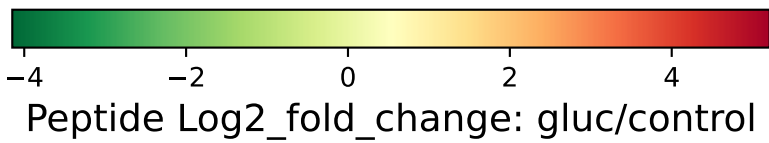
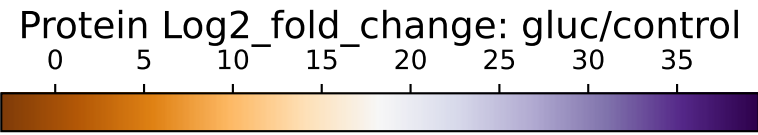
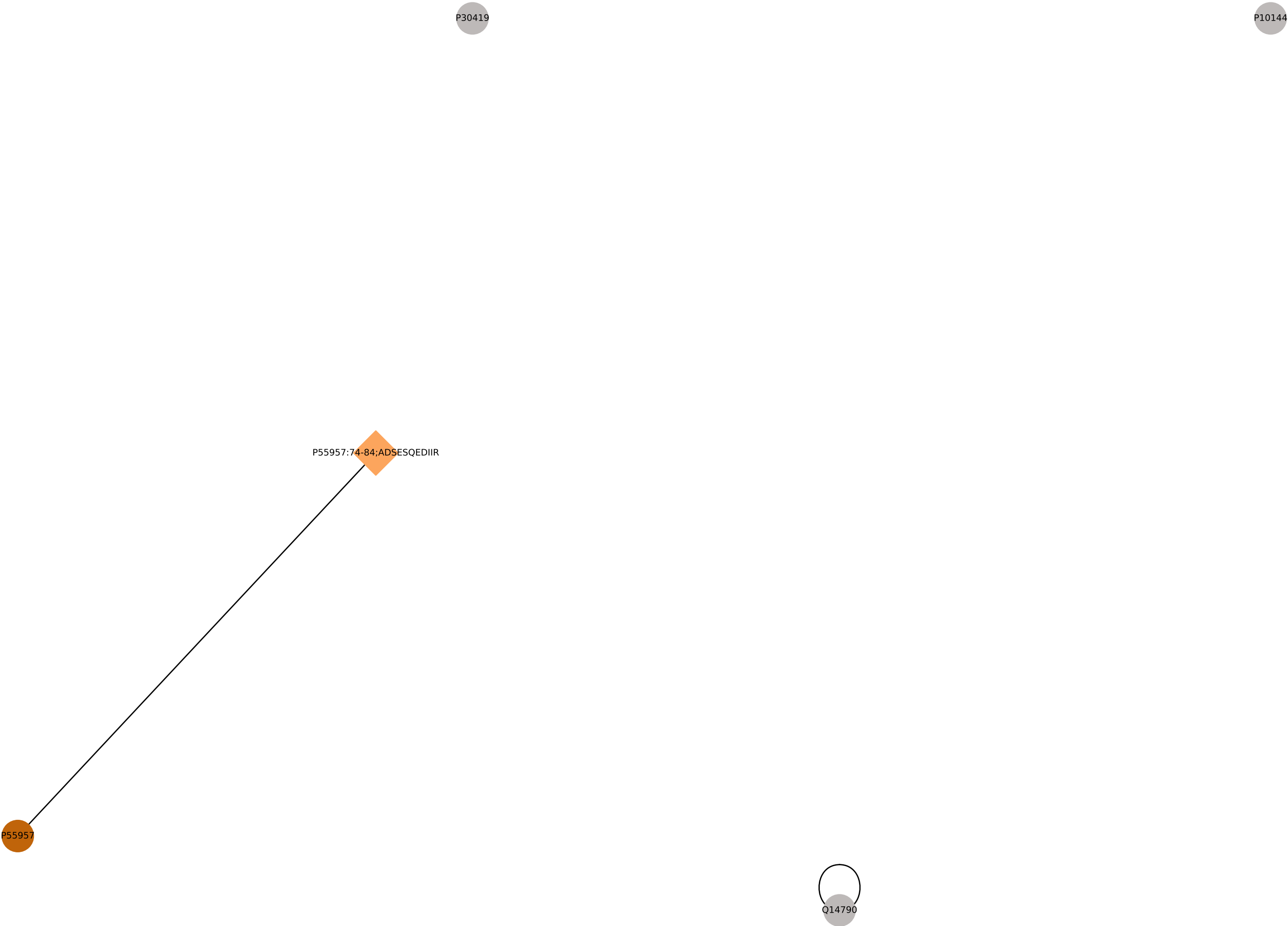
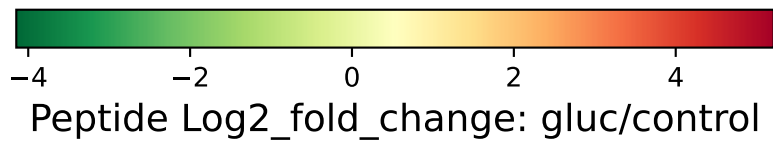
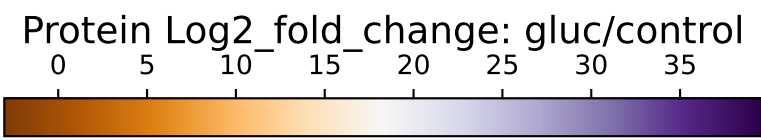
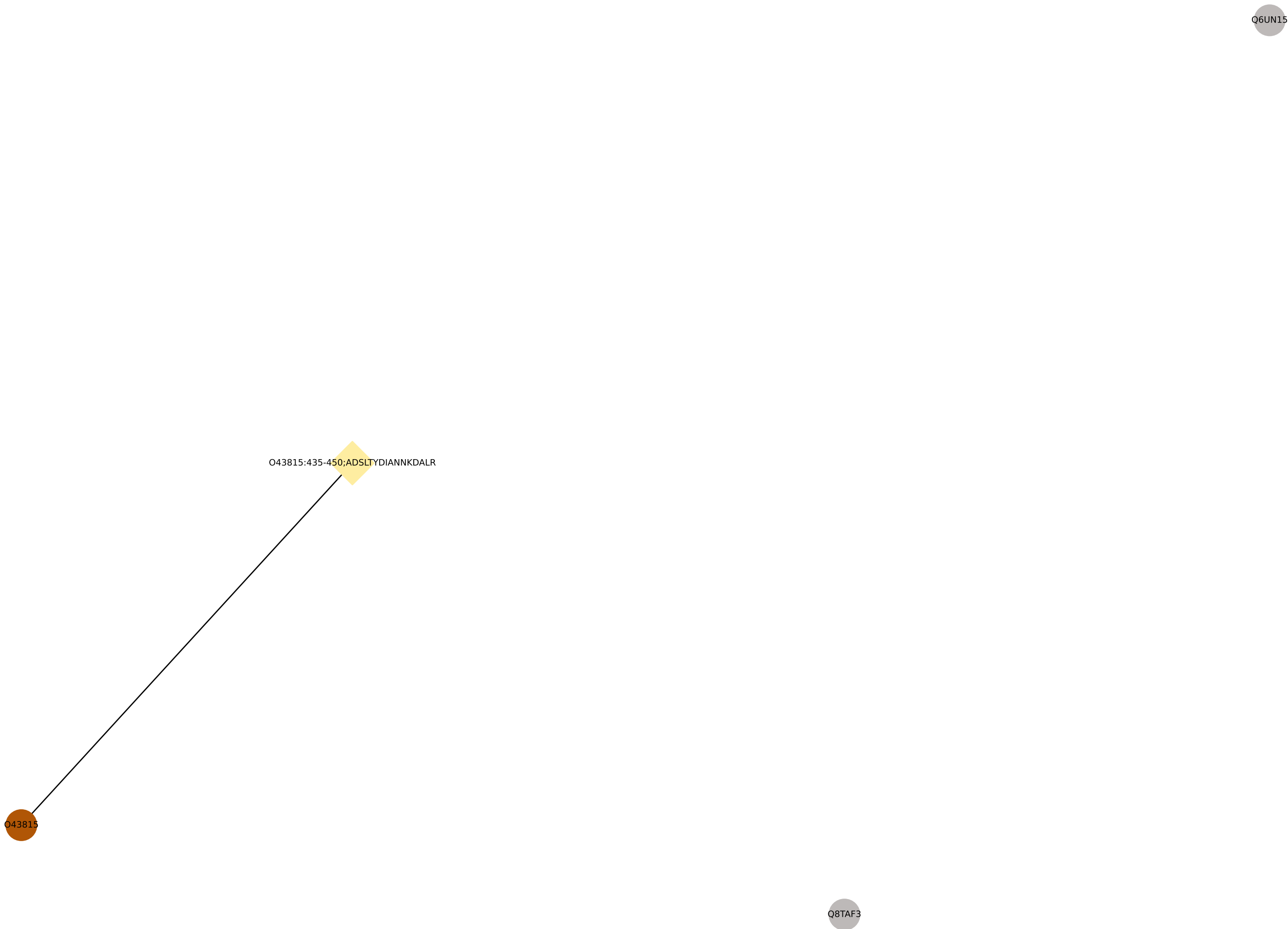


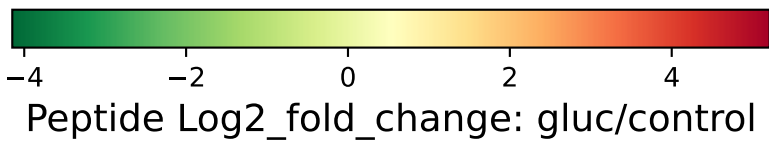
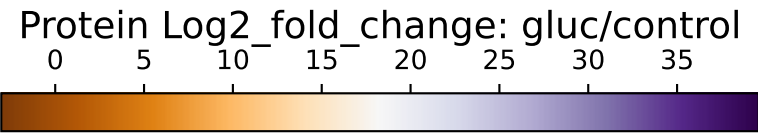
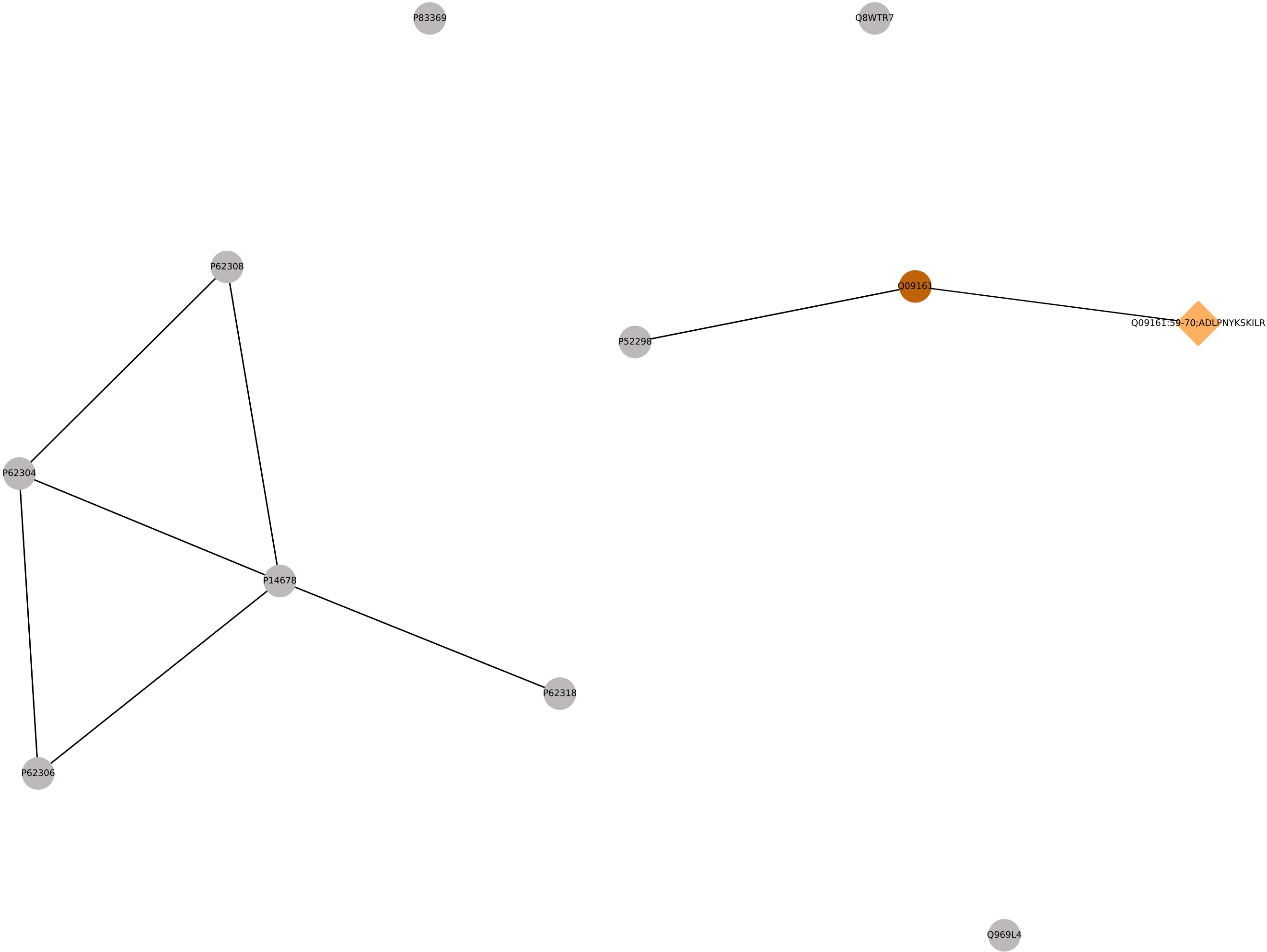
Log2\_fold\_change: gluc/control  
R-HSA-75108: Activation, myristoylation of BID and translocation to mitochondria  
p-value: 0.015117656007649471



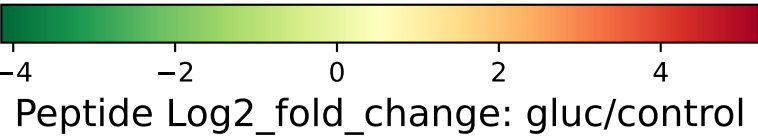
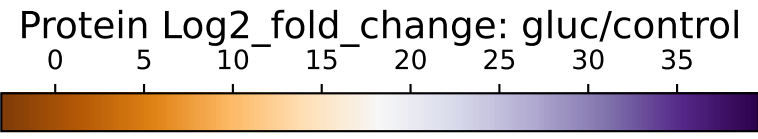
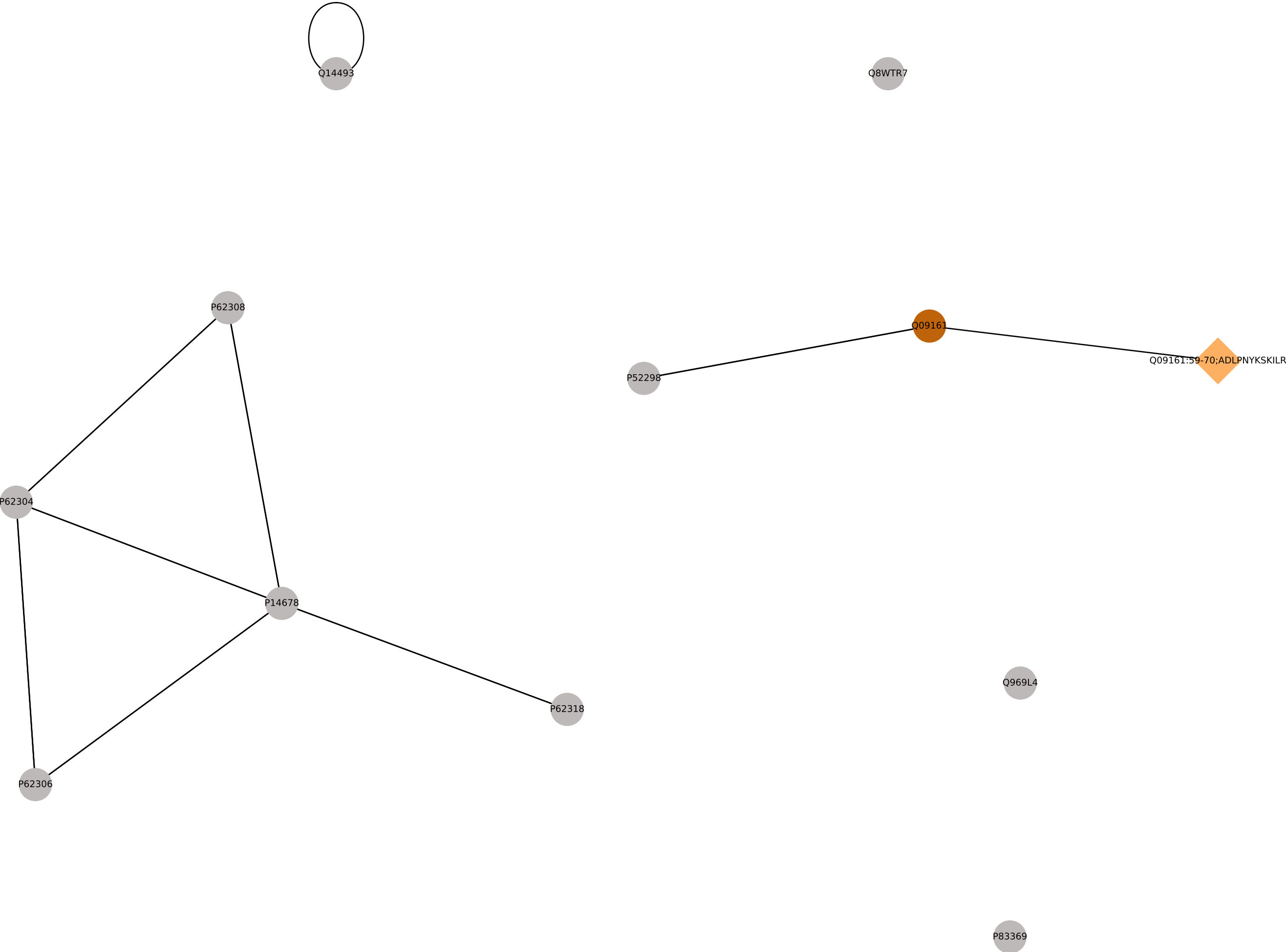
Log2\_fold\_change: gluc/control  
R-HSA-9673766: Signaling by cytosolic PDGFRA and PDGFRB fusion proteins  
p-value: 0.018861837637962453



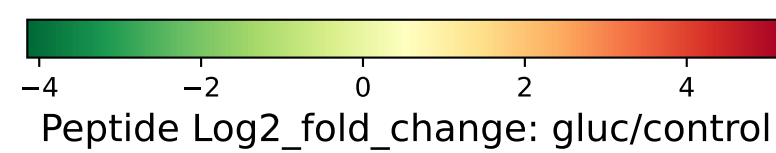
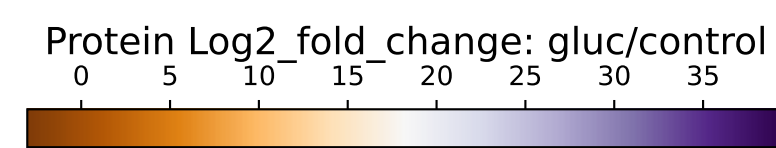
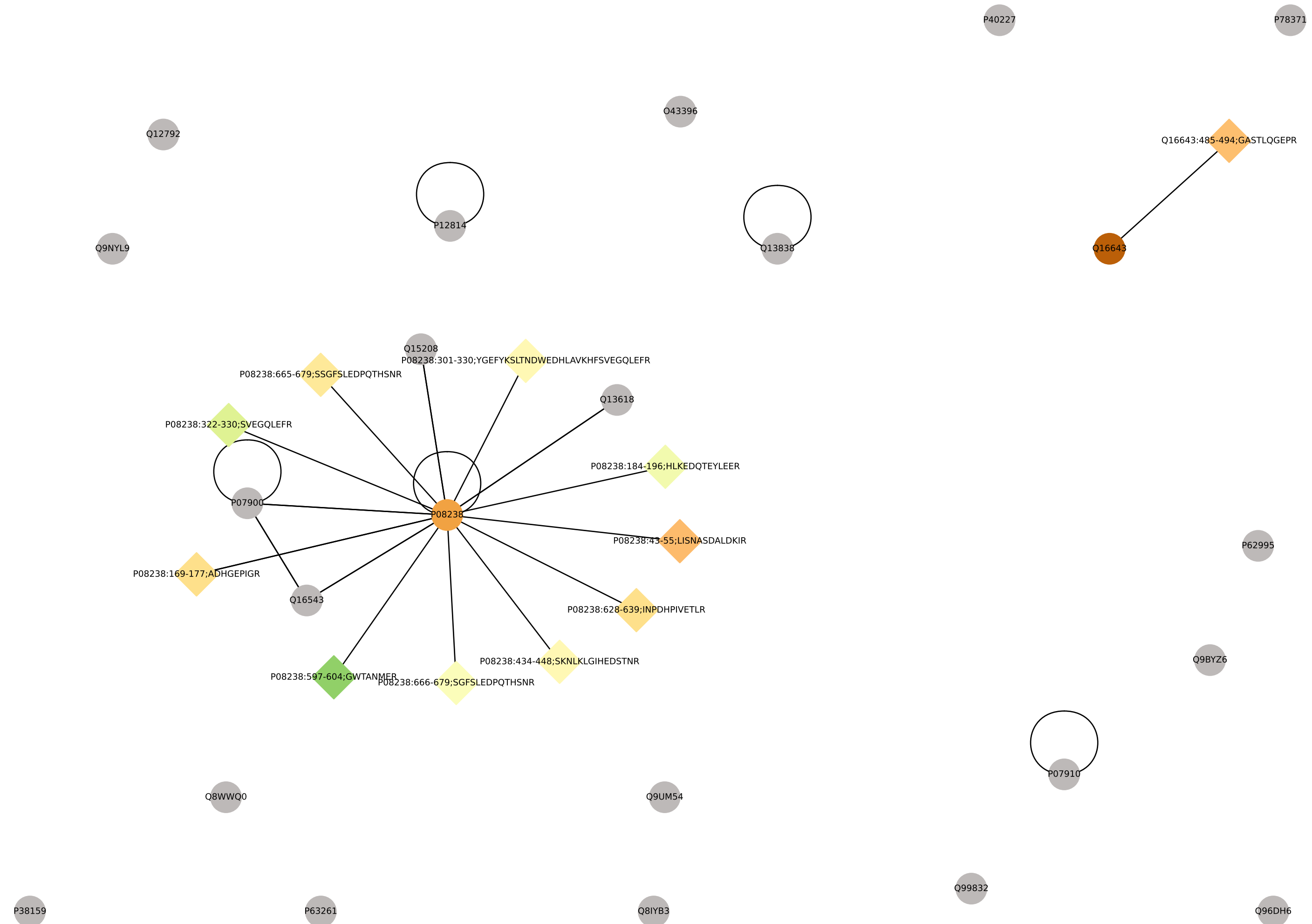
Log2\_fold\_change: gluc/control  
R-HSA-111367: SLBP independent Processing of Histone Pre-mRNAs  
p-value: 0.03737392638204595



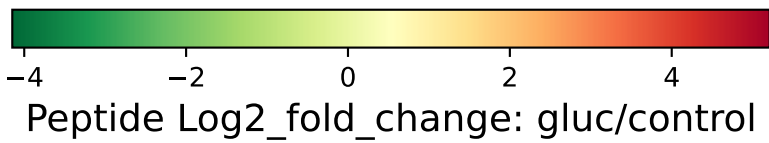
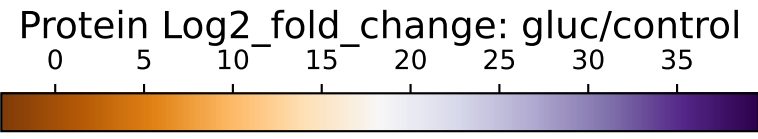
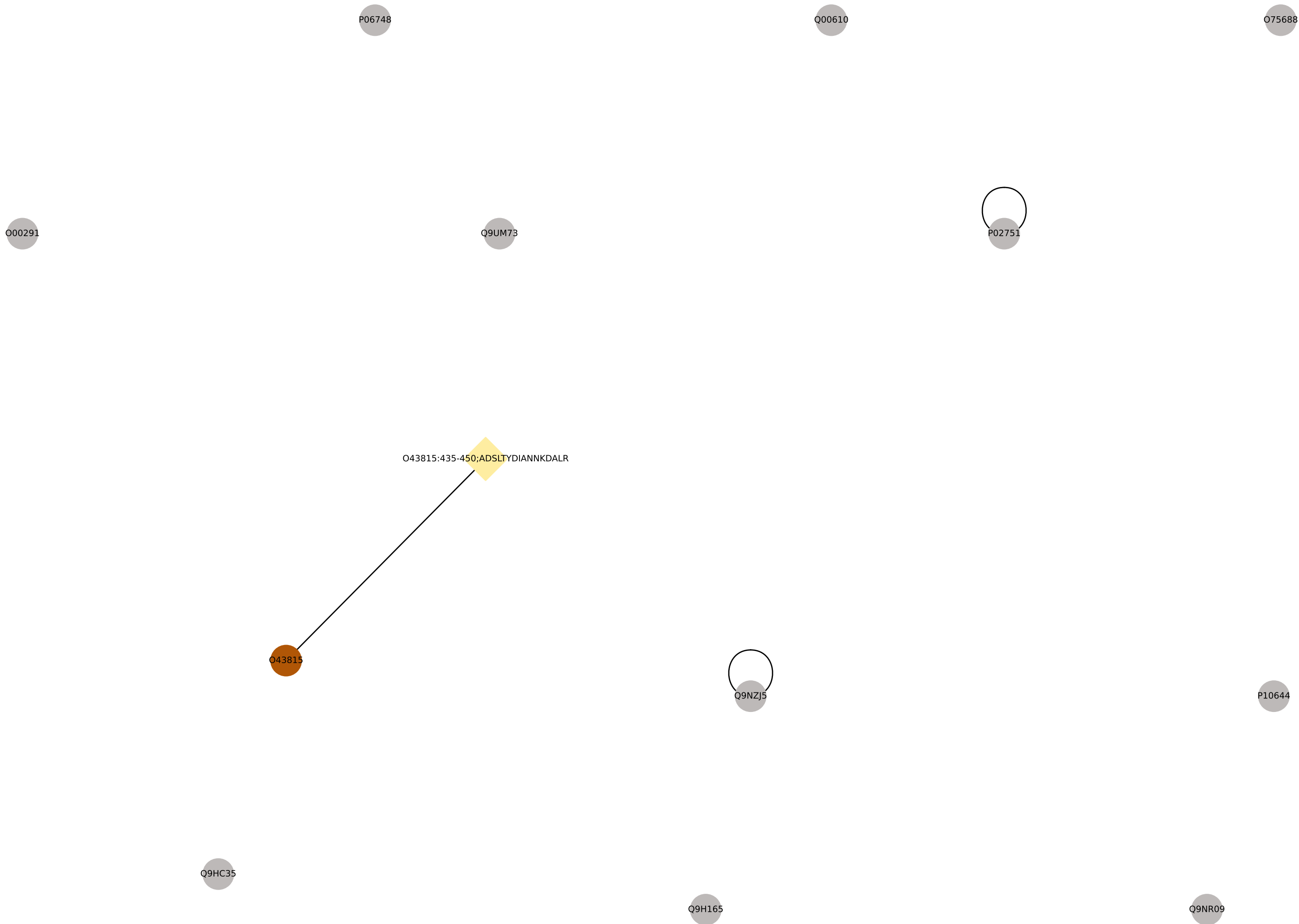
Log2\_fold\_change: gluc/control  
R-HSA-77588: SLBP Dependent Processing of Replication-Dependent Histone Pre-mRNAs  
p-value: 0.04103493715974438



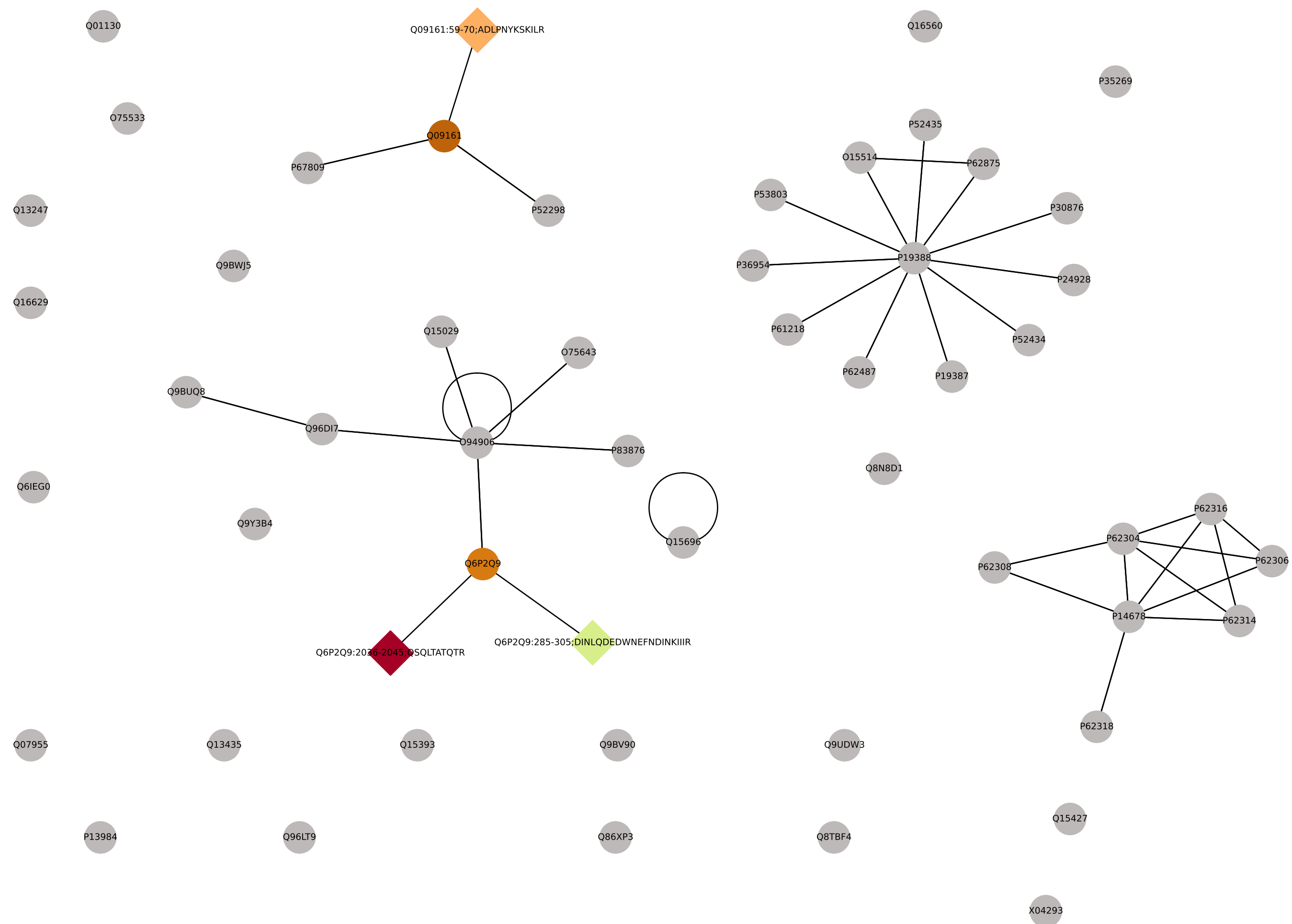
Log2\_fold\_change: gluc/control  
R-HSA-9013418: RHOBTB2 GTPase cycle  
p-value: 0.003870027935771292



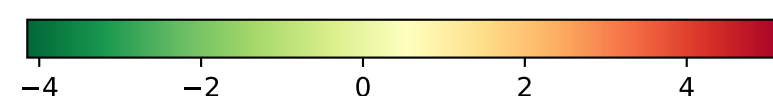
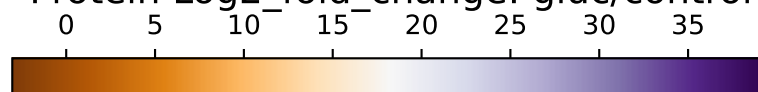
Log2\_fold\_change: gluc/control  
R-HSA-9700645: ALK mutants bind TKIs  
p-value: 0.04468226372702311



Log2\_fold\_change: gluc/control  
R-HSA-72165: mRNA Splicing - Minor Pathway  
p-value: 0.018216773752471993

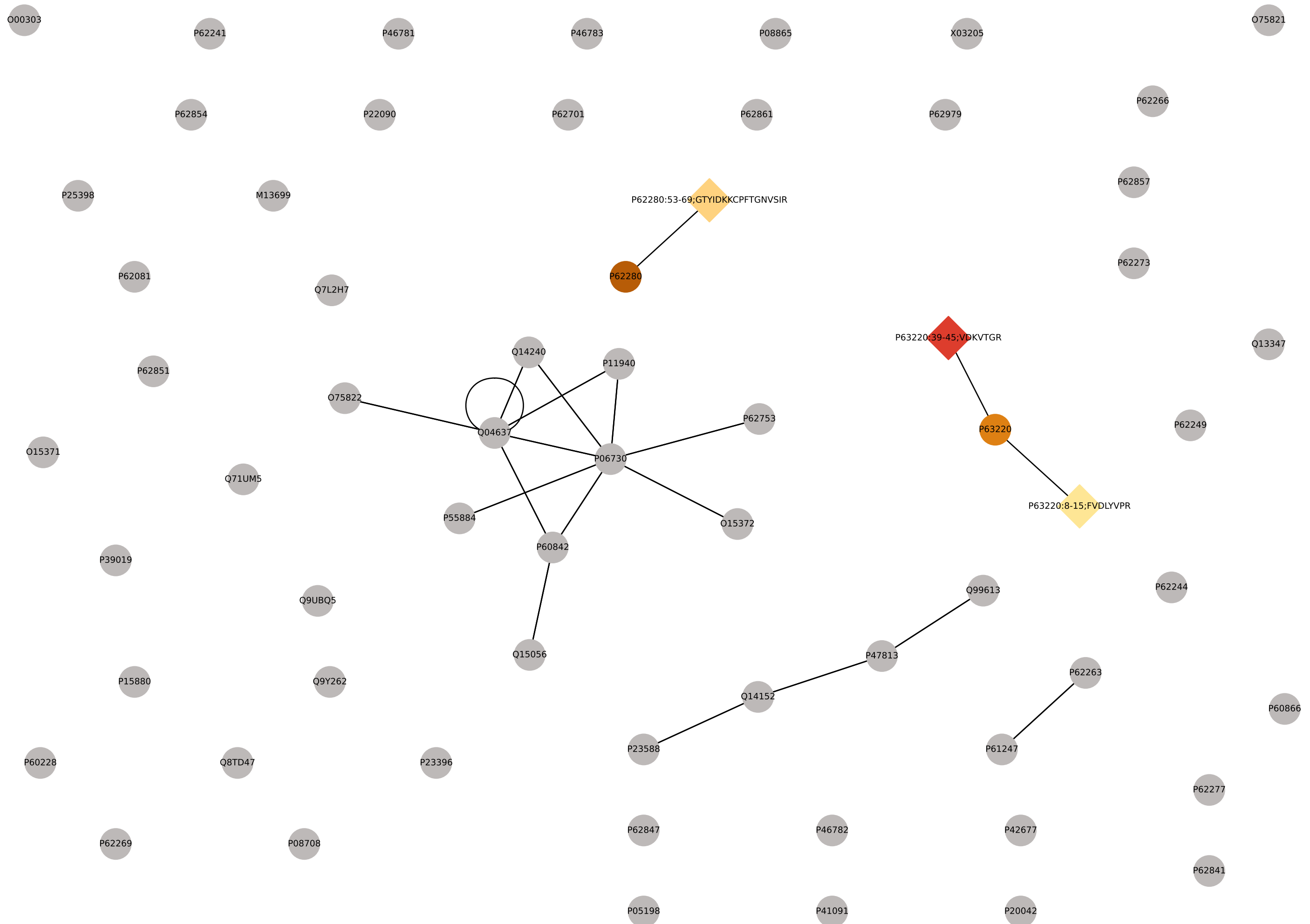


Protein Log2\_fold\_change: gluc/control



Peptide Log2\_fold\_change: gluc/control

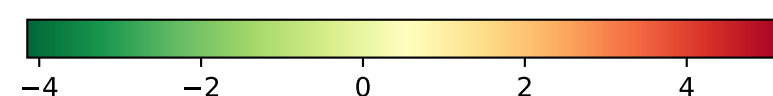
Log2\_fold\_change: gluc/control  
R-HSA-72649: Translation initiation complex formation  
p-value: 0.023554931232115828



Protein Log2\_fold\_change: gluc/control

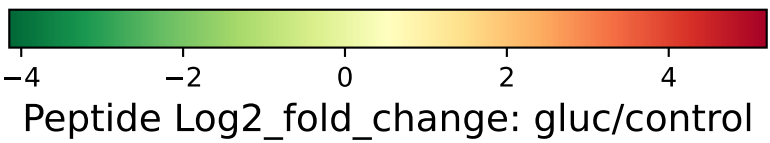
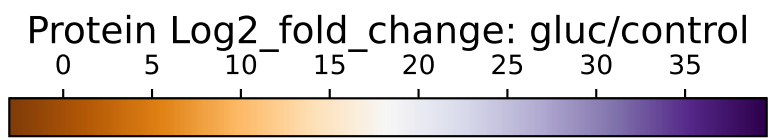
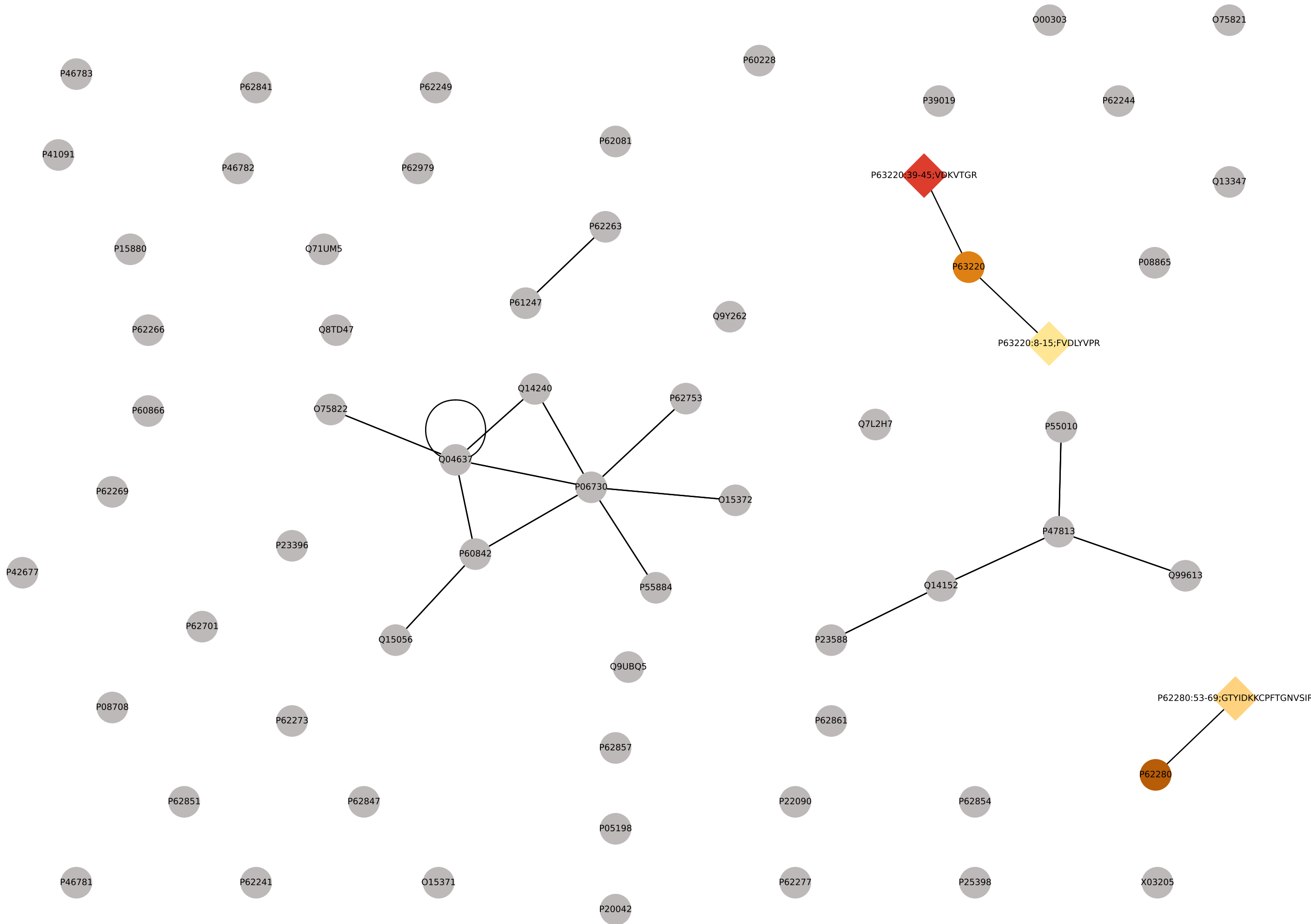


Peptide Log2\_fold\_change: gluc/control

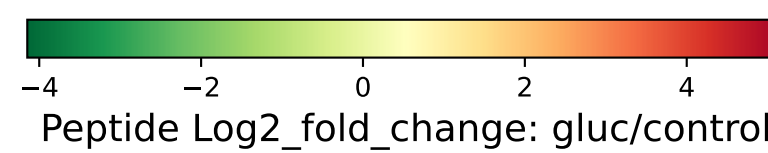
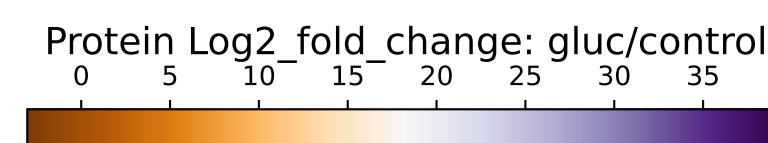
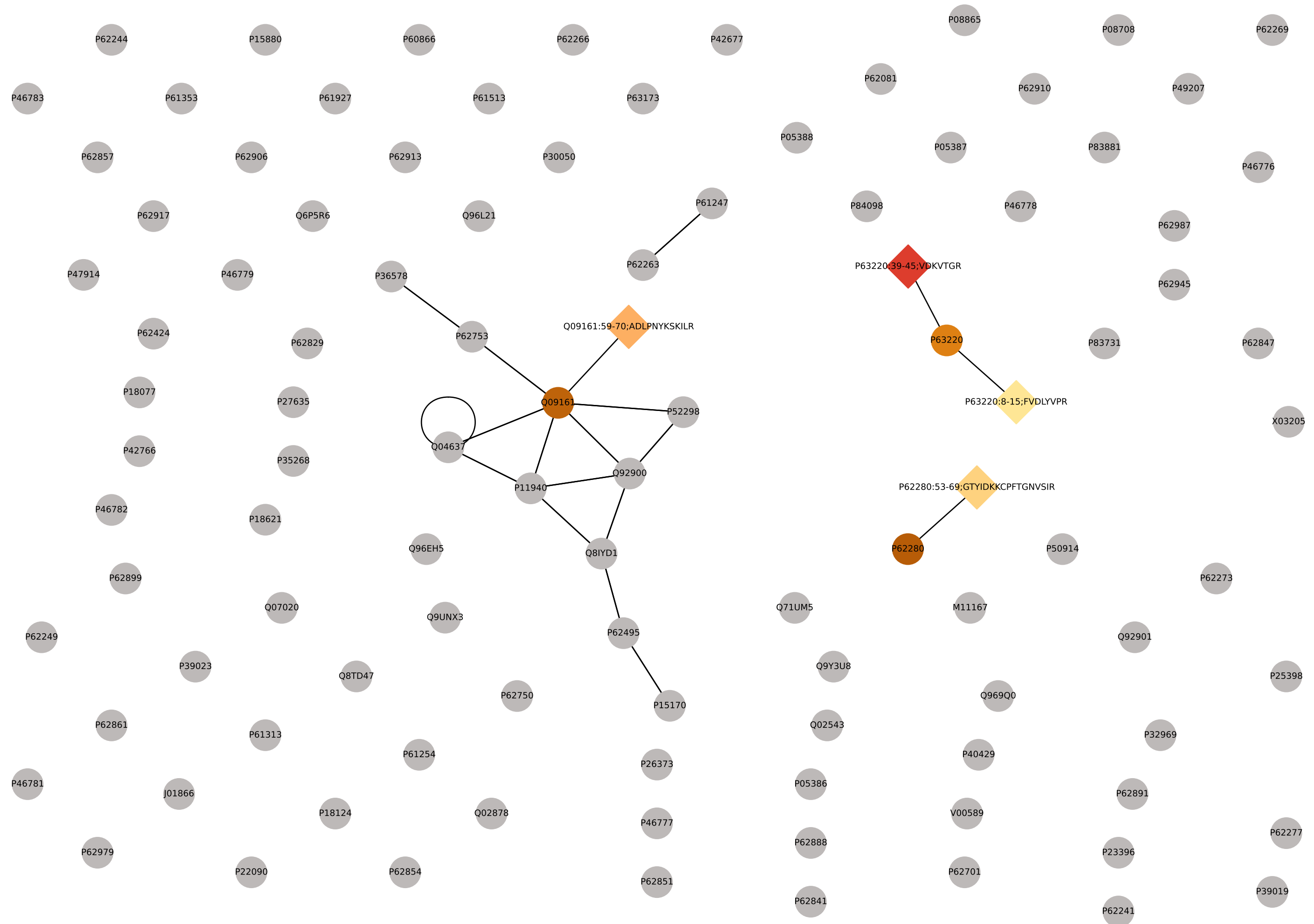




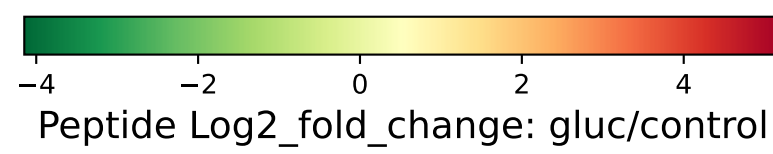
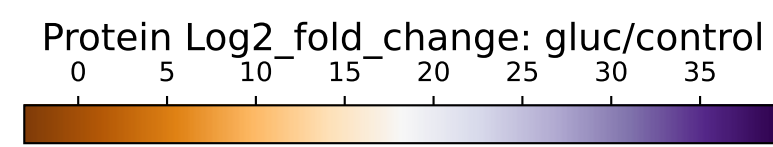
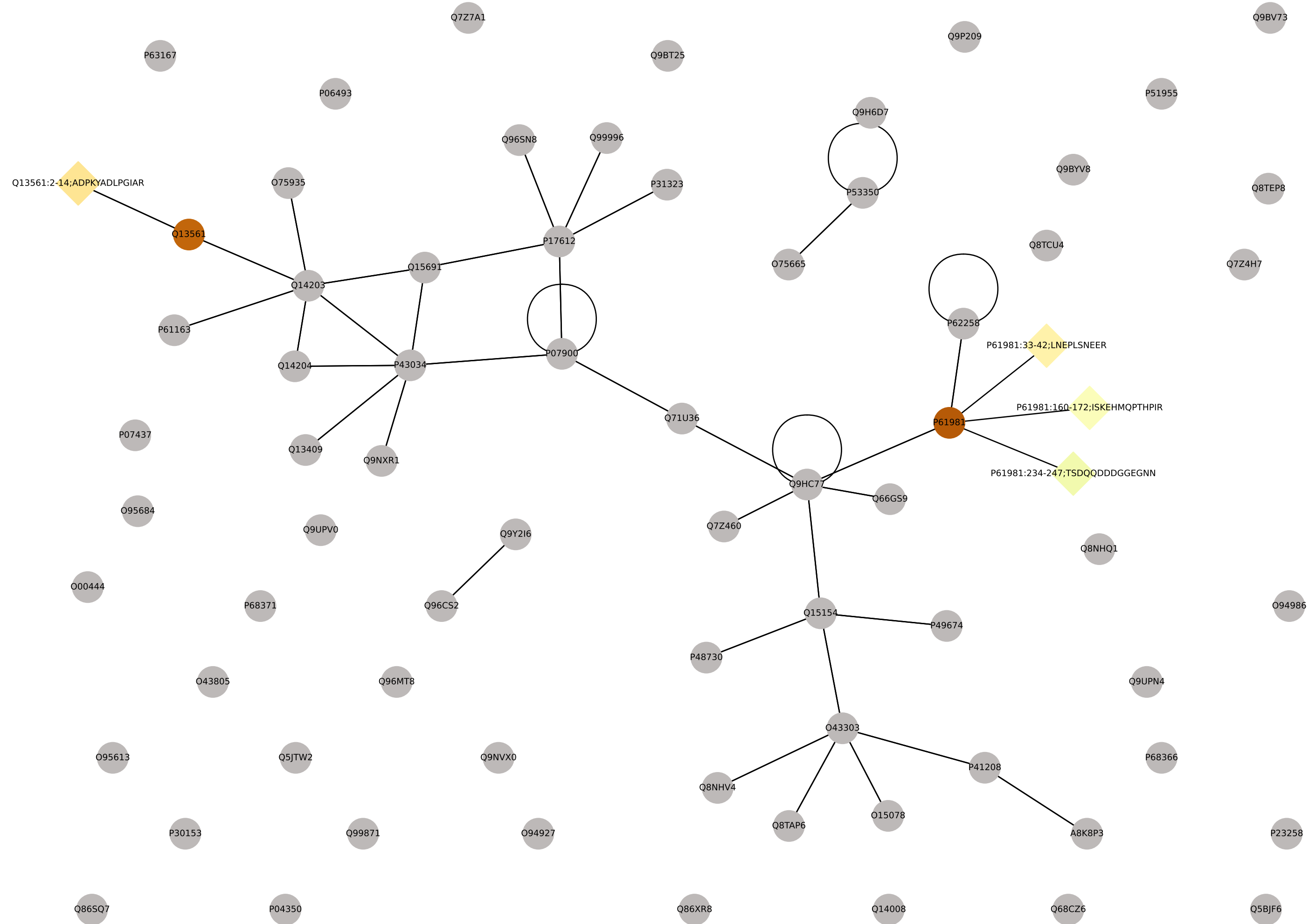
Log2\_fold\_change: gluc/control  
R-HSA-72702: Ribosomal scanning and start codon recognition  
p-value: 0.024978417082091475



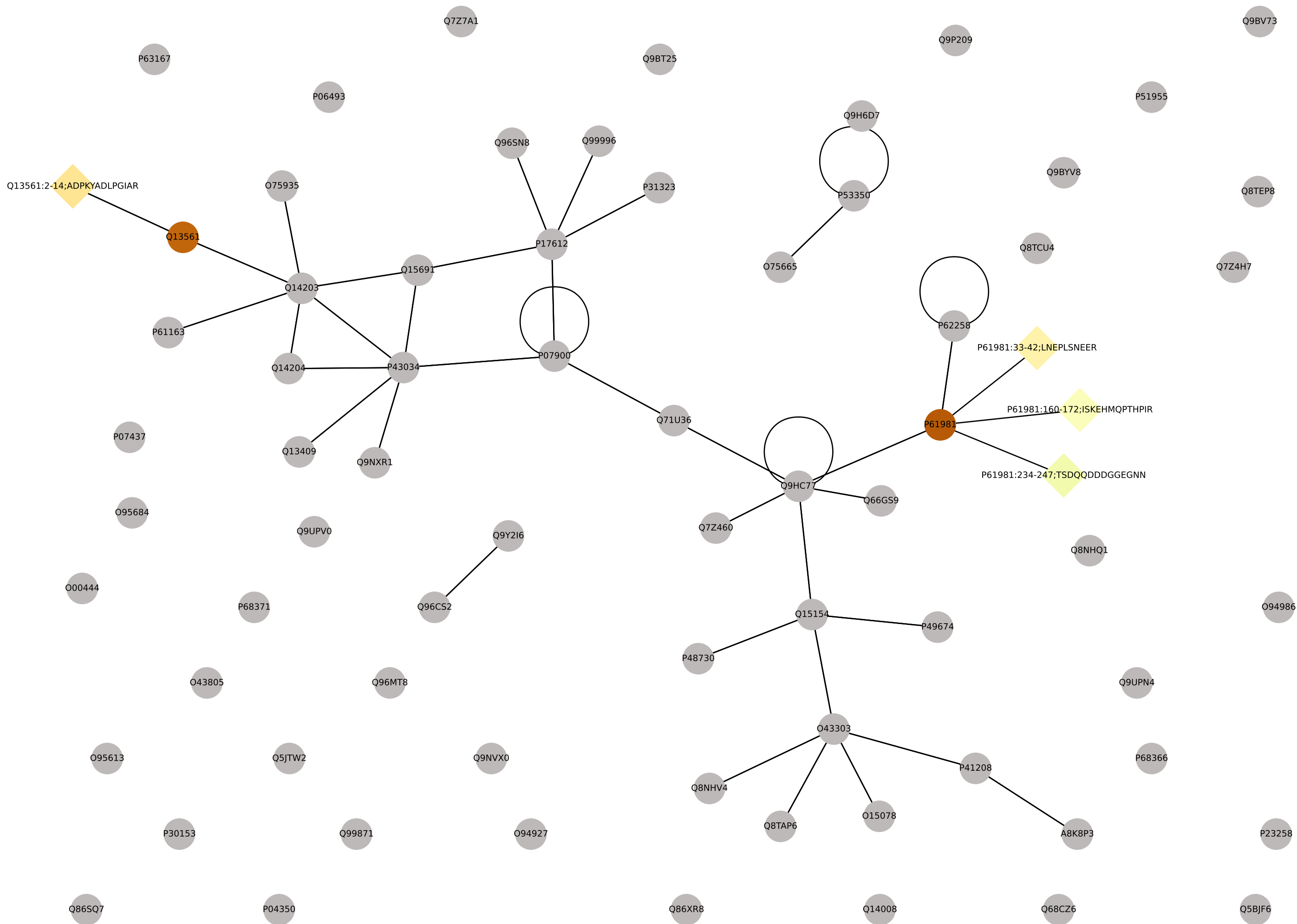
Log2\_fold\_change: gluc/control  
R-HSA-975956: Nonsense Mediated Decay (NMD) independent of the Exon Junction Complex (EJC)  
p-value: 0.006855569728165922



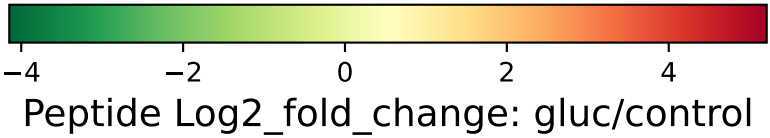
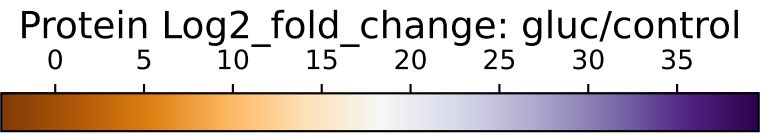
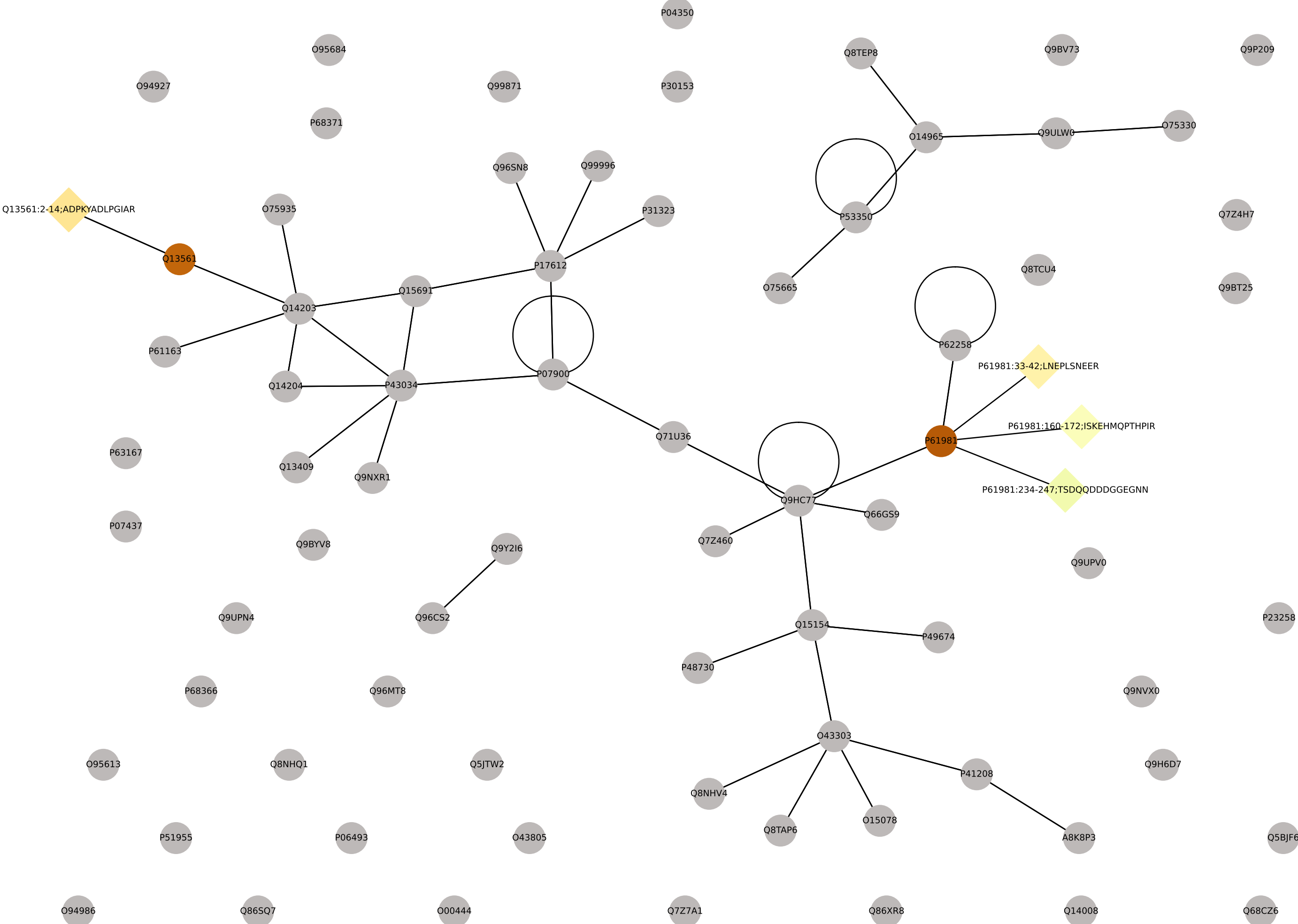
Log2\_fold\_change: gluc/control  
R-HSA-380284: Loss of proteins required for interphase microtubule organization from the centrosome  
p-value: 0.030227589686308676



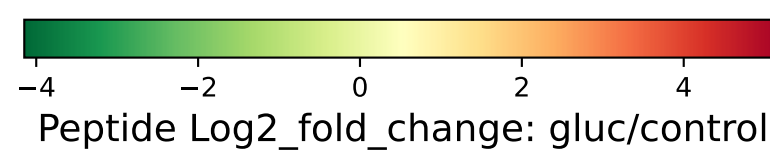
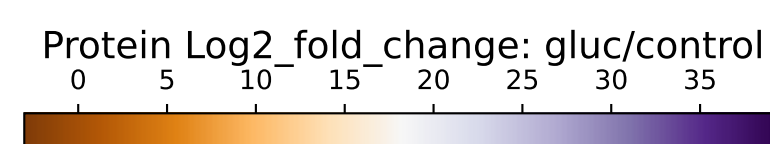
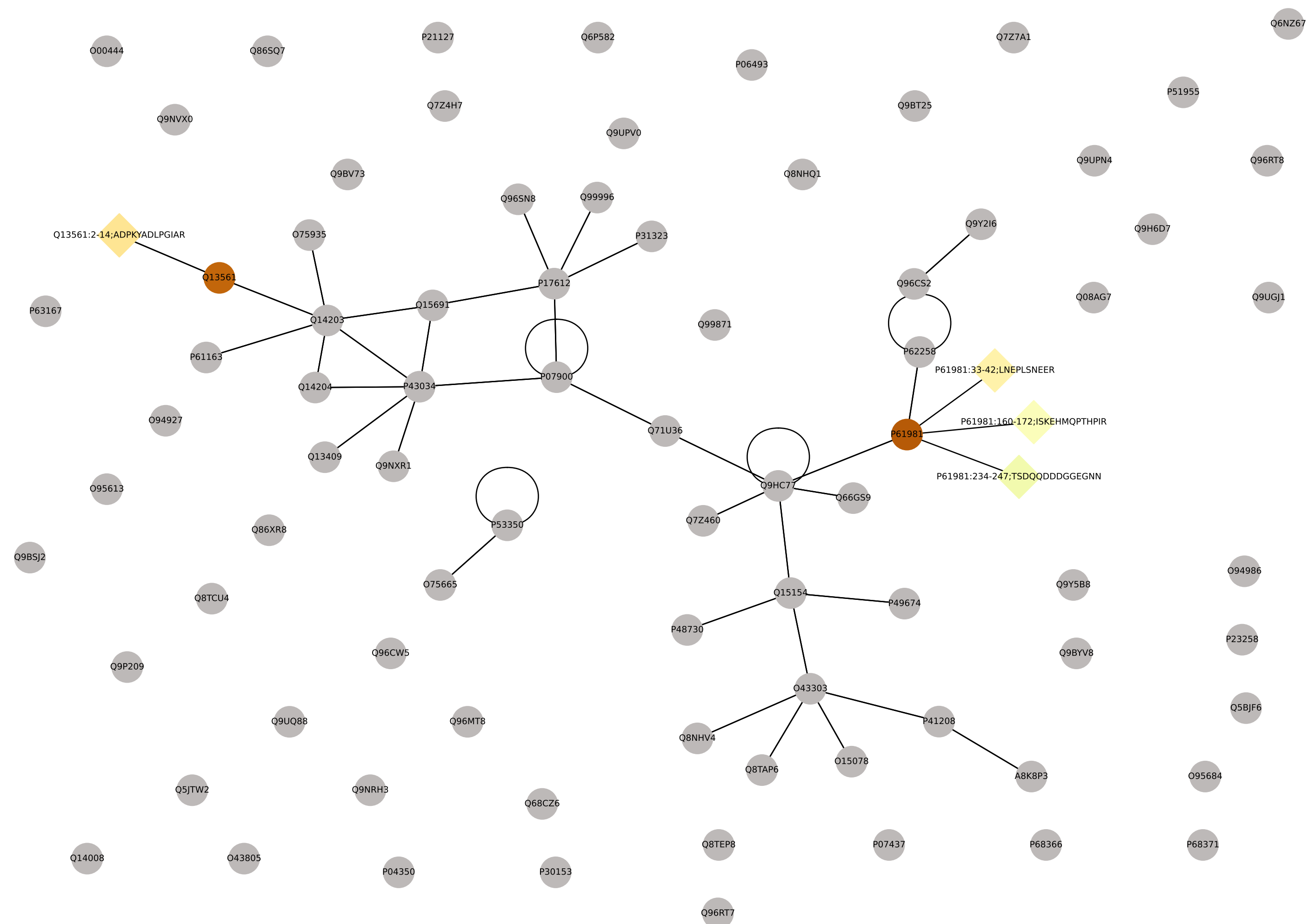
Log2\_fold\_change: gluc/control  
R-HSA-380259: Loss of Nlp from mitotic centrosomes  
p-value: 0.030227589686308676



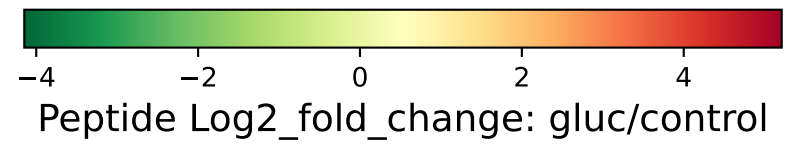
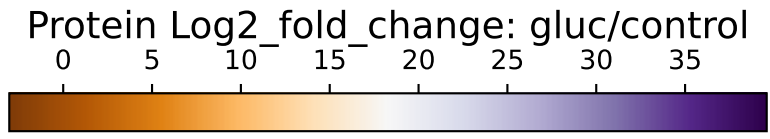
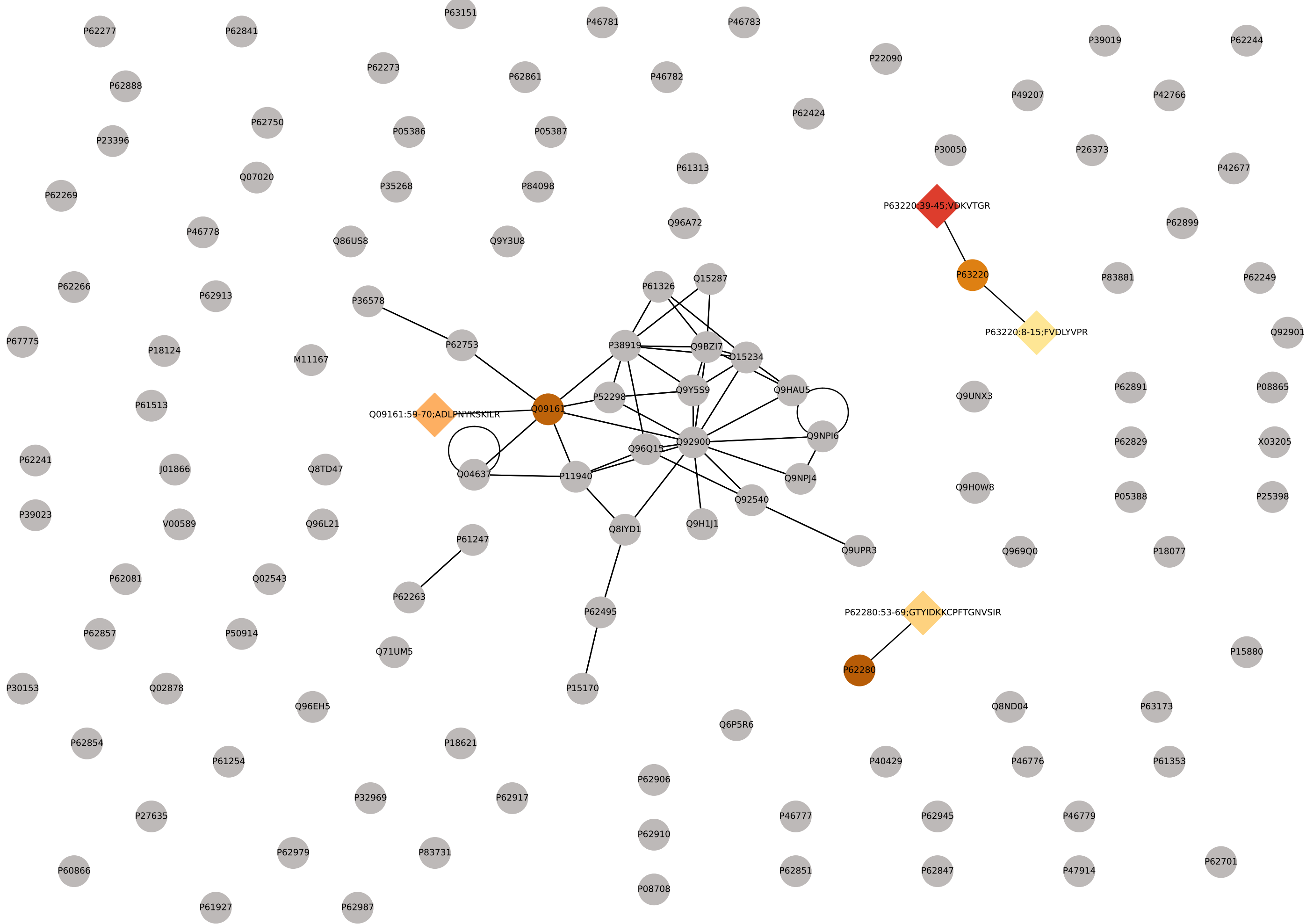
Log2\_fold\_change: gluc/control  
R-HSA-8854518: AURKA Activation by TPX2  
p-value: 0.032600049302680456



Log2\_fold\_change: gluc/control  
R-HSA-380270: Recruitment of mitotic centrosome proteins and complexes  
p-value: 0.0384084717376354

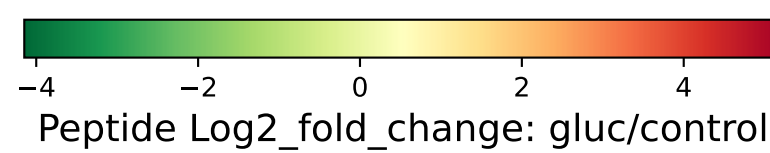
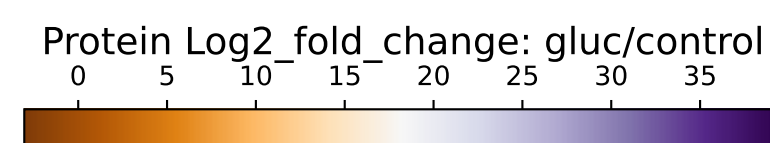
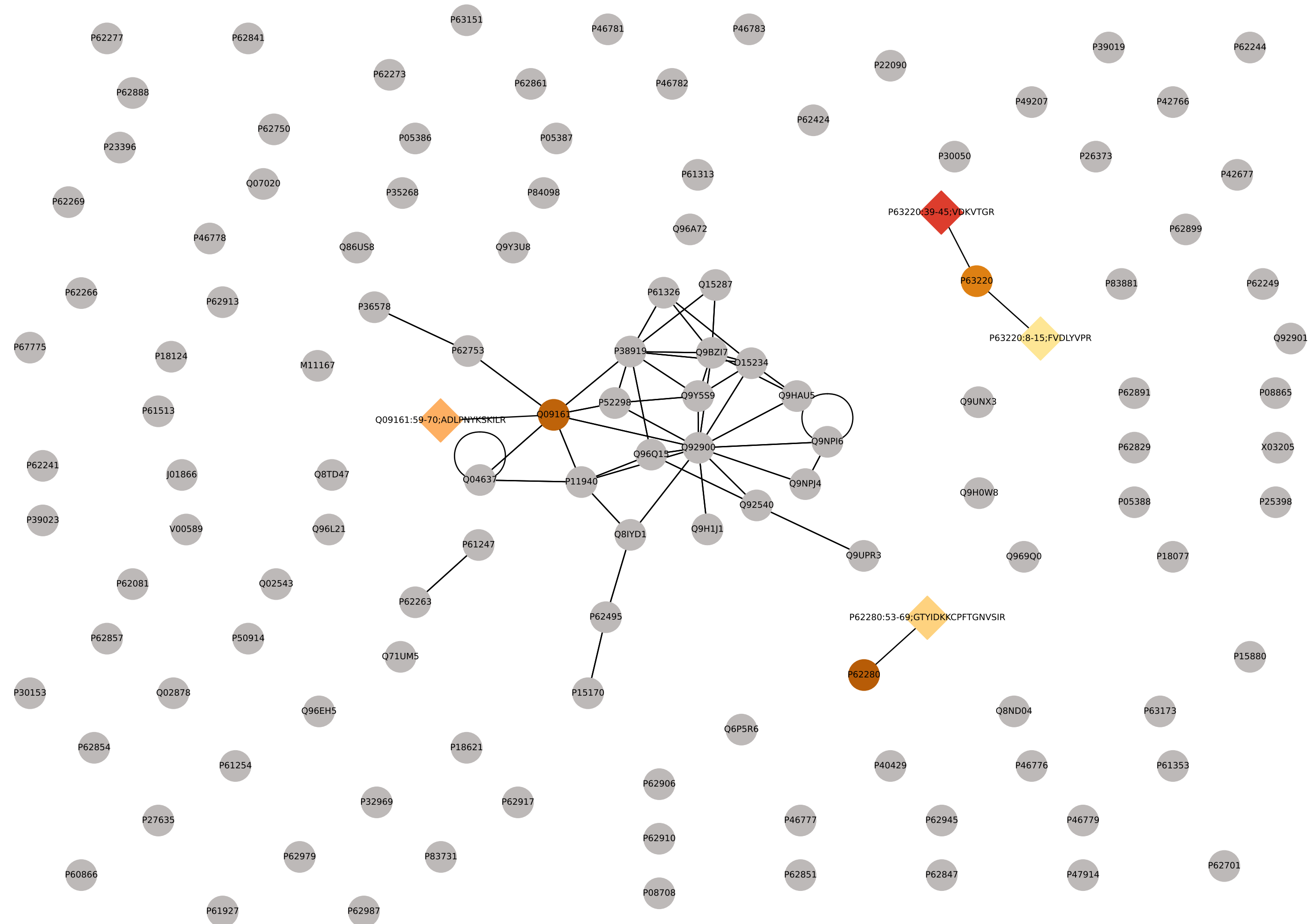


Log2\_fold\_change: gluc/control  
R-HSA-927802: Nonsense-Mediated Decay (NMD)  
p-value: 0.011931898374331729



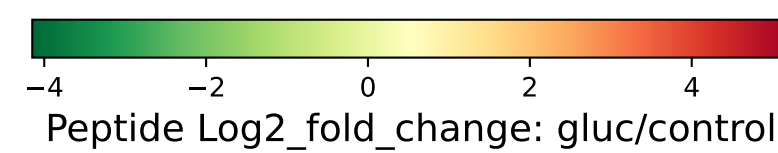
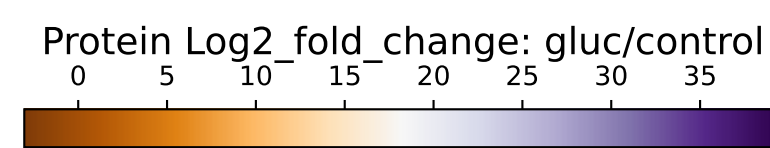
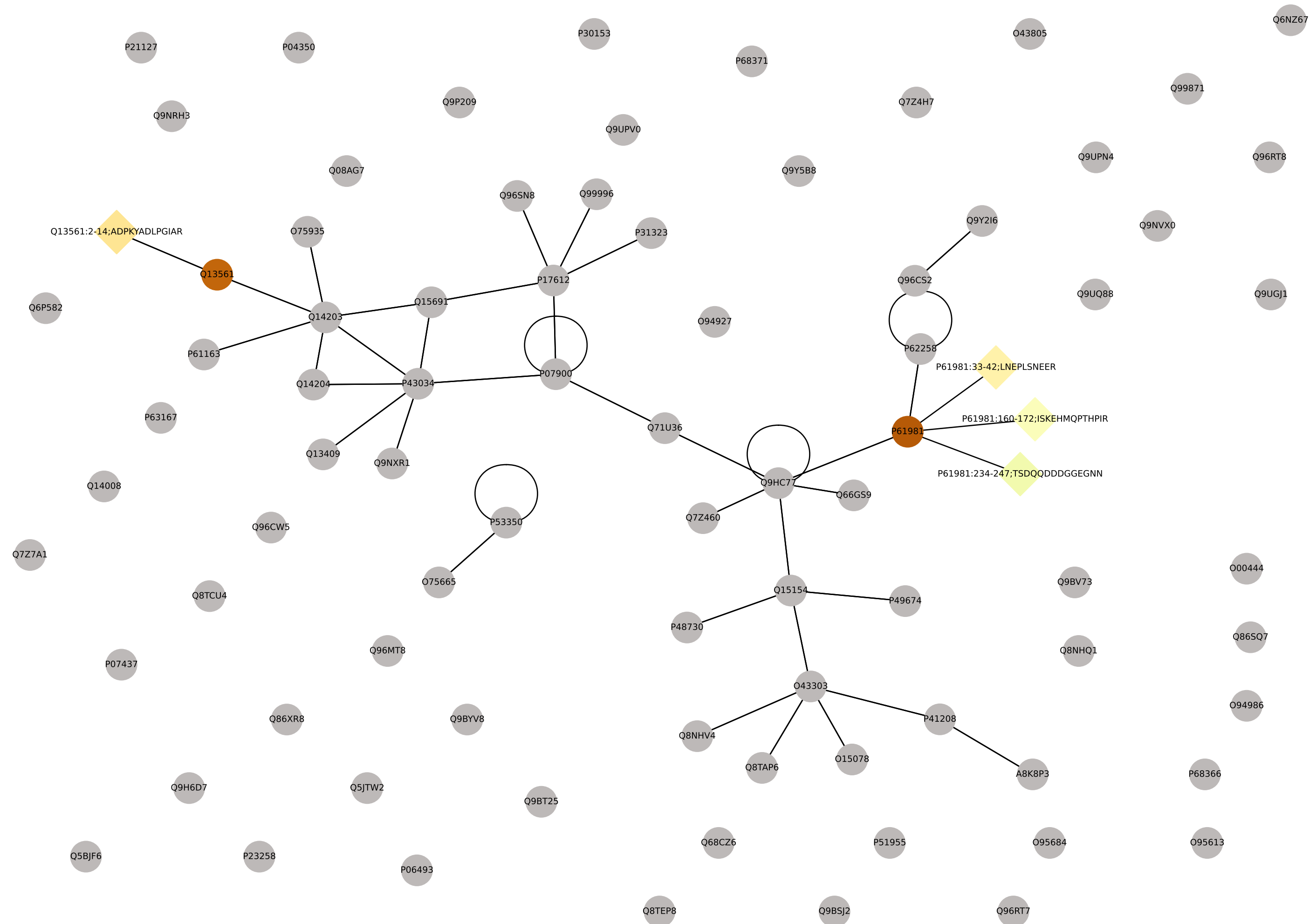


Log2\_fold\_change: gluc/control  
R-HSA-975957: Nonsense Mediated Decay (NMD) enhanced by the Exon Junction Complex (EJC)  
p-value: 0.011931898374331729

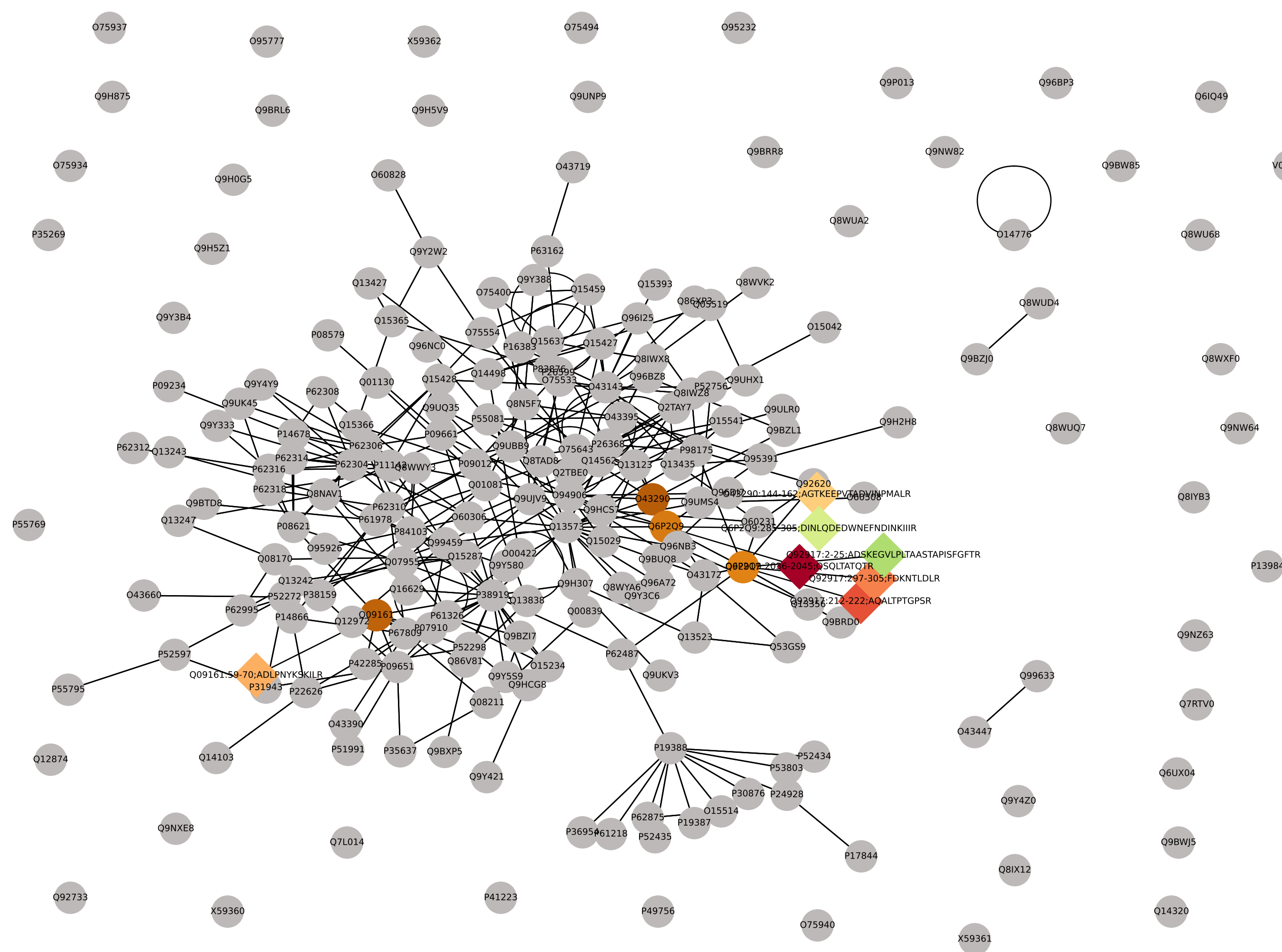




Log2\_fold\_change: gluc/control  
R-HSA-380287: Centrosome maturation  
p-value: 0.04013568745556928



Log2\_fold\_change: gluc/control  
R-HSA-72163: mRNA Splicing - Major Pathway  
p-value: 0.008907459881584257



Log2\_fold\_change: gluc/control  
R-HSA-72172: mRNA Splicing  
p-value: 0.01056574513476305

