Cor=0.23, p=5.8e−98, N=8,240

Scatter plot with x-axis "bulk RNA−seq DEG effect" and y-axis "scRNA−seq DEG effect".

Labeled points (upper region):
- CTLA4 (2.86e−03)
- HERPUD1 (5.22e−04)
- MGAT4A (1.41e−03)
- CLEC2B (1.39e−03)
- PRDM1 (1.95e−05)
- STX11 (1.21e−03)

Labeled points (lower region):
- LBH (4.35e−04)
- APBB1IP (7.84e−04)
- DYNLL1 (3.35e−04)
- LIMS1 (1.88e−03)
- EMP3 (1.28e−03)
- HSPA8 (2.82e−03)
- ICAM2 (1.93e−03)