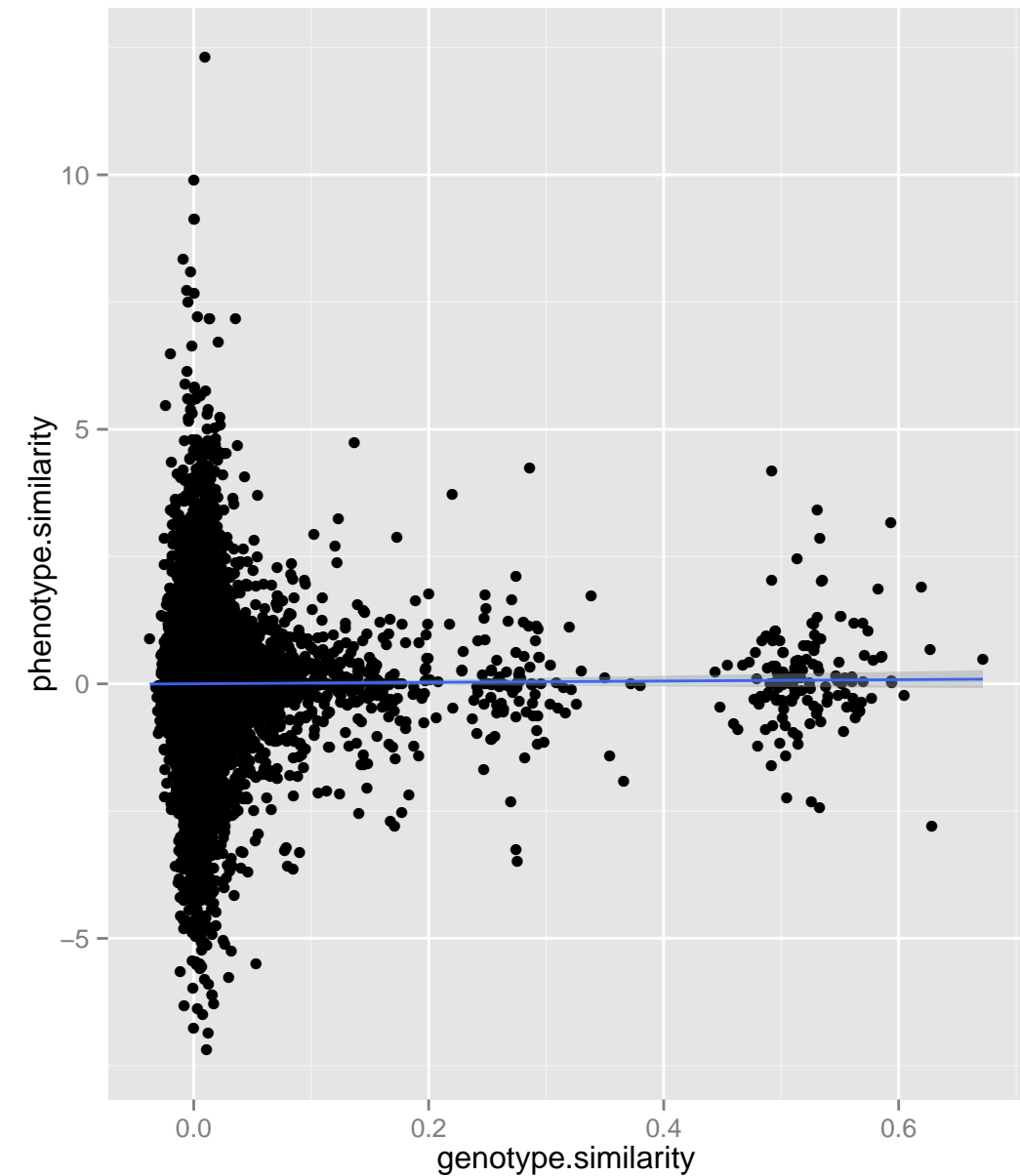
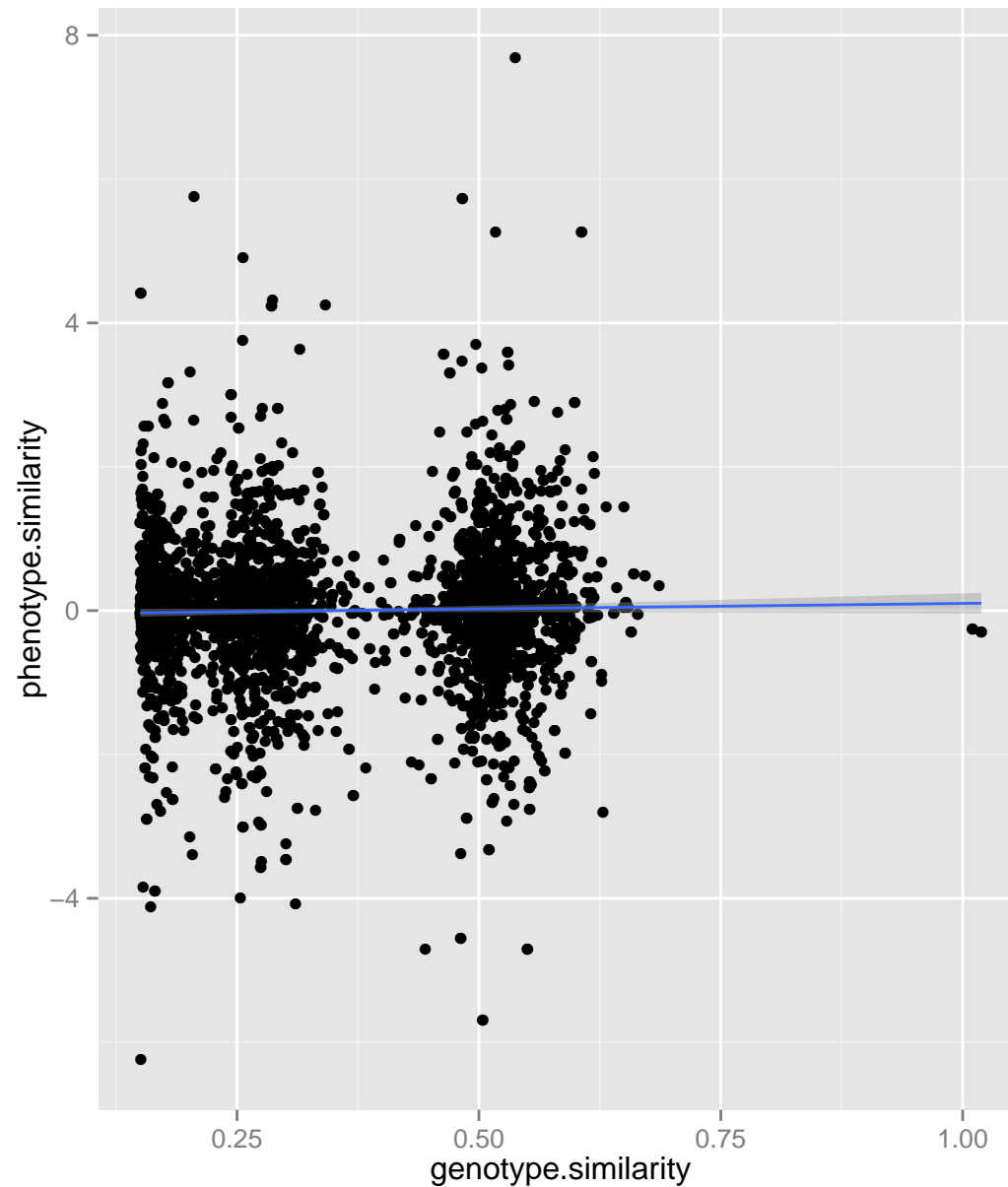


Phenotype vs genotype similarity: SCAP_meancorrect,
heri:0.0430847473386534



Phenotype vs genotype similarity: SCAP_meancorrect,
heri:0.191056479702018



Phenotype vs genotype similarity: SCAP_meancorrect,
heri:-0.0731564976648557

