TCGA–BRCA: Tumor vs Normal
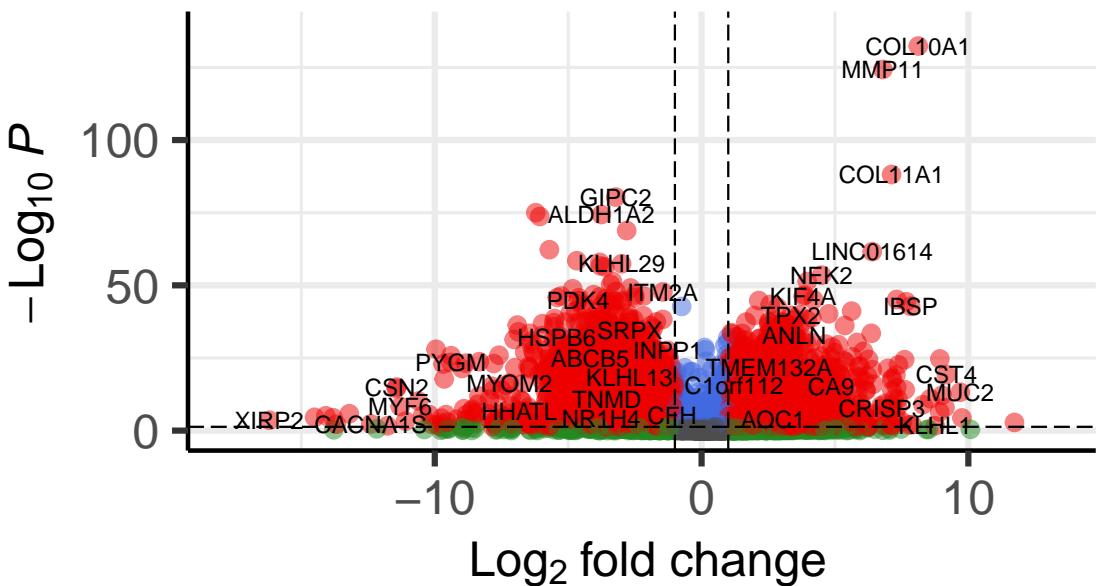
RNA−seq Differential Expression

Legend: NS, Log$_2$ FC, p−value, p−value and log$_2$ F

Y-axis: $-\mathrm{Log}_{10}\ P$ (values: 0, 50, 100)

X-axis: Log$_2$ fold change (values: −10, 0, 10)

Gene labels: COL10A1, MMP11, COL11A1, GIPC2, ALDH1A2, LINC01614, NEK2, KLHL29, KIF4A, IBSP, PDK4, ITM2A, TPX2, HSPB6, SRPX, ANLN, INPP1, PYGM, ABCB5, TMEM132A, CST4, CSN2, MYOM2, KLHL13, C1orf112, CA9, MUC2, MYF6, TNMD, CRISP3, XIRP2, CACNA1S, HHATL, NR1H4, CFH, AOC1, KLHL1

30 Tumor vs 30 Normal (TCGA−BRCA)