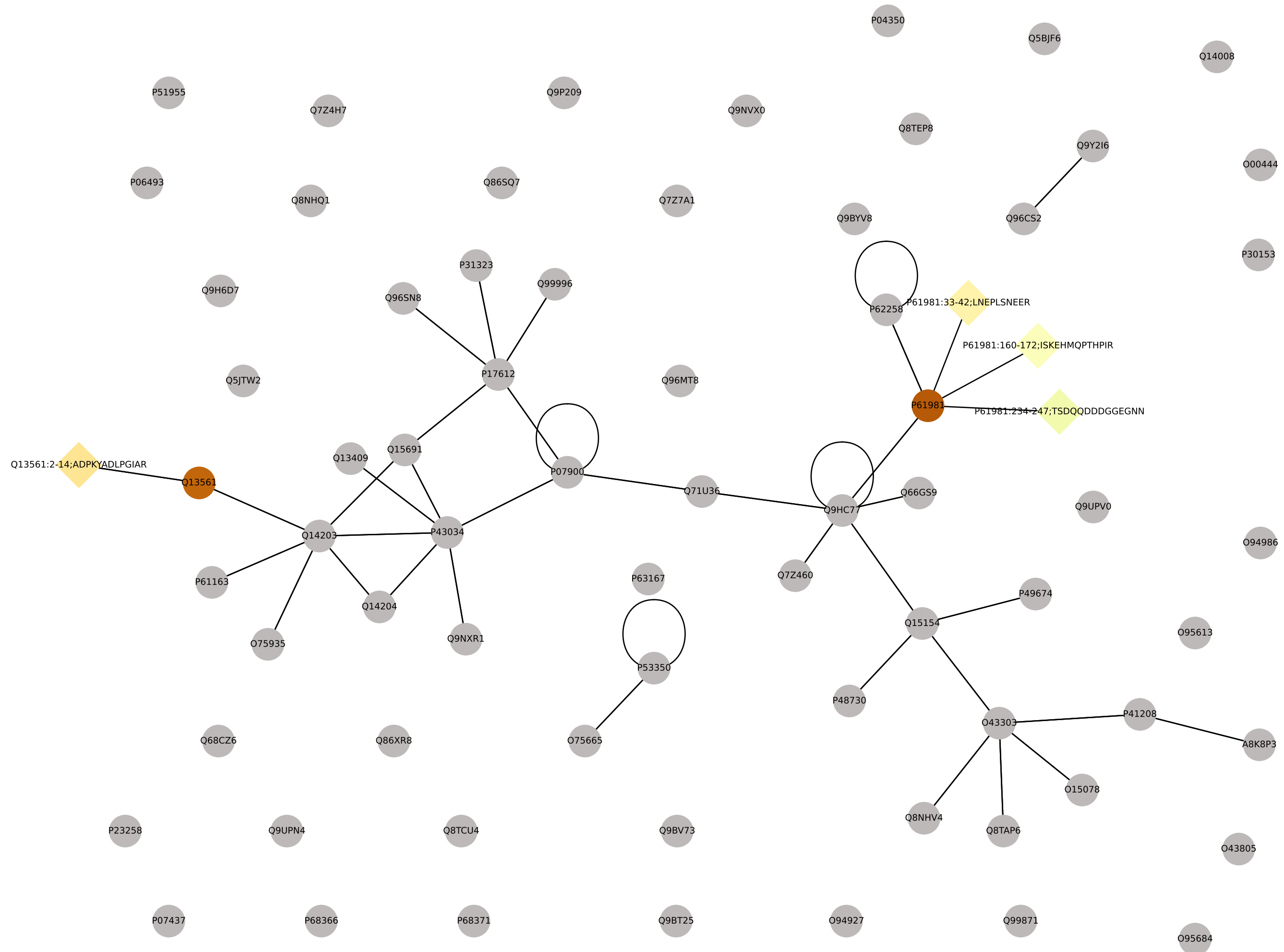
Log2\_fold\_change: gluc/control  
R-HSA-72702: Ribosomal scanning and start codon recognition  
p-value: 0.025157961205597323



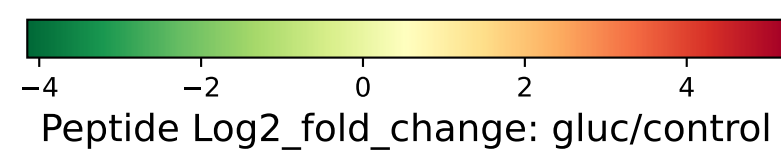
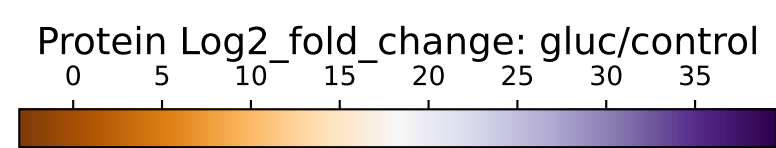
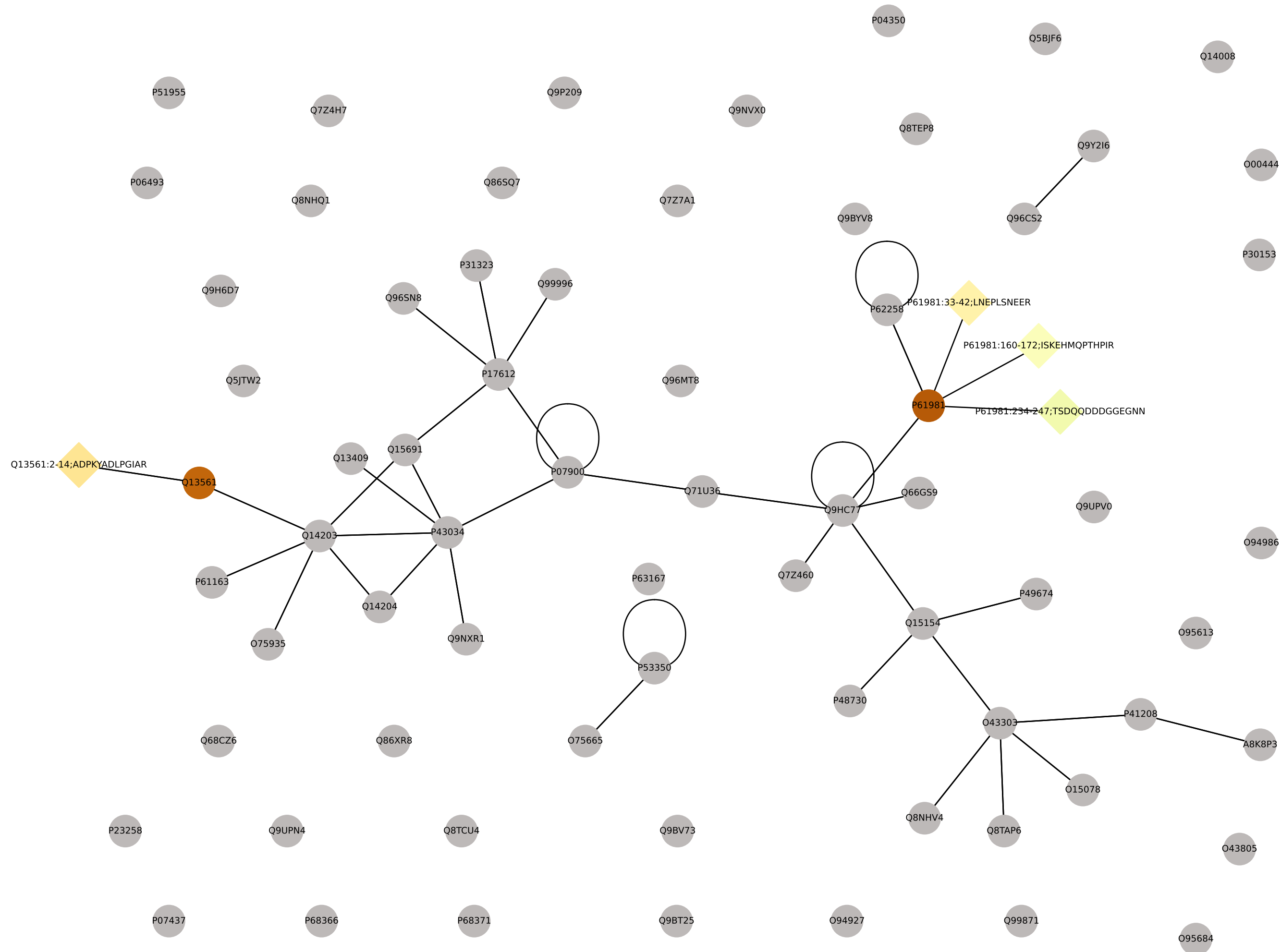
Log2\_fold\_change: gluc/control  
R-HSA-975956: Nonsense Mediated Decay (NMD) independent of the Exon Junction Complex (EJC)  
p-value: 0.0069285827914021825



Log2\_fold\_change: gluc/control  
R-HSA-380284: Loss of proteins required for interphase microtubule organization from the centrosome  
p-value: 0.030442907302830946

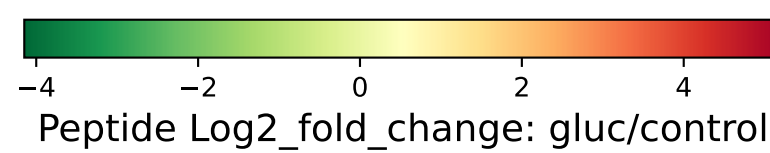
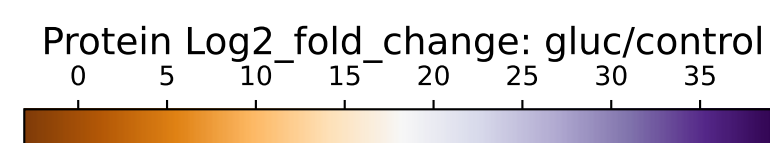
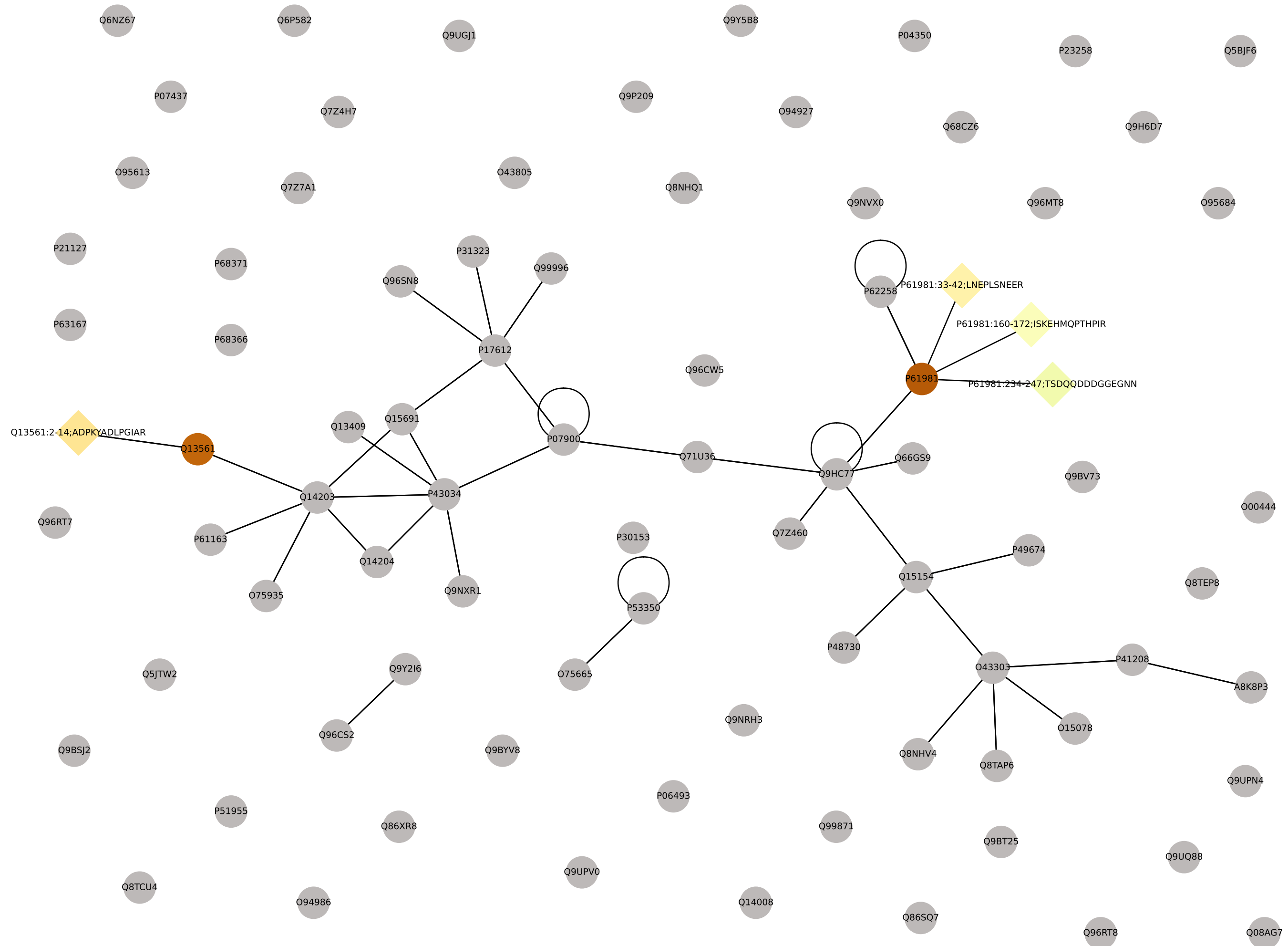


Log2\_fold\_change: gluc/control  
R-HSA-380259: Loss of Nlp from mitotic centrosomes  
p-value: 0.030442907302830946

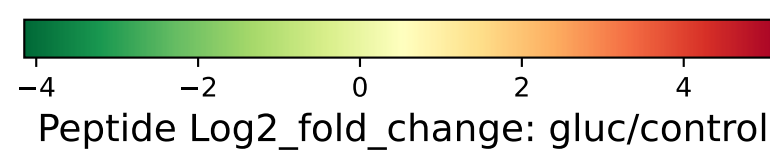
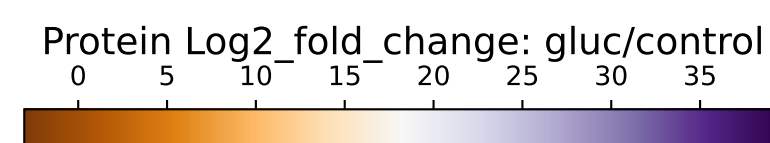
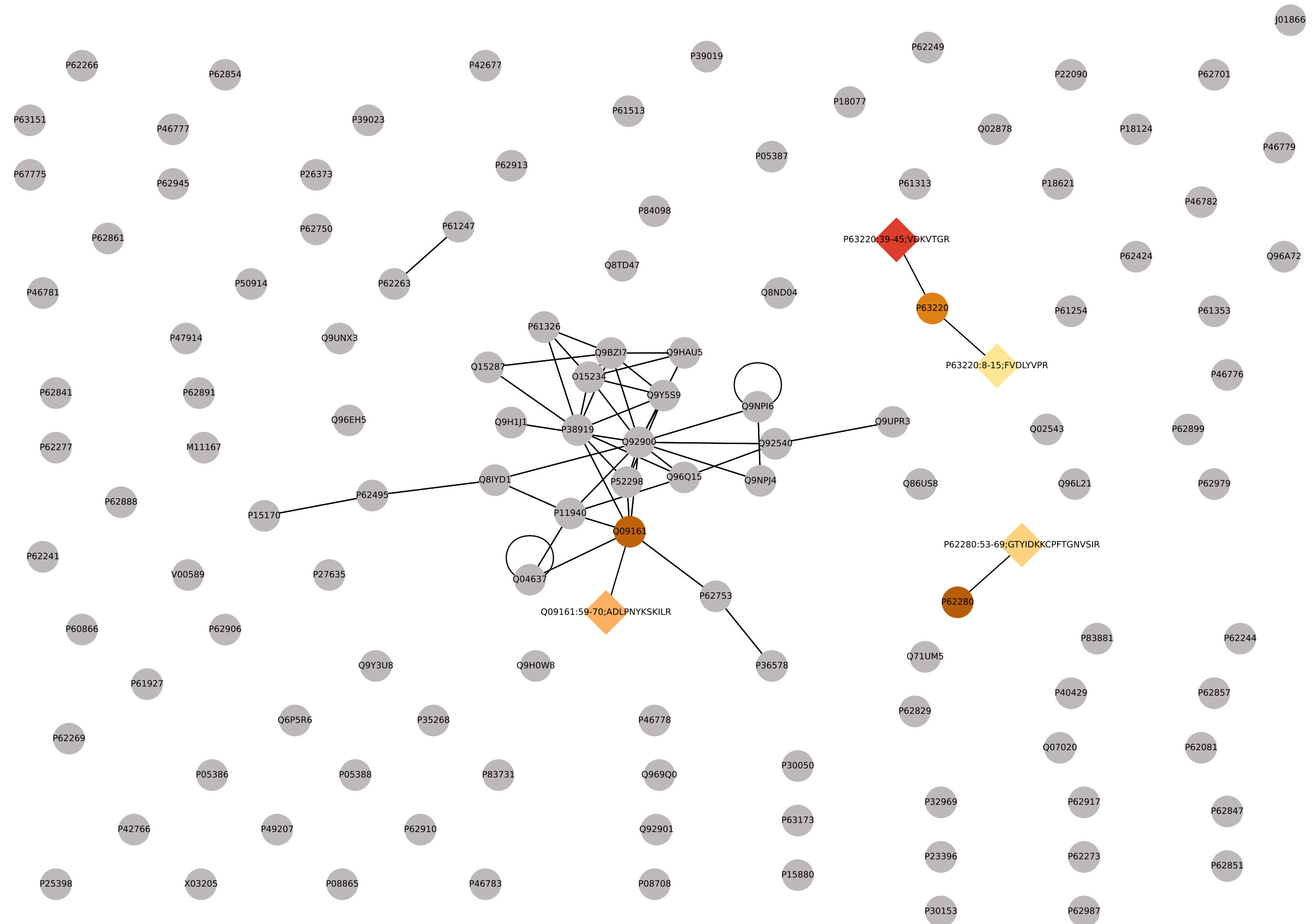




Log2\_fold\_change: gluc/control  
R-HSA-380270: Recruitment of mitotic centrosome proteins and complexes  
p-value: 0.038678528314858074

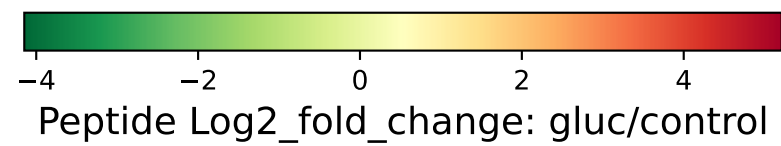
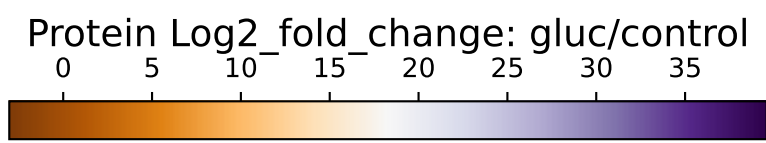
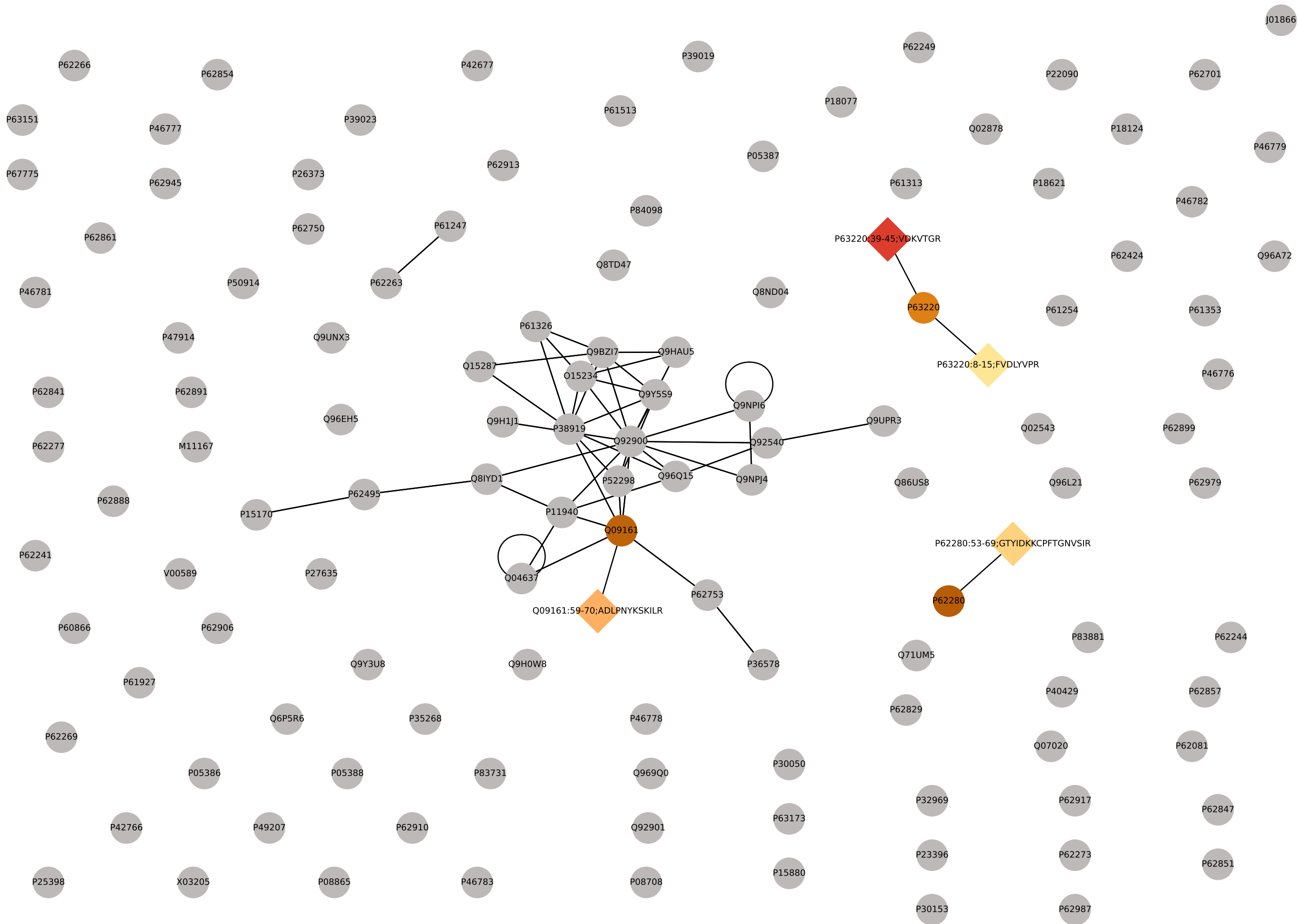


Log2\_fold\_change: gluc/control  
R-HSA-927802: Nonsense-Mediated Decay (NMD)  
p-value: 0.012056147753095936



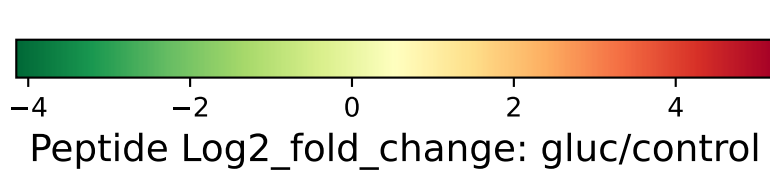
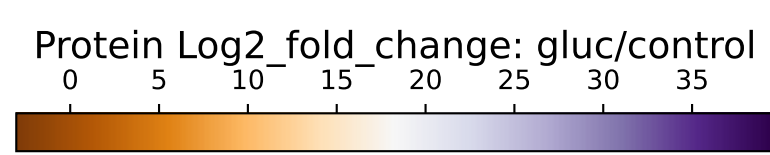
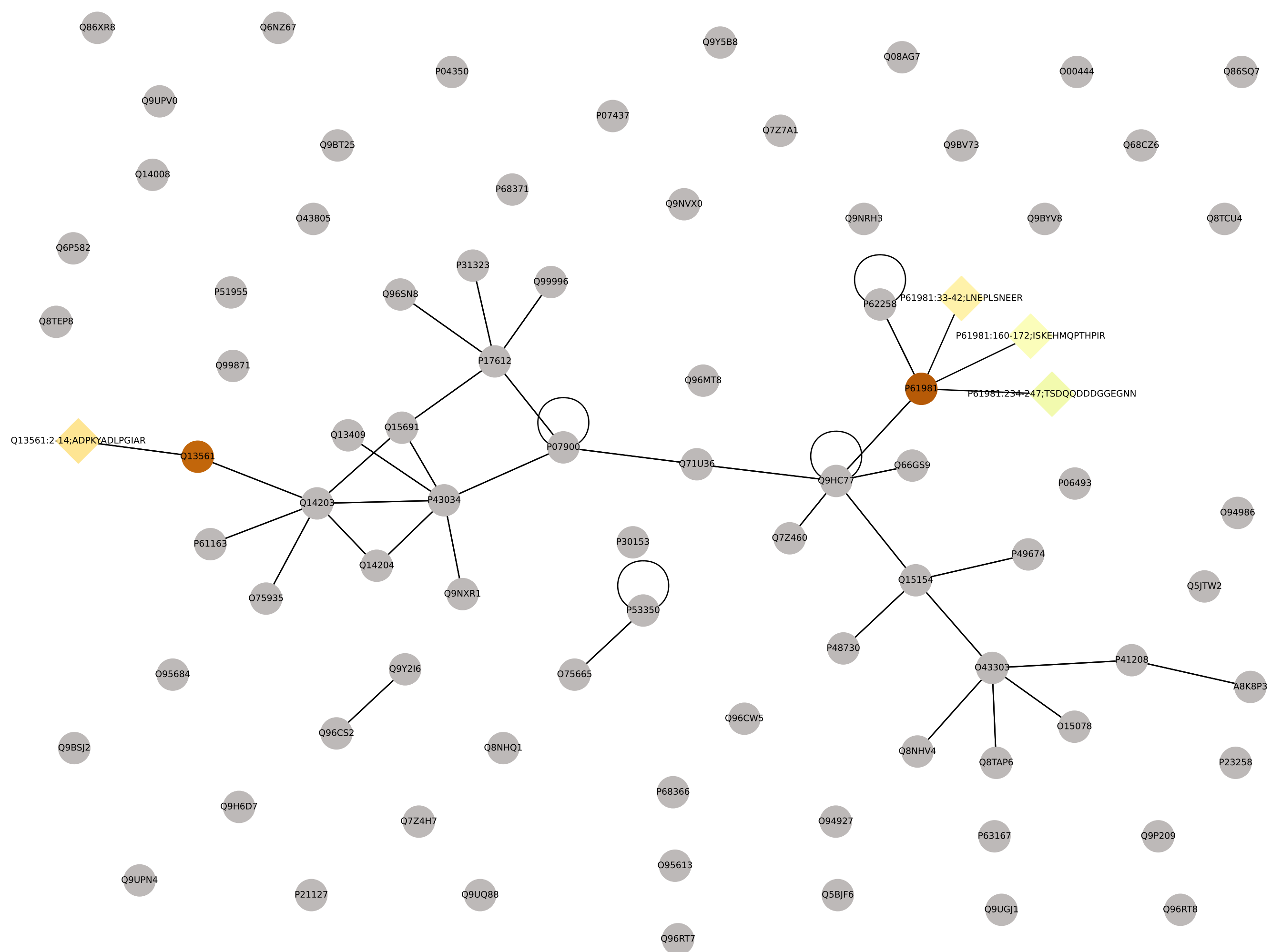


Log2\_fold\_change: gluc/control  
R-HSA-975957: Nonsense Mediated Decay (NMD) enhanced by the Exon Junction Complex (EJC)  
p-value: 0.012056147753095936





Log2\_fold\_change: gluc/control  
R-HSA-380287: Centrosome maturation  
p-value: 0.040417152198187645



Log2\_fold\_change: gluc/control  
R-HSA-72163: mRNA Splicing - Major Pathway  
p-value: 0.009025927255451371

