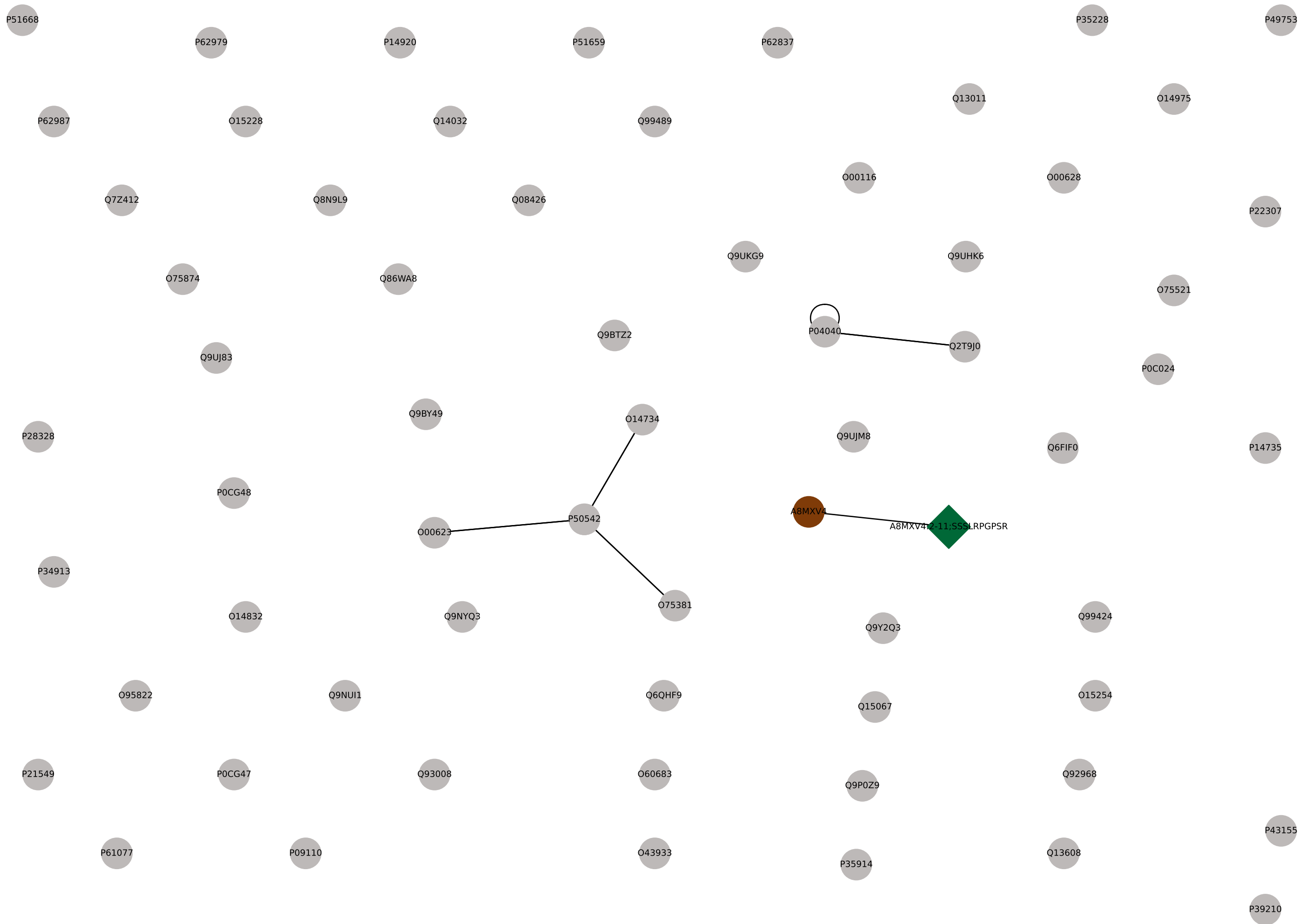


Log2 fold change: KO_untreated vs. KO_treated
R-HSA-9033241: Peroxisomal protein import
p-value: 0.004398634453781525



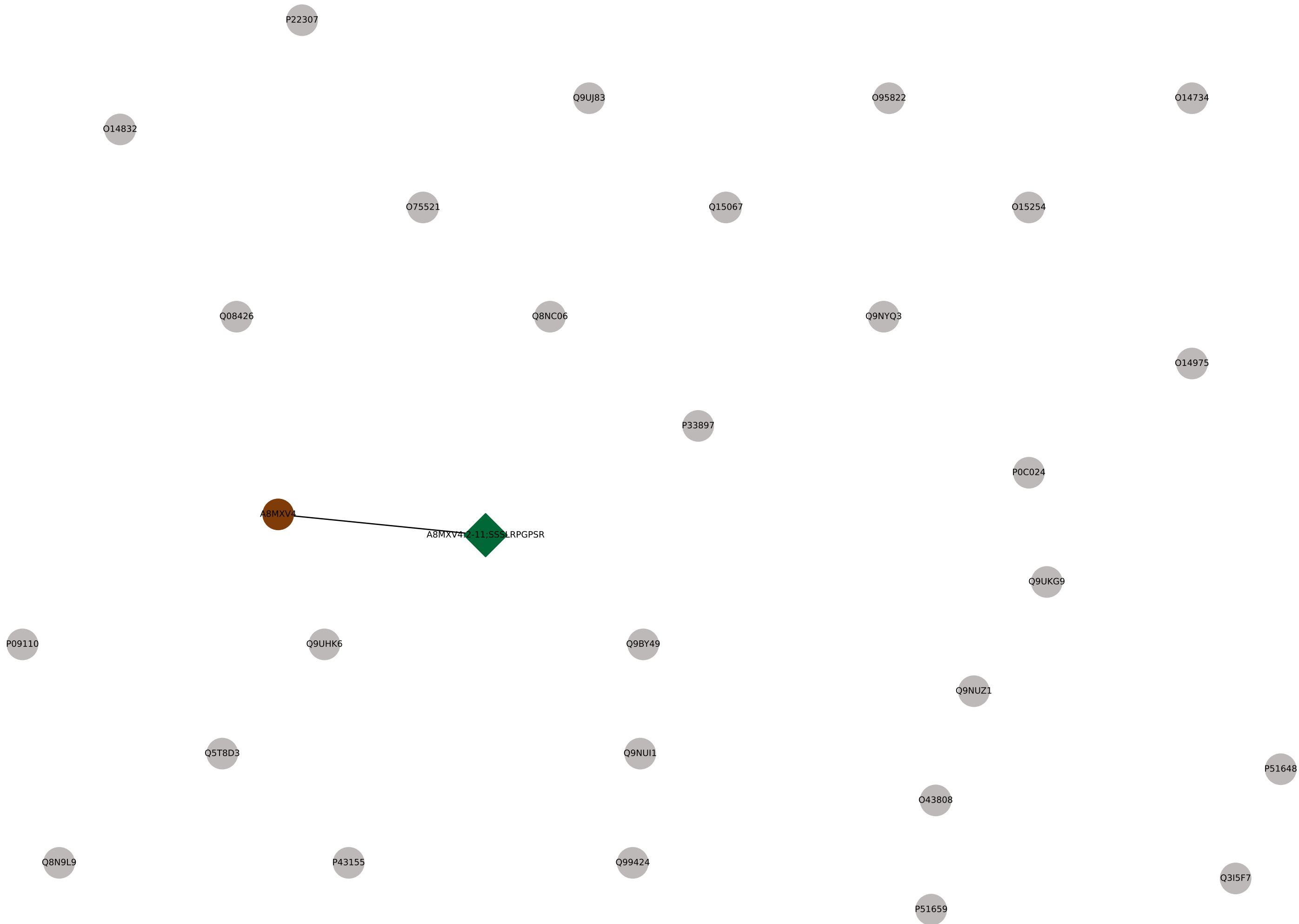
Protein Log2 fold change: KO_untreated vs. KO_treated



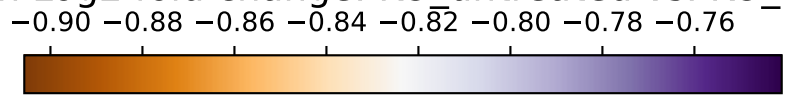
Peptide Log2 fold change: KO_untreated vs. KO_treated



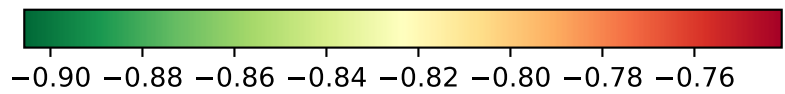
Log2 fold change: KO_untreated vs. KO_treated
R-HSA-390918: Peroxisomal lipid metabolism
p-value: 0.005908613445378186



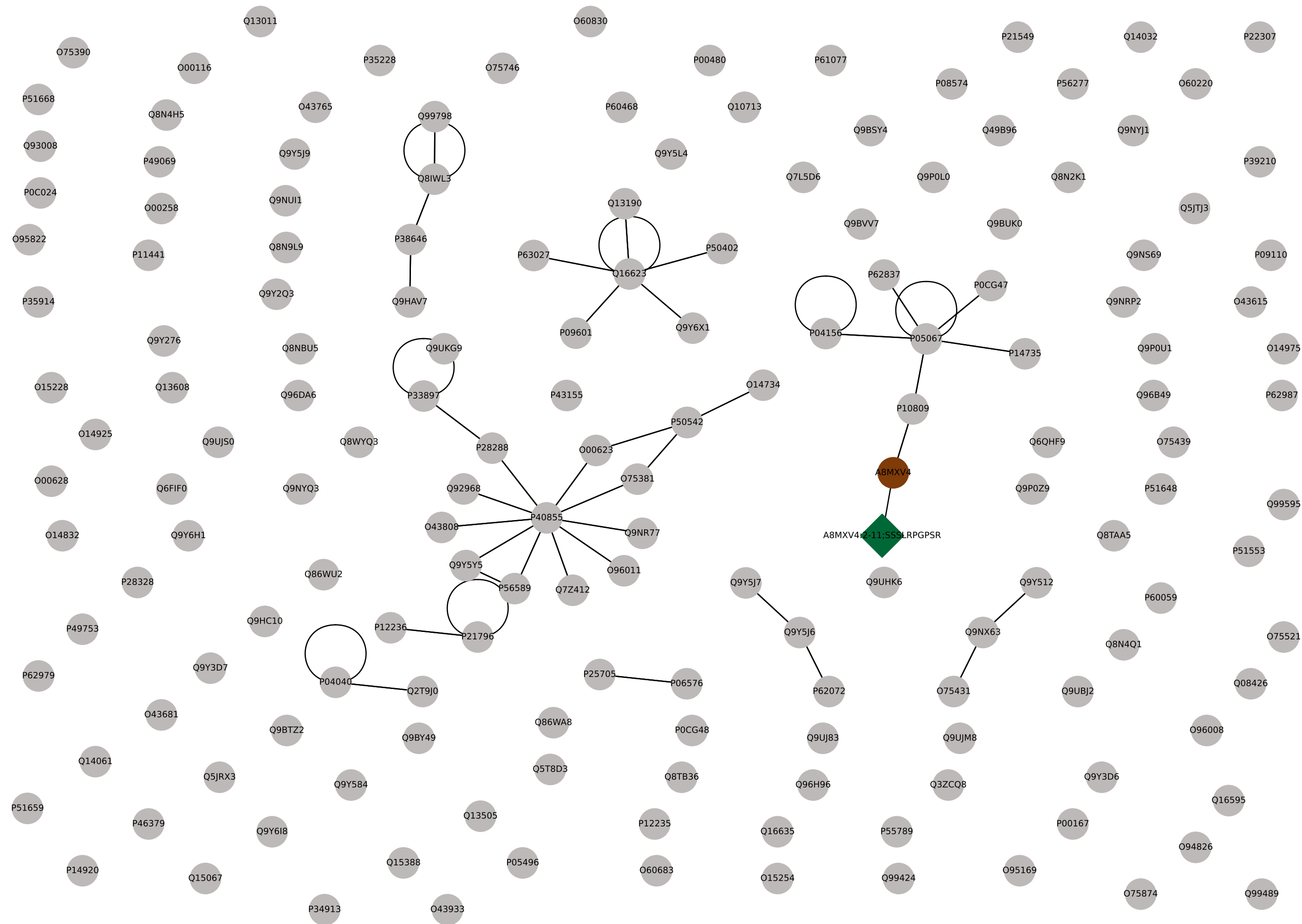
Protein Log2 fold change: KO_untreated vs. KO_treated



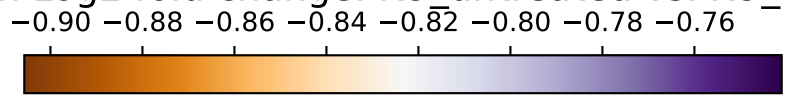
Peptide Log2 fold change: KO_untreated vs. KO_treated



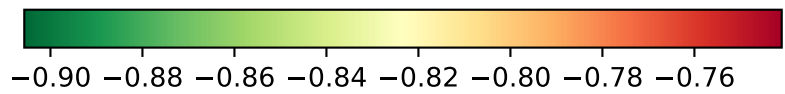
Log2 fold change: KO_untreated vs. KO_treated
R-HSA-9609507: Protein localization
p-value: 0.011160714285714302



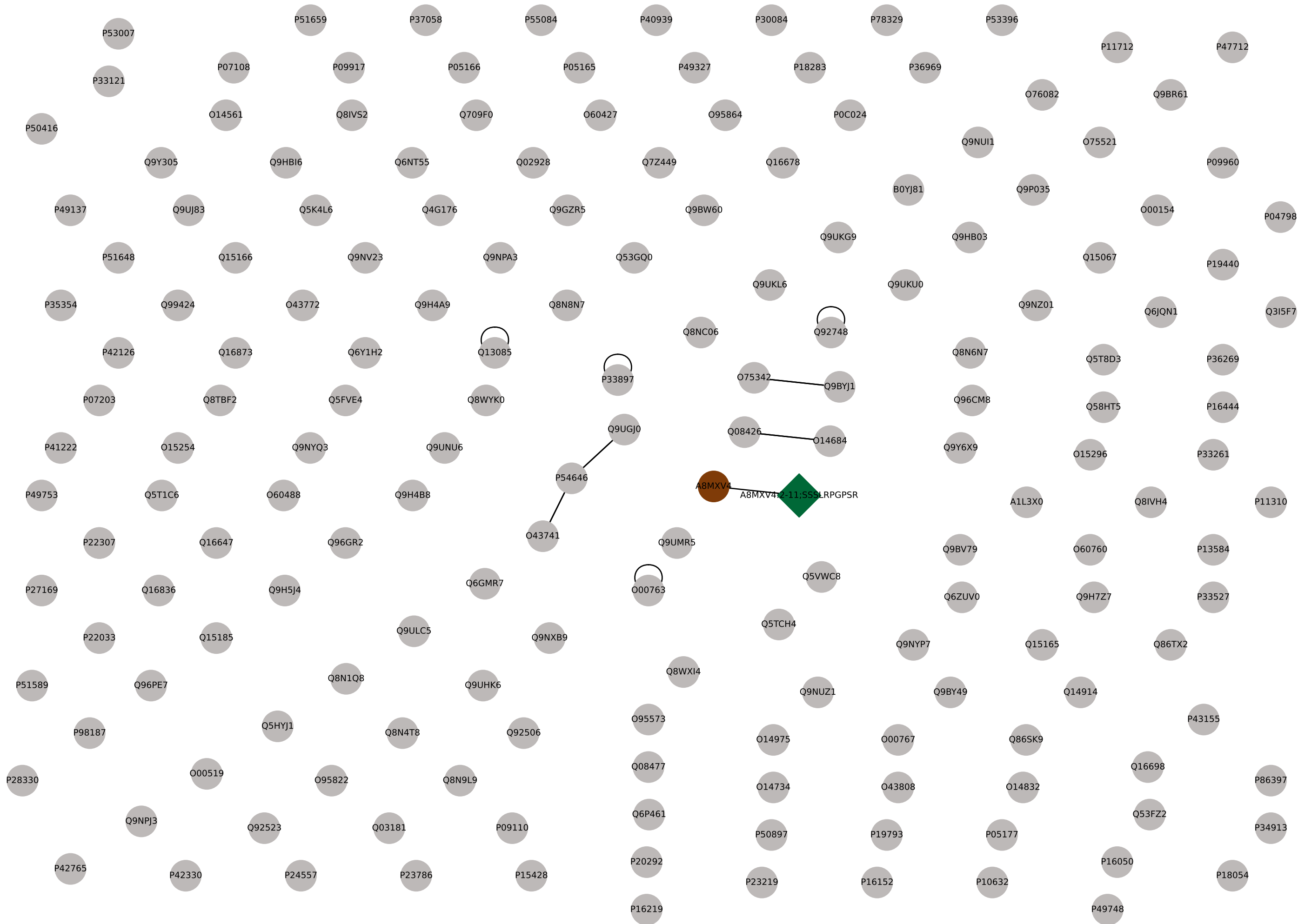
Protein Log2 fold change: KO_untreated vs. KO_treated



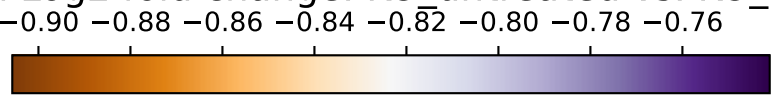
Peptide Log2 fold change: KO_untreated vs. KO_treated



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

