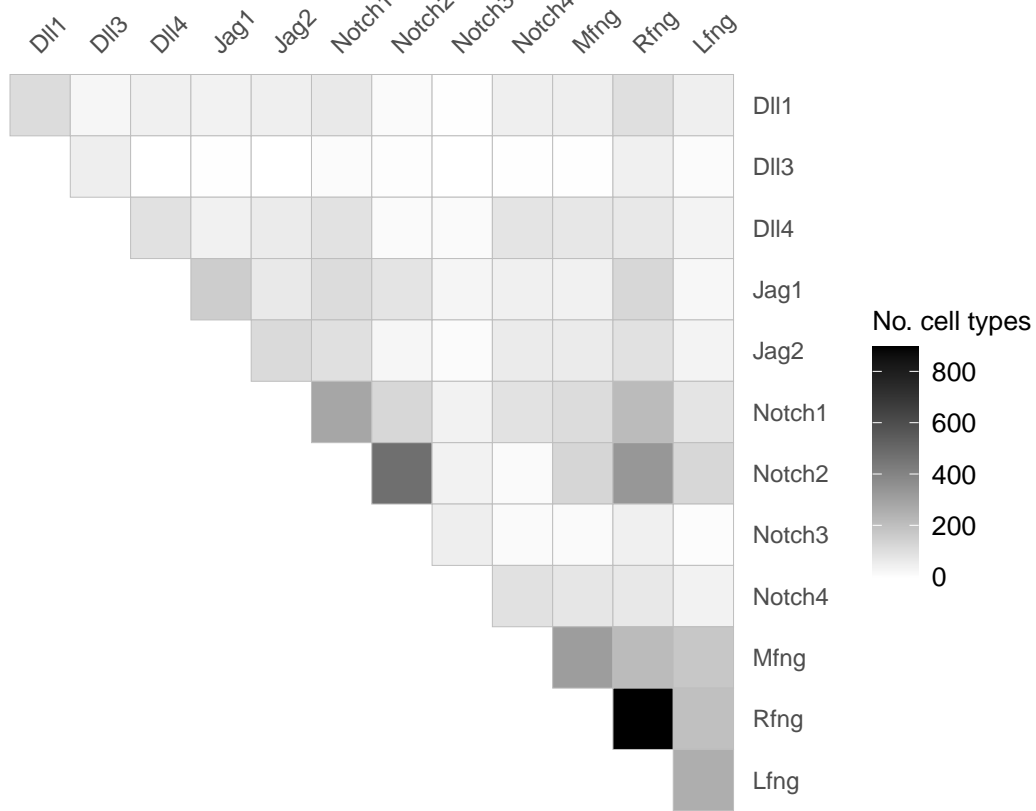


Notch (k_opt = 30)

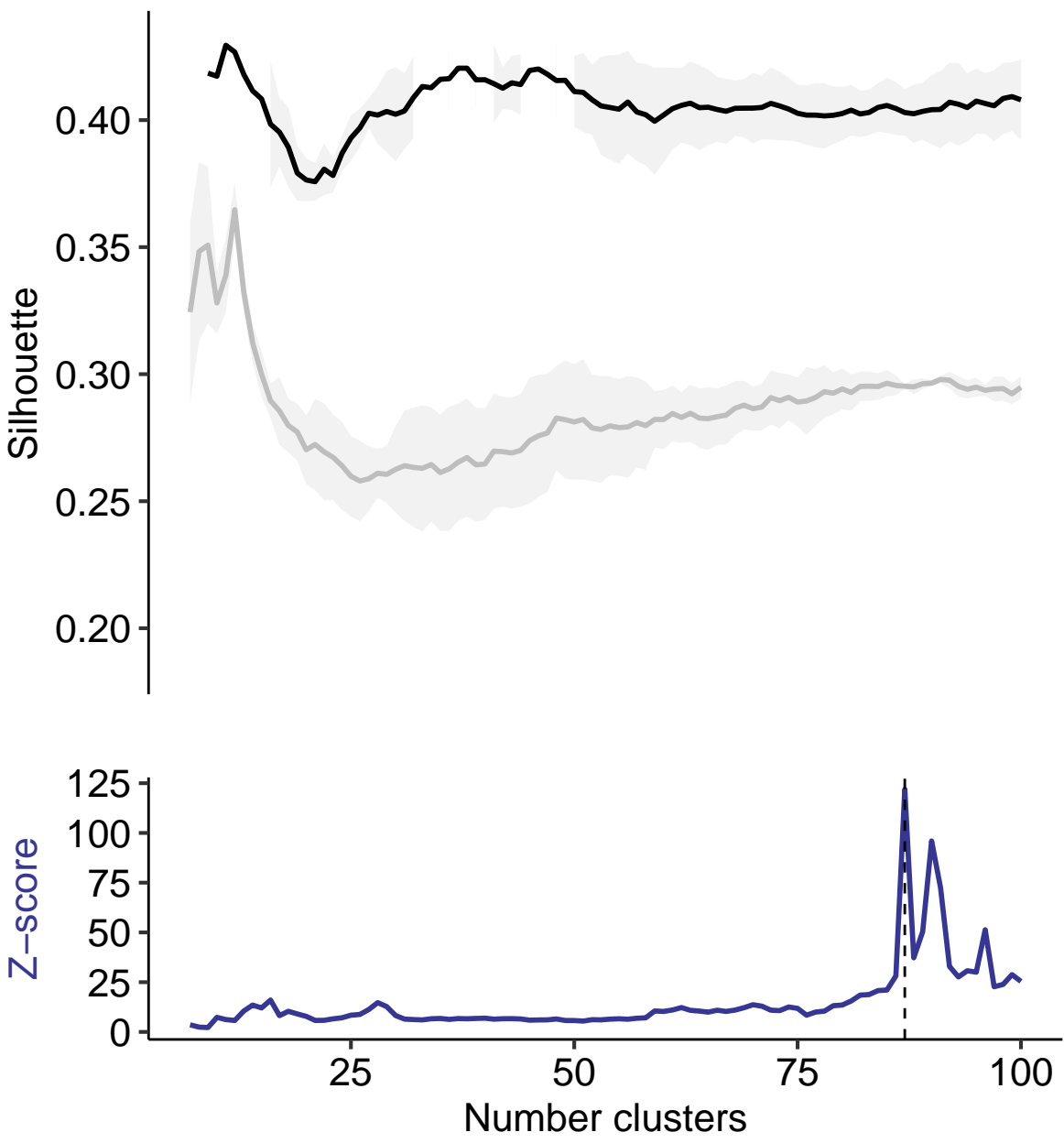
Pathway genes

Dll1	Jag2	Notch4
Dll3	Notch1	Mfng
Dll4	Notch2	Rfng
Jag1	Notch3	Lfng

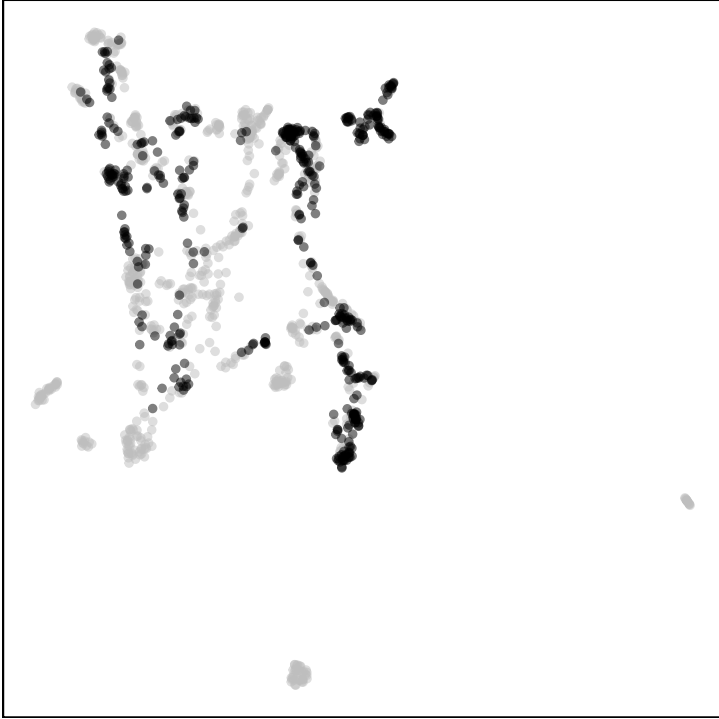
Gene Co-Expression Above Threshold



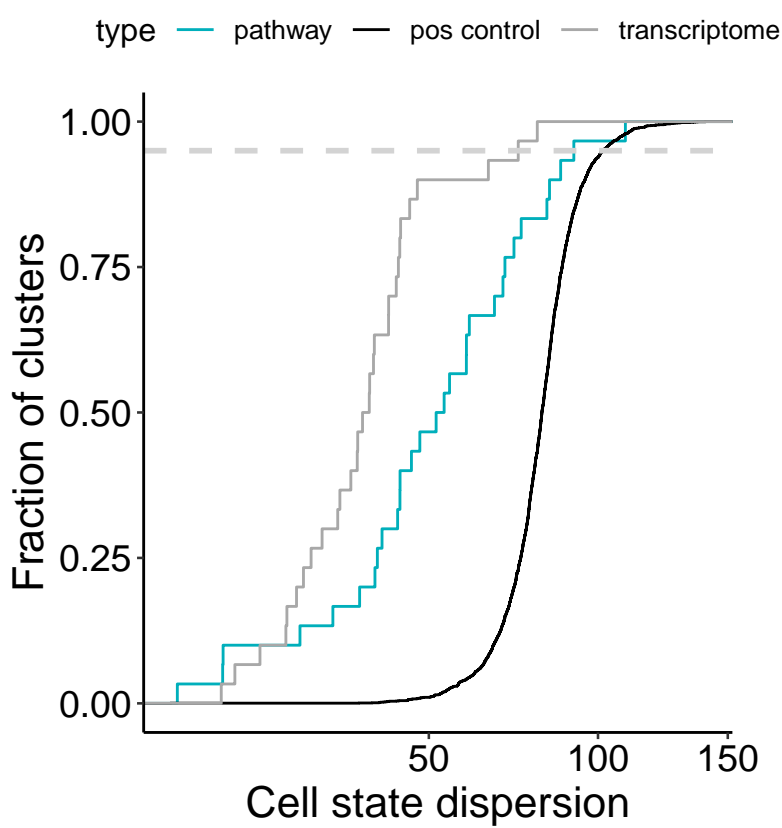
Silhouette and Z-score



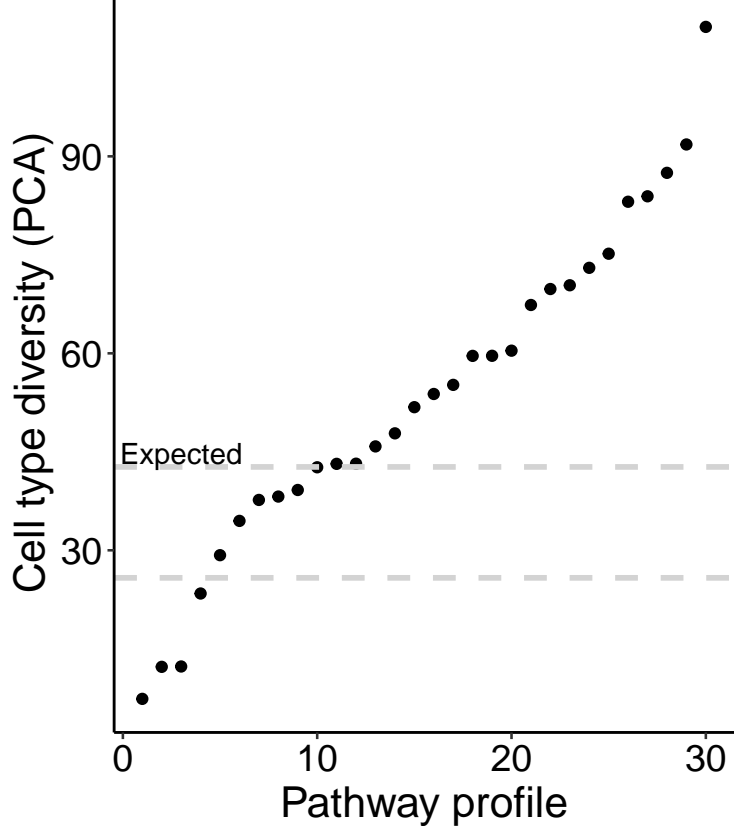
Global UMAP: Pathway ON



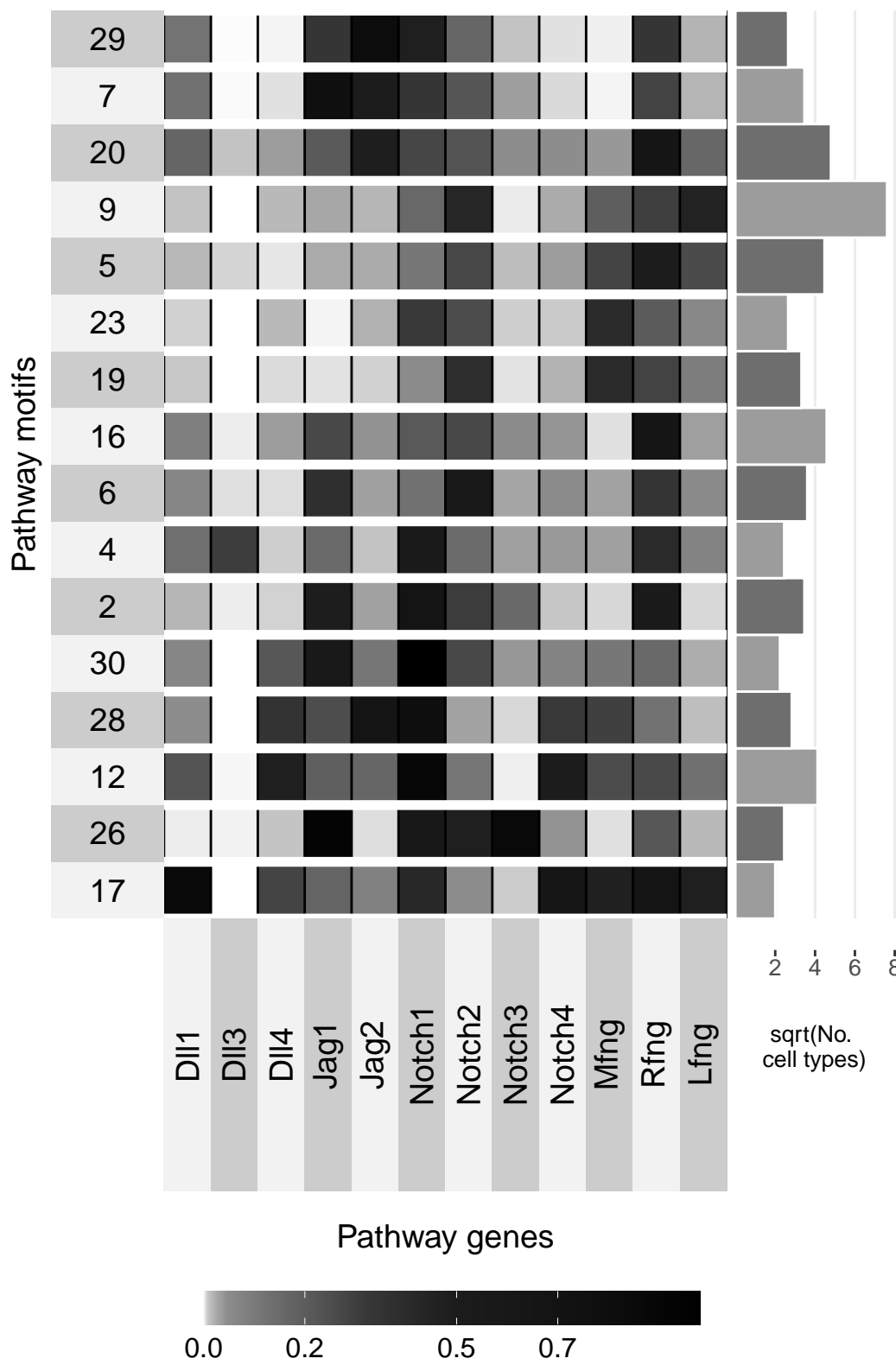
ECDF of Profile Dispersions



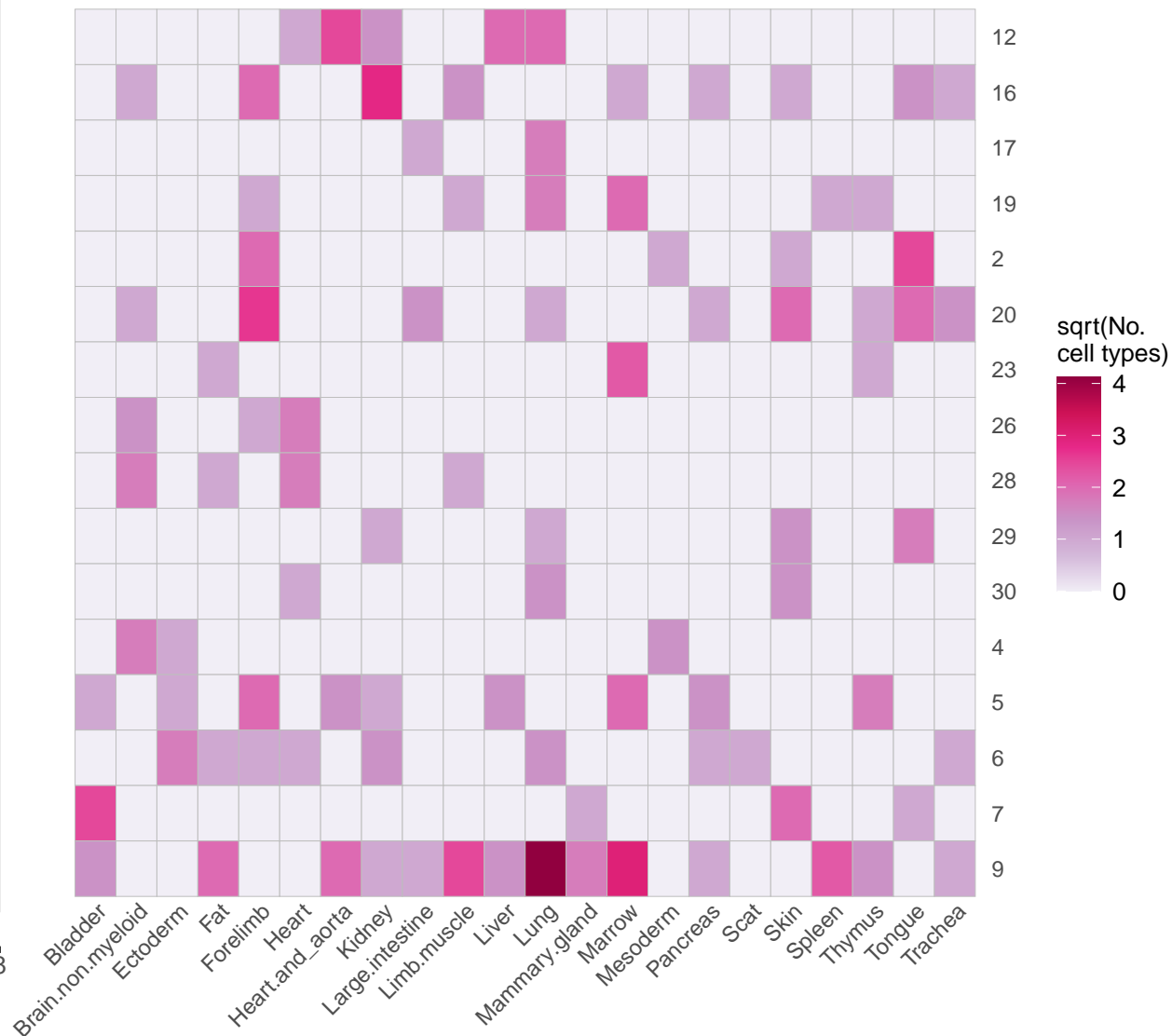
Profile Dispersion Ranking



Motifs, Dispersion >= 90th percentile



Motif Tissue Composition



Notch (k_opt = 30)

