condition_PARPi_vs_CTRL (BH <0.05)

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

Y-axis: Log$_{10}$ p adjusted

X-axis: Log$_2$ fold change

Labeled genes: GDPD2, KRT13, LCE1A, KRT3, PKP1, ALAS2, POU3F1, EGR3, PHYHIP, PPL, PGM5, S100A7, BTBD11

total = 22779 variables