Log2_fold_change: gluc/control
R-HSA-75108: Activation, myristolyation of BID and translocation to mitochondria
p-value: 0.015117656007649471

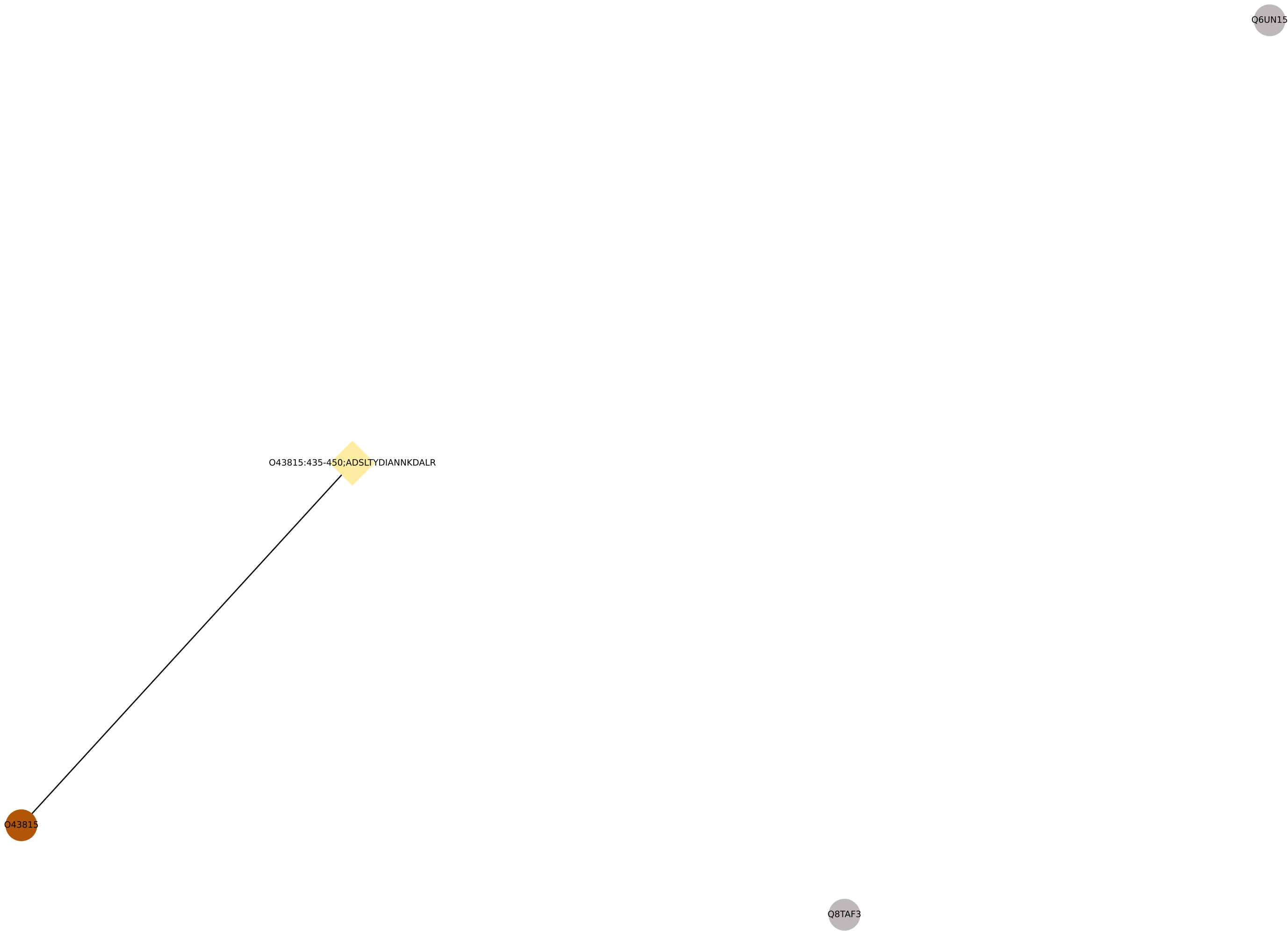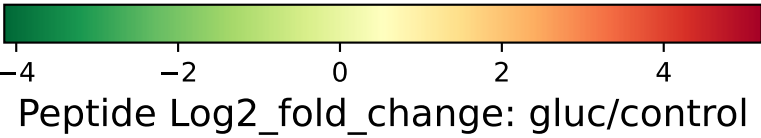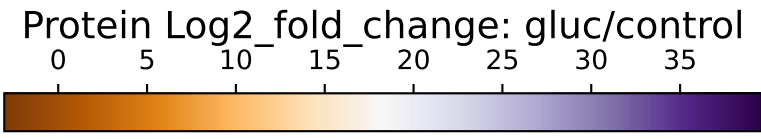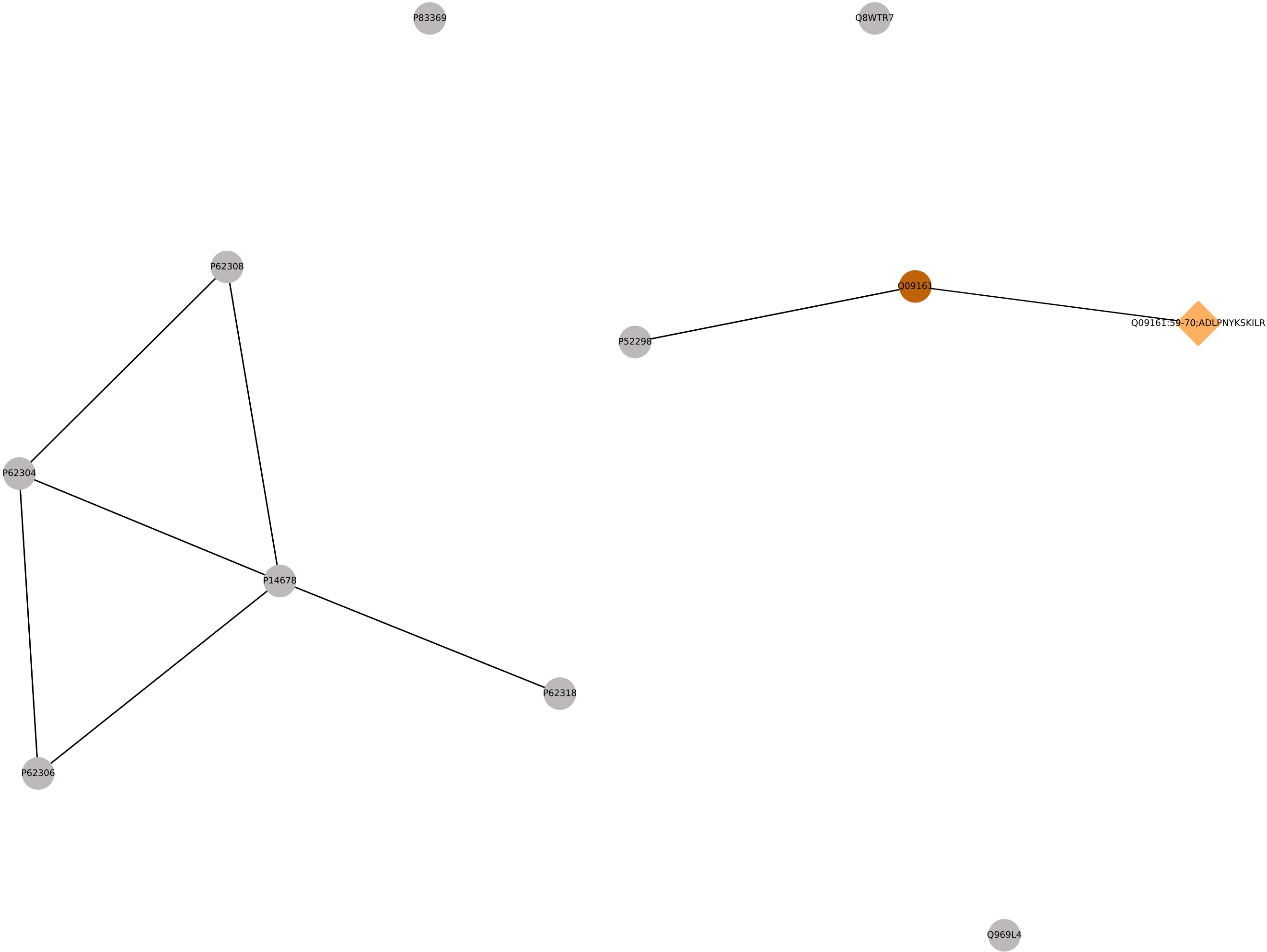Protein Log2_fold_change: gluc/control
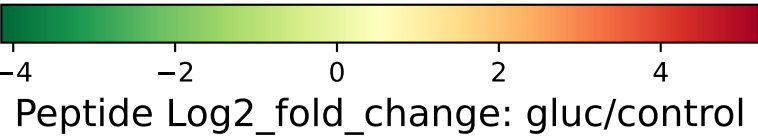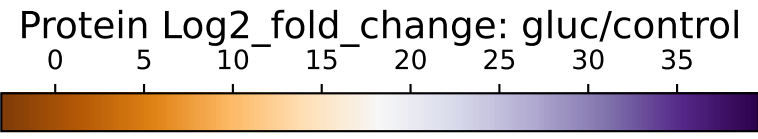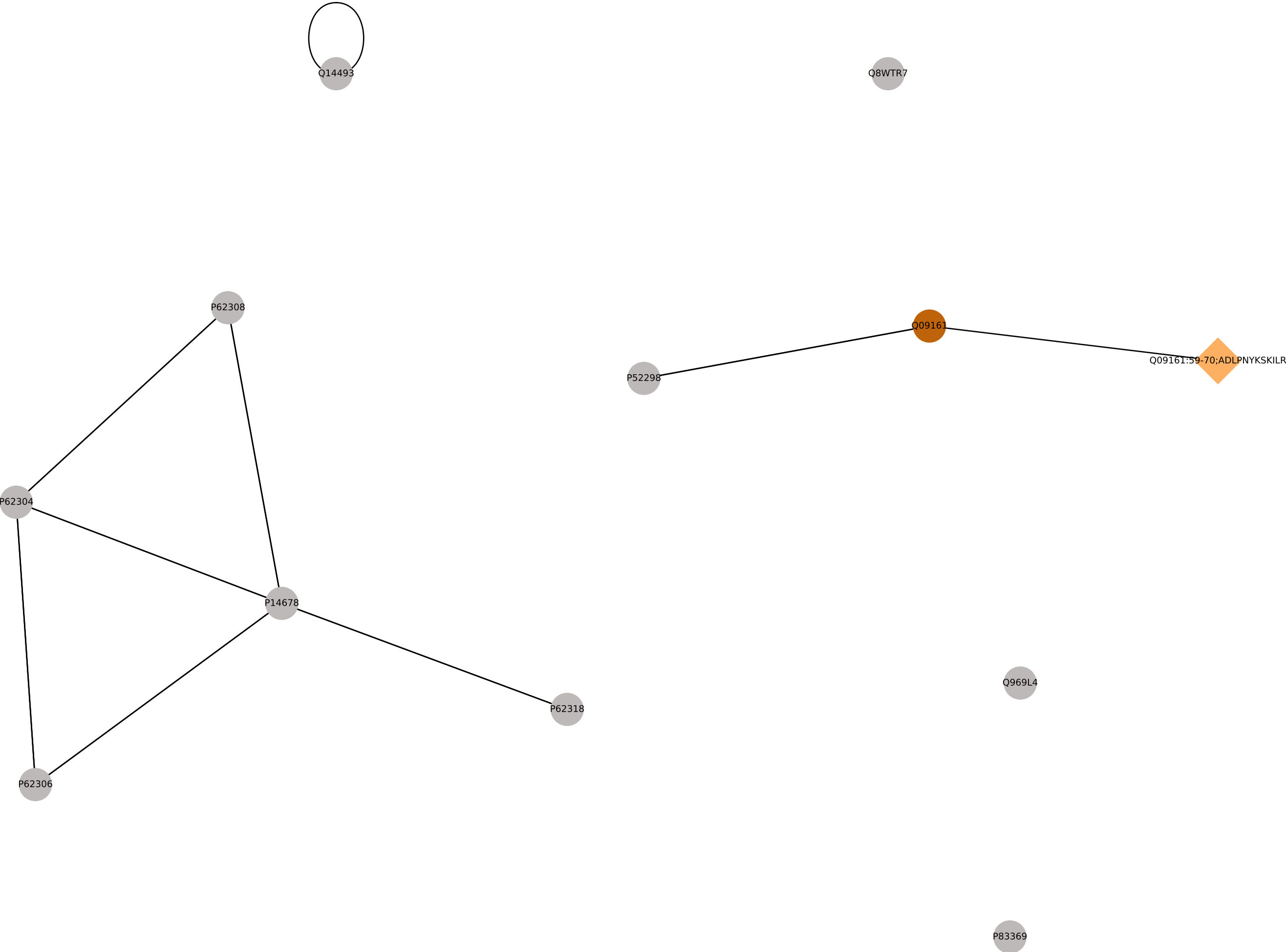
Peptide Log2_fold_change: gluc/control
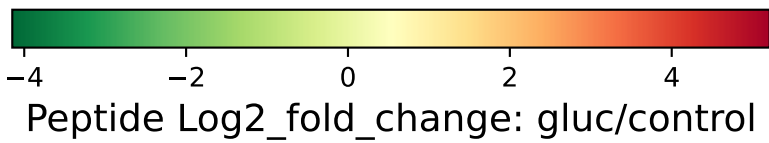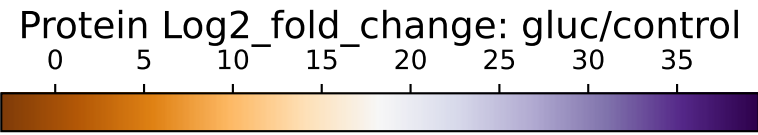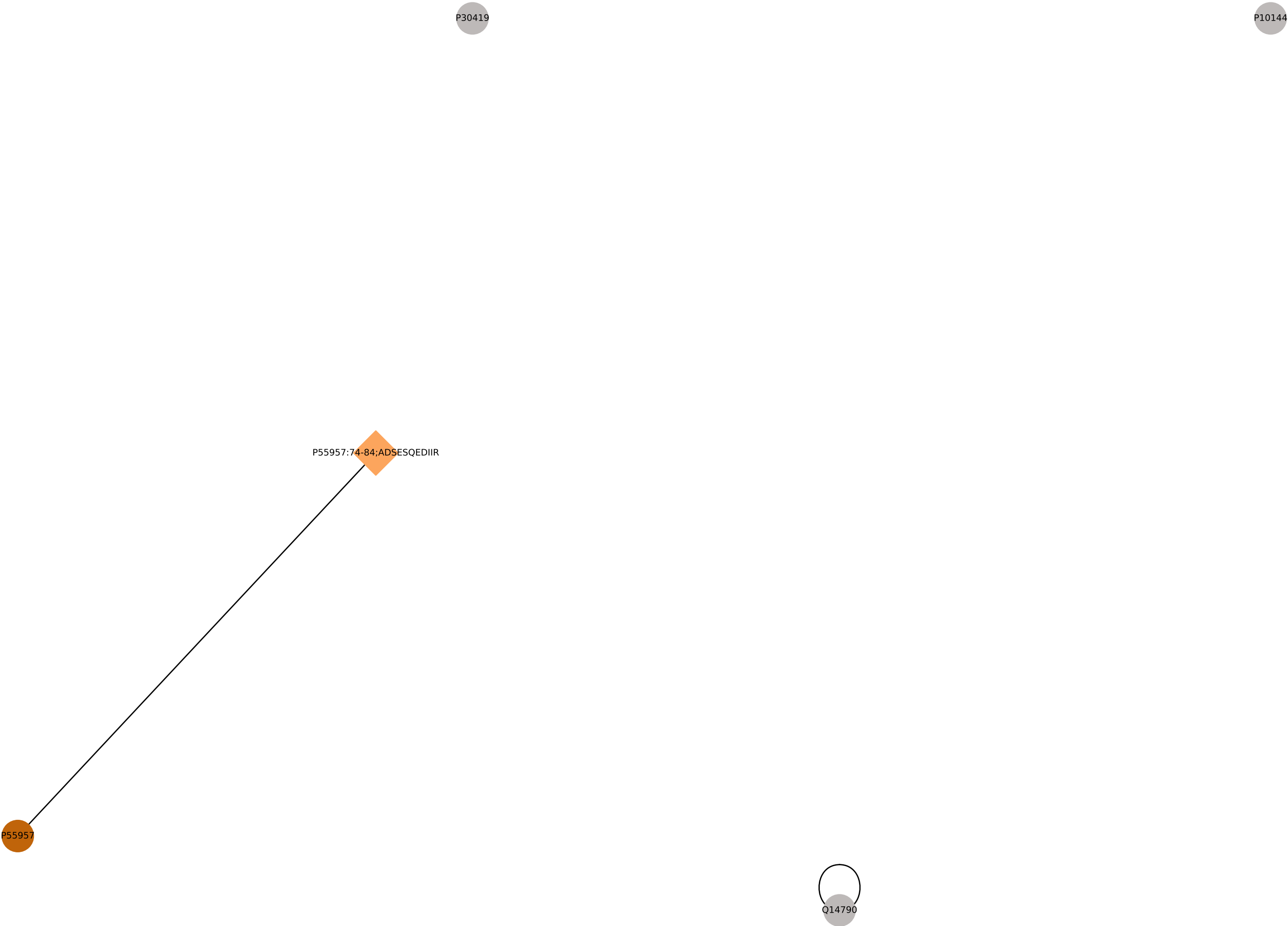
Log2_fold_change: gluc/control
R-HSA-9673766: Signaling by cytosolic PDGFRA and PDGFRB fusion proteins
p-value: 0.018861837637962453

Q6UN15

O43815:435-450;ADSLTYDIANNKDALR

O43815

Q8TAF3

Protein Log2_fold_change: gluc/control

0   5   10   15   20   25   30   35

Peptide Log2_fold_change: gluc/control

-4   -2   0   2   4
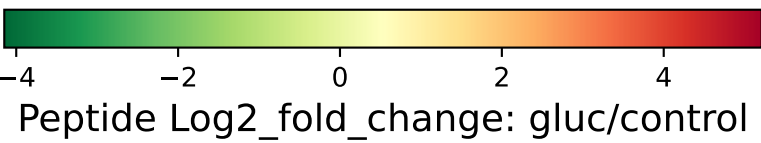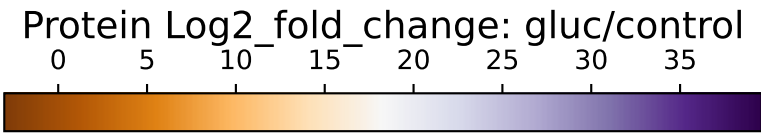
Log2_fold_change: gluc/control
R-HSA-111367: SLBP independent Processing of Histone Pre-mRNAs
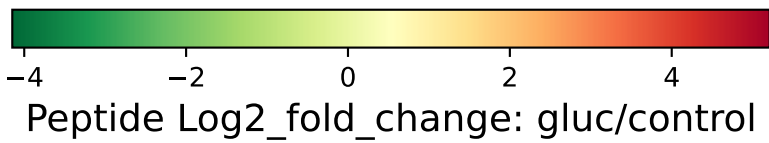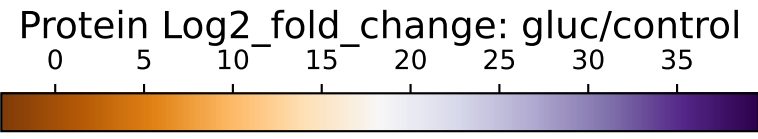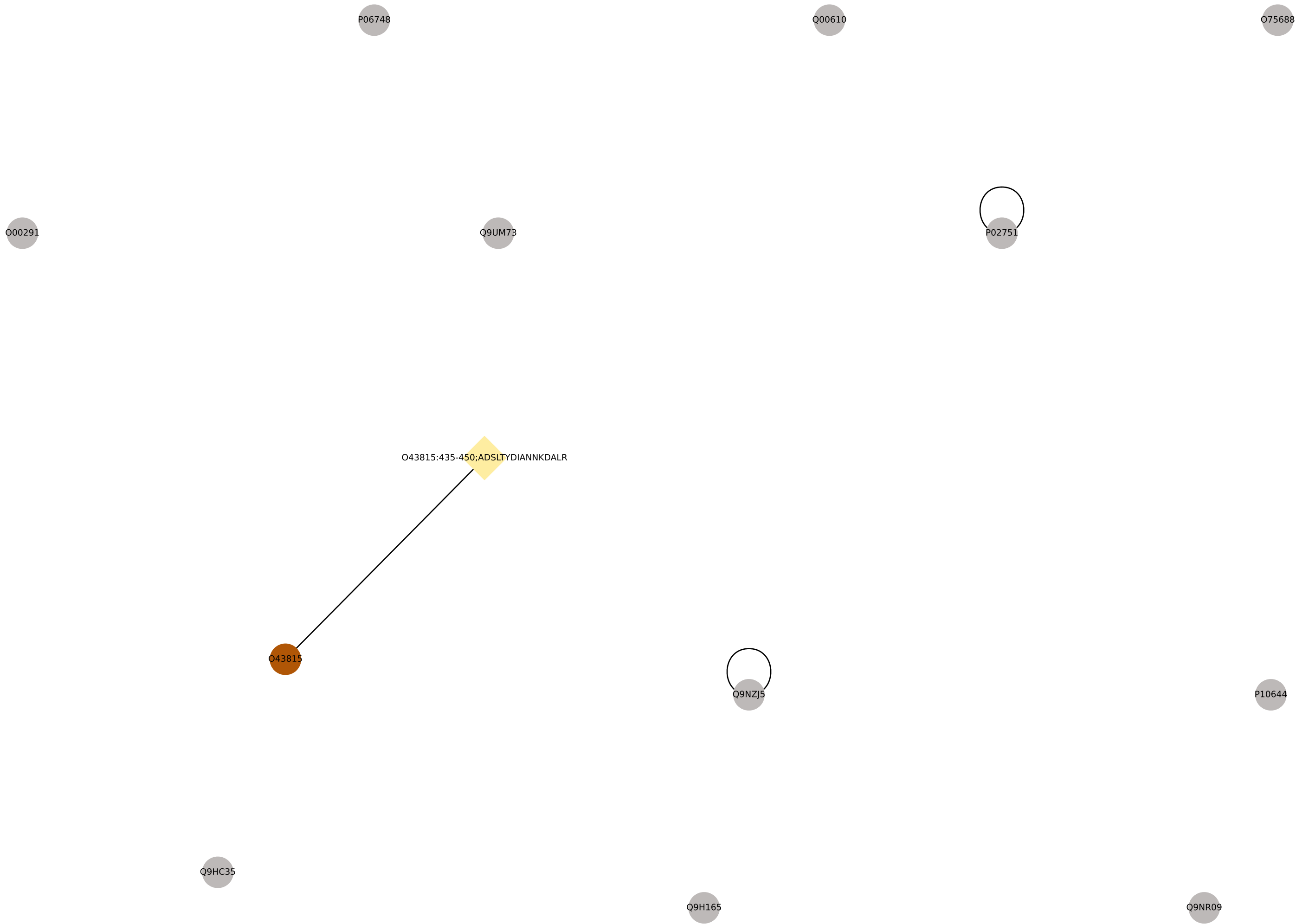p-value: 0.03737392638204595

Log2_fold_change: gluc/control
R-HSA-77588: SLBP Dependent Processing of Replication-Dependent Histone Pre-mRNAs
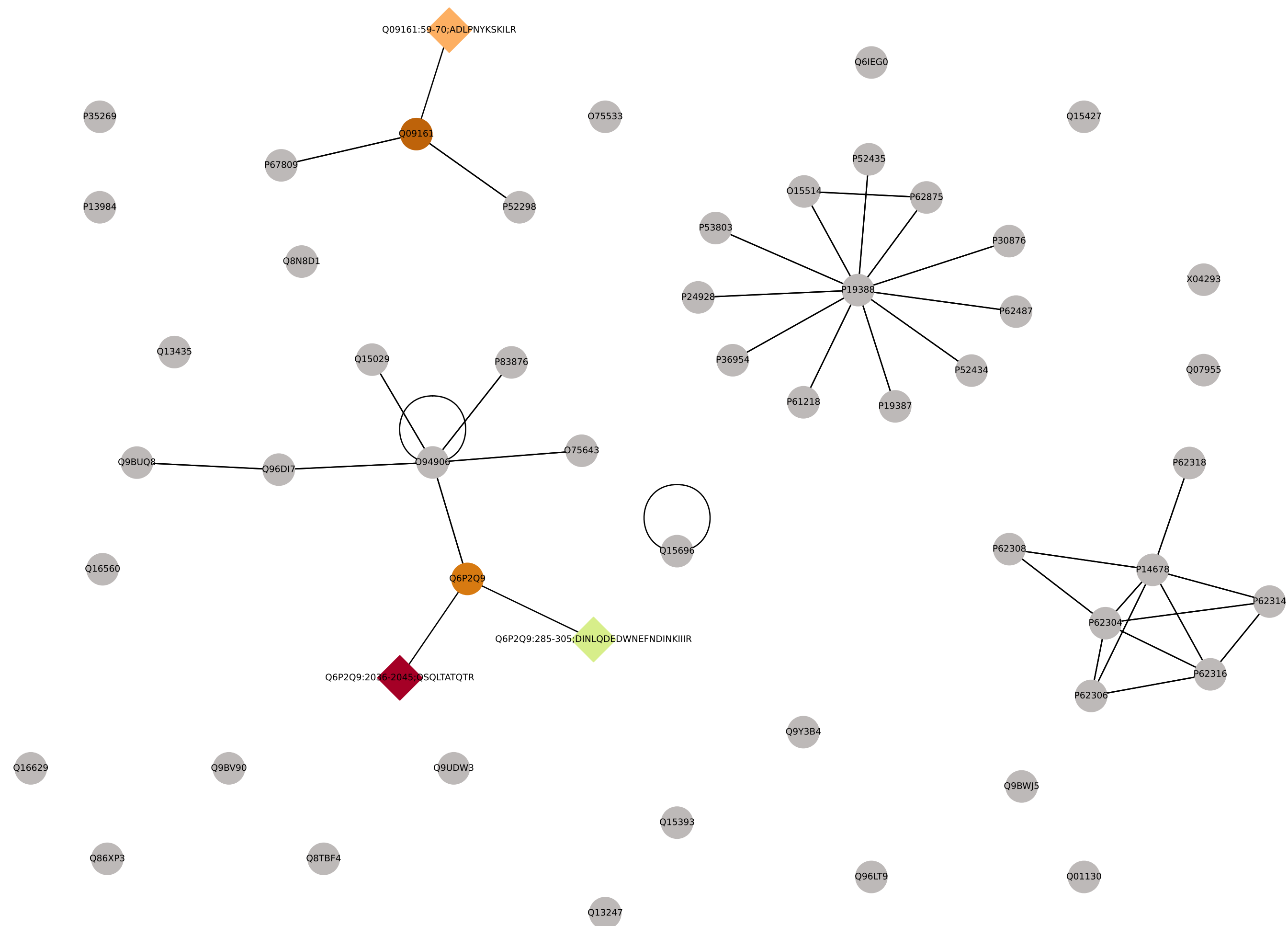p-value: 0.04103493715974438

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control

Log2_fold_change: gluc/control
R-HSA-9013418: RHOBTB2 GTPase cycle
p-value: 0.003870027935771292

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control
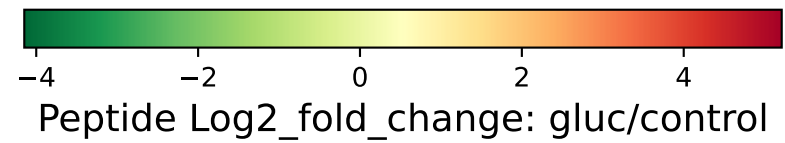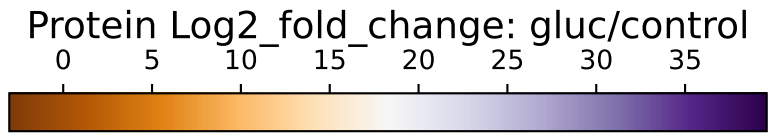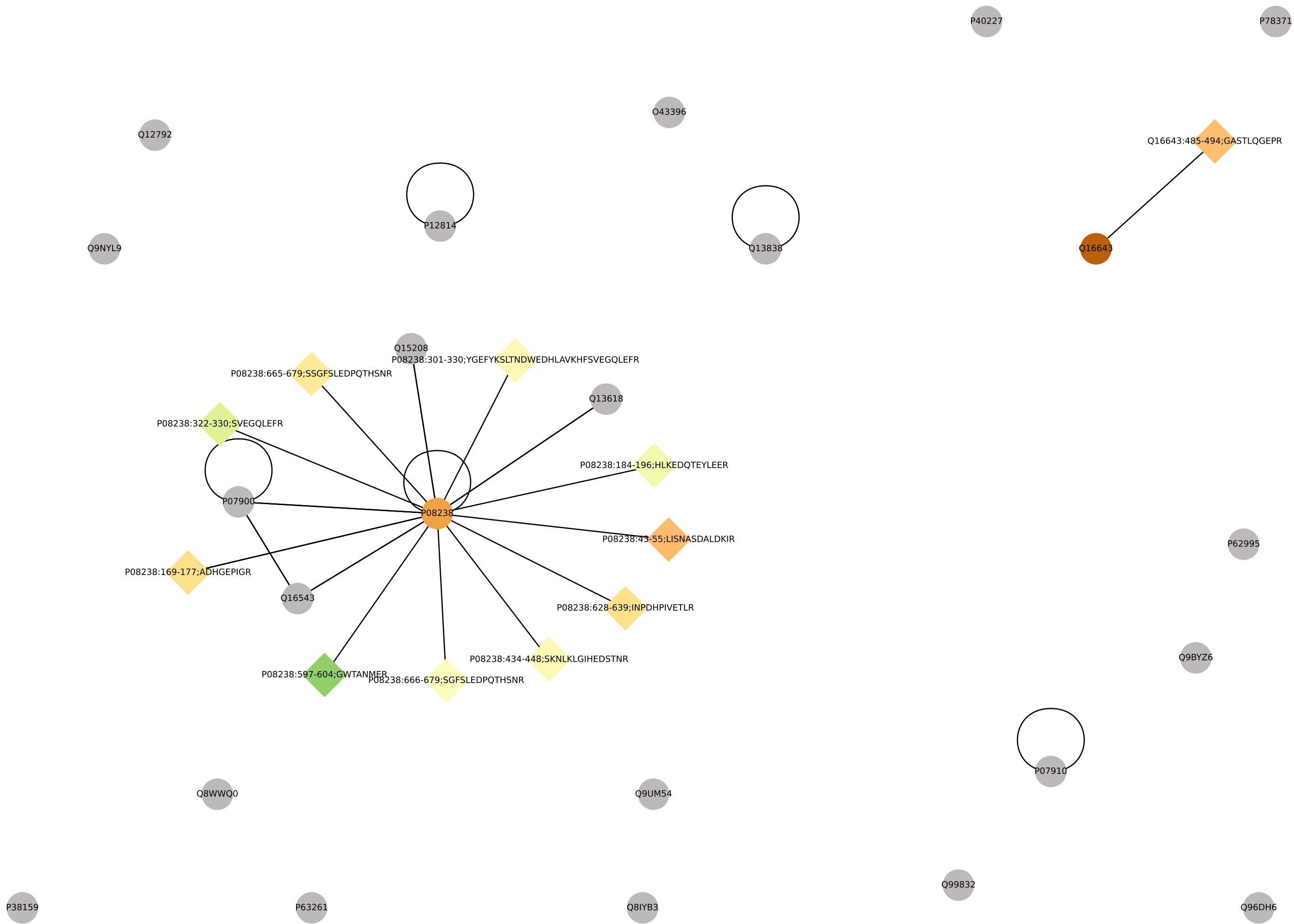
Log2_fold_change: gluc/control
R-HSA-9700645: ALK mutants bind TKIs
p-value: 0.04468226372702311

O43815:435-450:ADSLTYDIANNKDALR

Protein Log2_fold_change: gluc/control
0   5   10   15   20   25   30   35

Peptide Log2_fold_change: gluc/control
-4   -2   0   2   4

Log2_fold_change: gluc/control
R-HSA-72165: mRNA Splicing - Minor Pathway
p-value: 0.018216773752471993
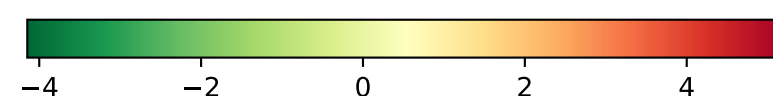
Log2_fold_change: gluc/control
R-HSA-72649: Translation initiation complex formation
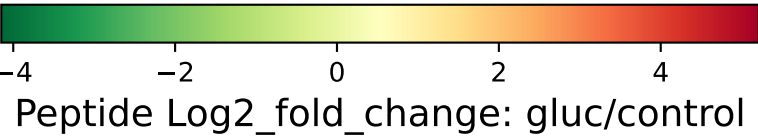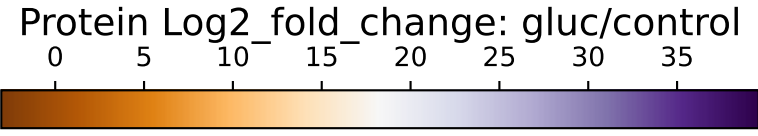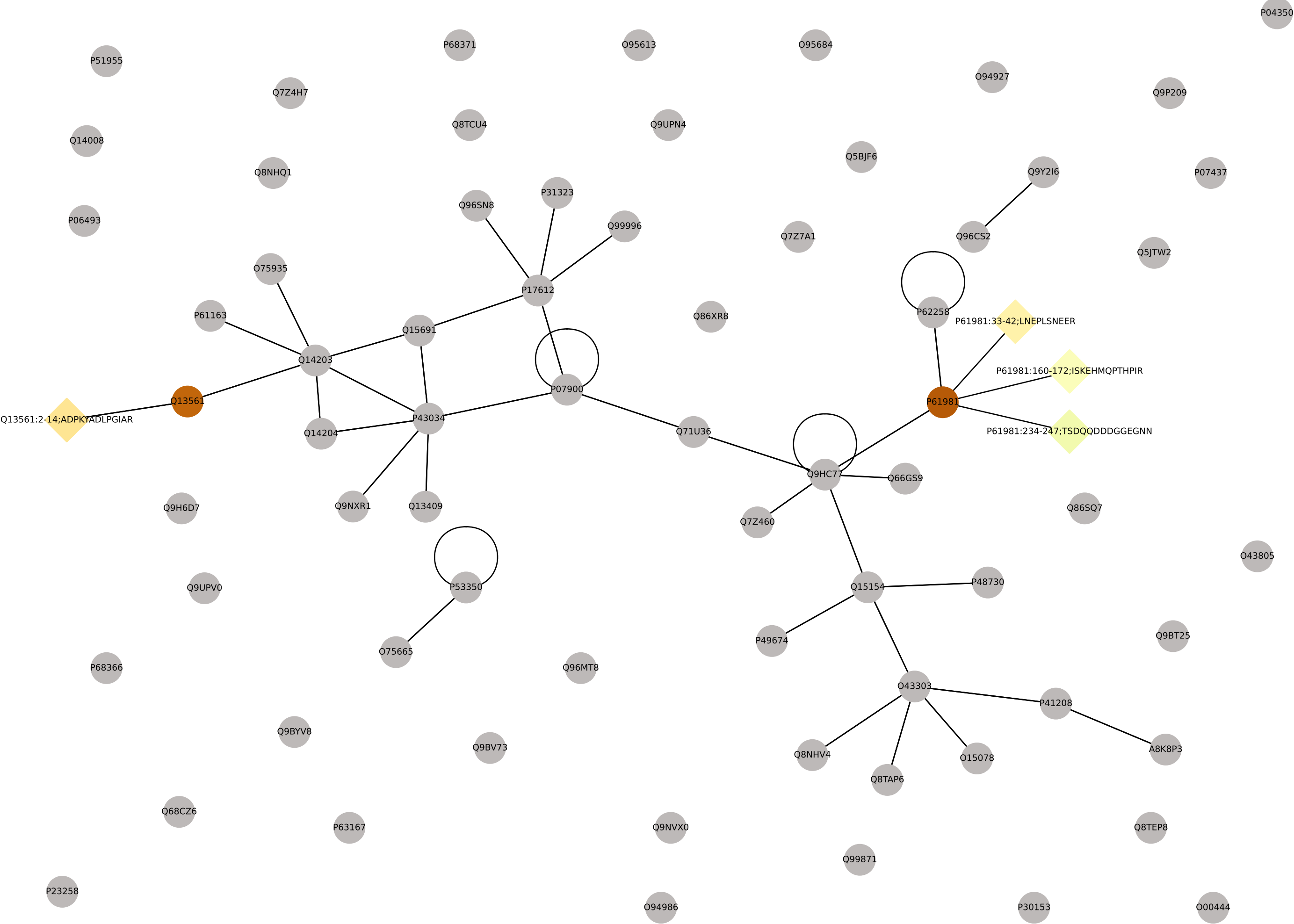p-value: 0.023554931232115828
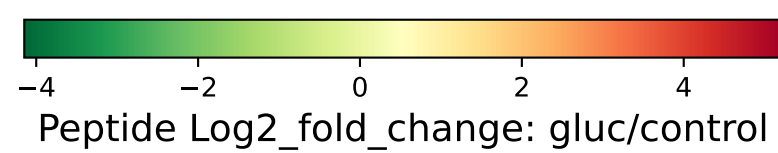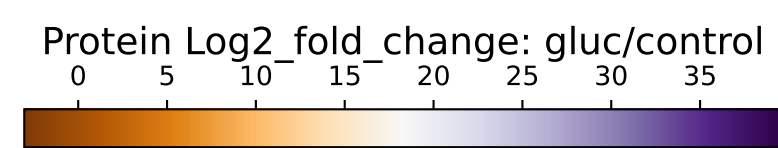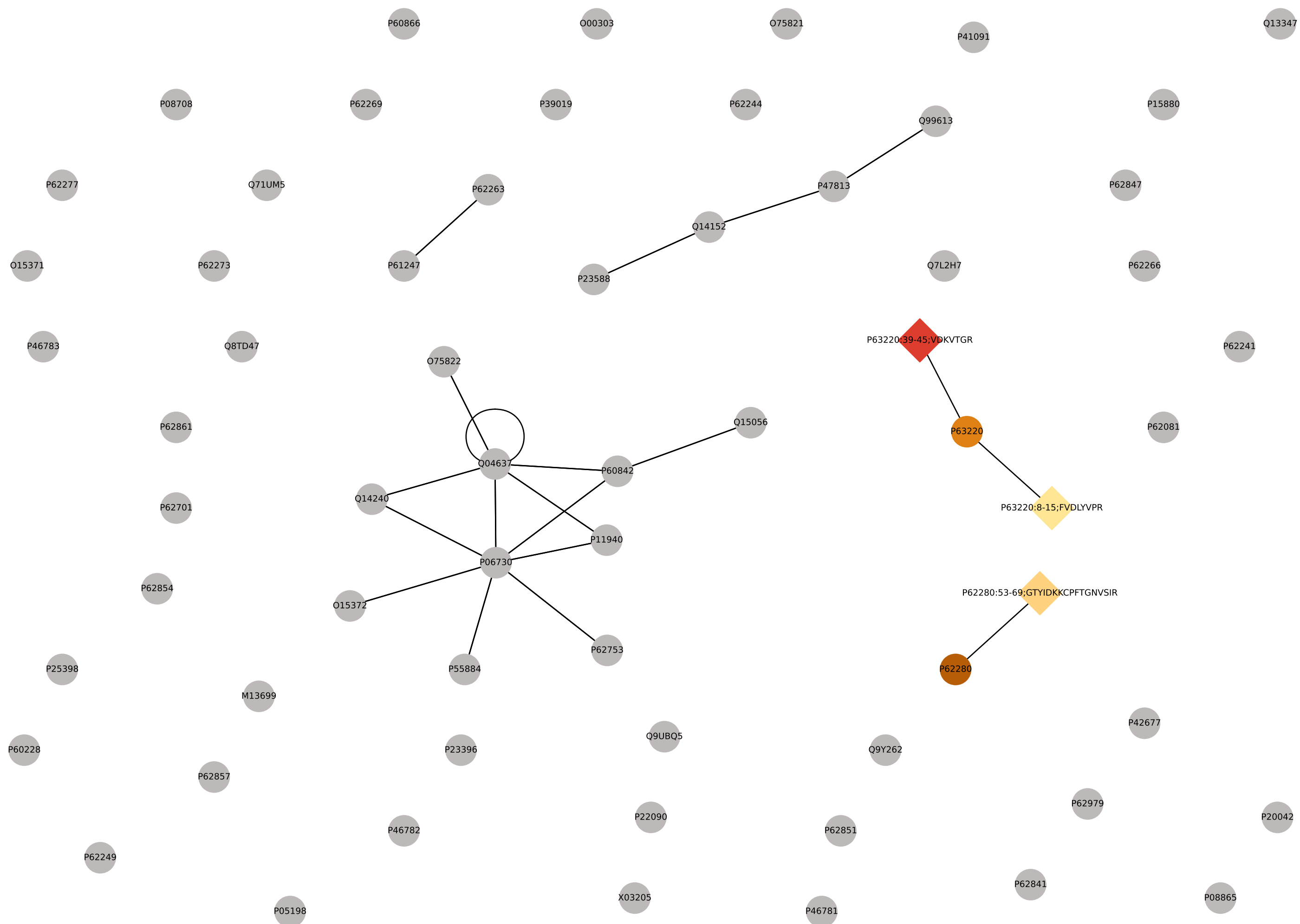
Log2_fold_change: gluc/control
R-HSA-72702: Ribosomal scanning and start codon recognition
p-value: 0.024978417082091475

Protein Log2_fold_change: gluc/control

0 5 10 15 20 25 30 35

Peptide Log2_fold_change: gluc/control

-4 -2 0 2 4
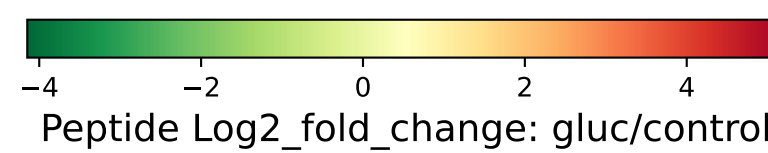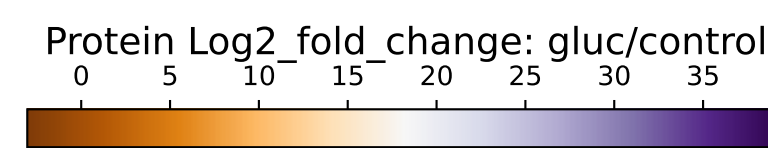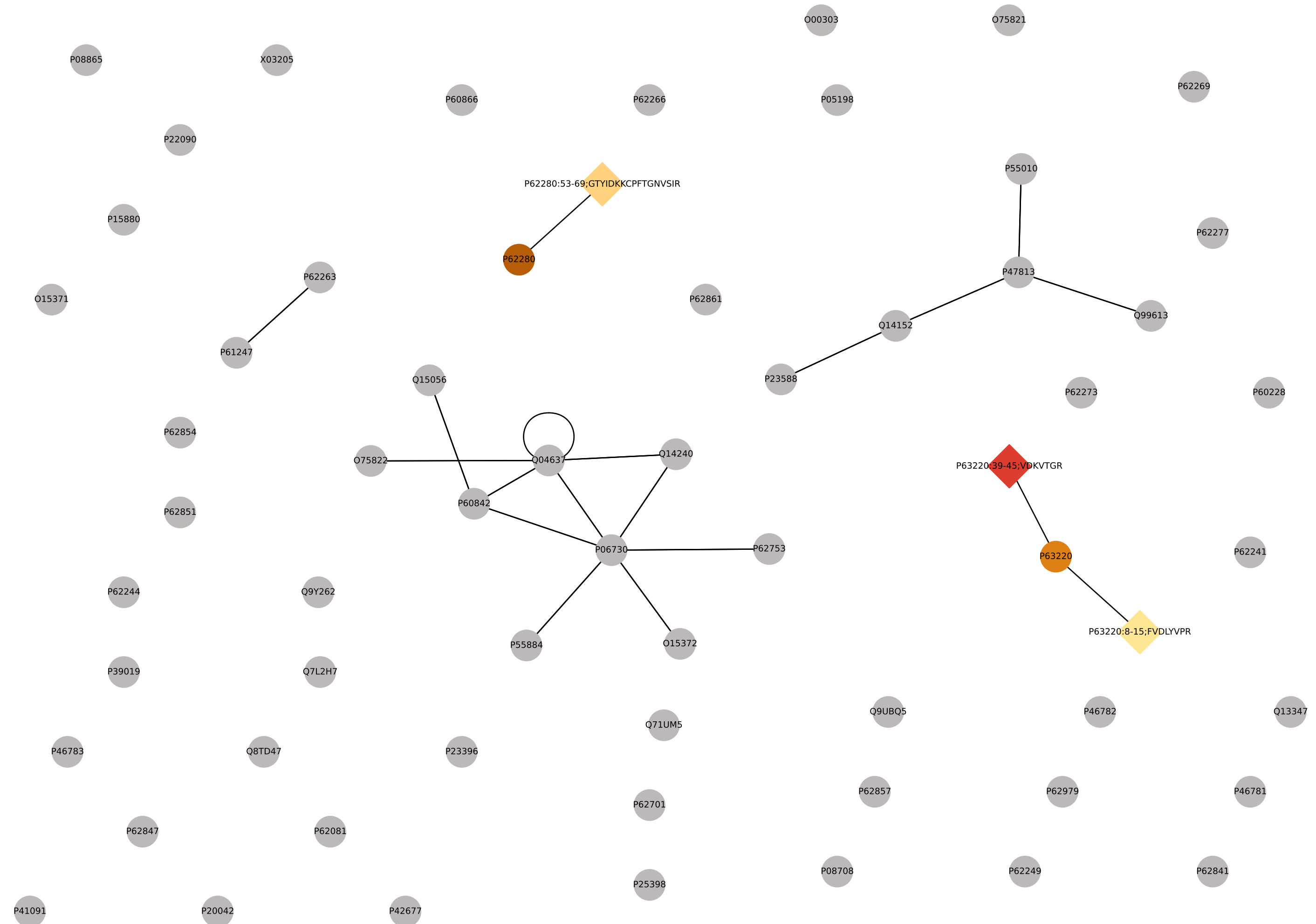
Log2_fold_change: gluc/control
R-HSA-975956: Nonsense Mediated Decay (NMD) independent of the Exon Junction Complex (EJC)
p-value: 0.006855569728165922

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control
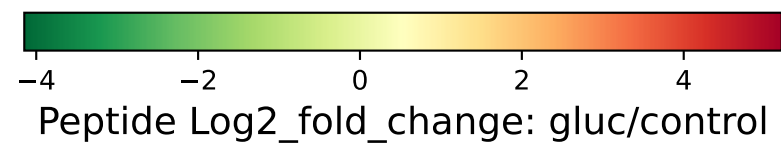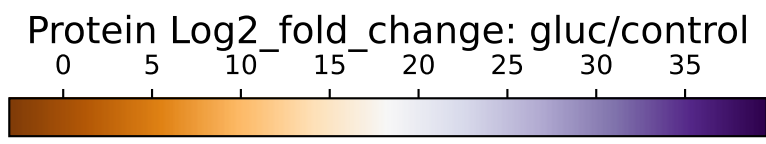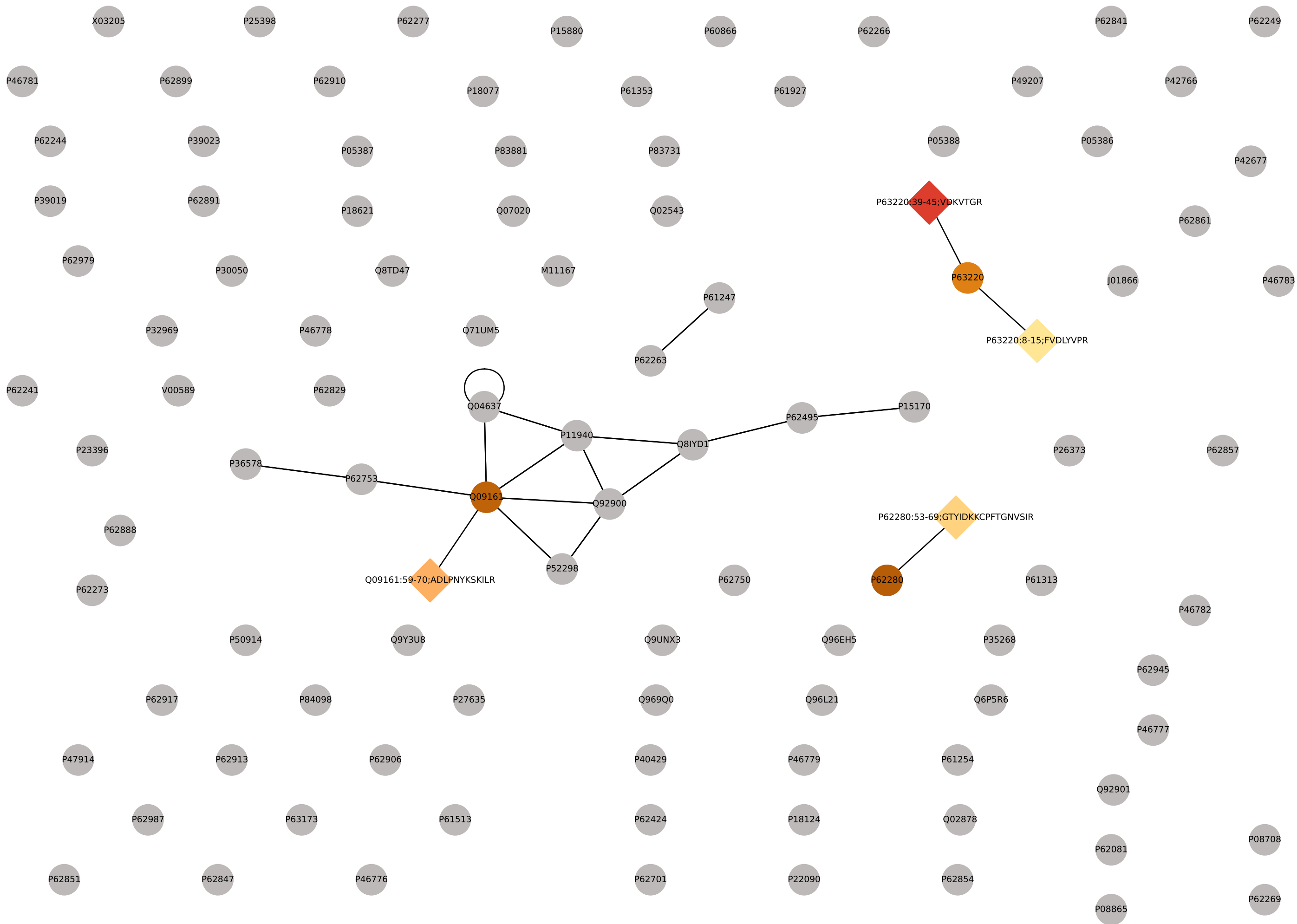
Log2_fold_change: gluc/control
R-HSA-380284: Loss of proteins required for interphase microtubule organization from the centrosome
p-value: 0.030227589686308676

Protein Log2_fold_change: gluc/control
0   5   10   15   20   25   30   35

Peptide Log2_fold_change: gluc/control
-4   -2   0   2   4

Log2_fold_change: gluc/control
R-HSA-380259: Loss of Nlp from mitotic centrosomes
p-value: 0.030227589686308676

Protein Log2_fold_change: gluc/control

0   5   10   15   20   25   30   35

Peptide Log2_fold_change: gluc/control

-4   -2   0   2   4

Log2_fold_change: gluc/control
R-HSA-8854518: AURKA Activation by TPX2
p-value: 0.032600049302680456

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control

Log2_fold_change: gluc/control
R-HSA-380270: Recruitment of mitotic centrosome proteins and complexes
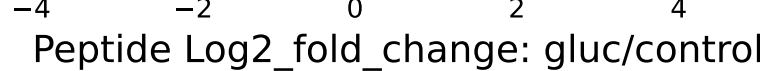p-value: 0.0384084717376354

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control

Log2_fold_change: gluc/control
R-HSA-927802: Nonsense-Mediated Decay (NMD)
p-value: 0.011931898374331729

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control

Log2_fold_change: gluc/control
R-HSA-975957: Nonsense Mediated Decay (NMD) enhanced by the Exon Junction Complex (EJC)
p-value: 0.011931898374331729

Protein Log2_fold_change: gluc/control

Peptide Log2_fold_change: gluc/control

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
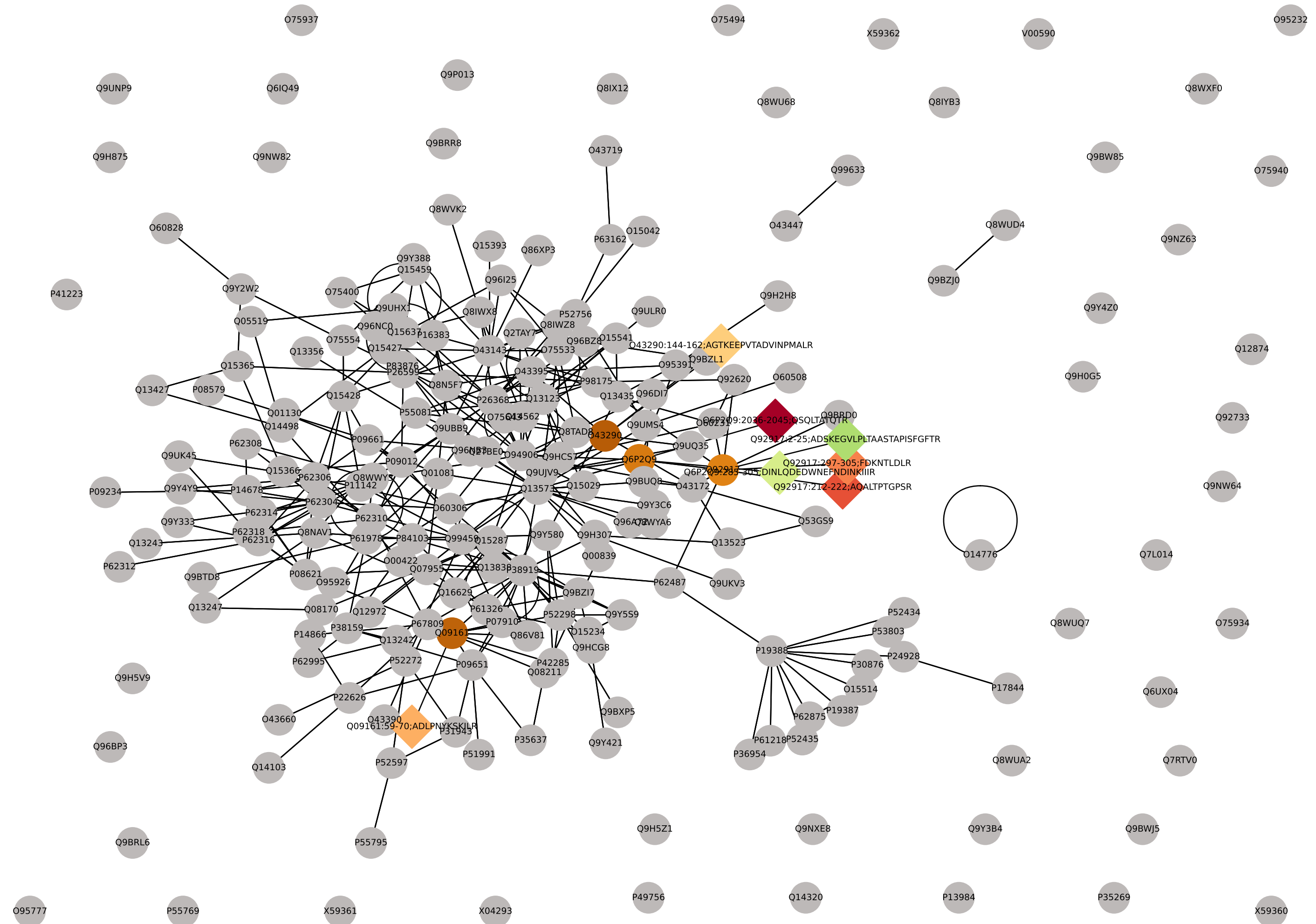
Protein Log2_fold_change: gluc/control
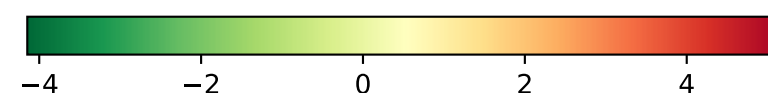
Peptide Log2_fold_change: gluc/control

Log2_fold_change: gluc/control
R-HSA-72172: mRNA Splicing
p-value: 0.01056574513476305