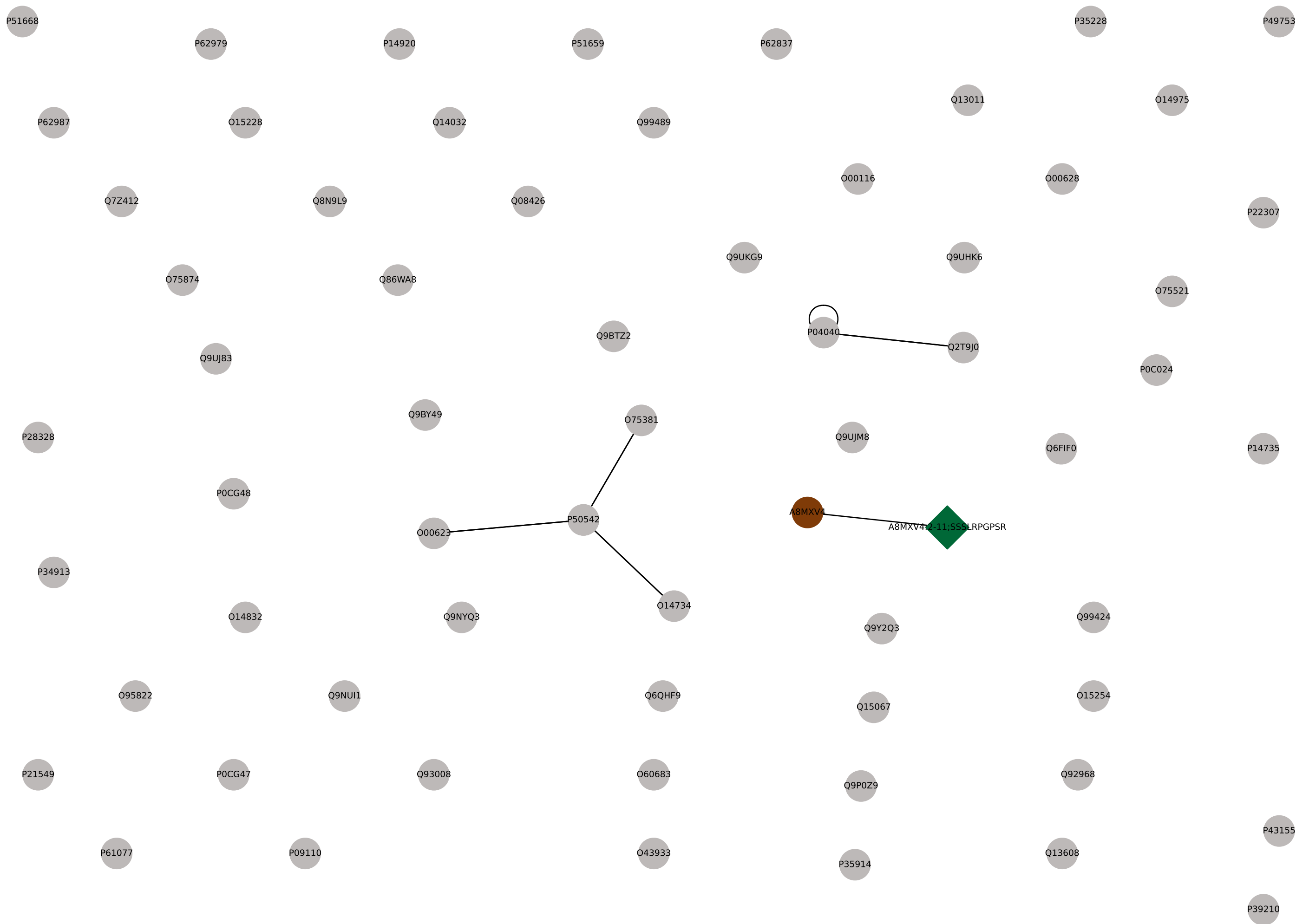


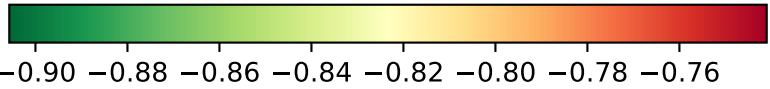
Log2 fold change: KO\_untreated vs. KO\_treated  
R-HSA-9033241: Peroxisomal protein import  
p-value: 0.004398634453781525



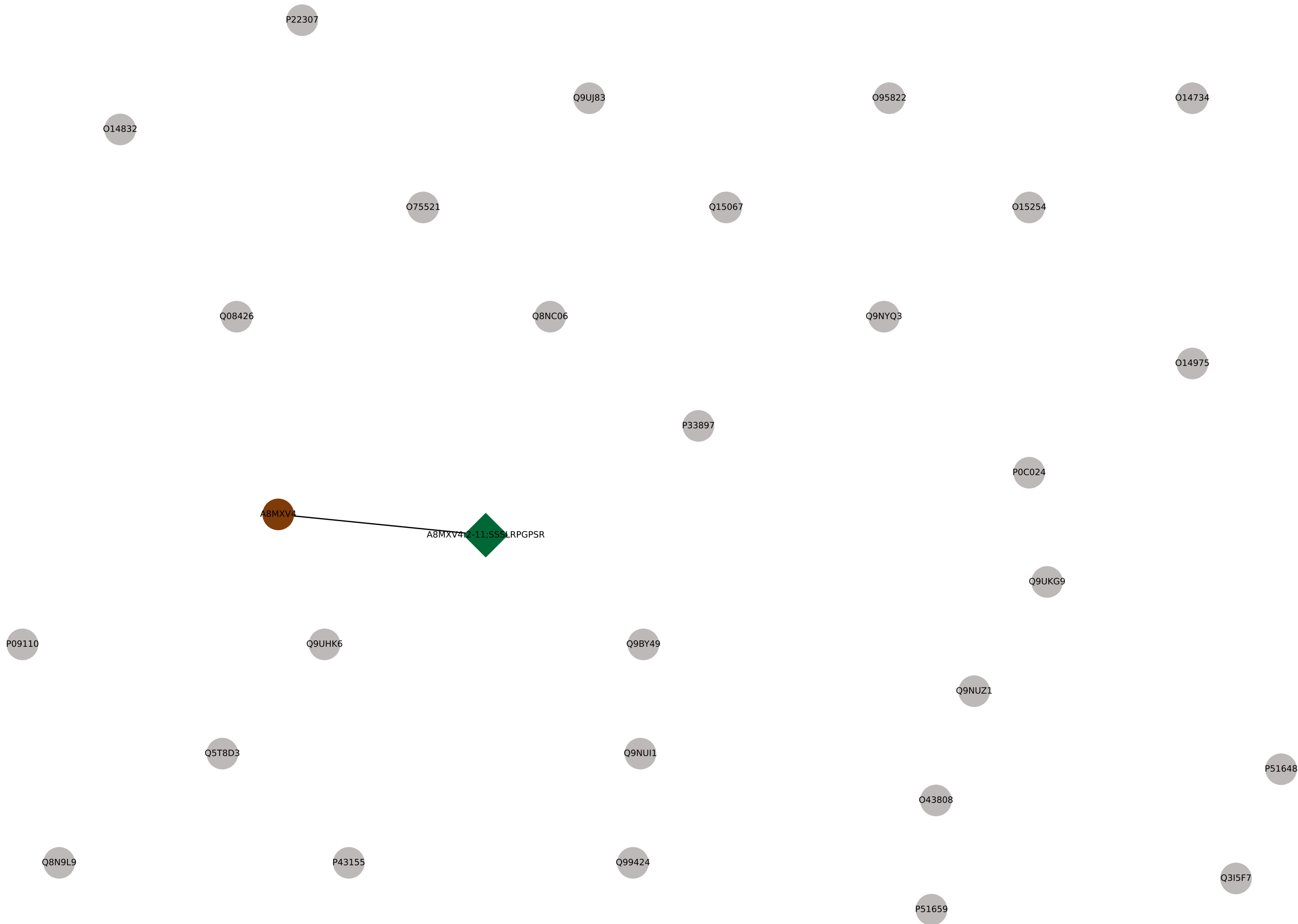
Protein Log2 fold change: KO\_untreated vs. KO\_treated



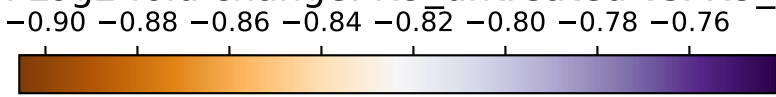
Peptide Log2 fold change: KO\_untreated vs. KO\_treated



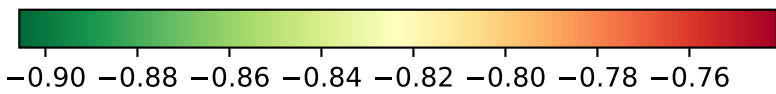
Log2 fold change: KO\_untreated vs. KO\_treated  
R-HSA-390918: Peroxisomal lipid metabolism  
p-value: 0.005908613445378186



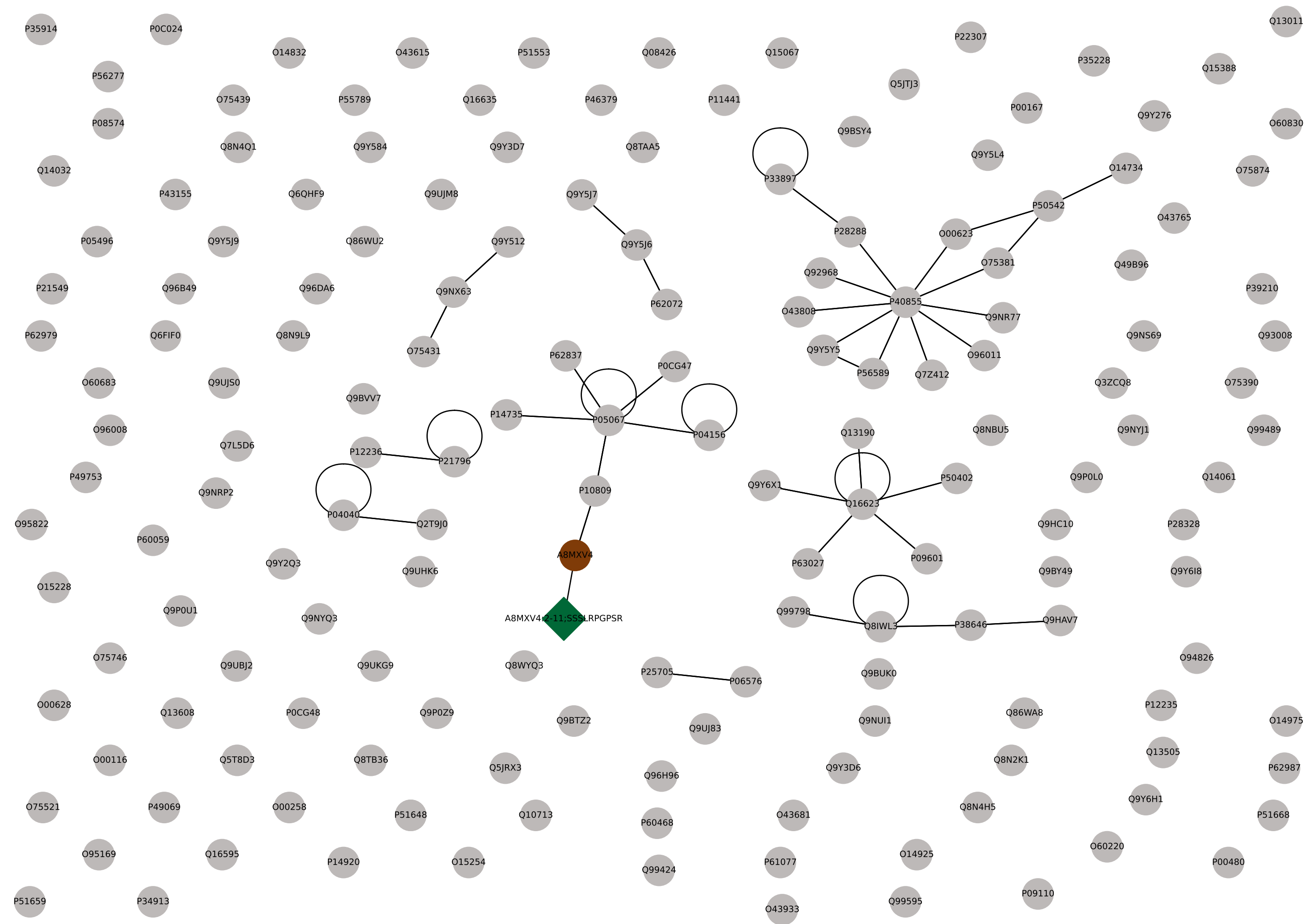
Protein Log2 fold change: KO\_untreated vs. KO\_treated



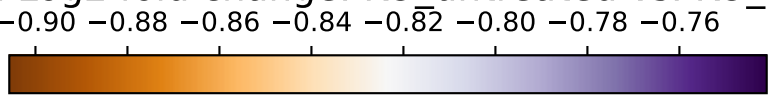
Peptide Log2 fold change: KO\_untreated vs. KO\_treated



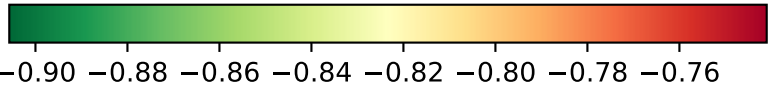
Log2 fold change: KO\_untreated vs. KO\_treated  
R-HSA-9609507: Protein localization  
p-value: 0.011160714285714302



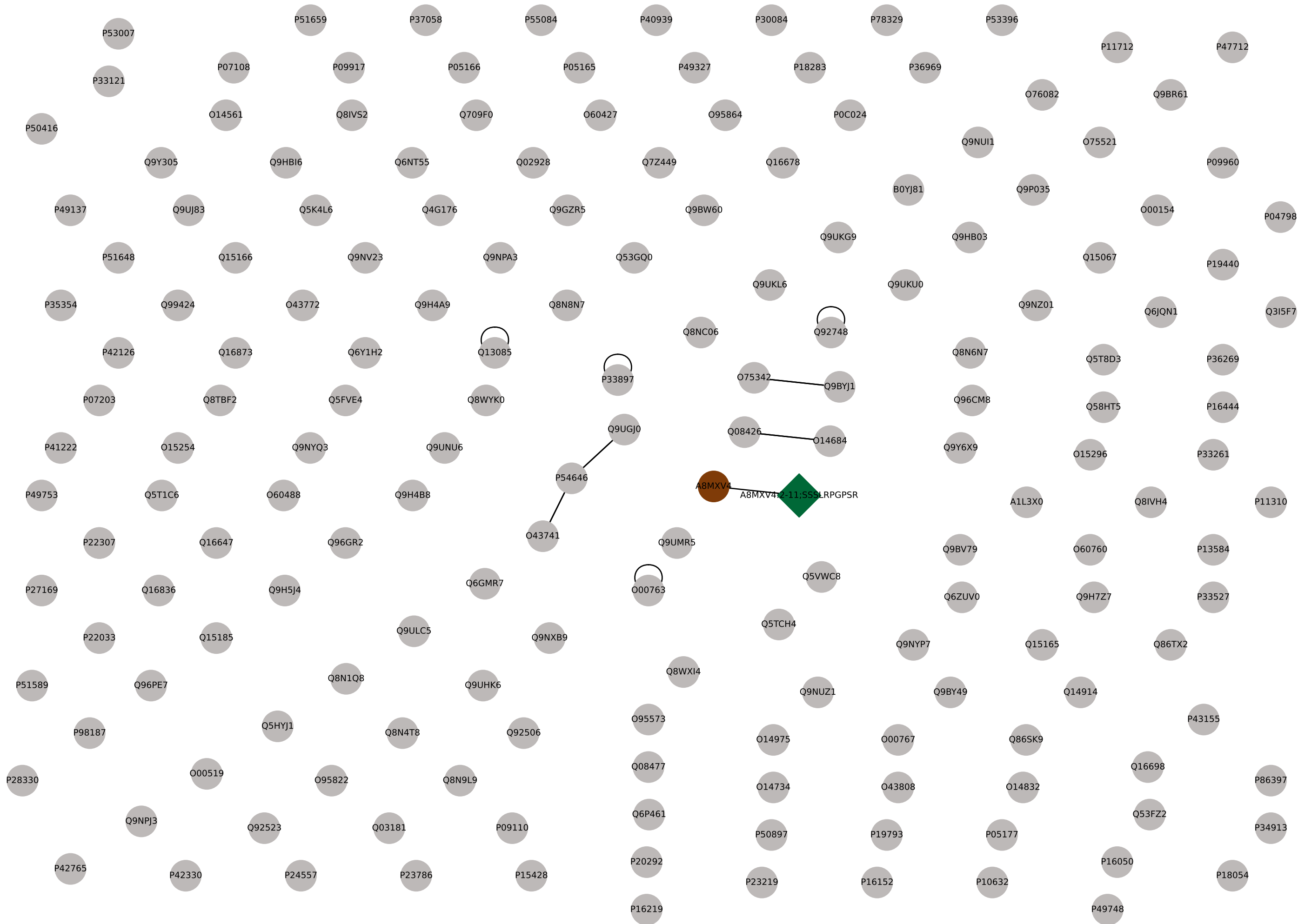
Protein Log2 fold change: KO\_untreated vs. KO\_treated



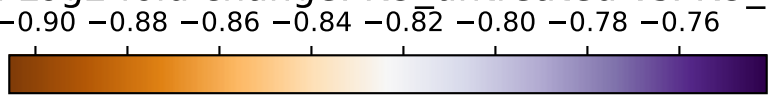
Peptide Log2 fold change: KO\_untreated vs. KO\_treated



Log2 fold change: KO\_untreated vs. KO\_treated  
R-HSA-8978868: Fatty acid metabolism  
p-value: 0.028295693277310963



Protein Log2 fold change: KO\_untreated vs. KO\_treated



Peptide Log2 fold change: KO\_untreated vs. KO\_treated

