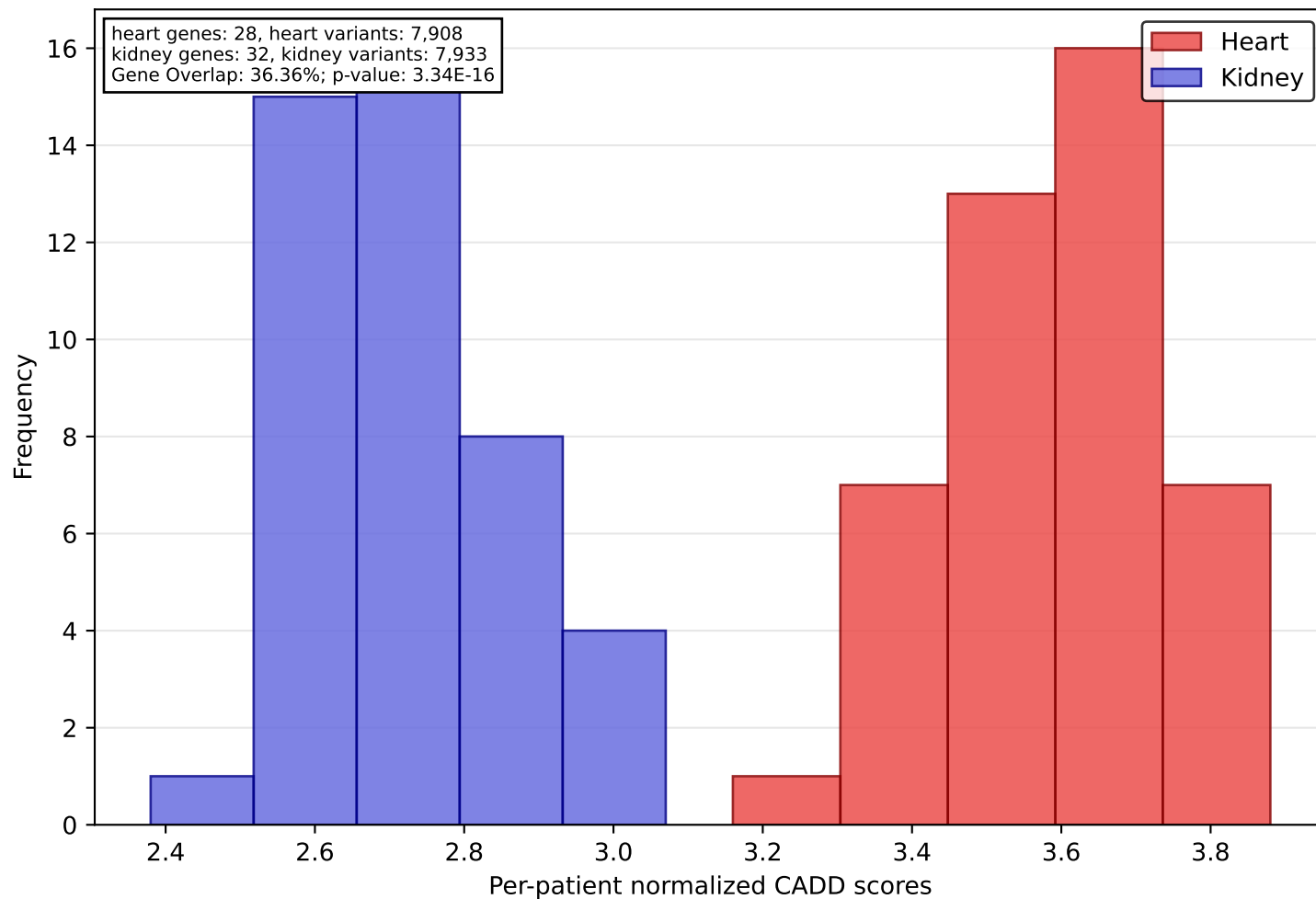
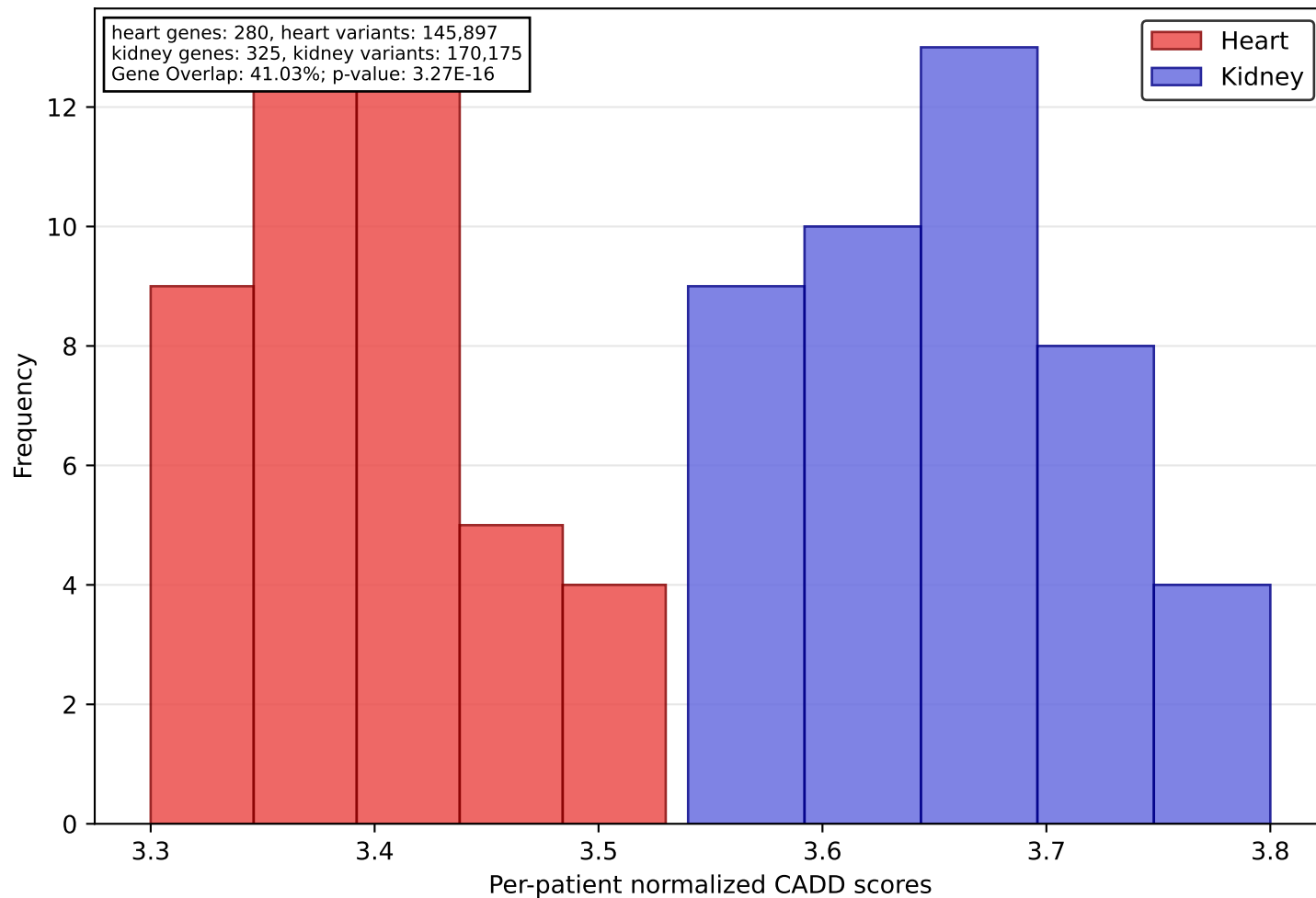


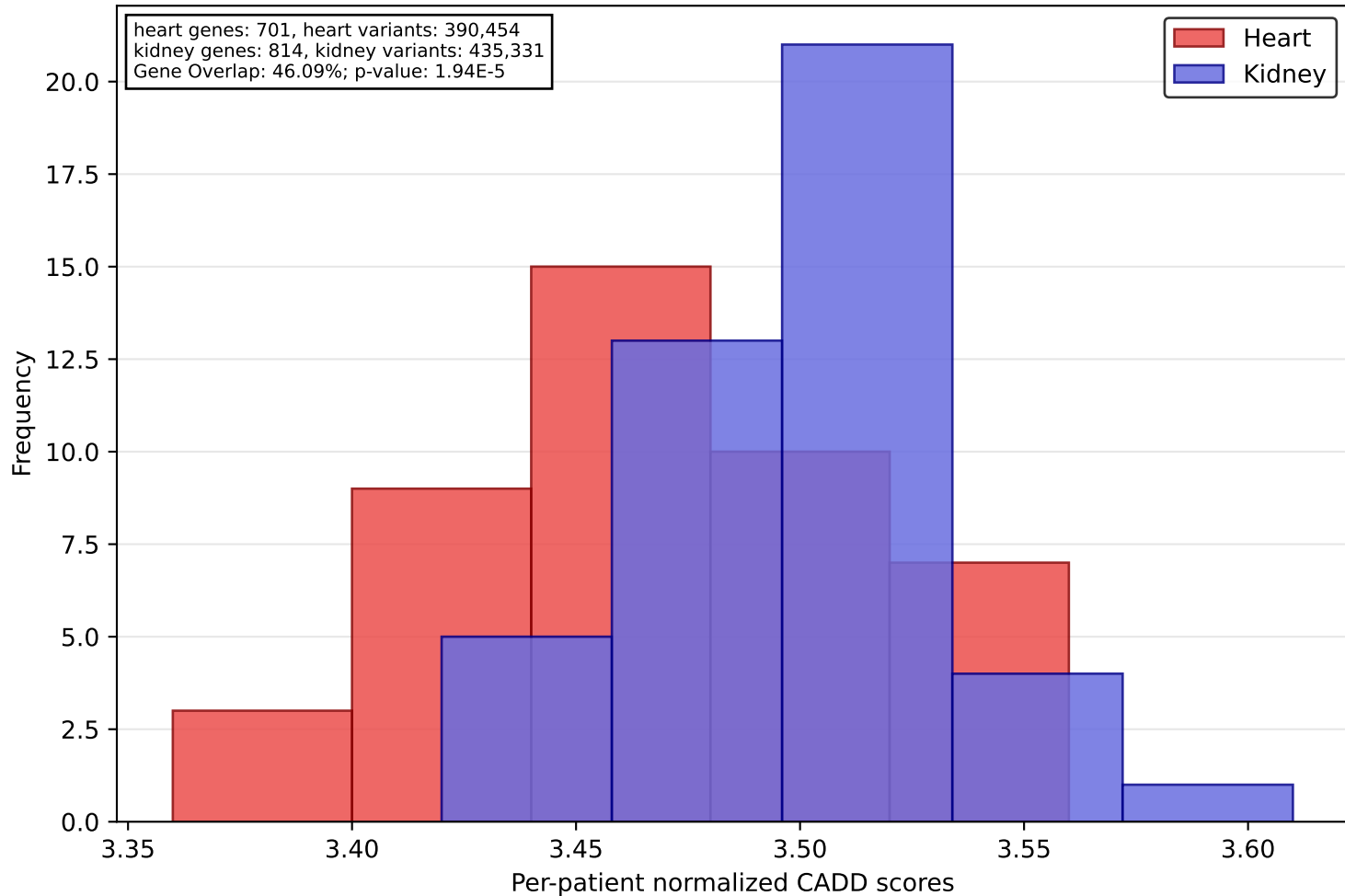
Top 0.1% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



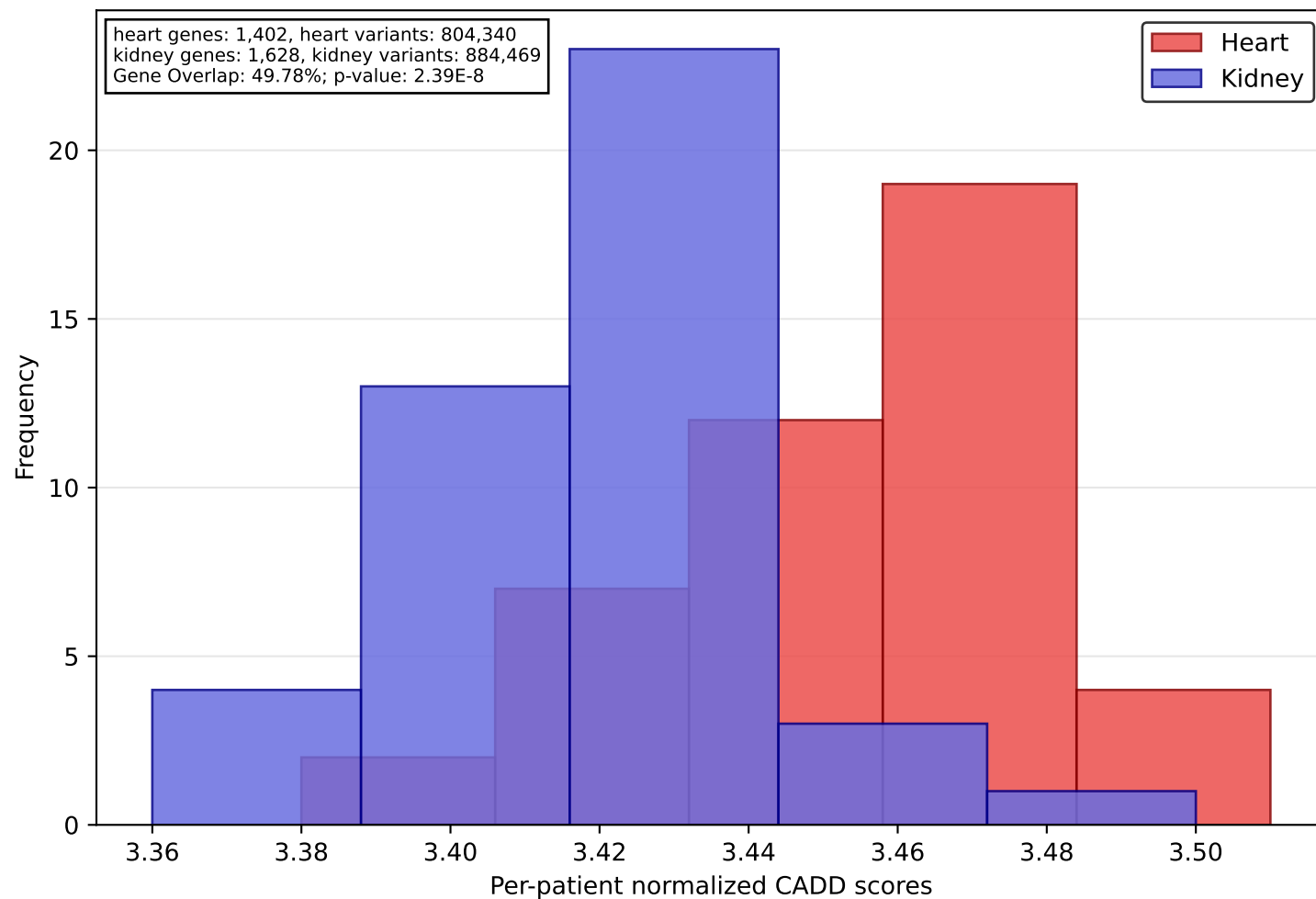
Top 1.0% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



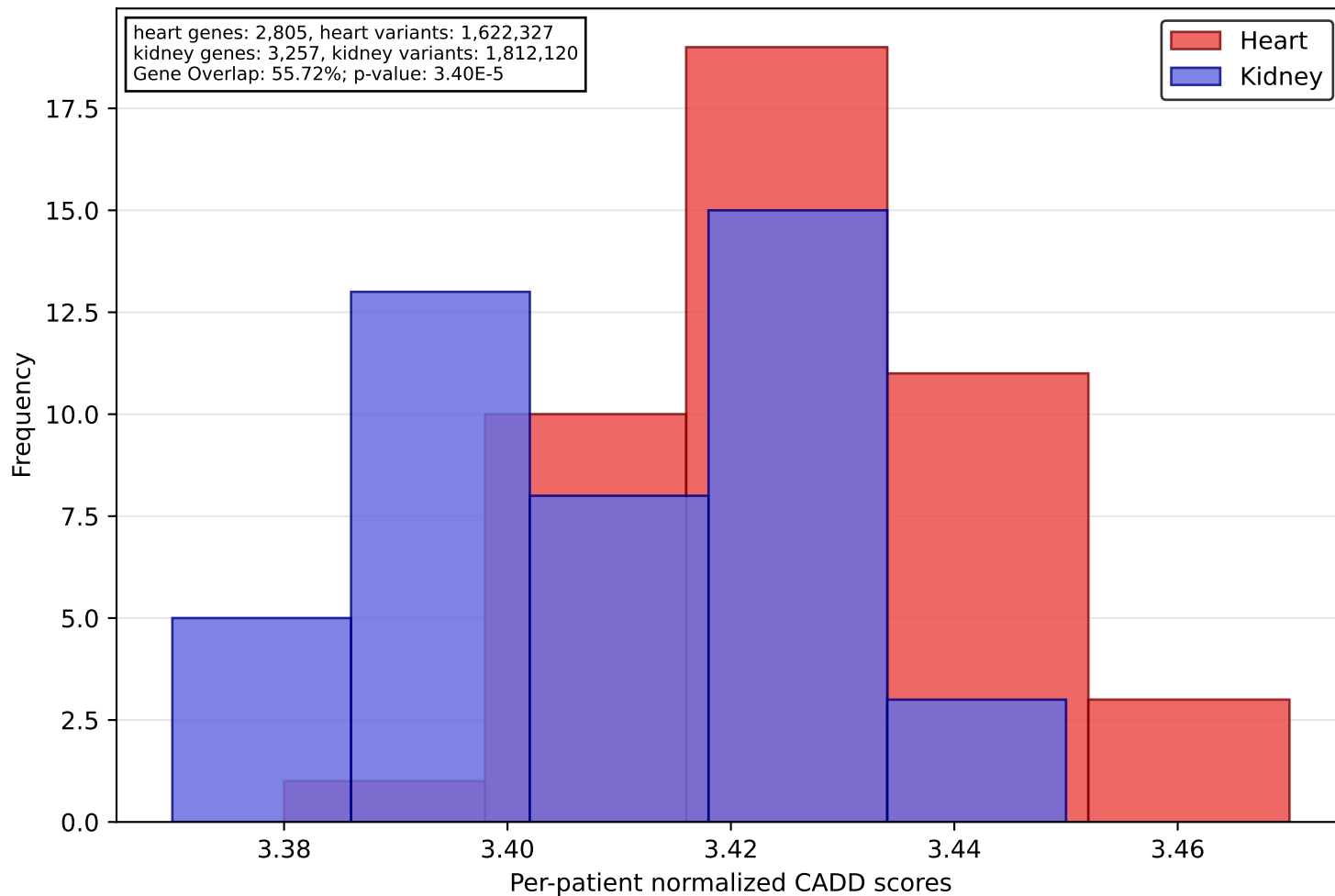
Top 2.5% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



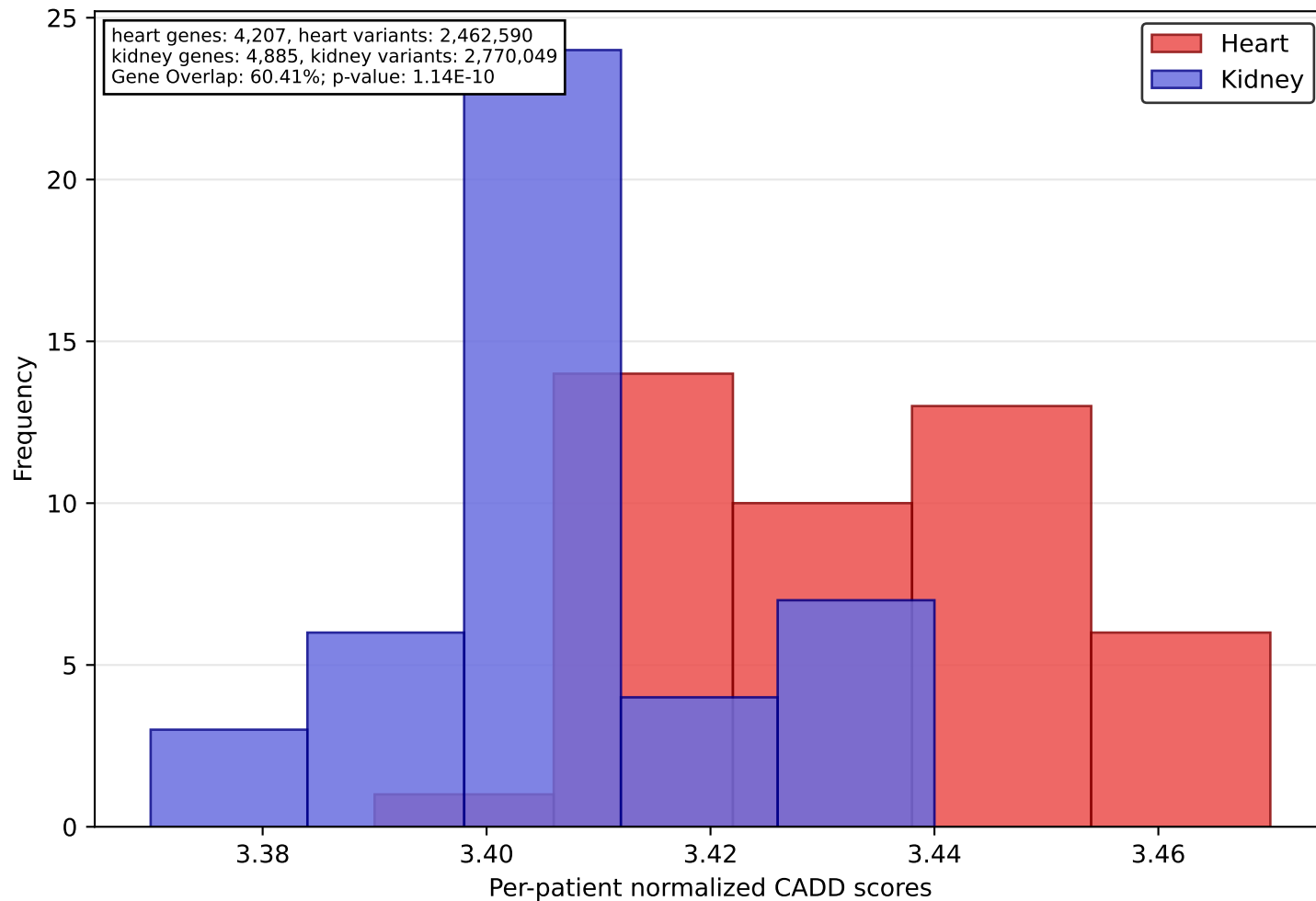
Top 5.0% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



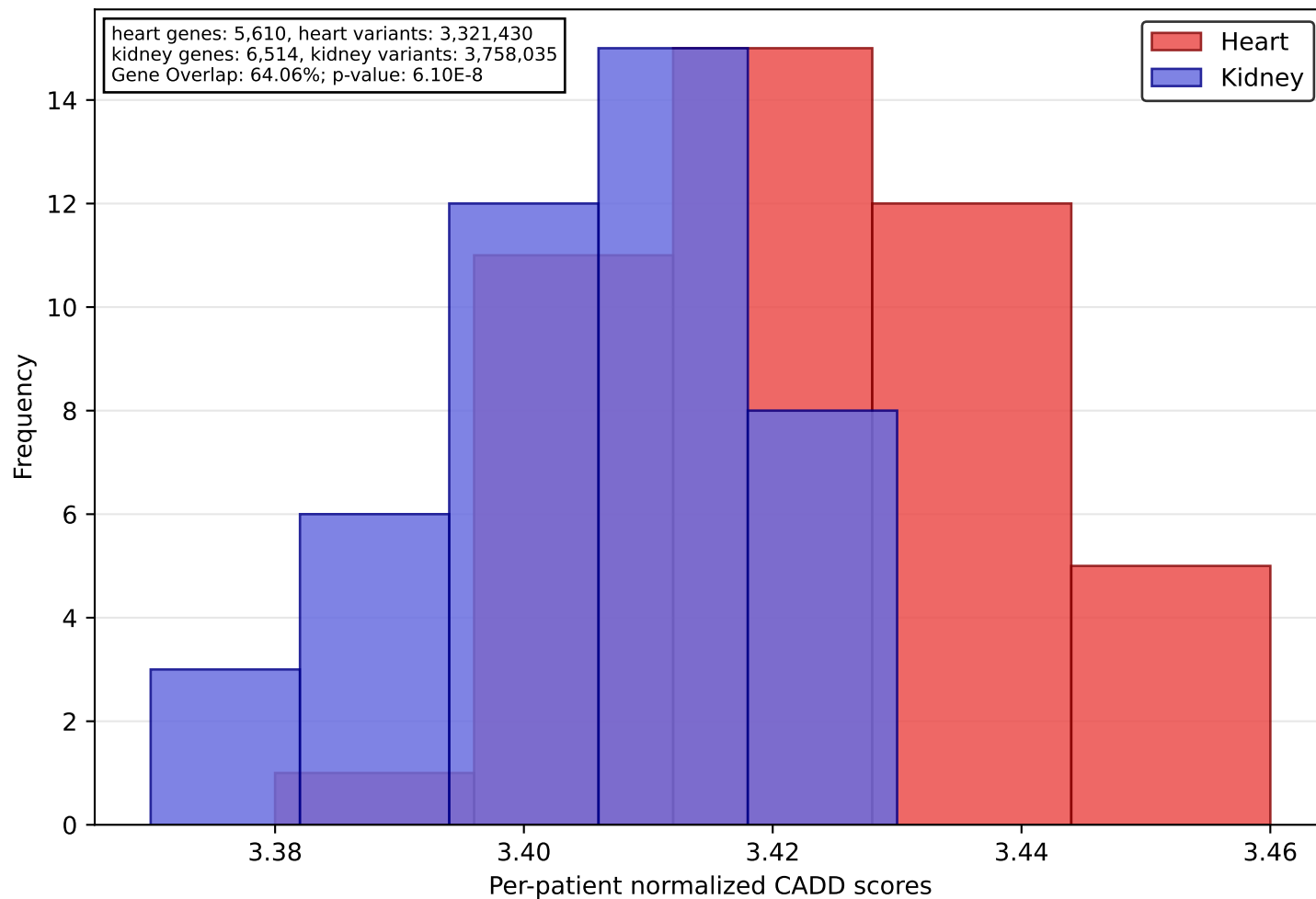
Top 10.0% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



Top 15.0% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



Top 20.0% expressed genes from GTEx heart and kidney data
were used to filter variants from the KUTD cohort



Summary of KUTD analysis using GTEx tissue datasets

