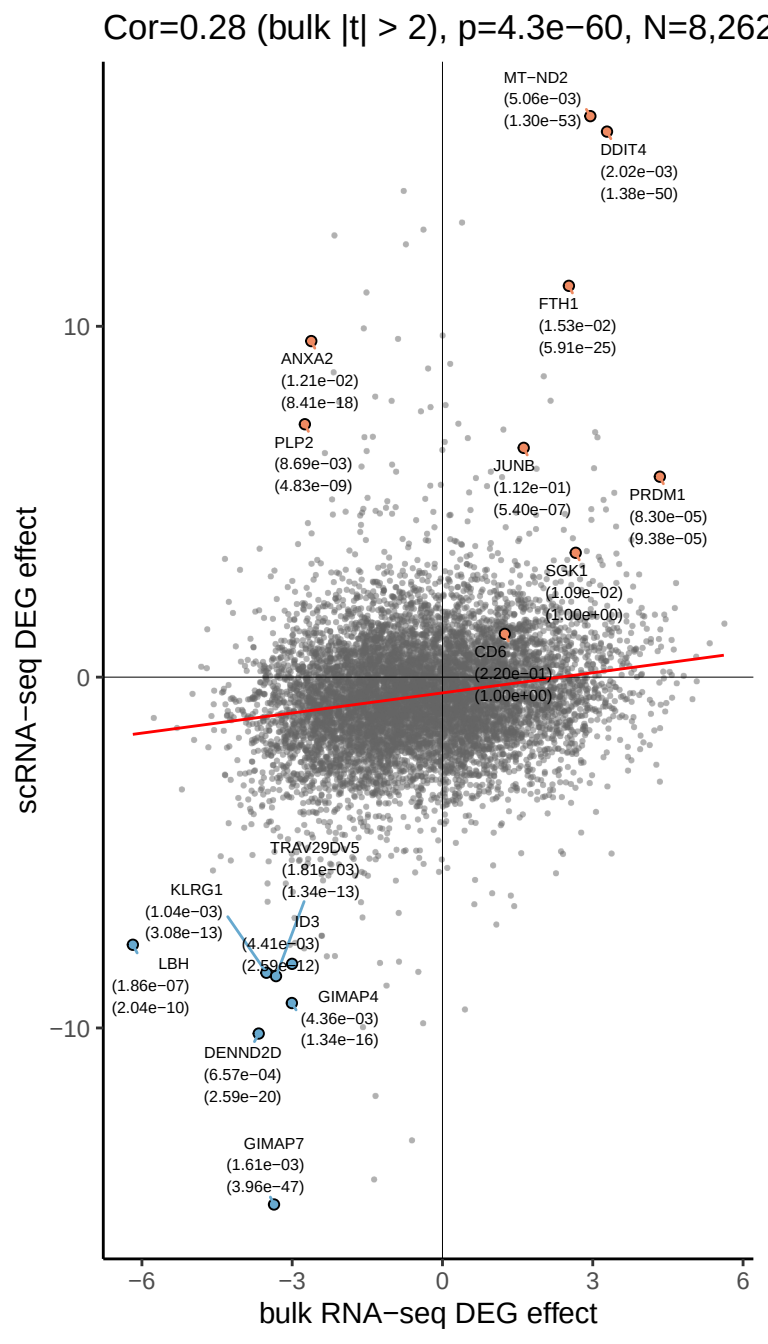
Cor=0.28 (bulk |t| > 2), p=4.3e−60, N=8,262
MT−ND2
(5.06e−03)
(1.30e−53)
DDIT4
(2.02e−03)
(1.38e−50)
FTH1
(1.53e−02)
(5.91e−25)
ANXA2
(1.21e−02)
(8.41e−18)
PLP2
(8.69e−03)
(4.83e−09)
JUNB
(1.12e−01)
(5.40e−07)
PRDM1
(8.30e−05)
(9.38e−05)
SGK1
(1.09e−02)
(1.00e+00)
CD6
(2.20e−01)
(1.00e+00)
TRAV29DV5
(1.81e−03)
(1.34e−13)
KLRG1
(1.04e−03)
(3.08e−13)
ID3
(4.41e−03)
(2.59e−12)
LBH
(1.86e−07)
(2.04e−10)
GIMAP4
(4.36e−03)
(1.34e−16)
DENND2D
(6.57e−04)
(2.59e−20)
GIMAP7
(1.61e−03)
(3.96e−47)
10
0
−10
−6
−3
0
3
6
scRNA−seq DEG effect
bulk RNA−seq DEG effect