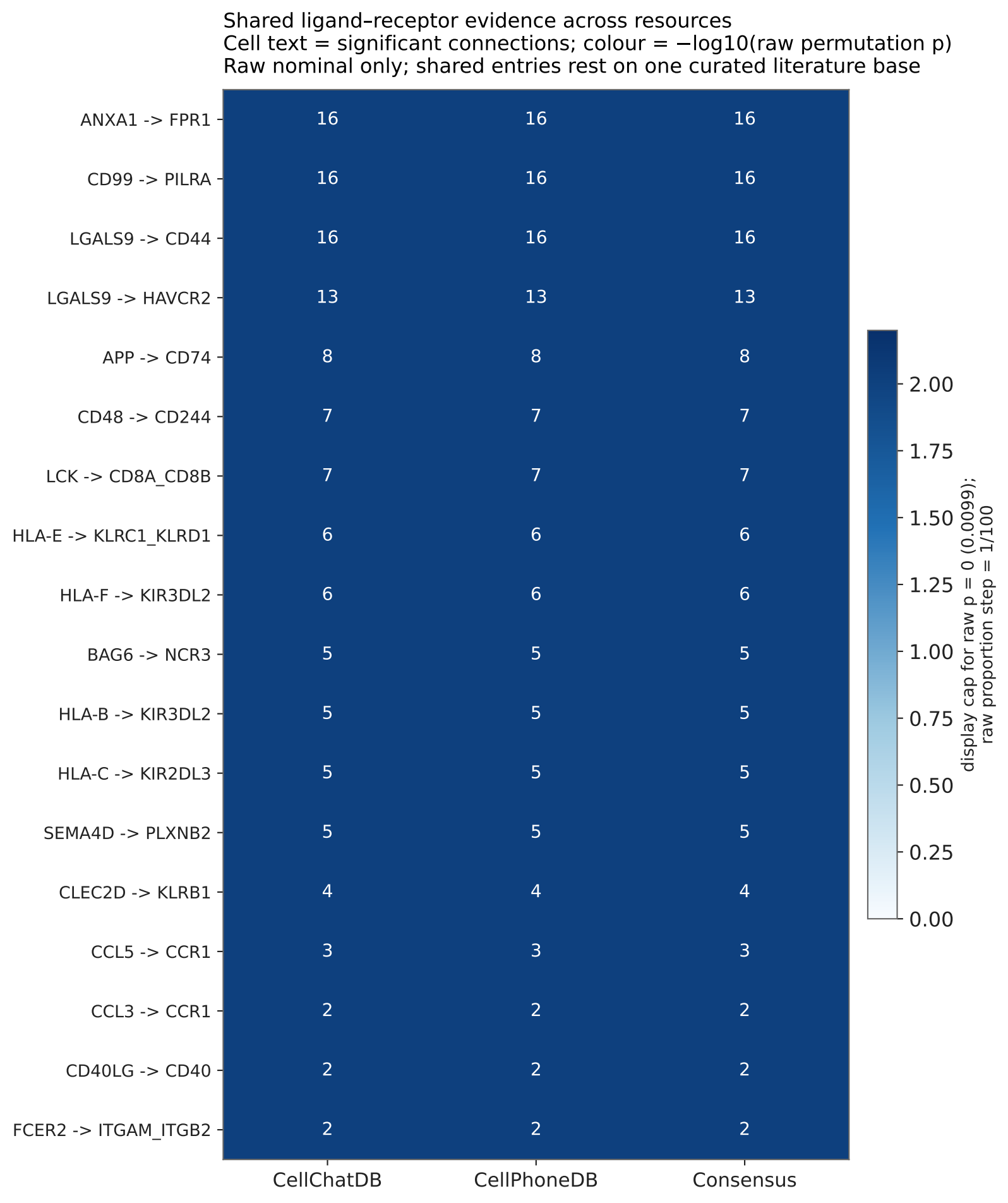

Shared ligand–receptor evidence across resources
Cell text = significant connections; colour = −log10(raw permutation p)
Raw nominal only; shared entries rest on one curated literature base
ANXA1 -> FPR1
CD99 -> PILRA
LGALS9 -> CD44
LGALS9 -> HAVCR2
APP -> CD74
CD48 -> CD244
LCK -> CD8A_CD8B
HLA-E -> KLRC1_KLRD1
HLA-F -> KIR3DL2
BAG6 -> NCR3
HLA-B -> KIR3DL2
HLA-C -> KIR2DL3
SEMA4D -> PLXNB2
CLEC2D -> KLRB1
CCL5 -> CCR1
CCL3 -> CCR1
CD40LG -> CD40
FCER2 -> ITGAM_ITGB2
16 16 16
16 16 16
16 16 16
13 13 13
8 8 8
7 7 7
7 7 7
6 6 6
6 6 6
5 5 5
5 5 5
5 5 5
5 5 5
4 4 4
3 3 3
2 2 2
2 2 2
2 2 2
CellChatDB
CellPhoneDB
Consensus
display cap for raw p = 0 (0.0099);
raw proportion step = 1/100
2.00
1.75
1.50
1.25
1.00
0.75
0.50
0.25
0.00