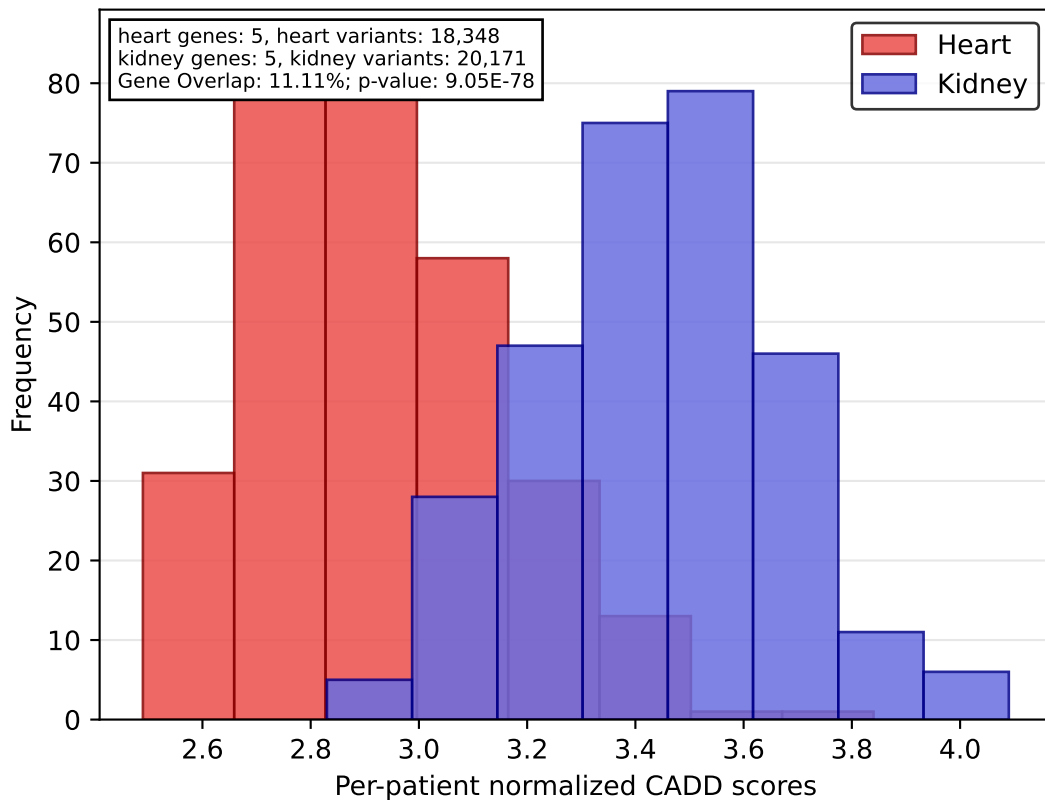
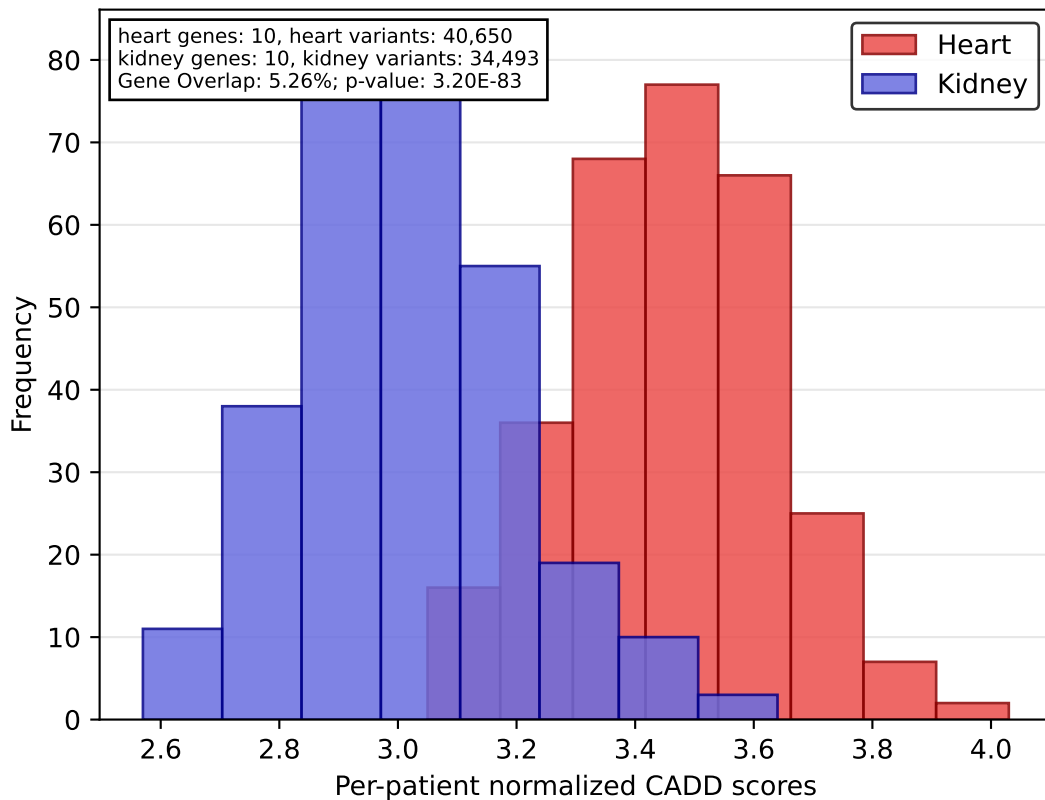


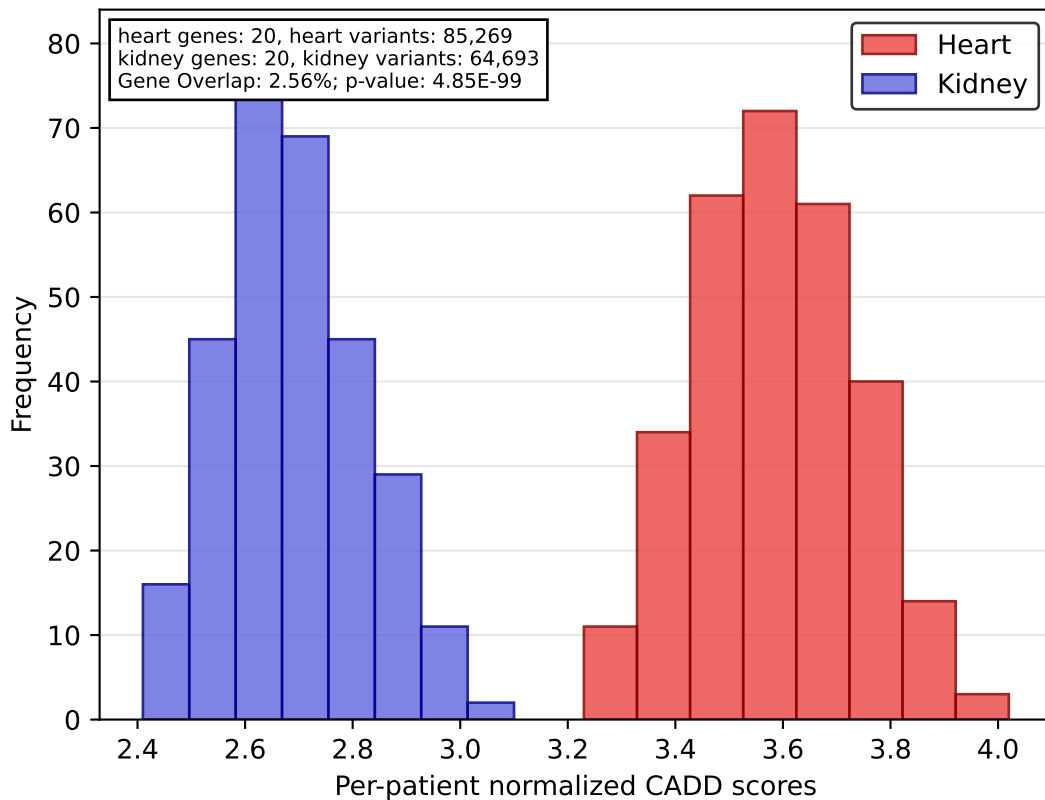
Top 5 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



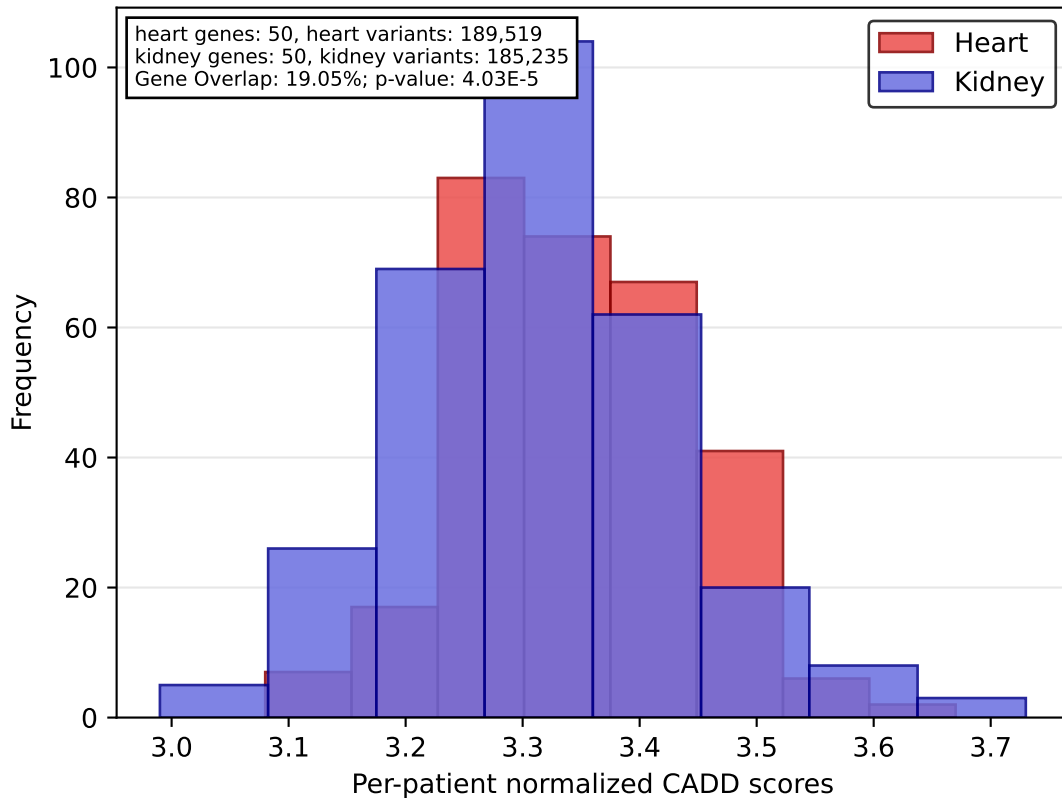
Top 10 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



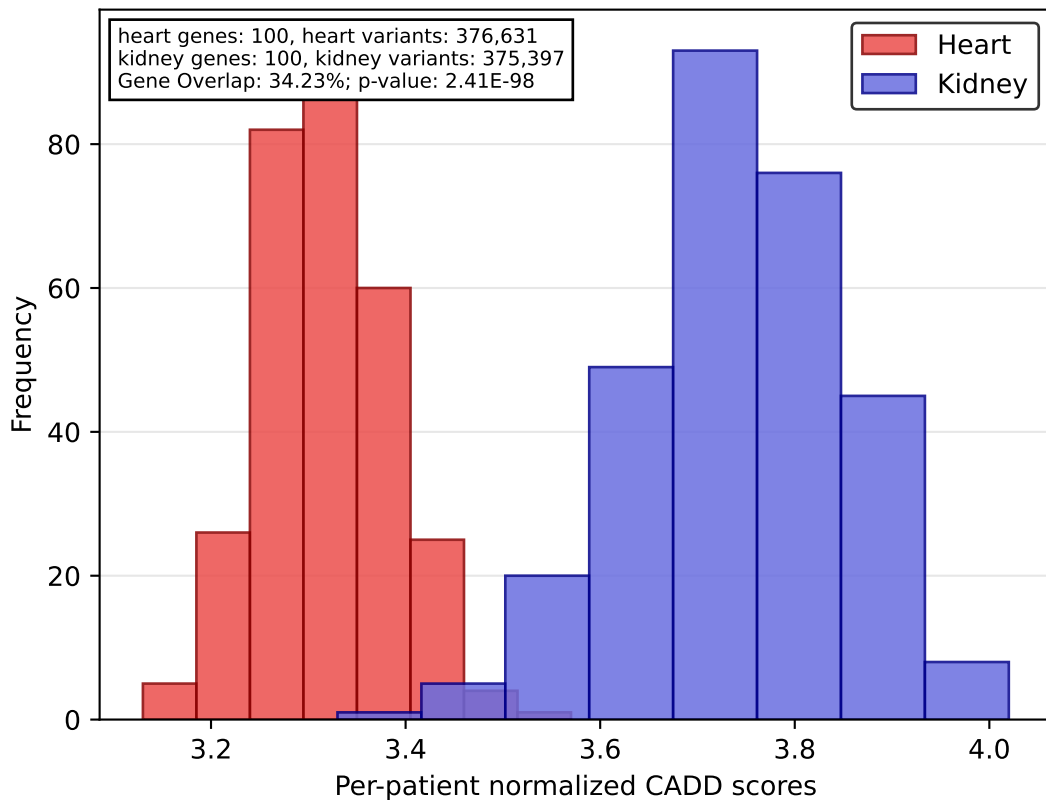
Top 20 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



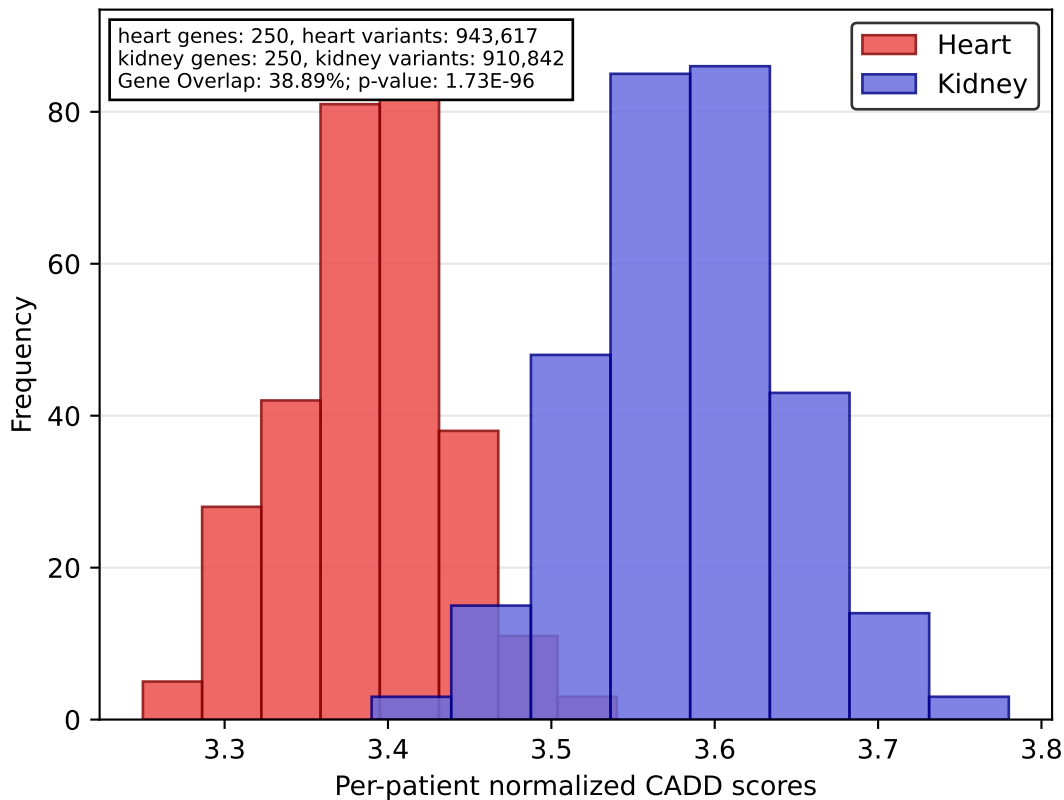
Top 50 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



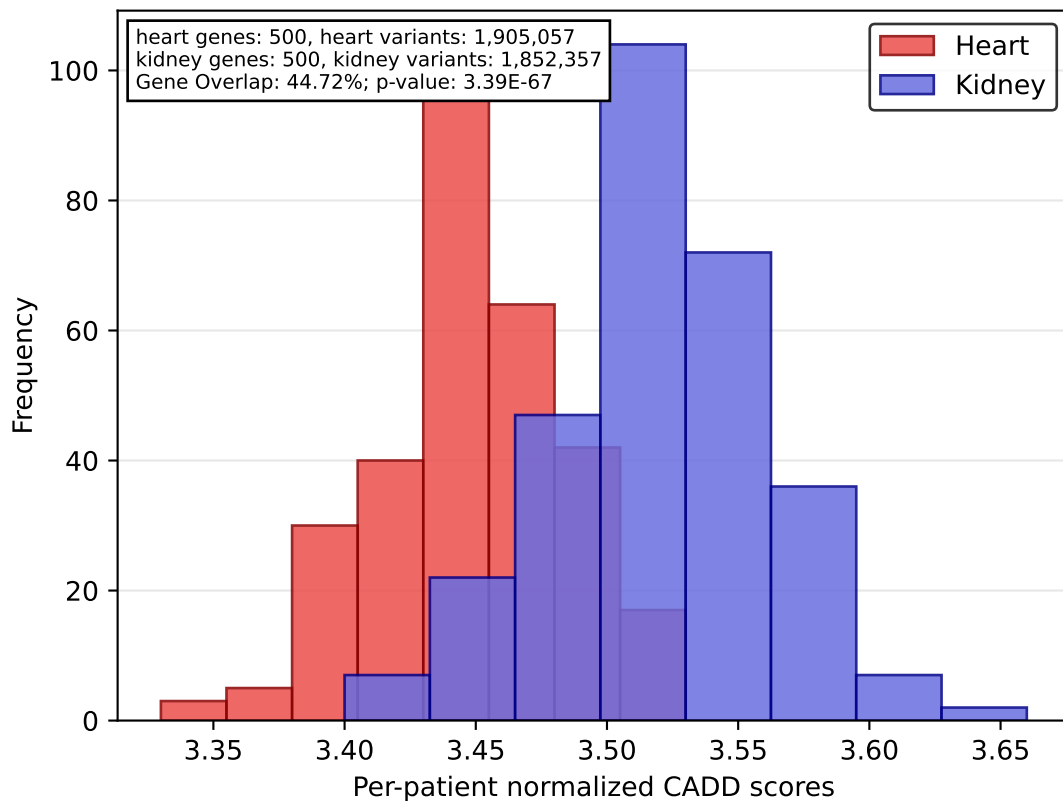
Top 100 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



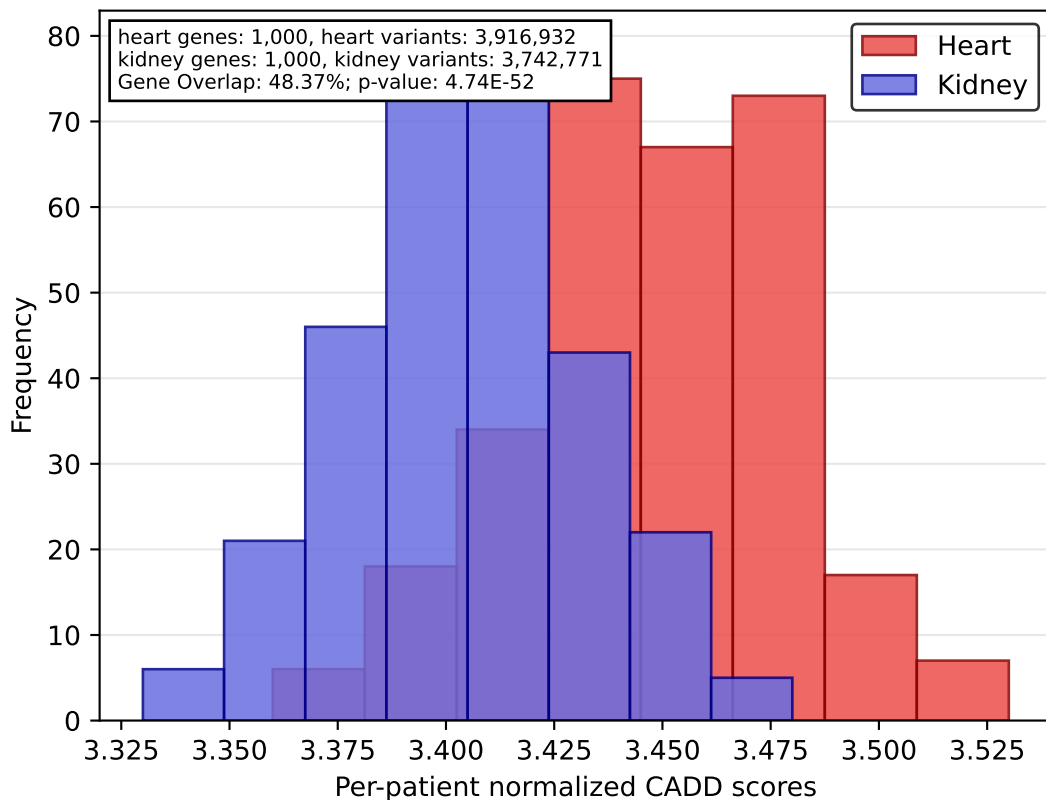
Top 250 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



Top 500 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



Top 1000 expressed genes from GTEx heart and kidney data were used to filter variants from the CHD cohort



Summary of CHD analysis using GTEx tissue datasets

