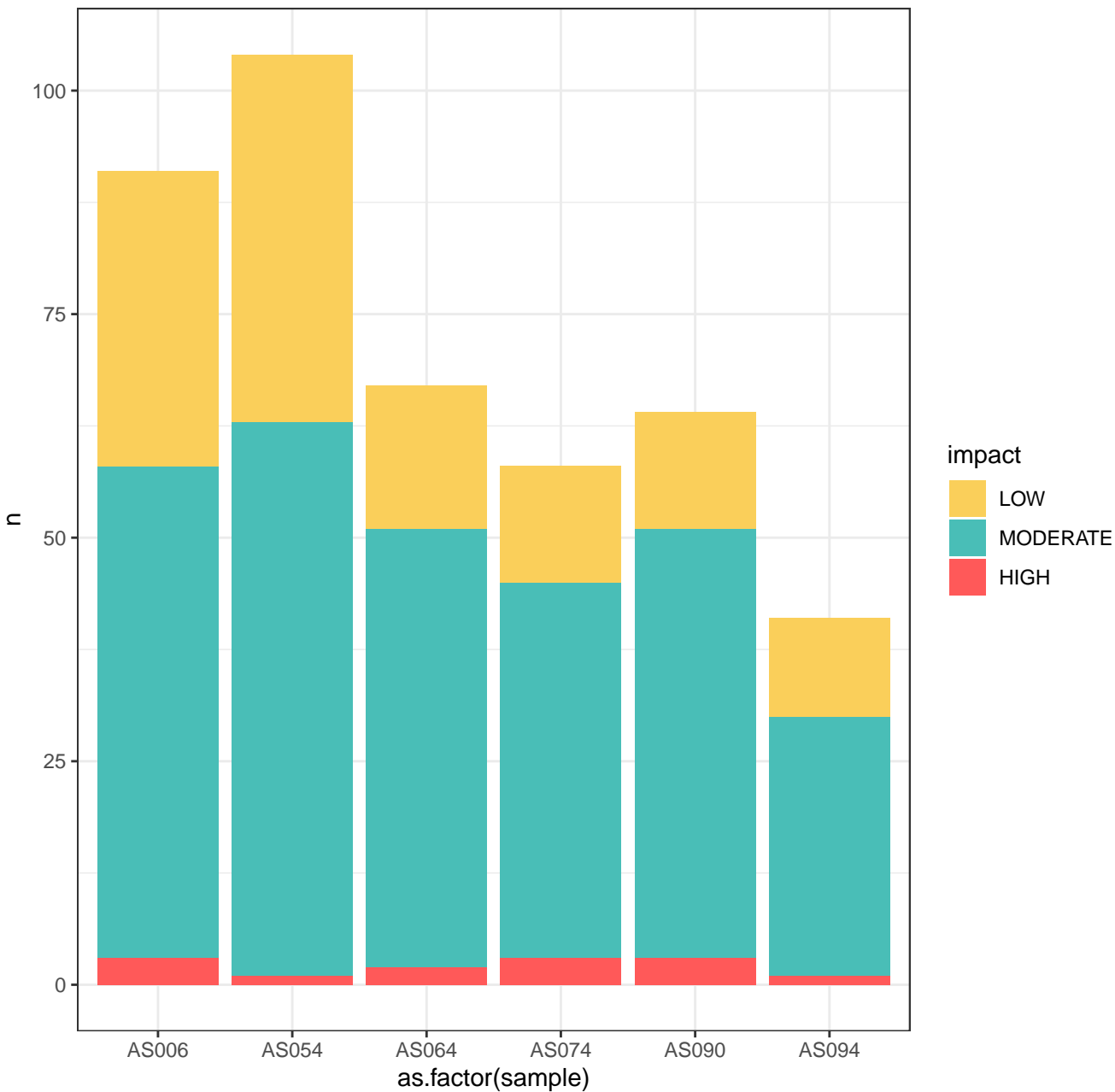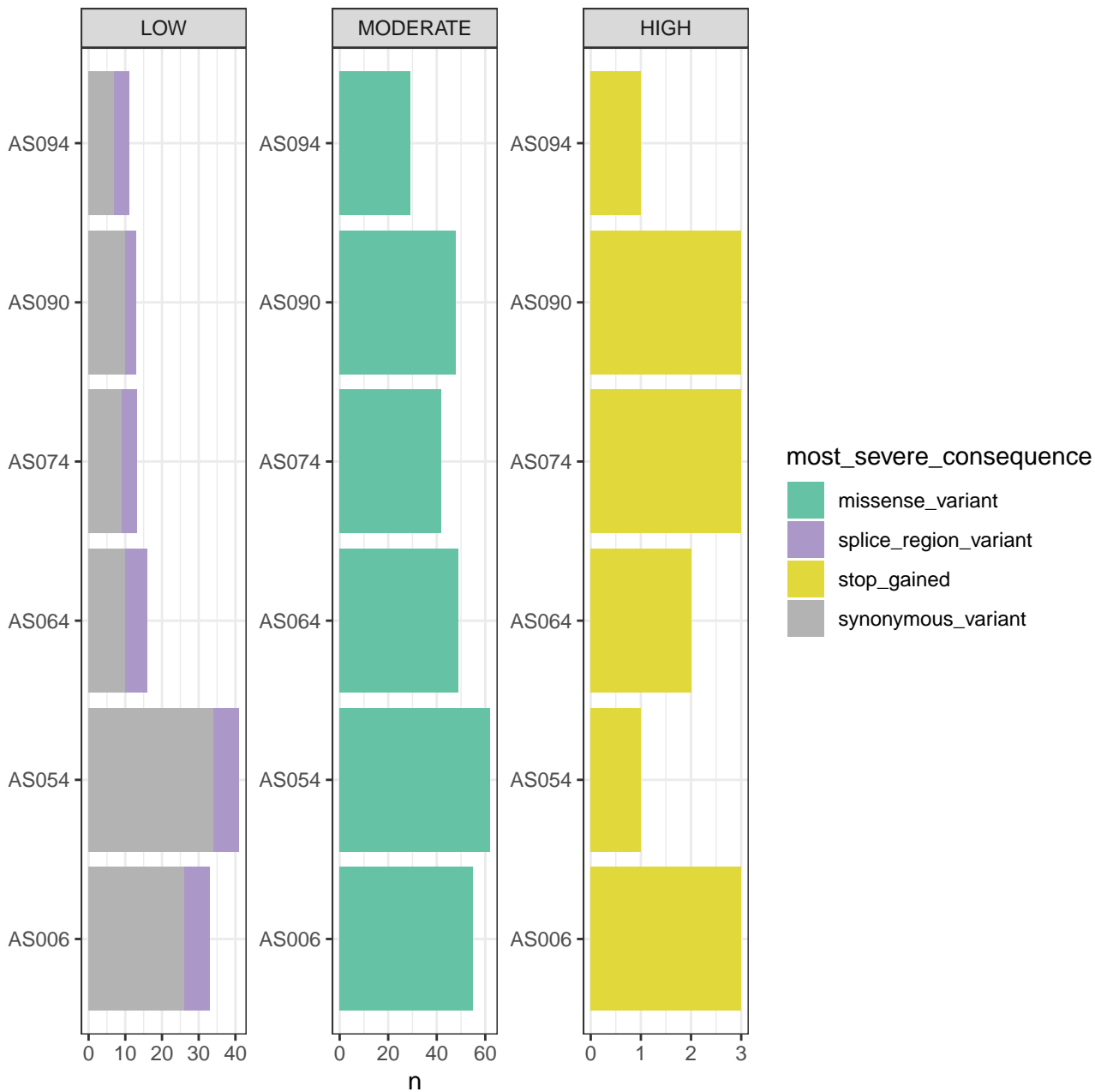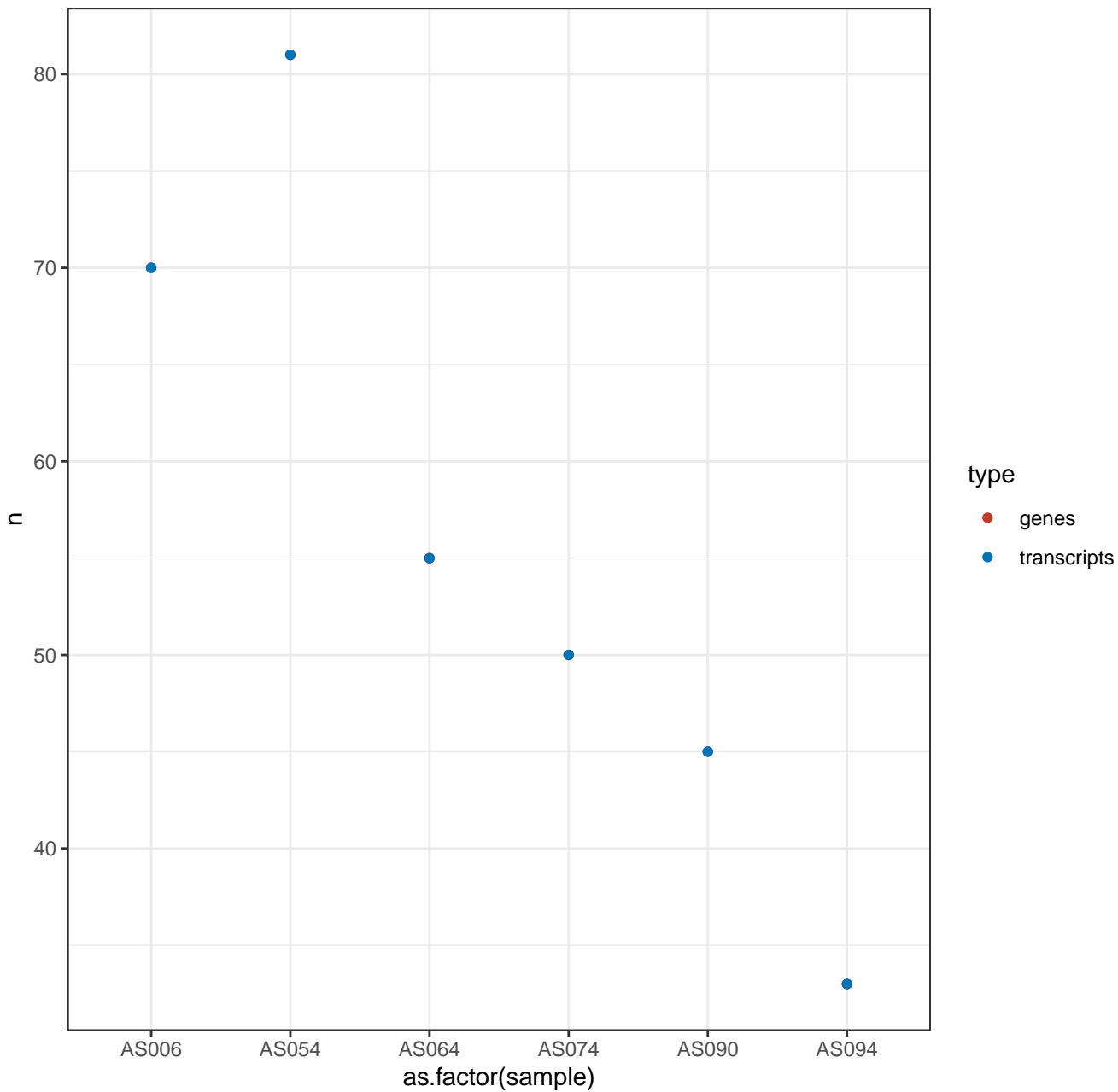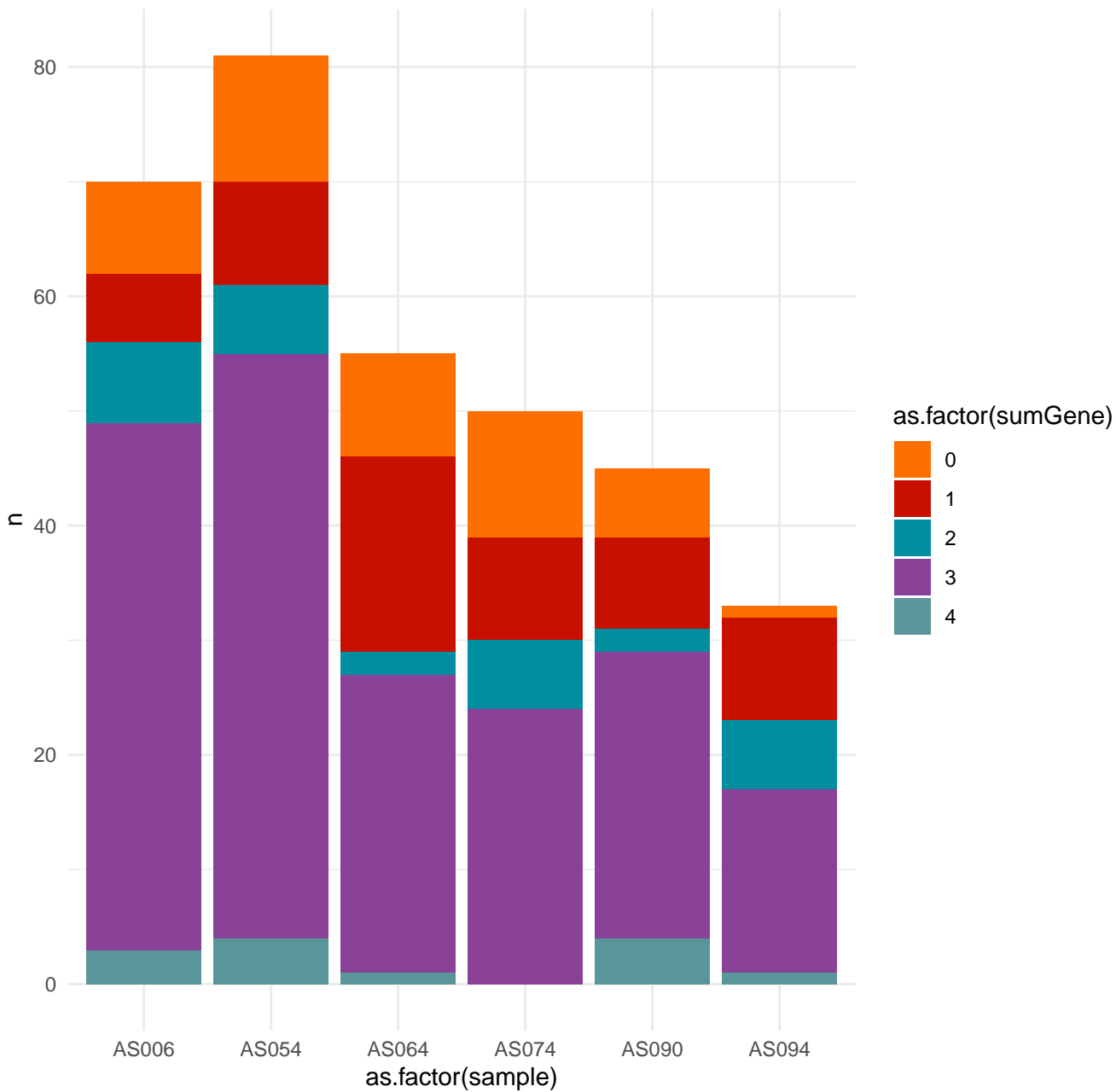f4 – Unique variants per sample
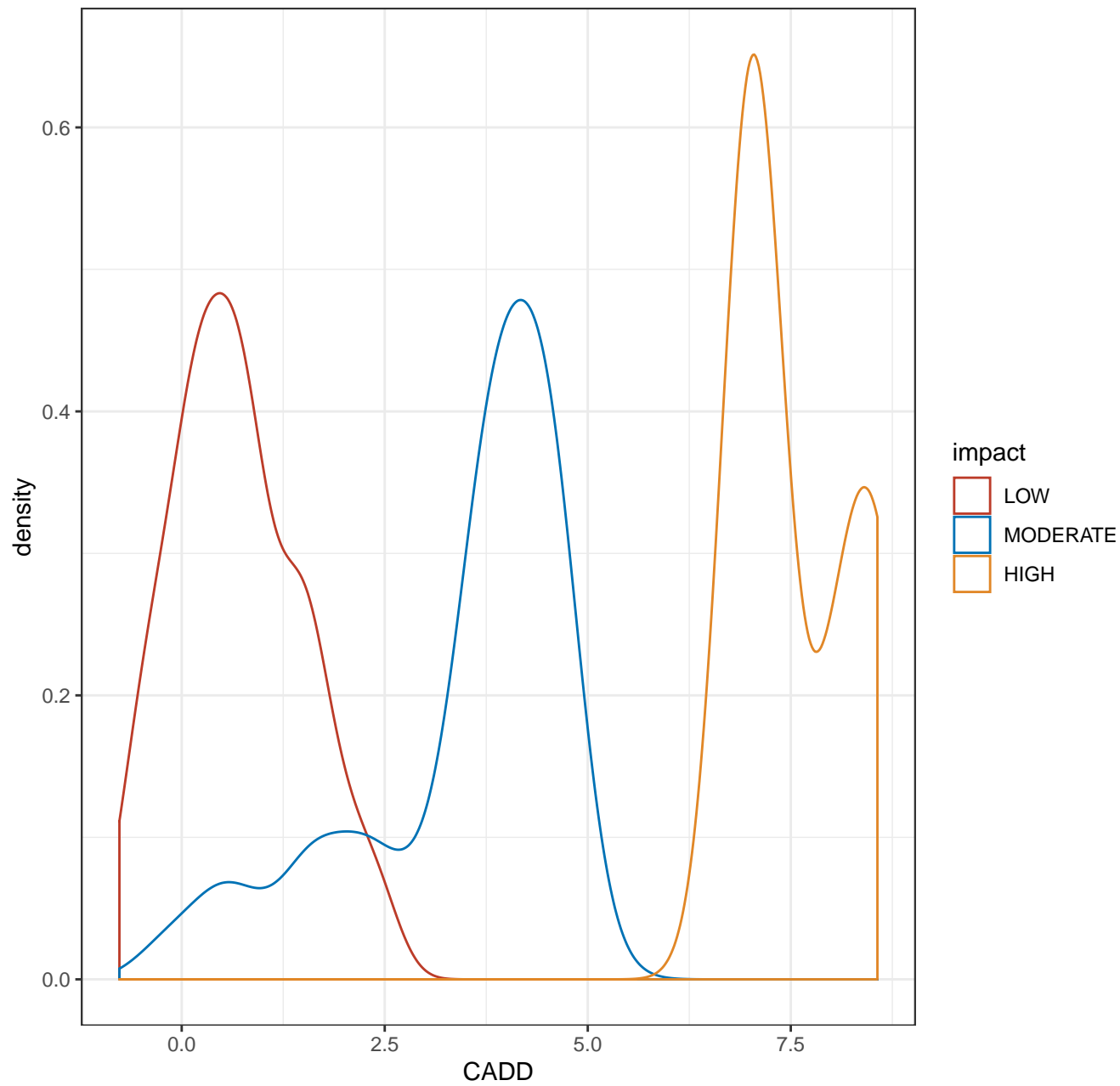
f4 – most_severe_consequence
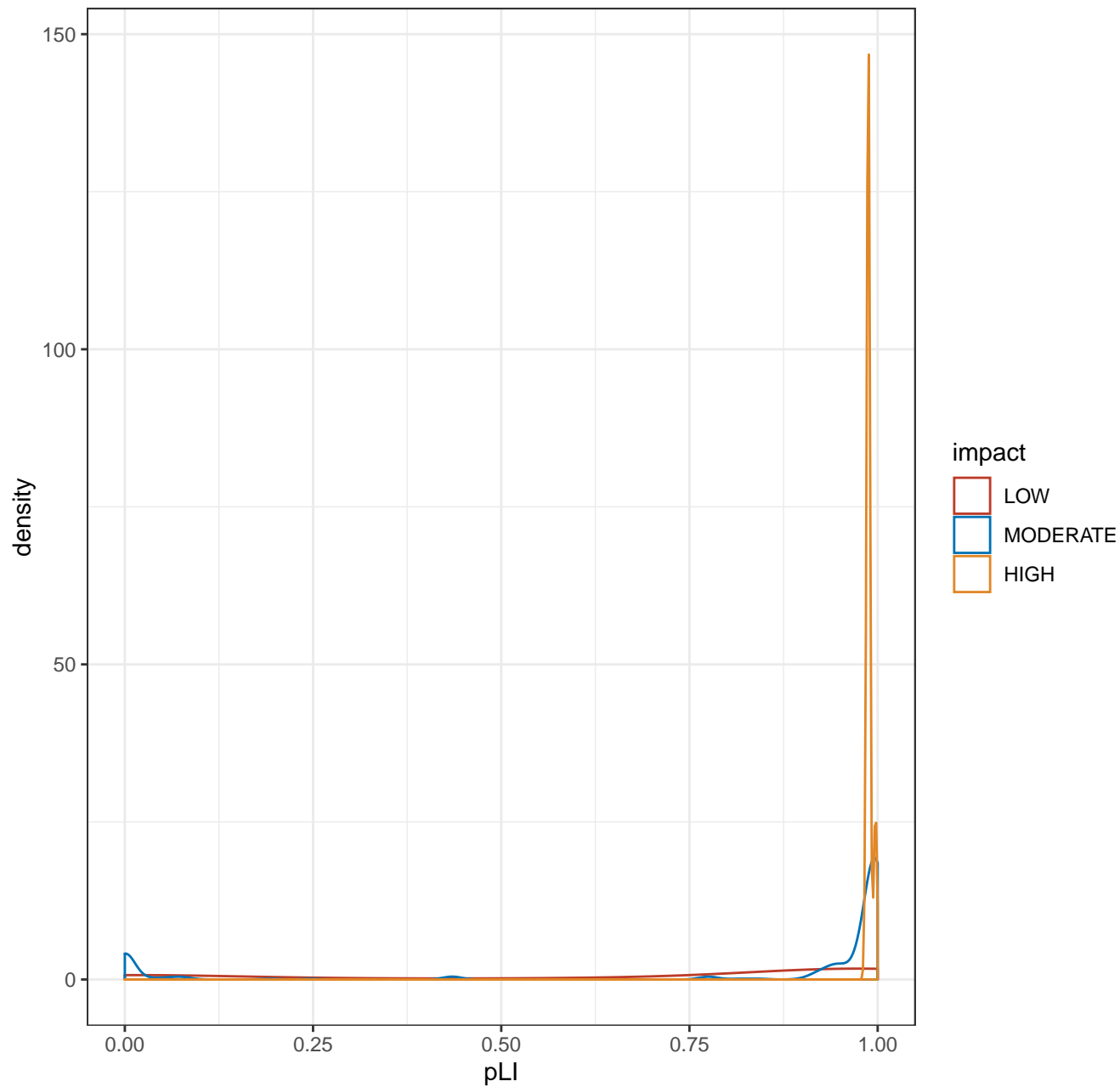
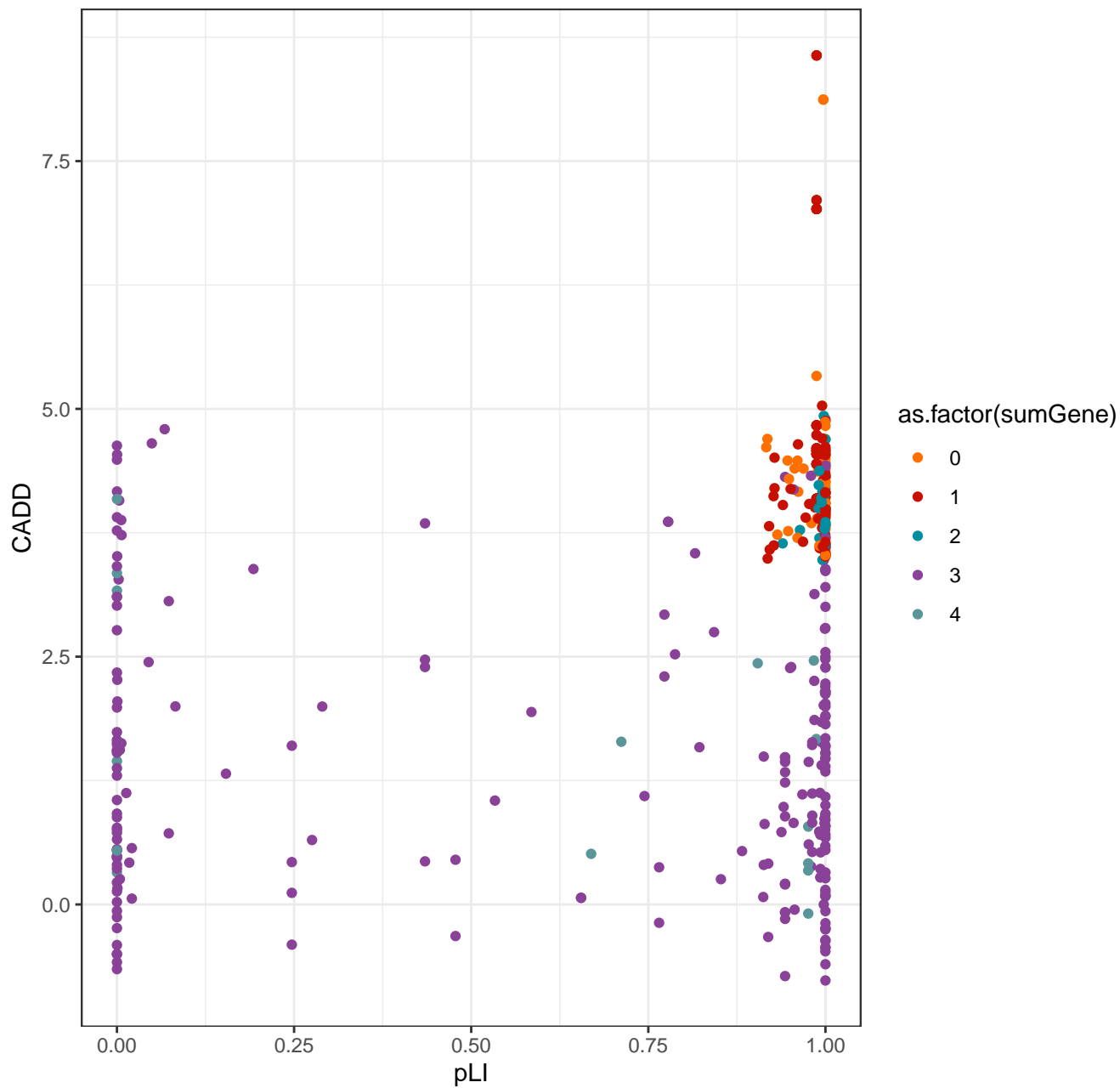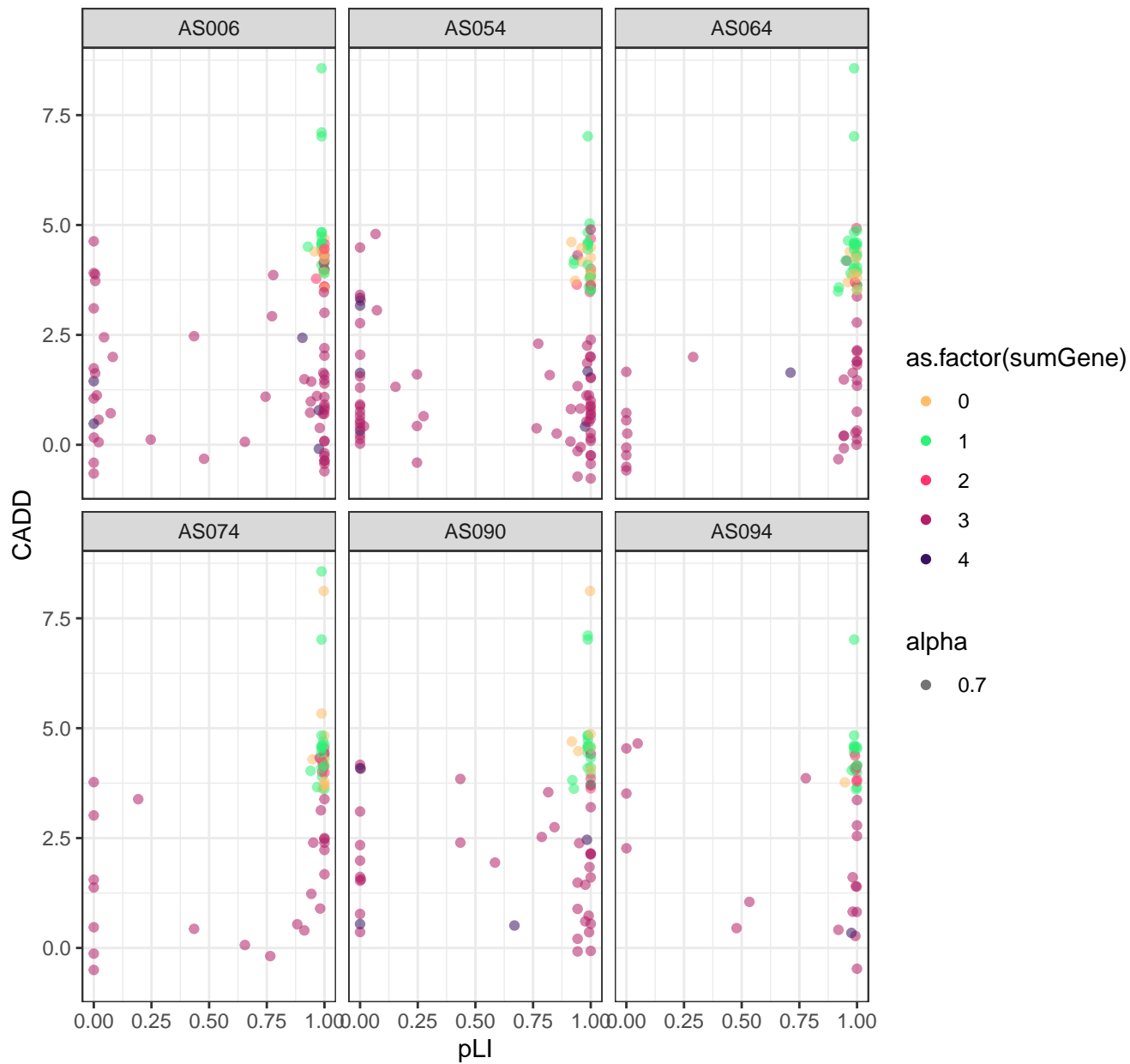f4 – Genes and transcripts

f4 – Genes in lists

f4 – CADD

f4 − pLI

f4 – pLI CADD sumGene

f4pLI, CADD, gene lists
unique sites with data =373

f4 – Genes shared among samples