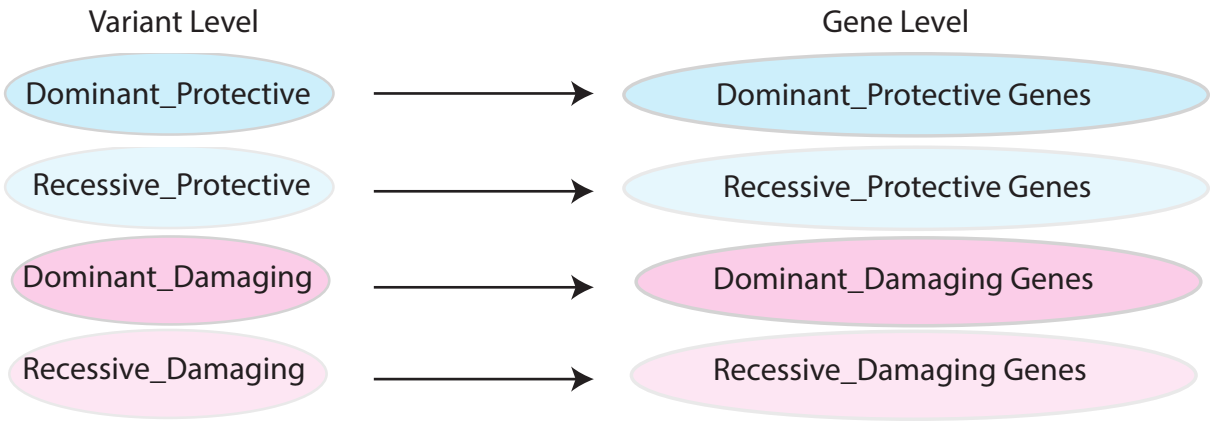
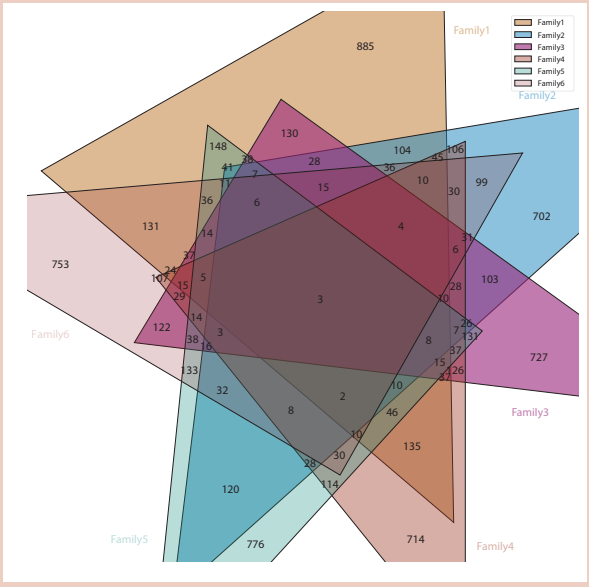
Gene Level Analysis

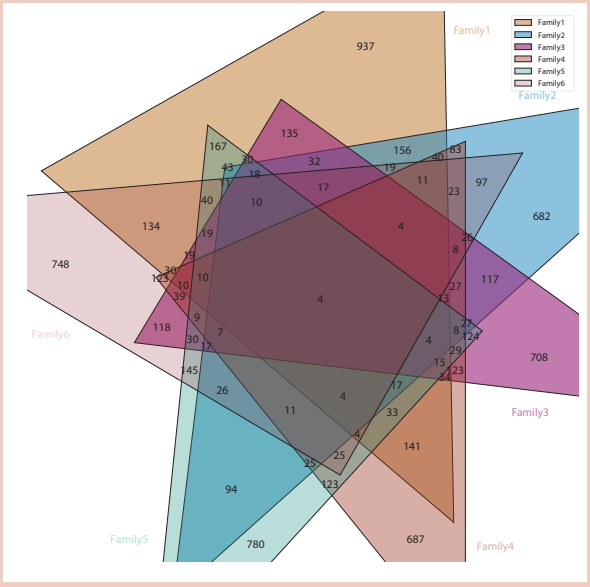
For this analysis, we are going to abstract the variants out to the gene level per family, and then run intersections. It only makes sense to perform this at the HIGH/MED level, not cis-regulatory



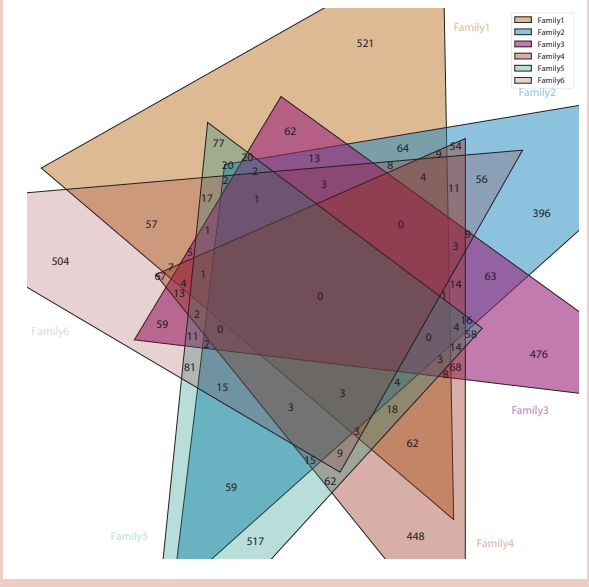
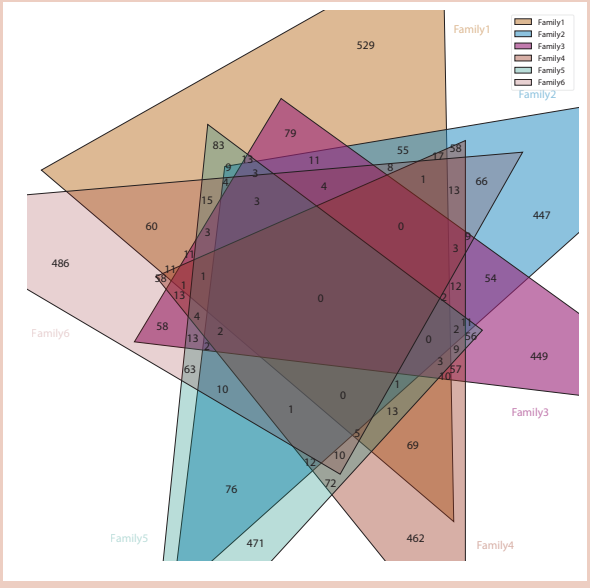
Dominant Protective



Dominant Damaging



Recessive Protective



Recessive Damaging