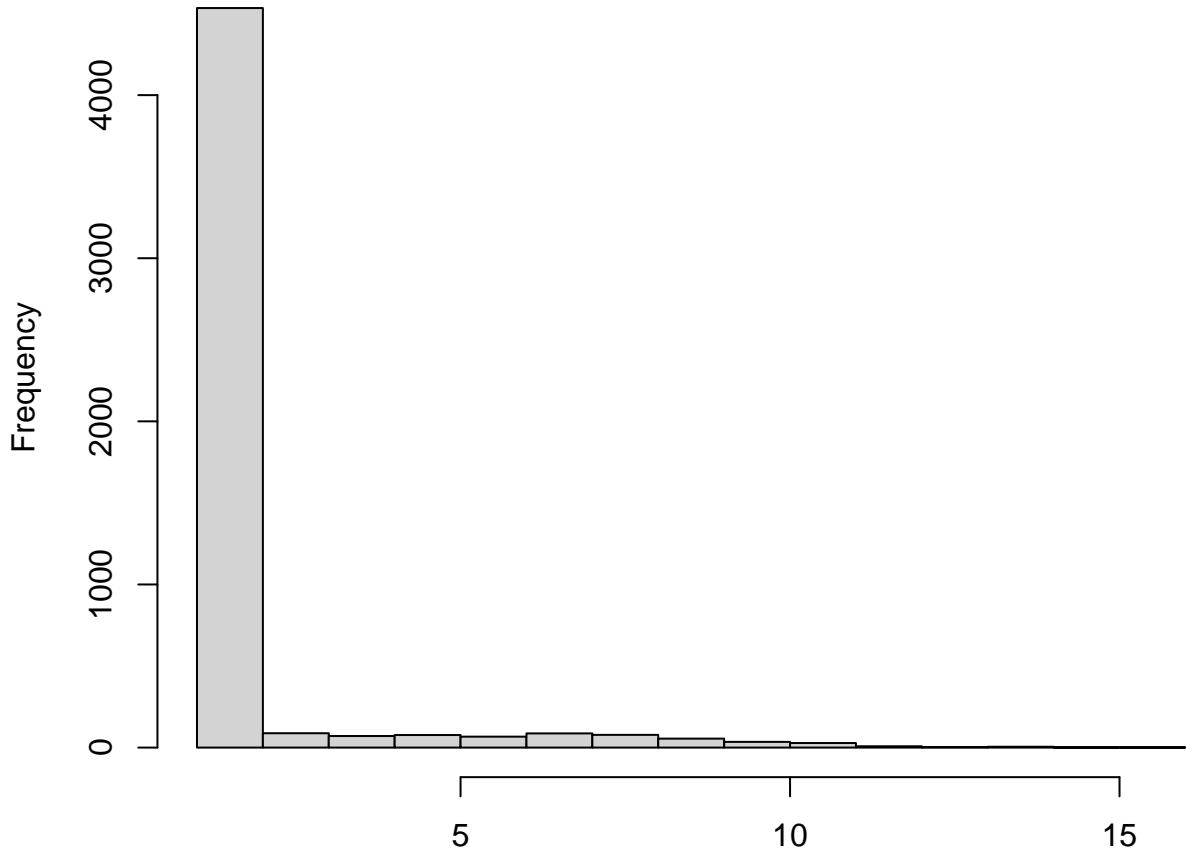
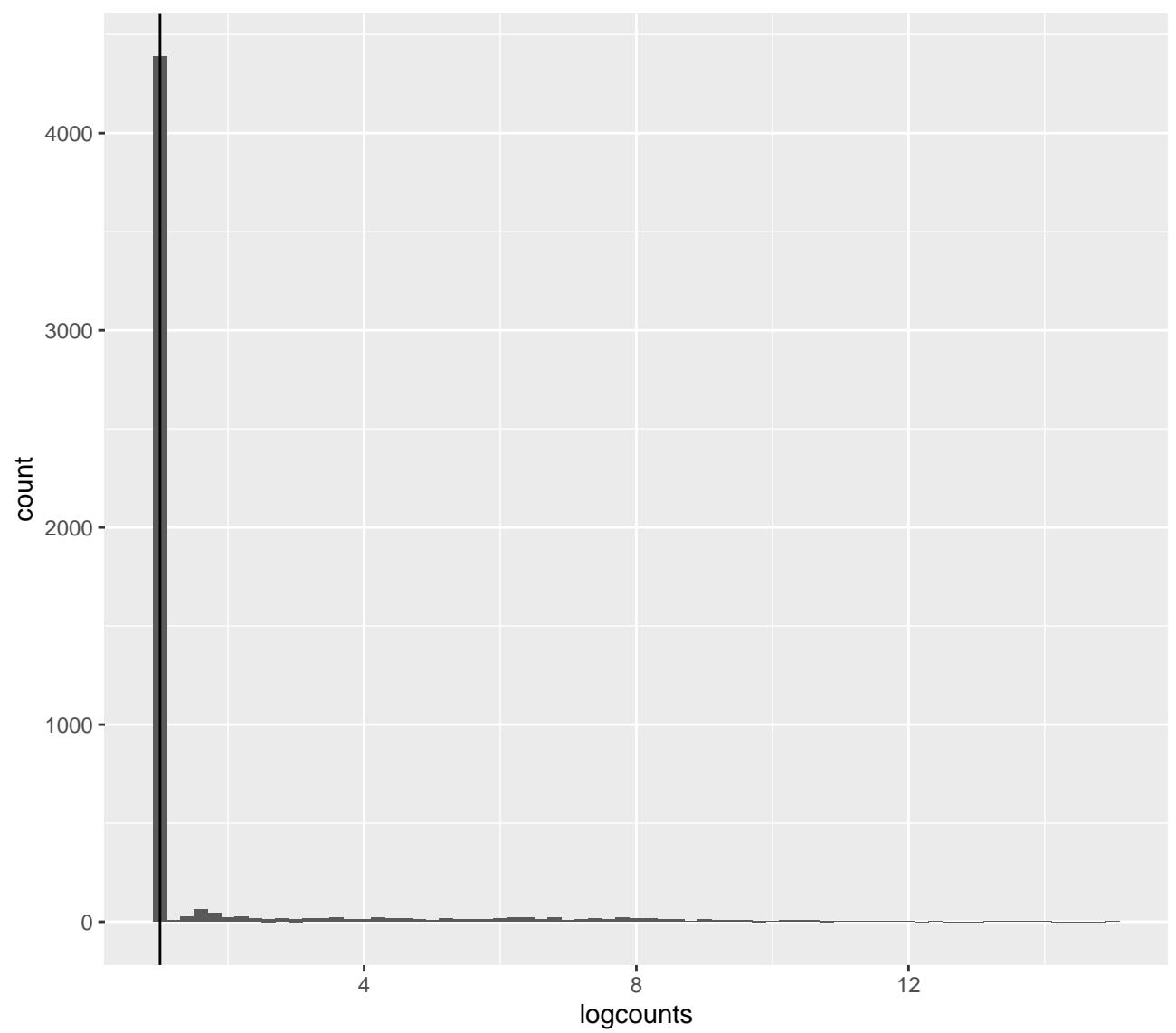
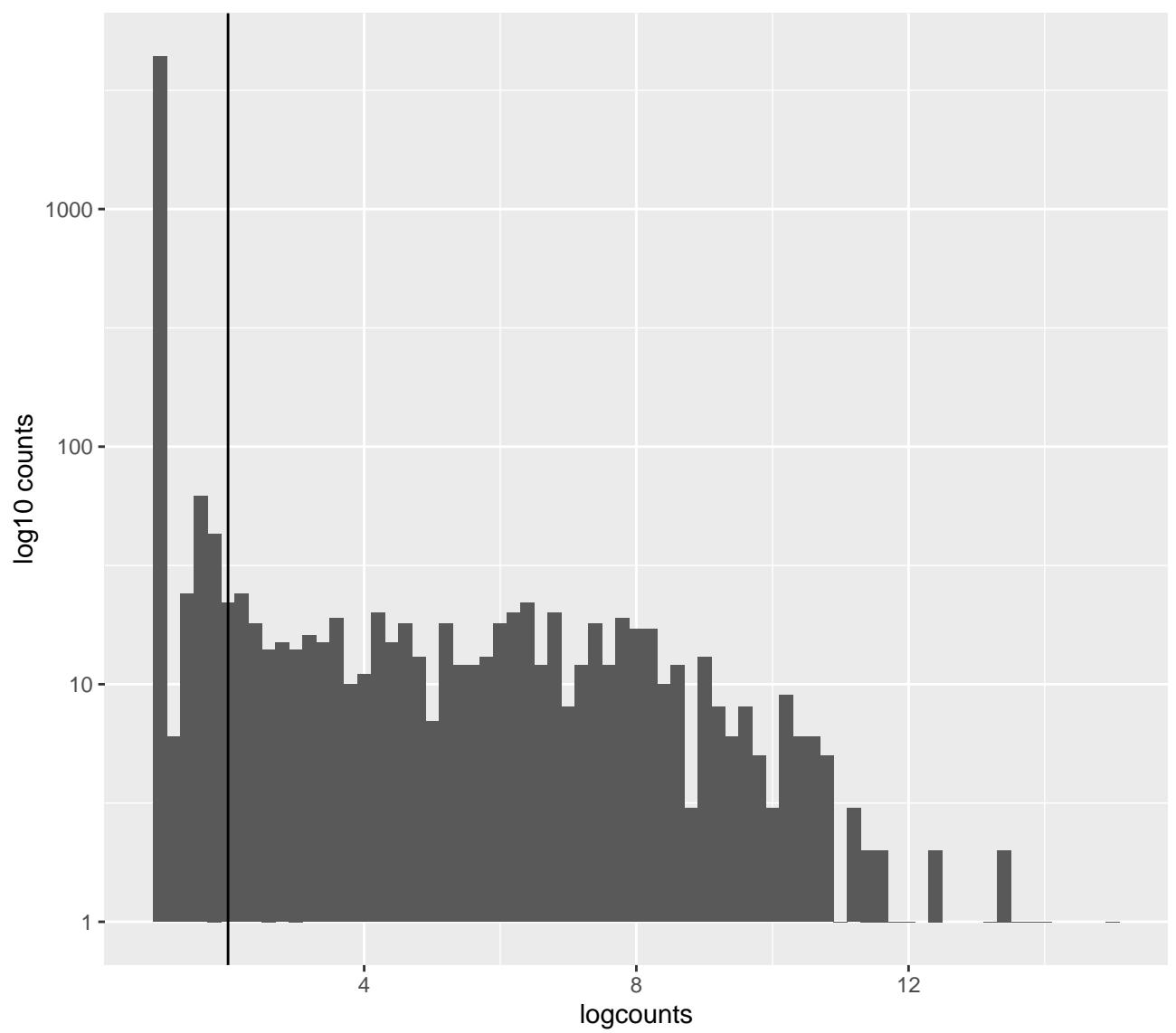


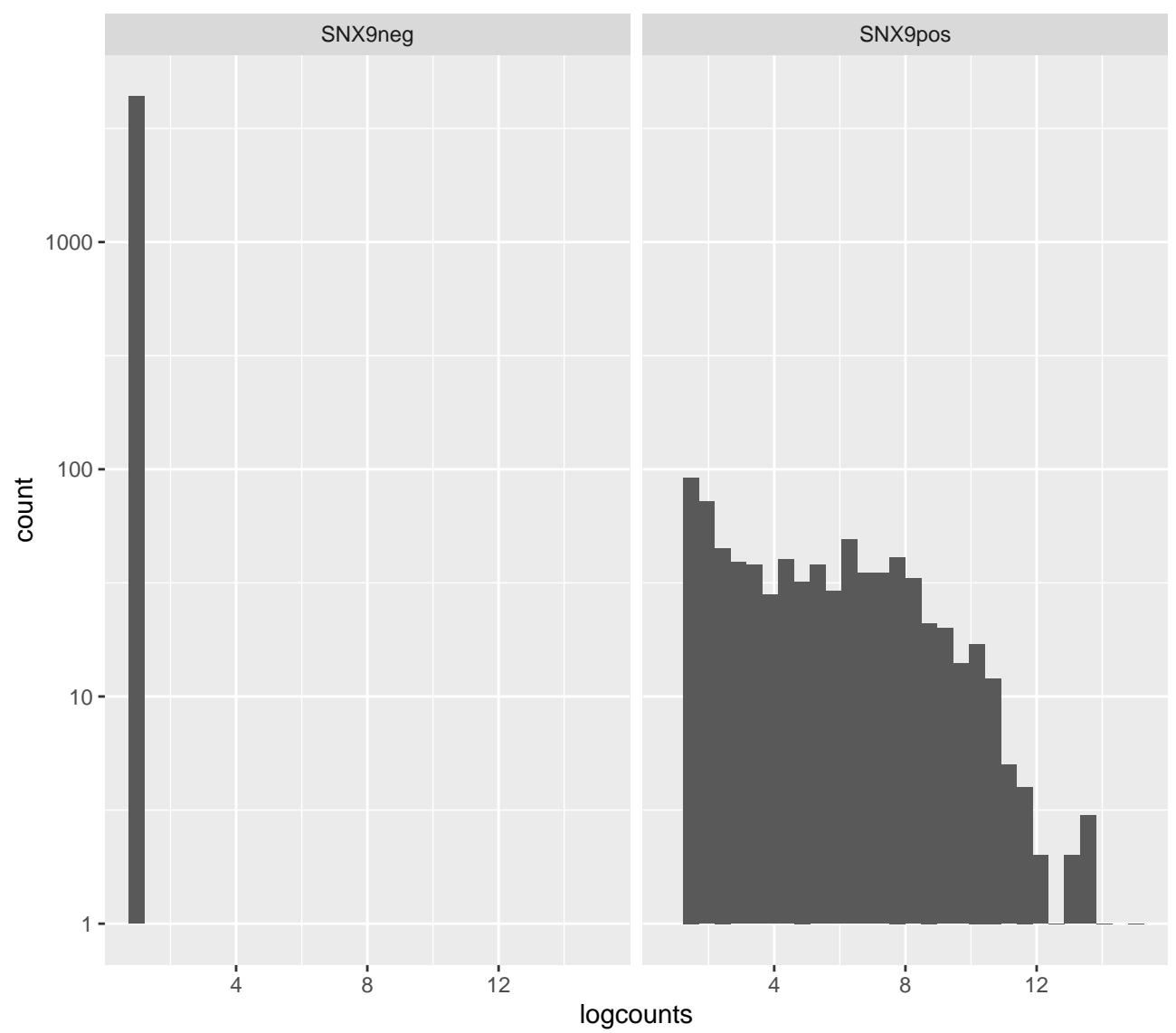
histogram of logcounts(sce_CD8T)[rowData(sce_CD8T)\$GeneSymbol == "S



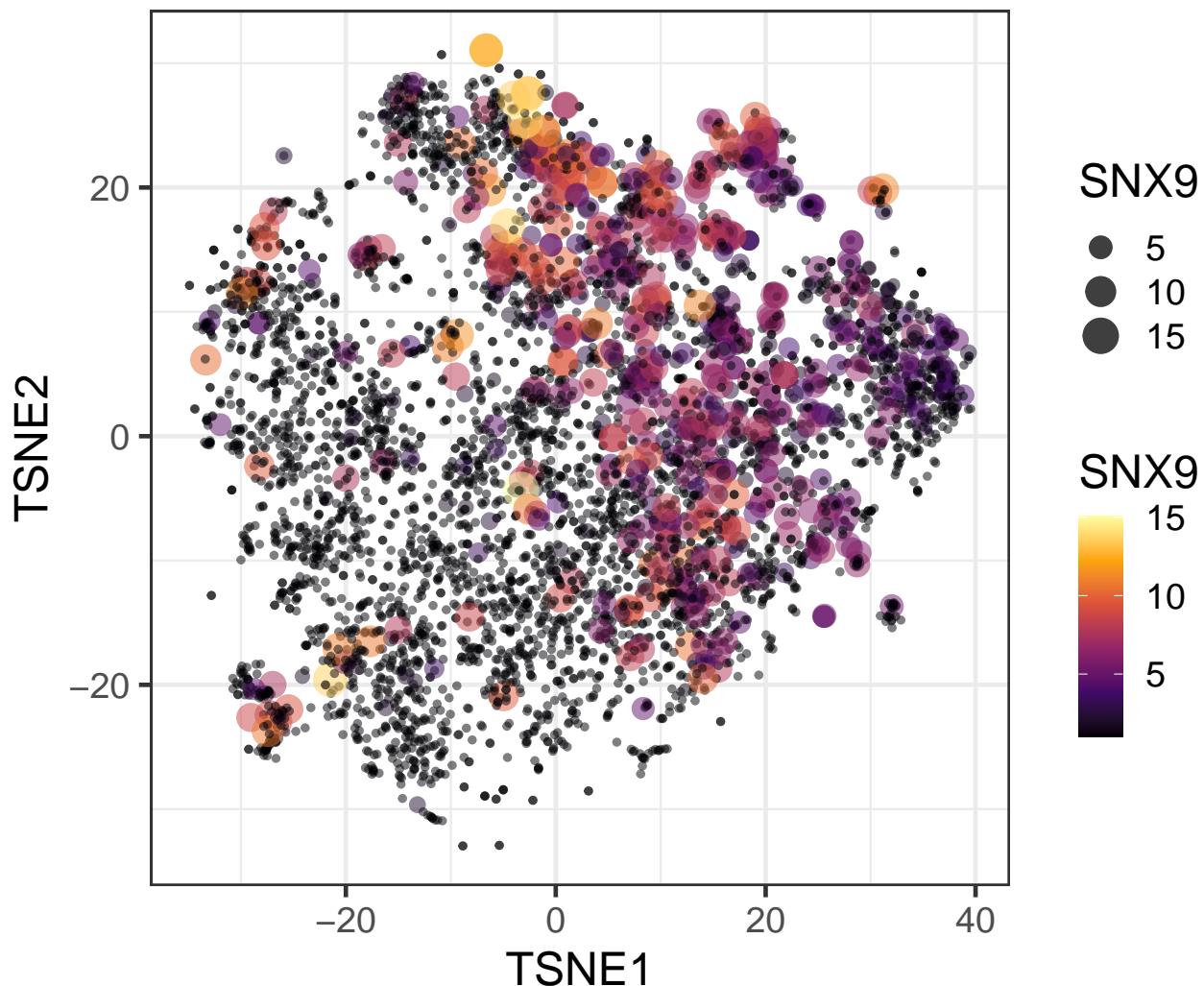
logcounts(sce_CD8T)[rowData(sce_CD8T)\$GeneSymbol == "SNX9",]



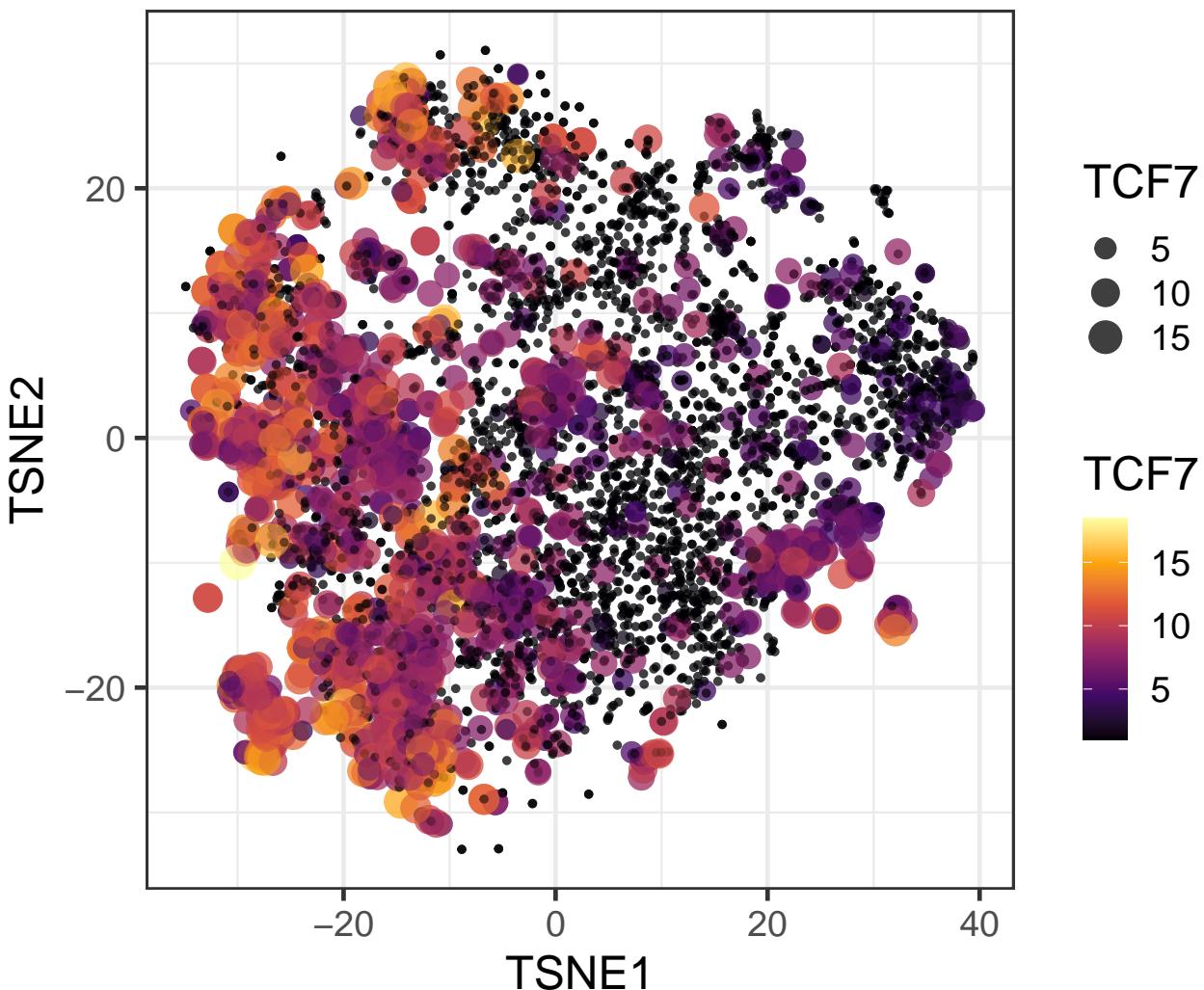




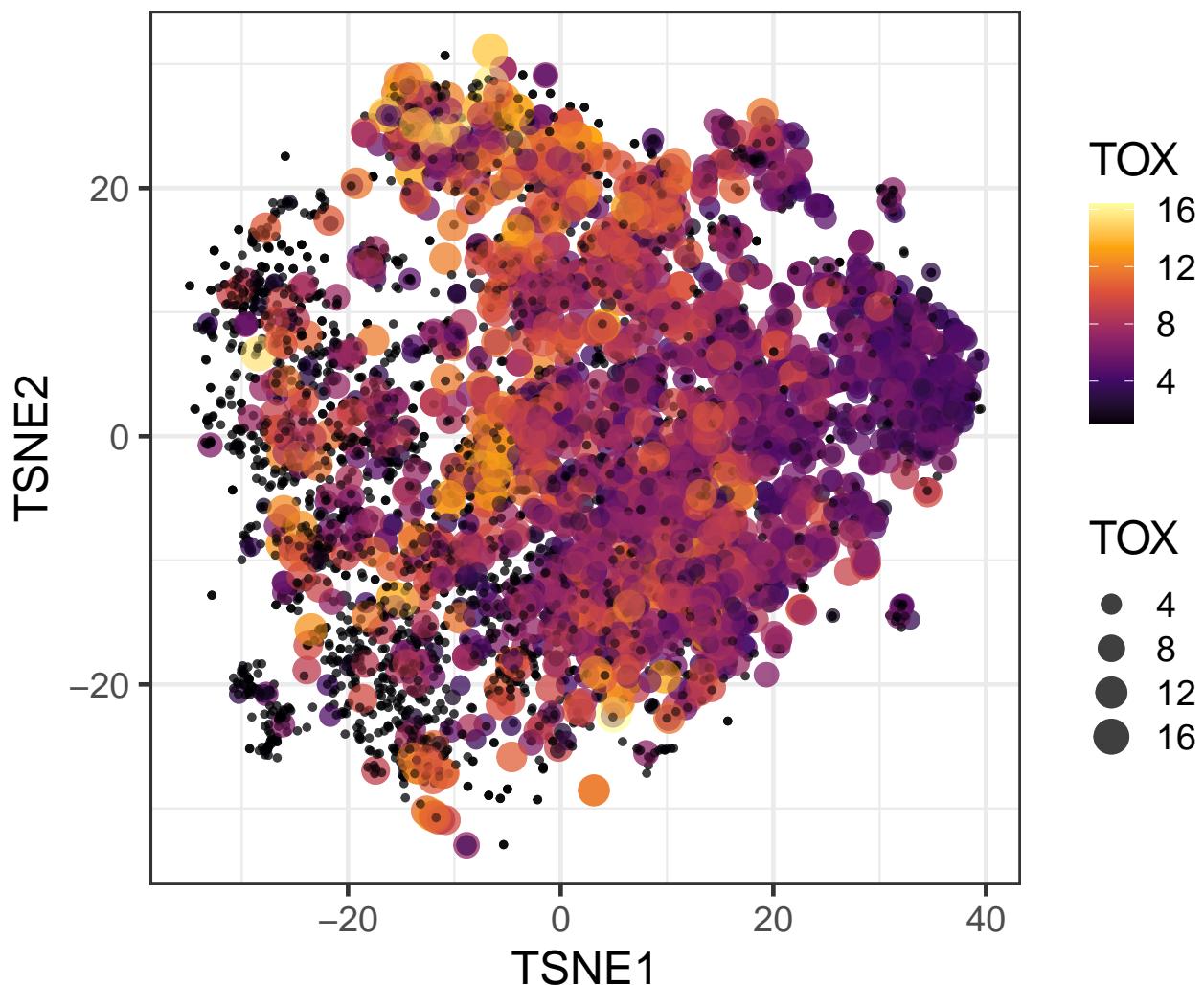
SNX9 in CD8+ T cells



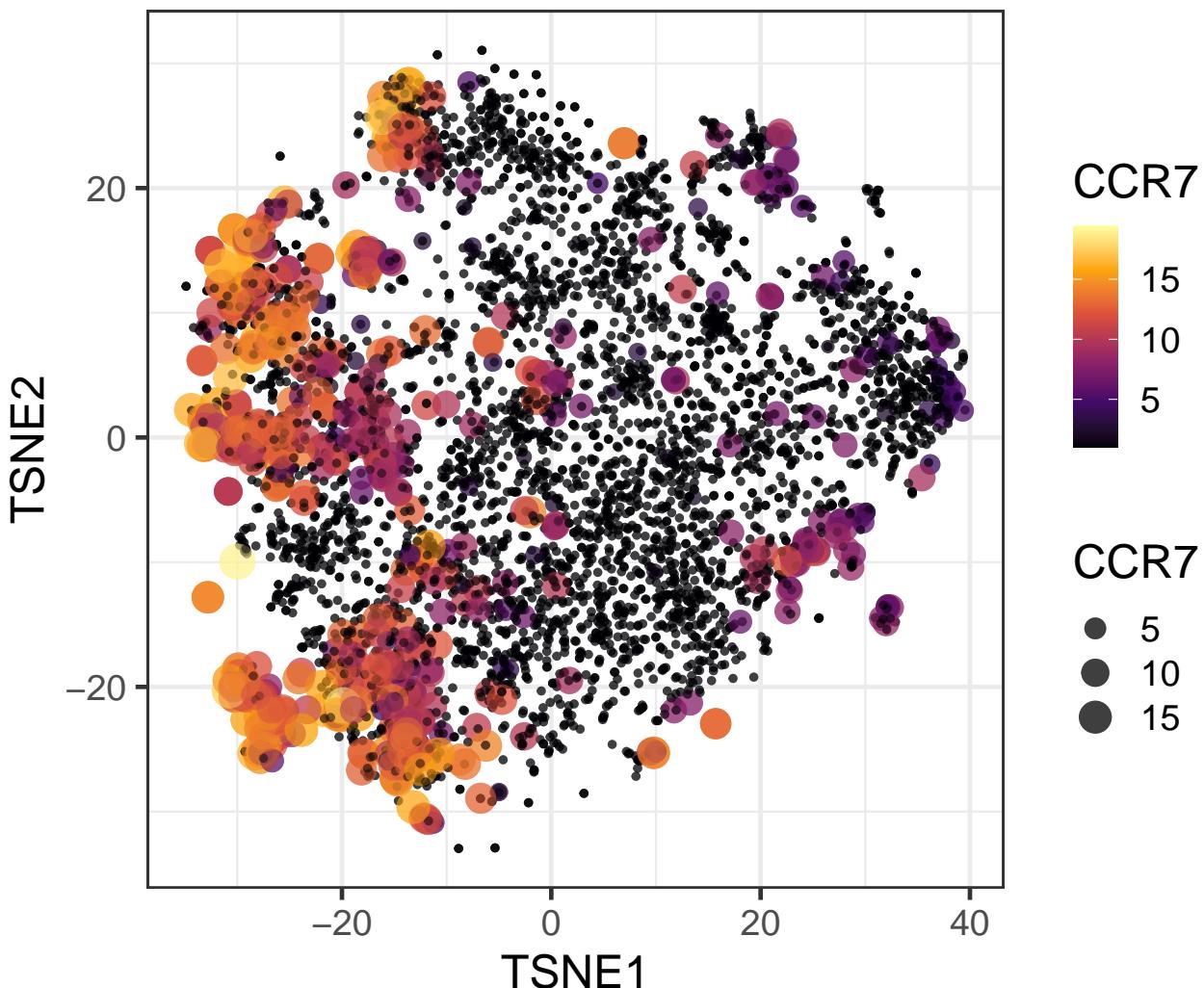
TCF7 in CD8+ T cells



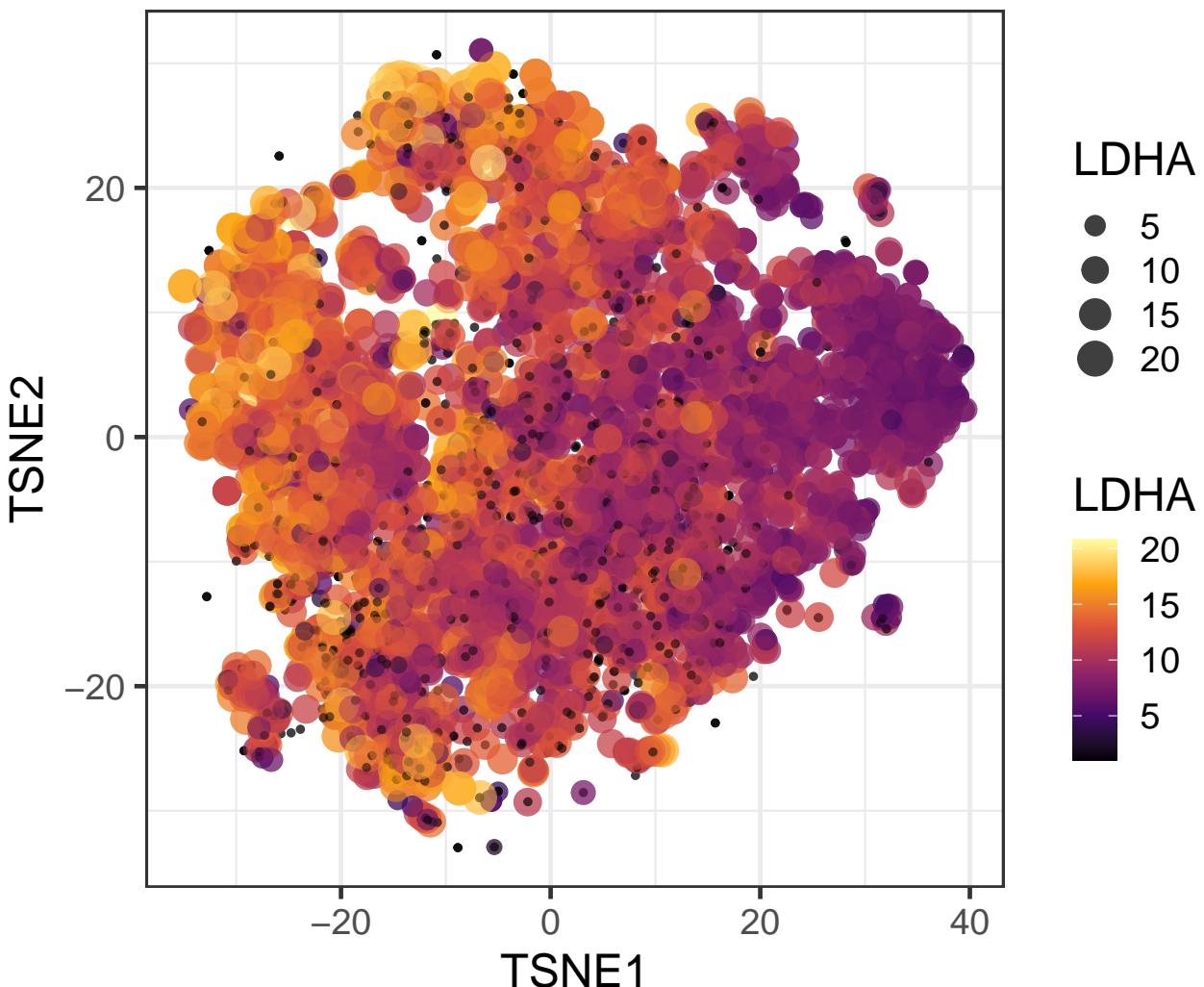
TOX in CD8+ T cells



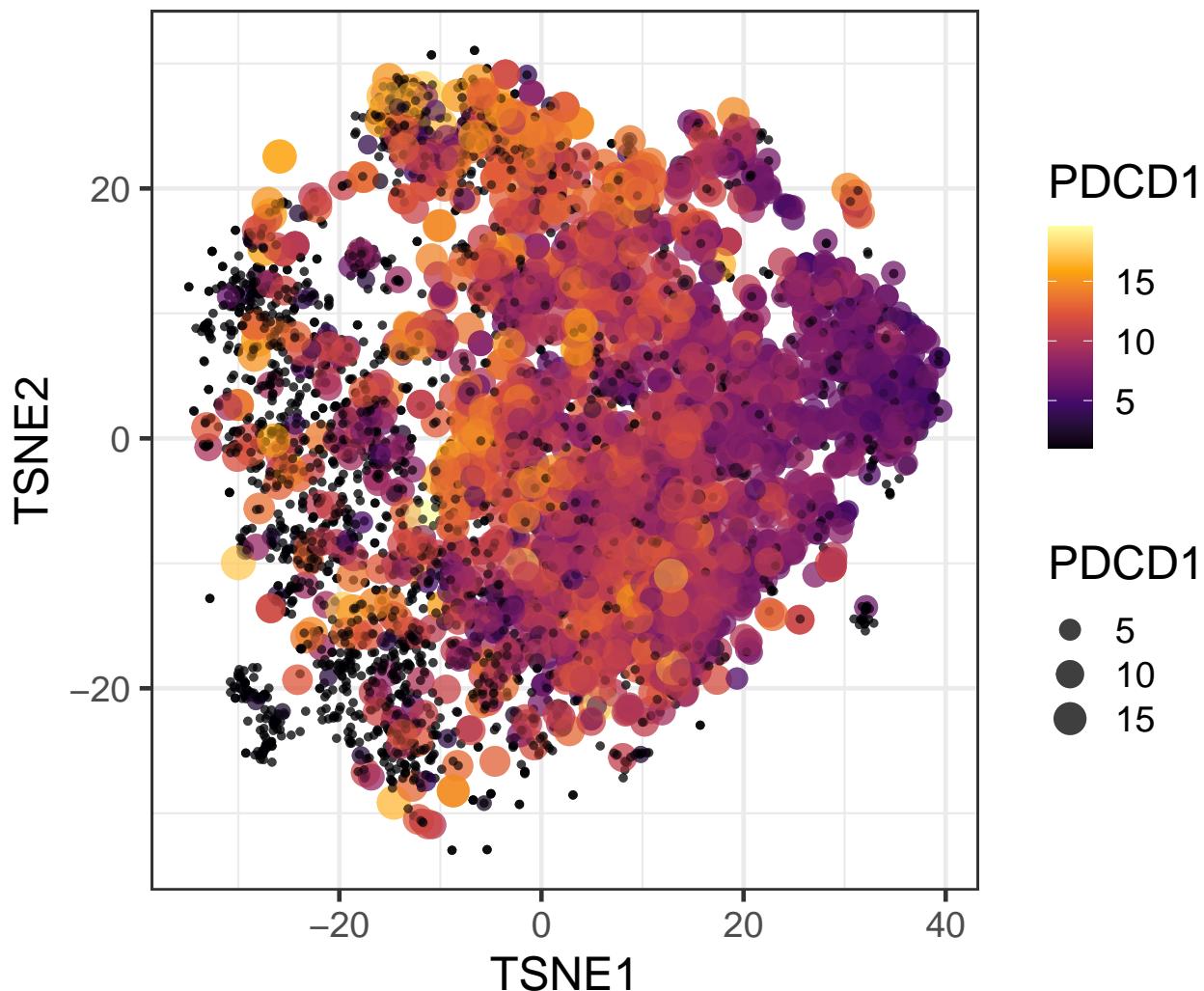
CCR7 in CD8+ T cells



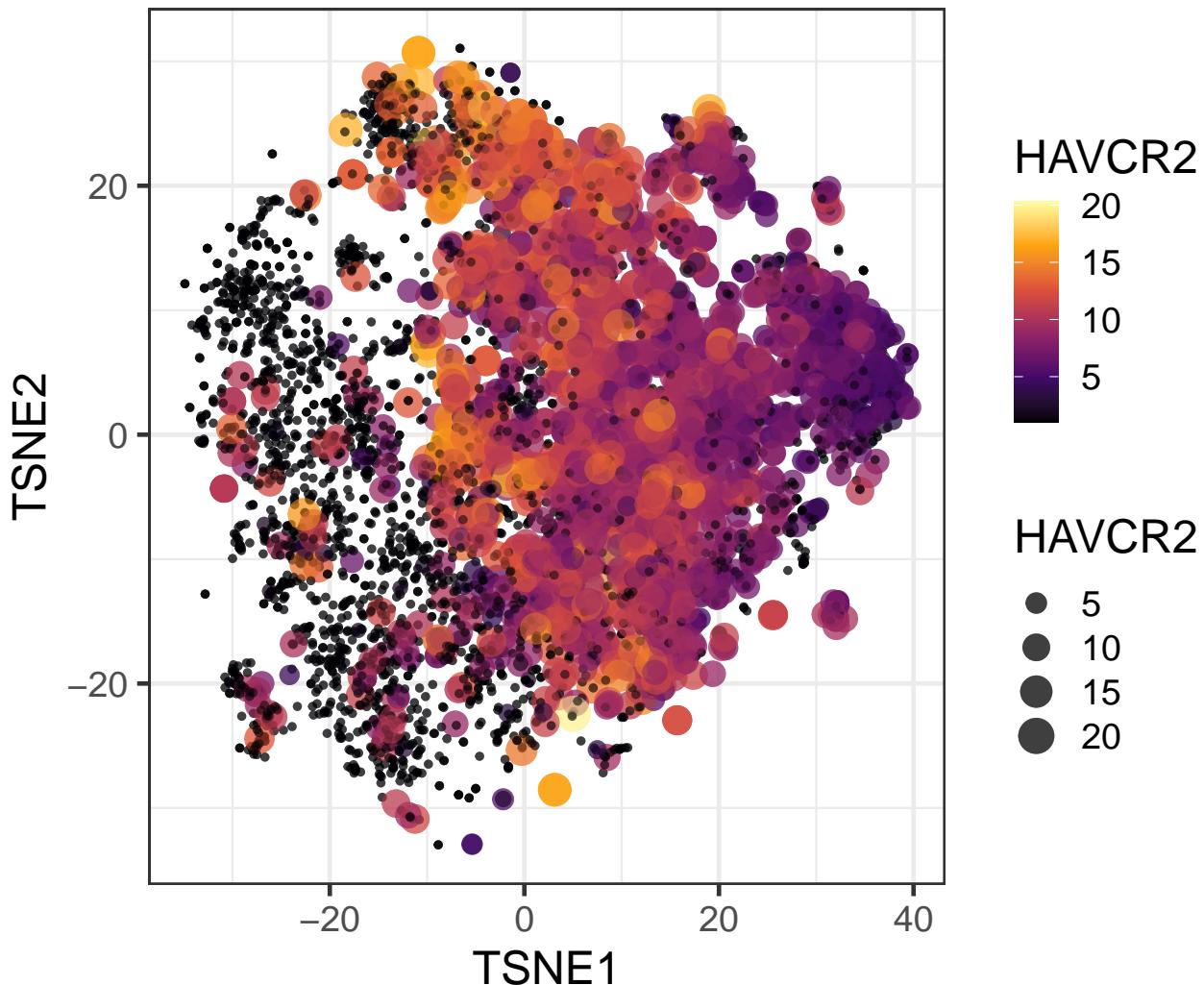
LDHA in CD8+ T cells



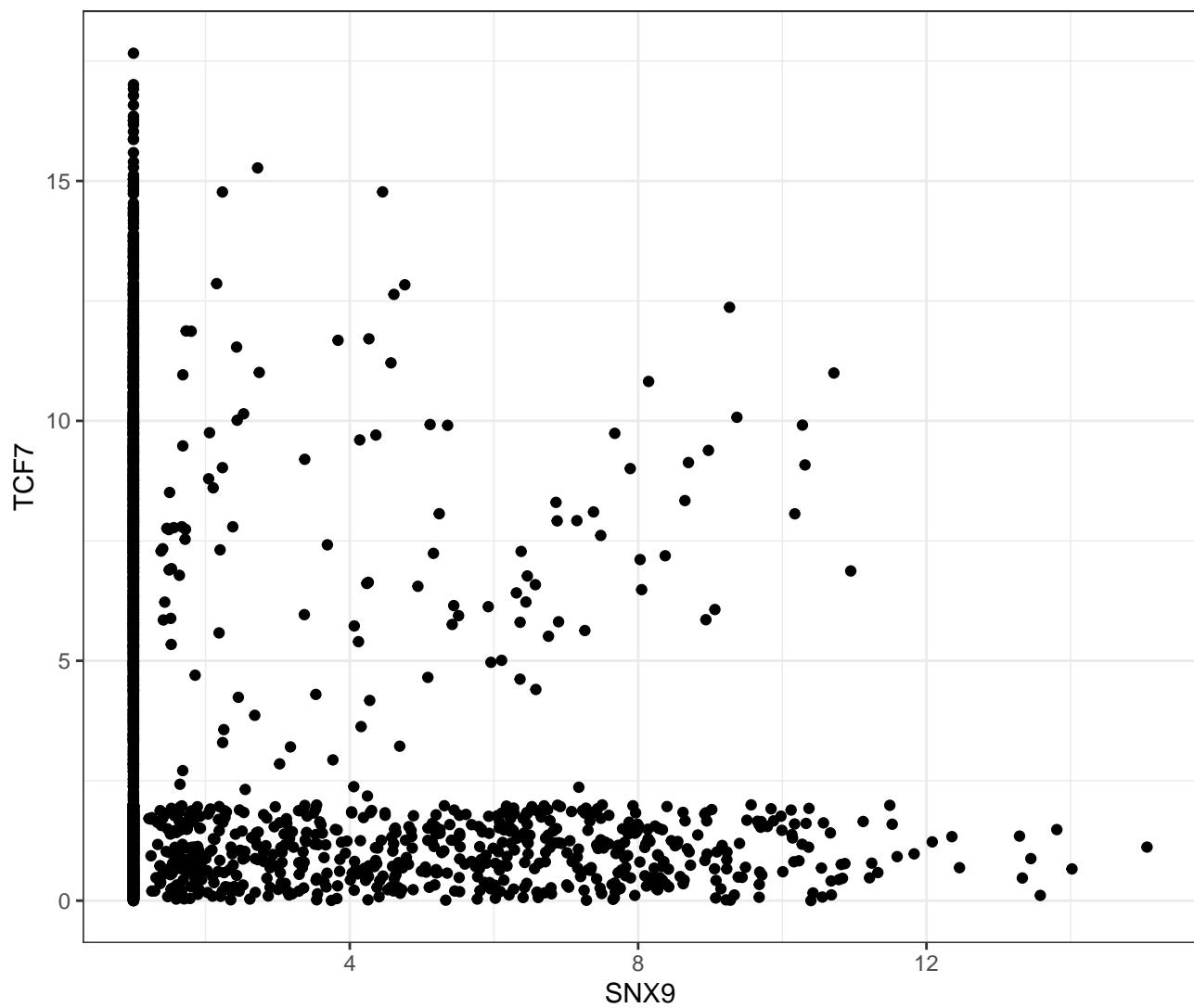
PDCD1 in CD8+ T cells



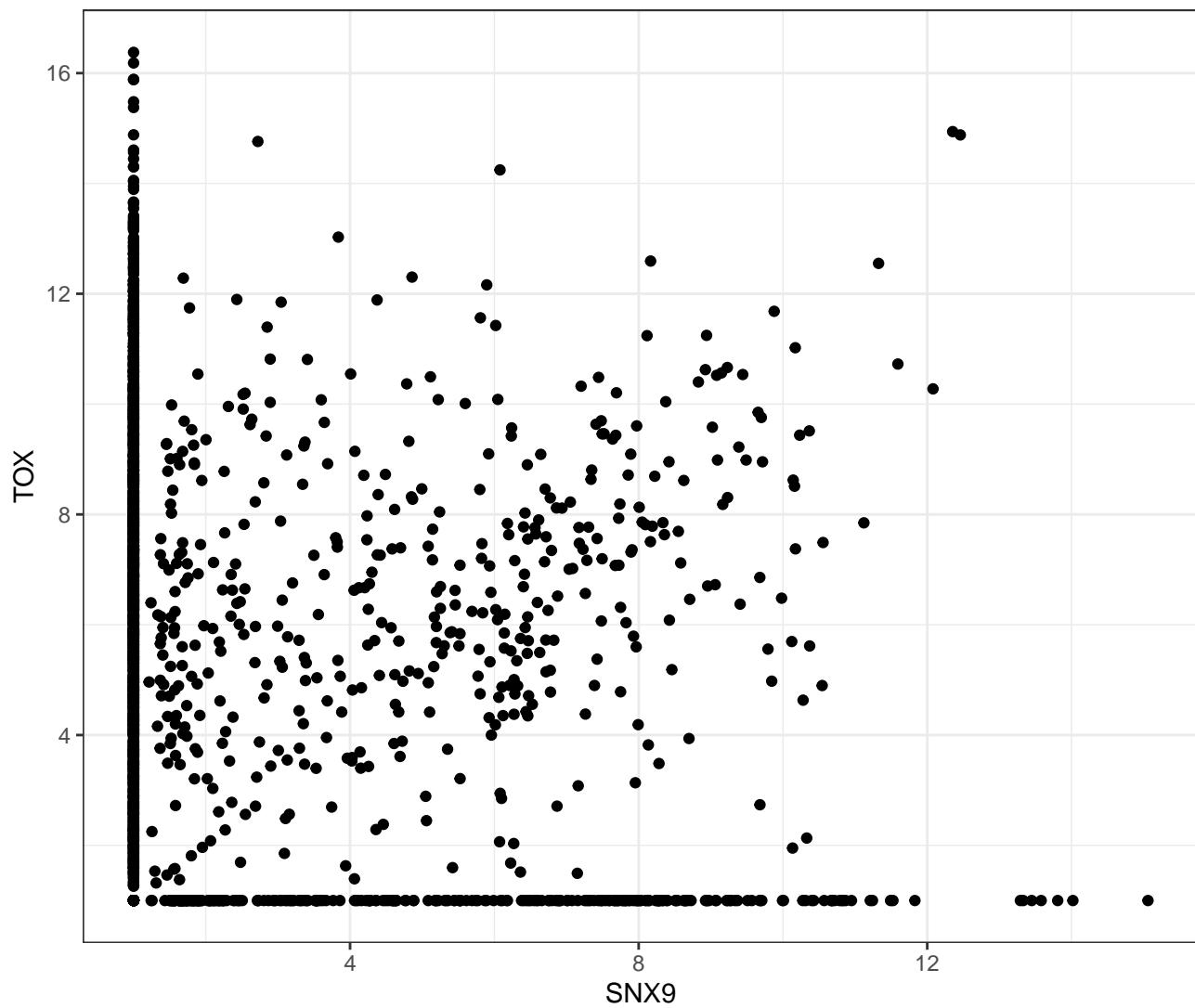
HAVCR2 in CD8+ T cells



SNX9 vs TCF7 in CD8+ T cells

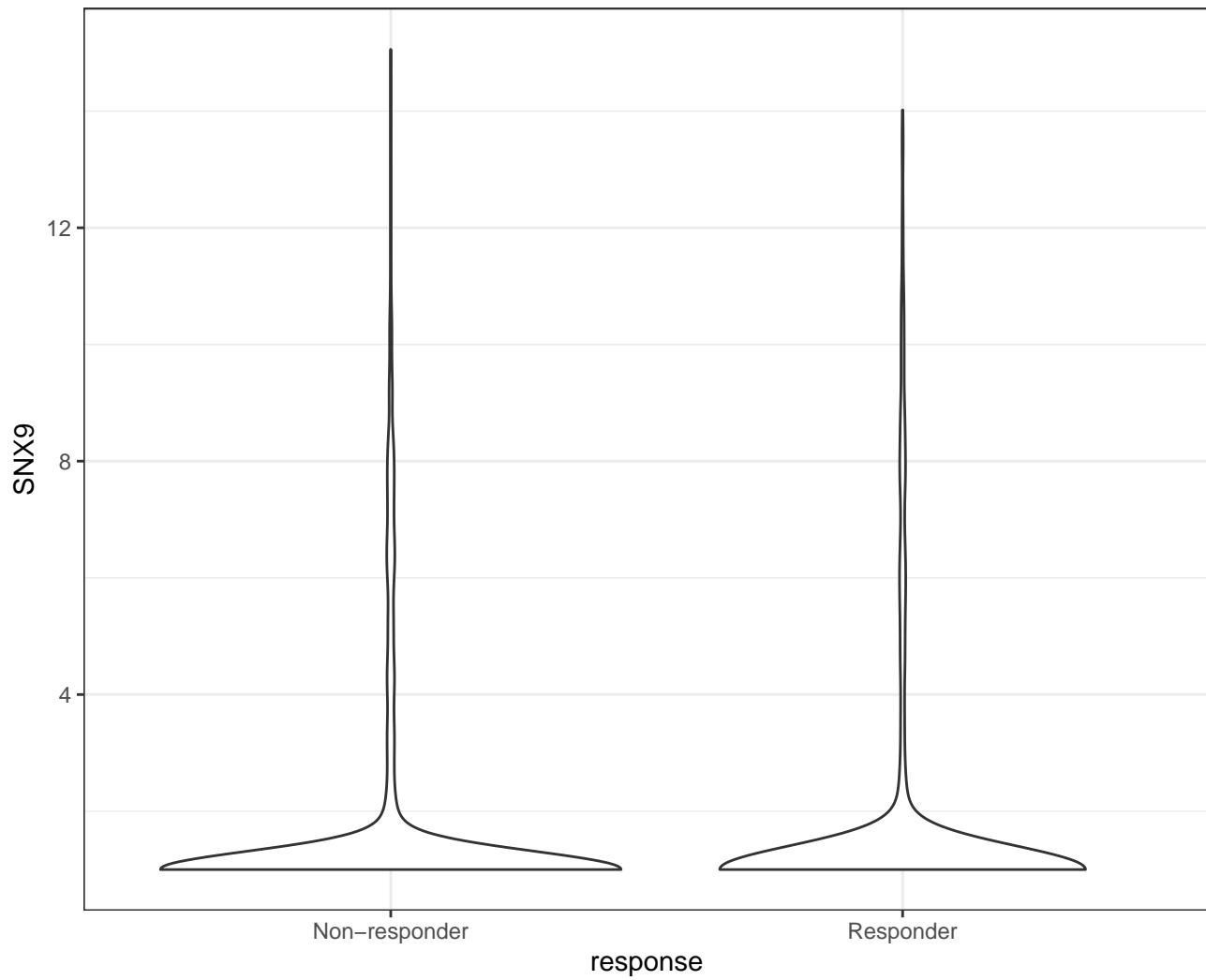


SNX9 vs TOX in CD8+ T cells



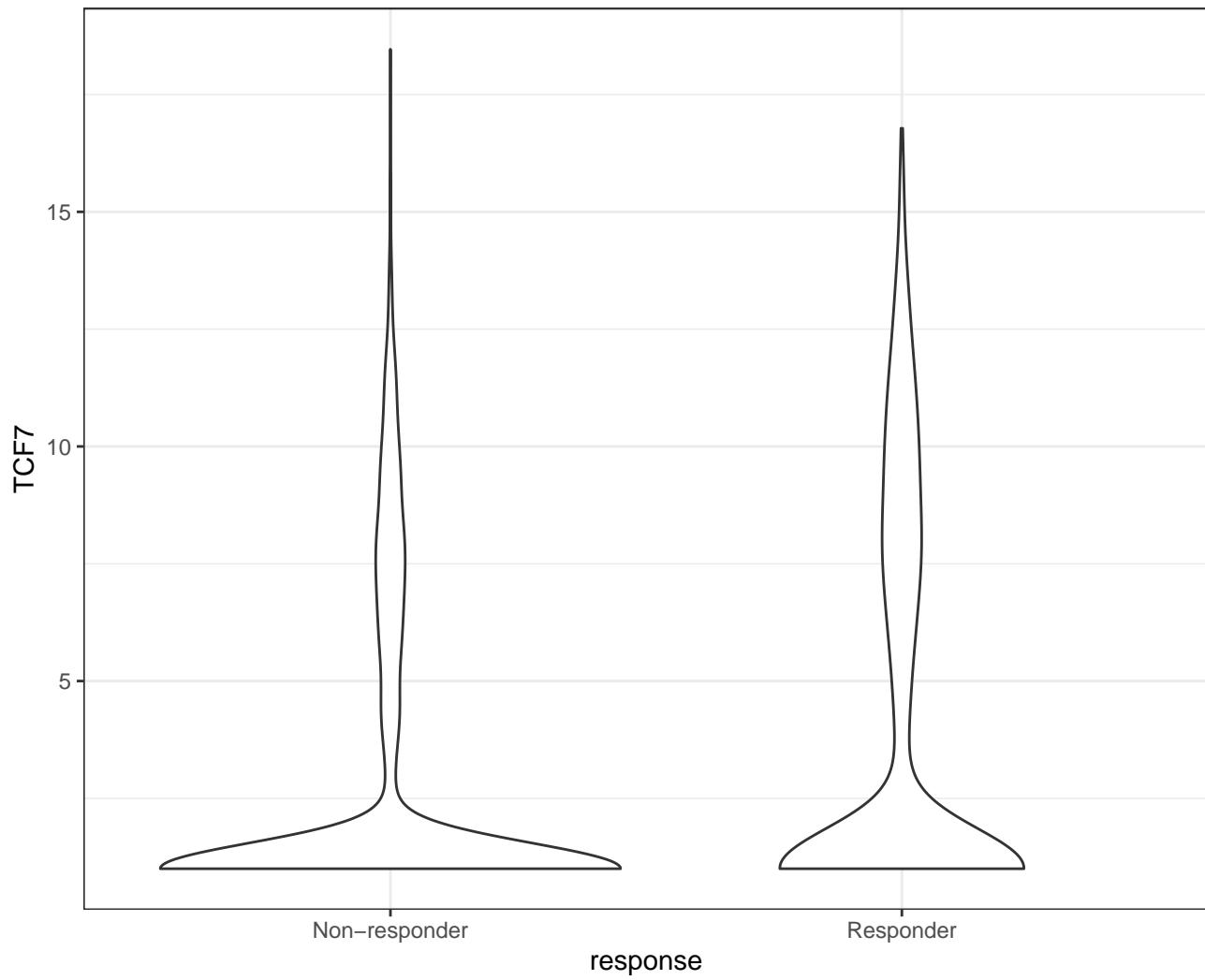
Response vs SNX9 in CD8+

wilcoxon p-value = 2.98e-02



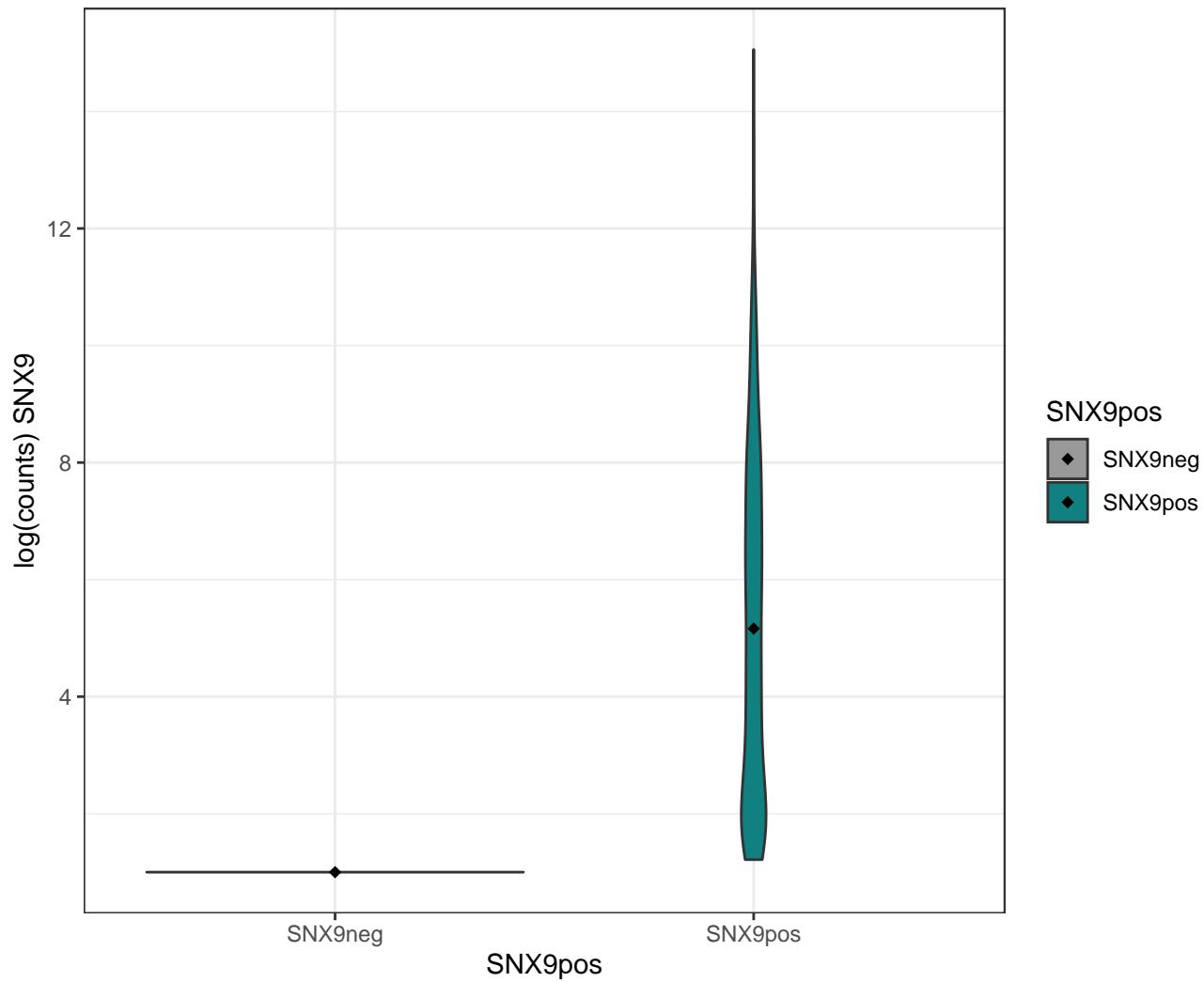
Response vs TCF7 in CD8+

wilcoxon p-value = 2.82e-28



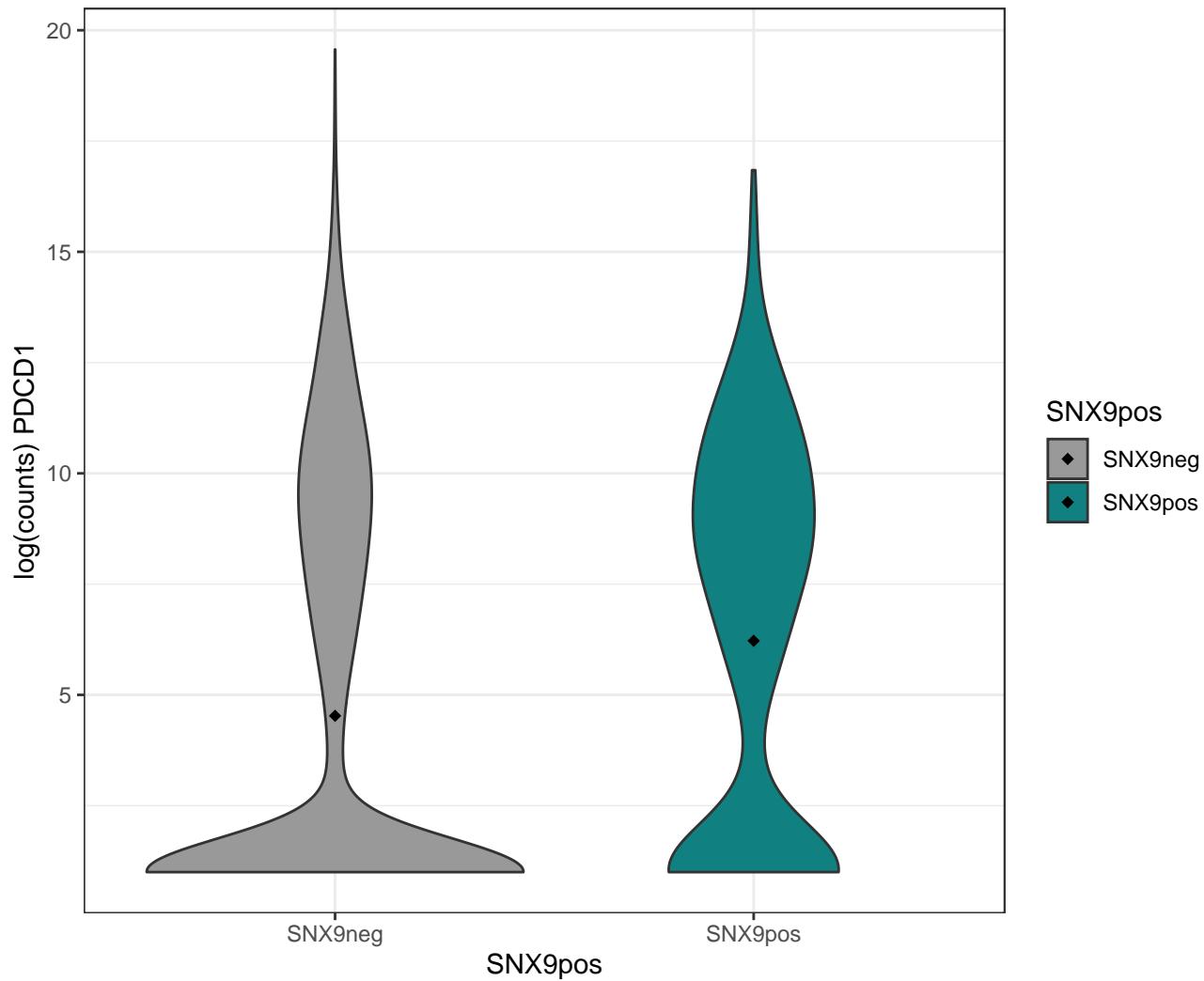
SNX9 expression in CD8+

wilcoxon p-value = 0.00e+00



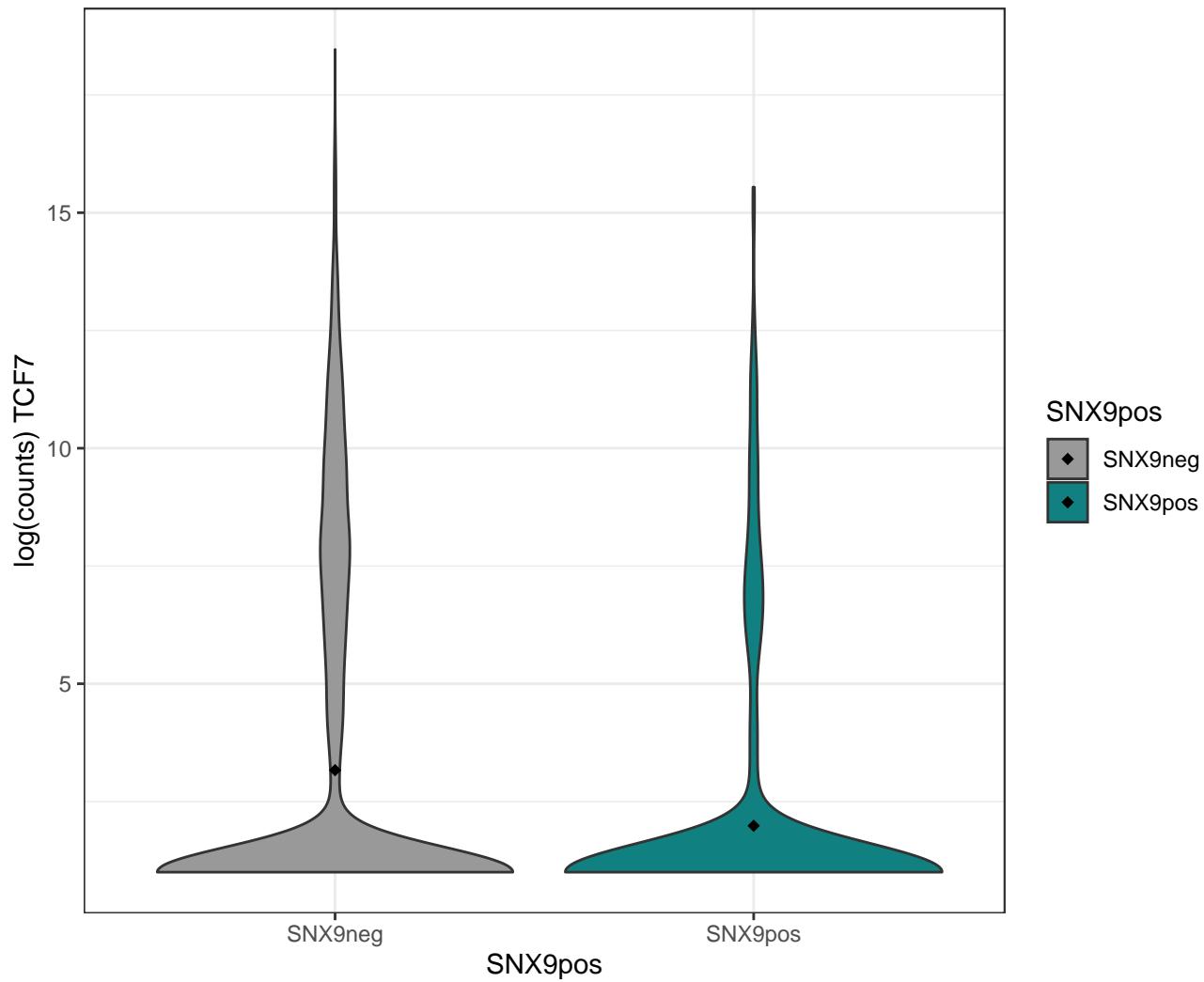
PD_CD1 expression in CD8+

wilcoxon p-value = 1.85e-24



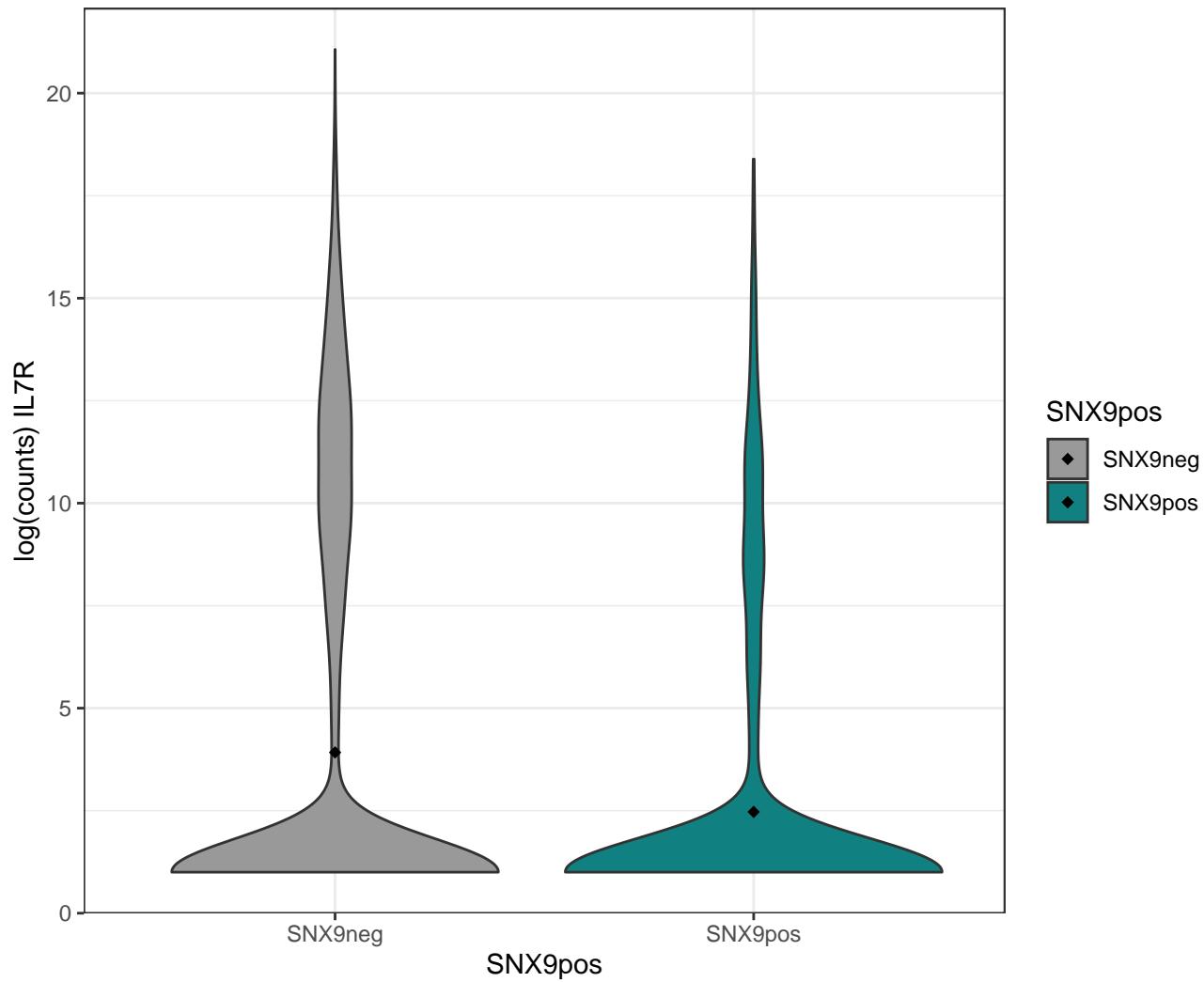
TCF7 expression in CD8+

wilcoxon p-value = 1.82e-18



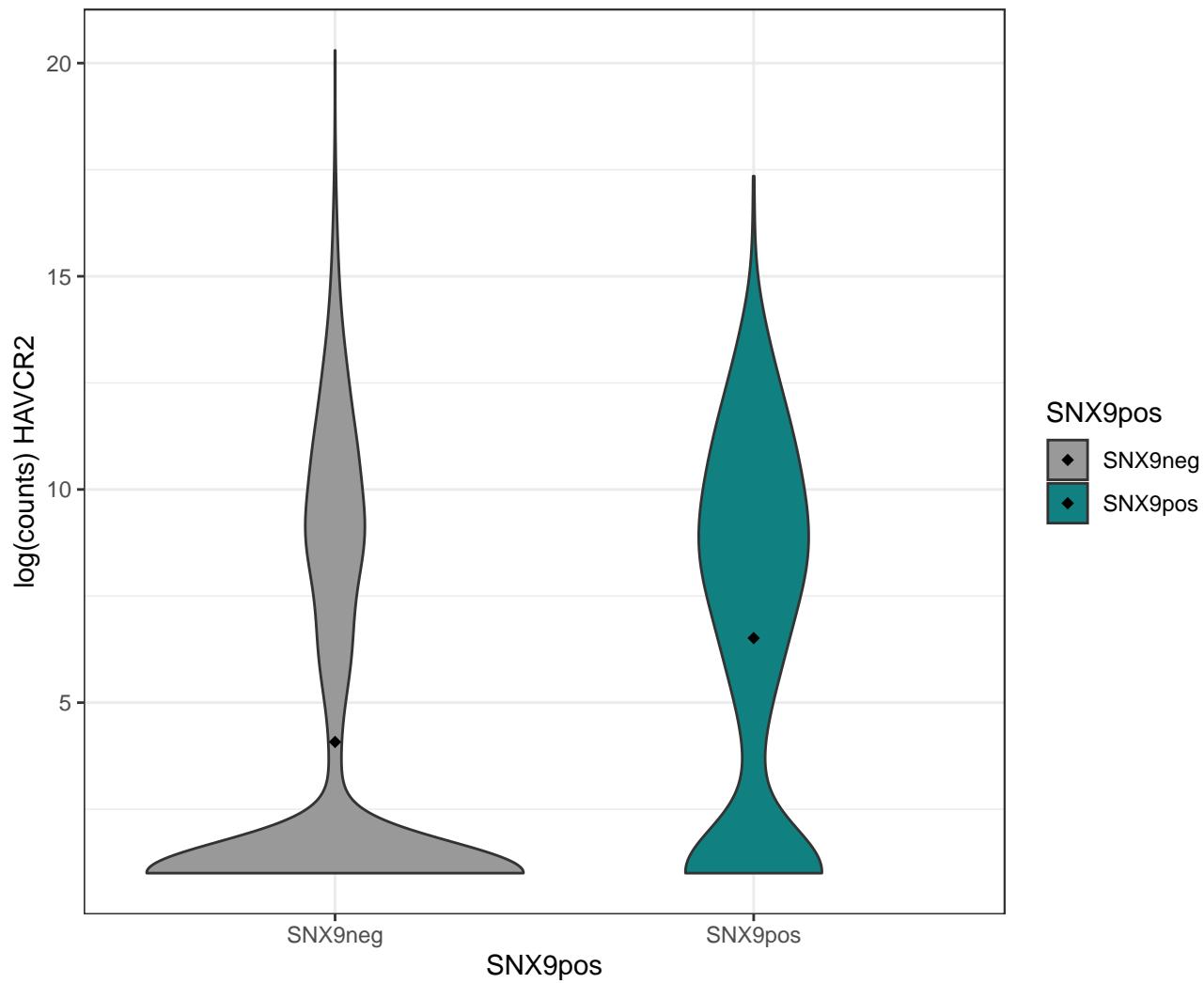
IL7R expression in CD8+

wilcoxon p-value = 9.09e-13



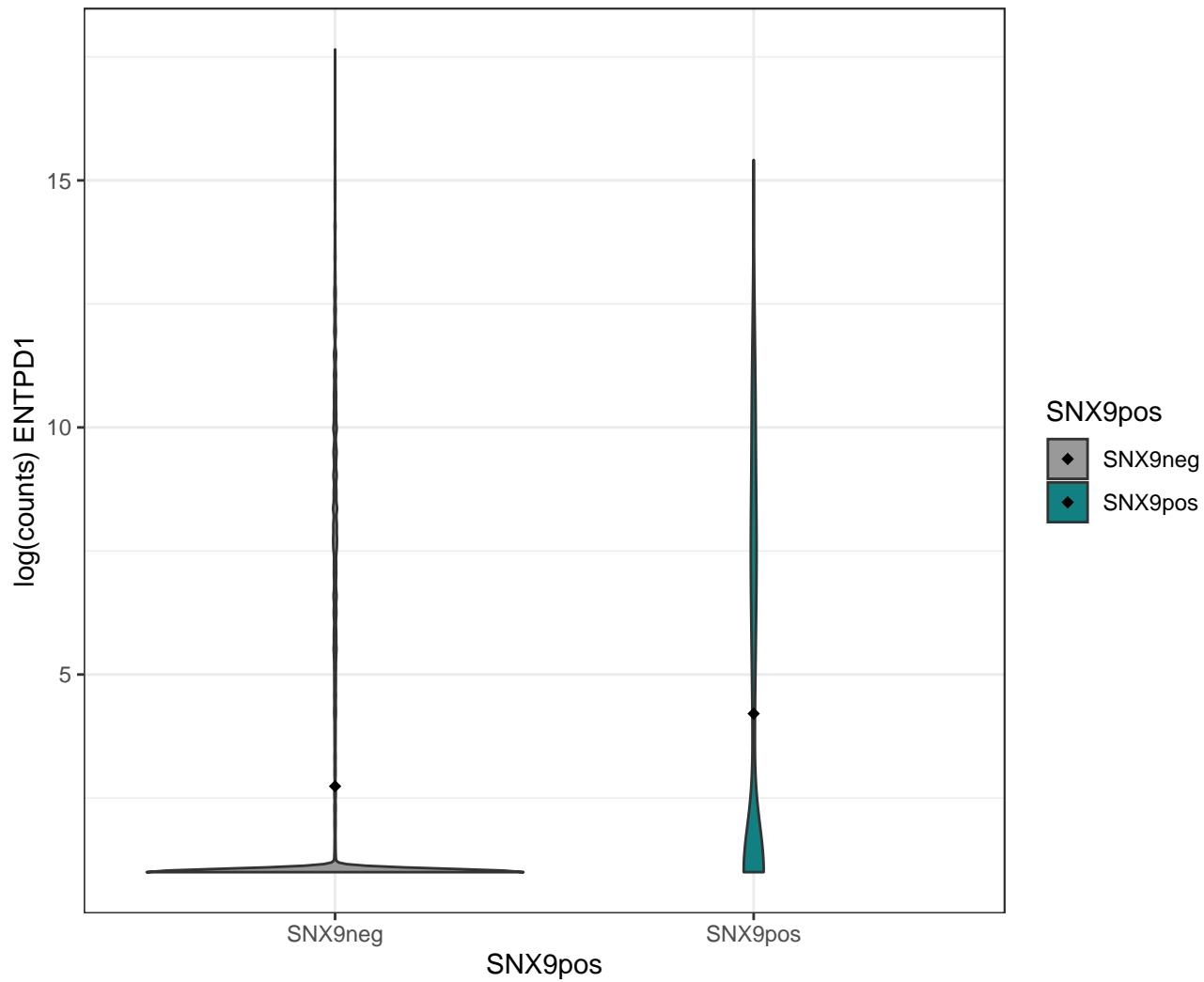
HAVCR2 expression in CD8+

wilcoxon p-value = 8.96e-50



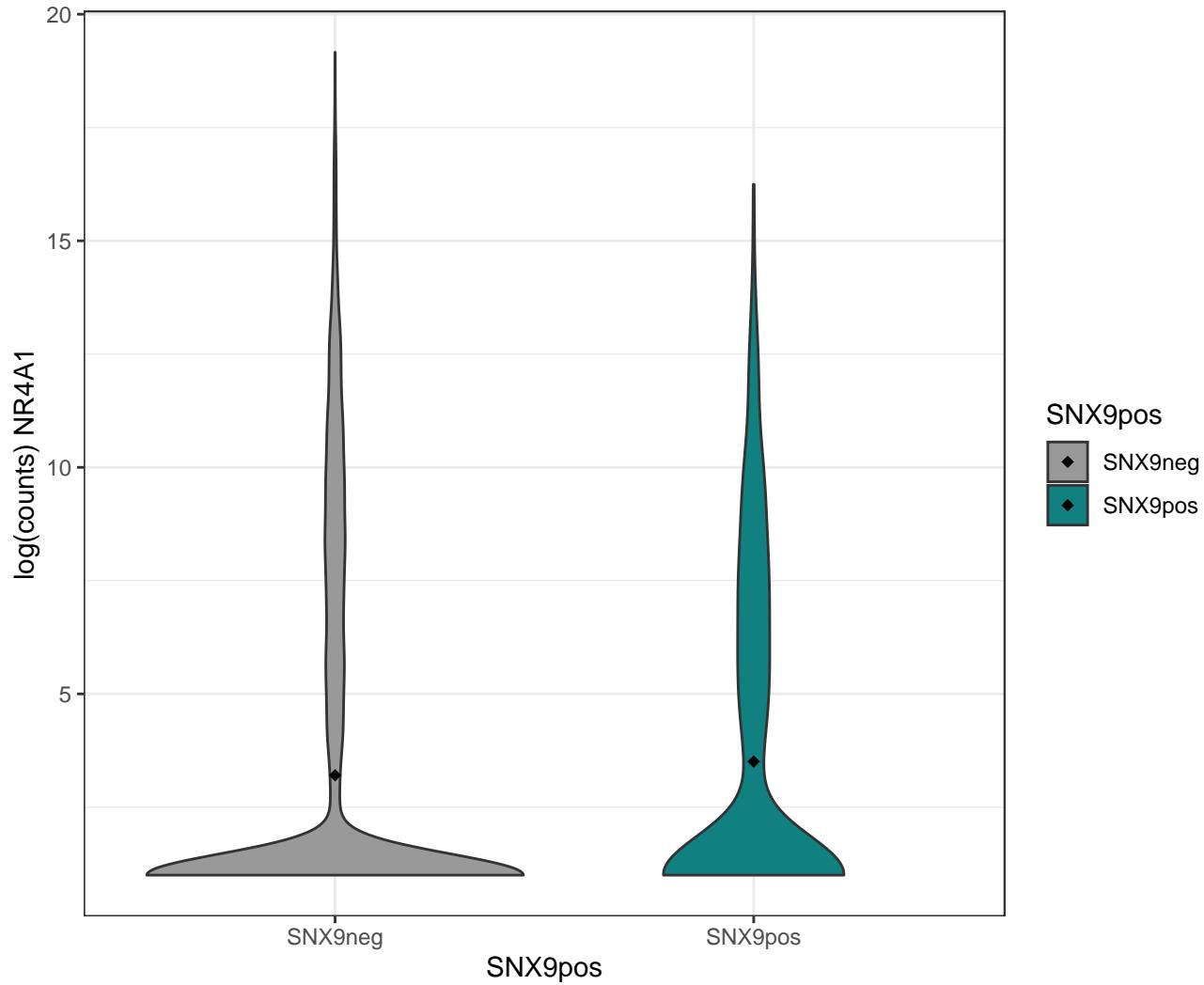
ENTPD1 expression in CD8+

wilcoxon p-value = 9.00e-34



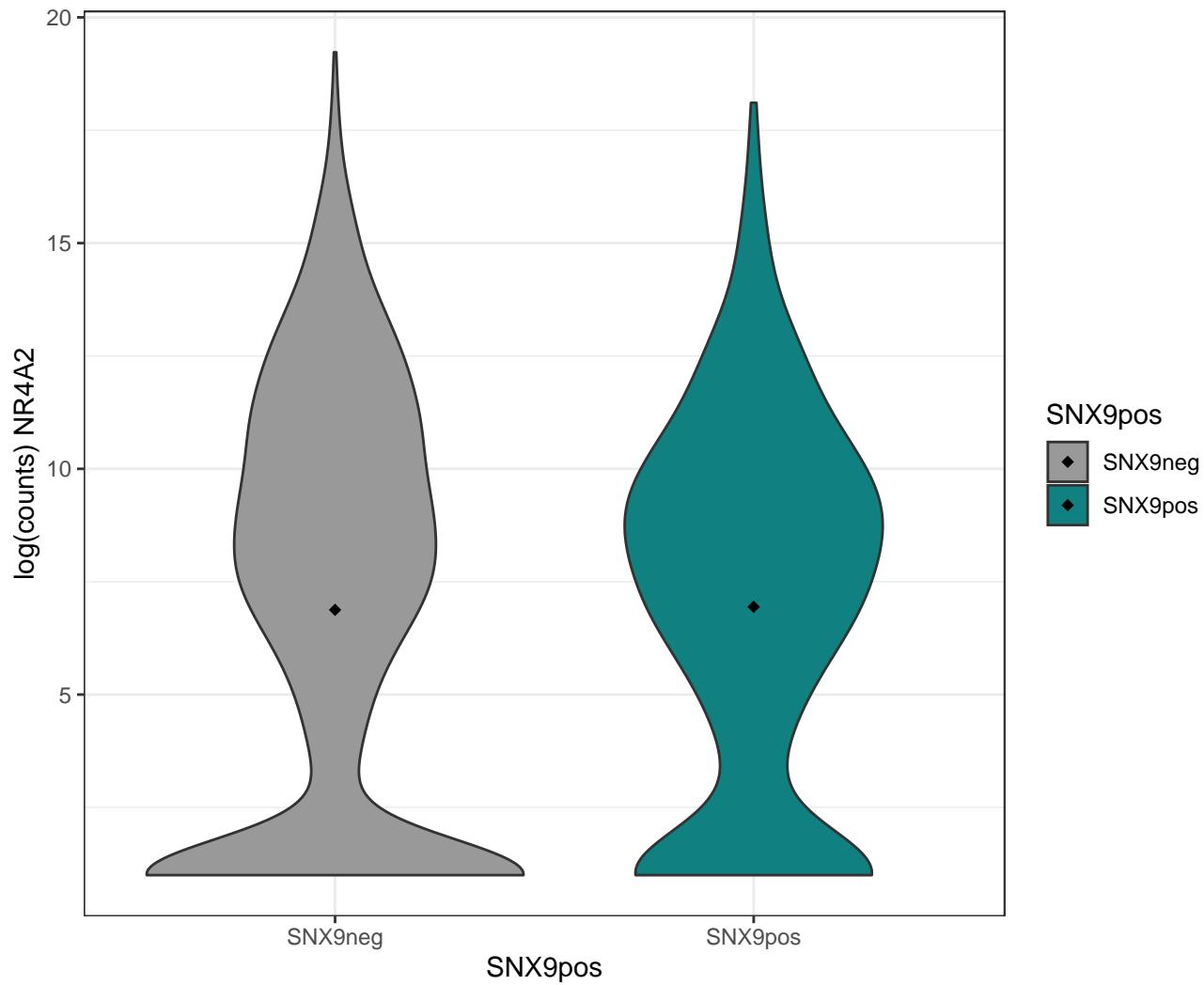
NR4A1 expression in CD8+

wilcoxon p-value = 9.74e-05



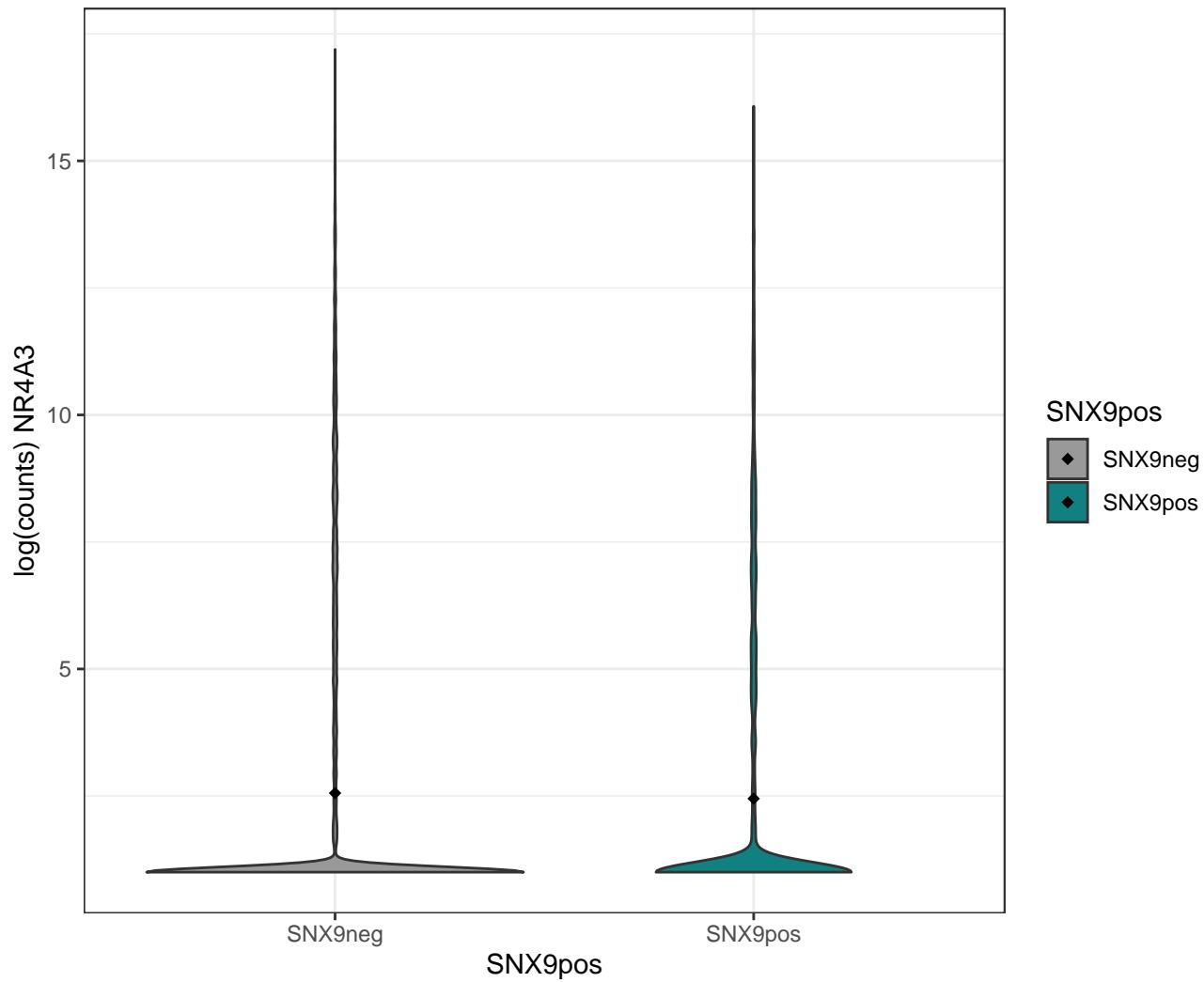
NR4A2 expression in CD8+

wilcoxon p-value = 9.09e-01



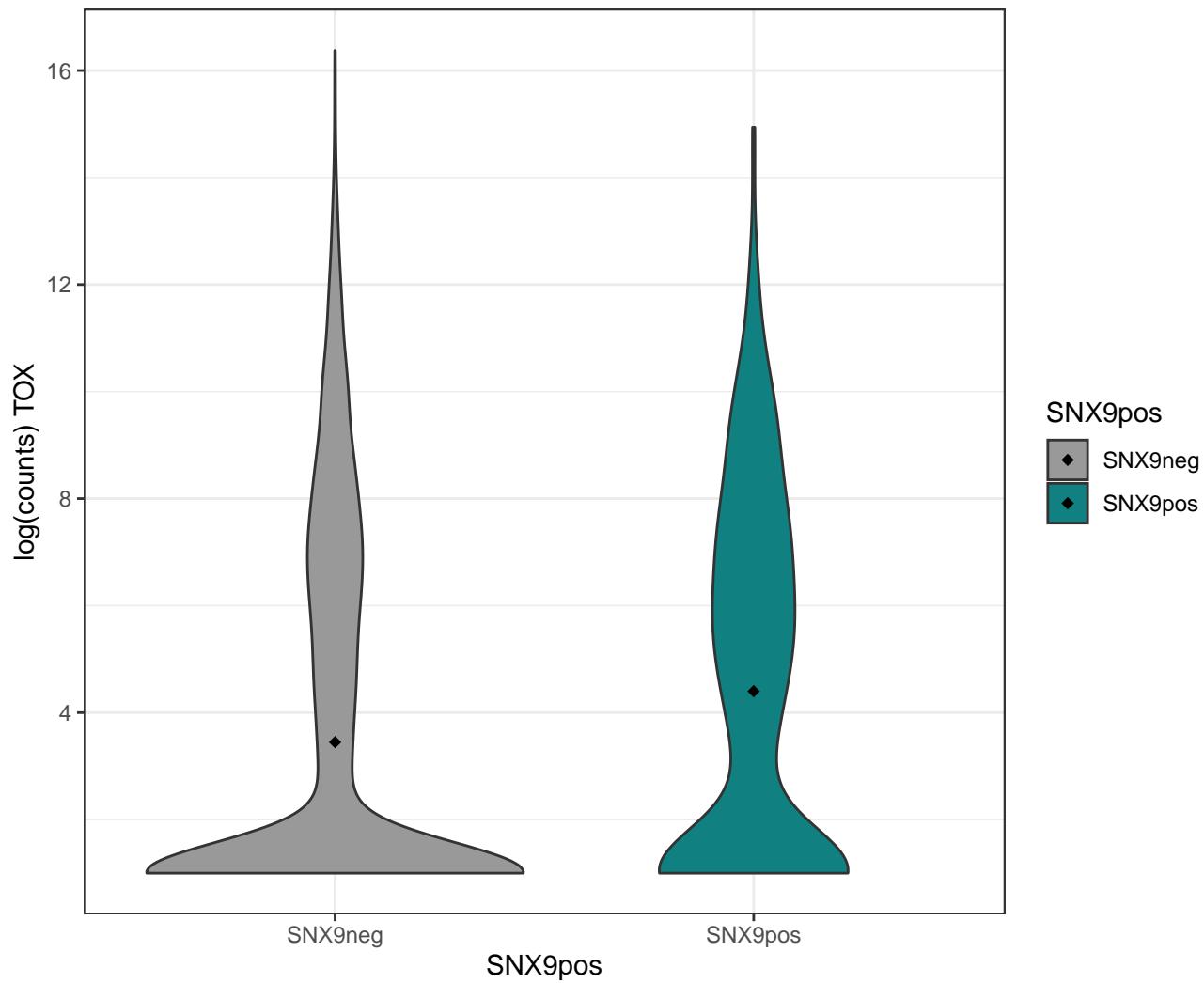
NR4A3 expression in CD8+

wilcoxon p-value = 9.27e-01



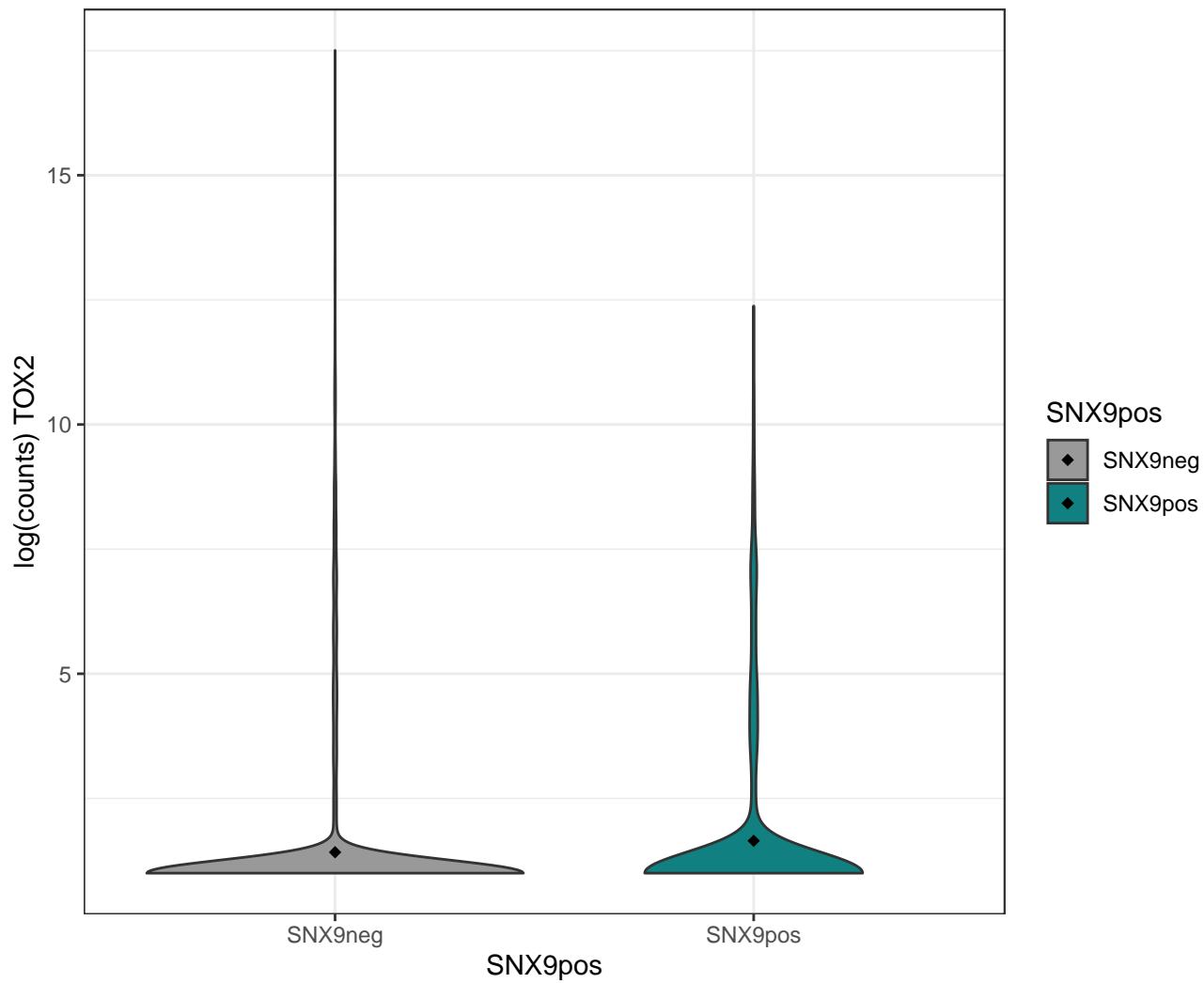
TOX expression in CD8+

wilcoxon p-value = 8.49e-17



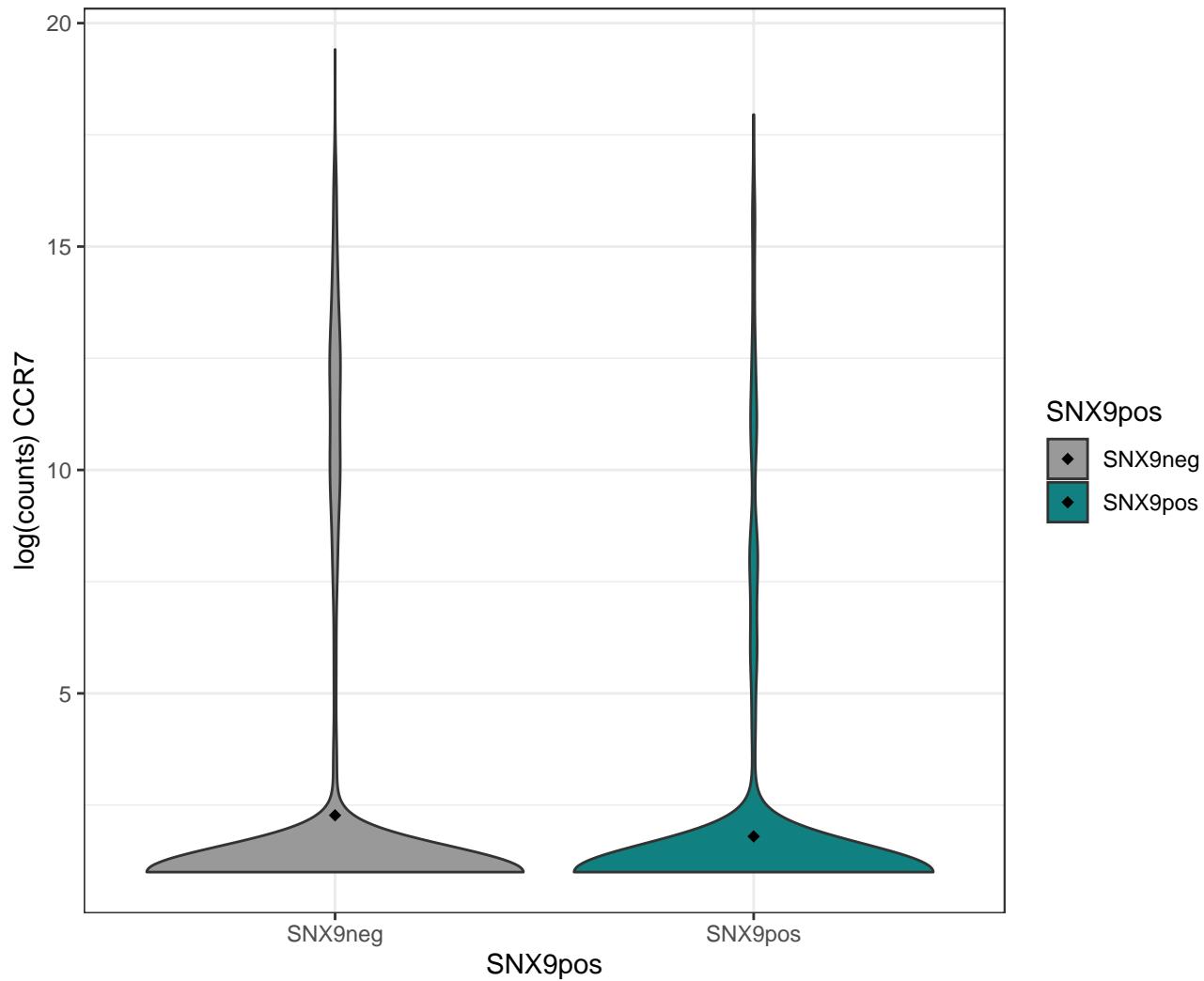
TOX2 expression in CD8+

wilcoxon p-value = 1.04e-06



CCR7 expression in CD8+

wilcoxon p-value = 1.02e-02



LDHA expression in CD8+

wilcoxon p-value = 4.50e-01

