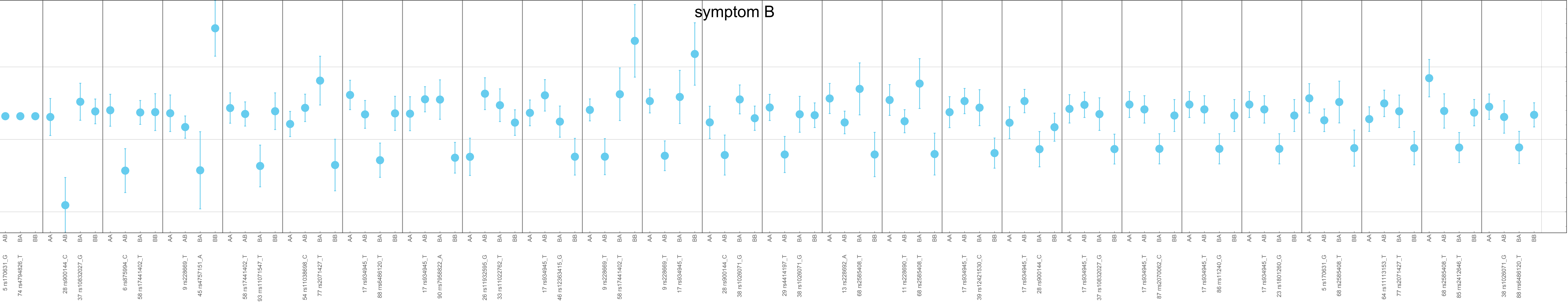


freq(symptom | genes alleles)



gene pair