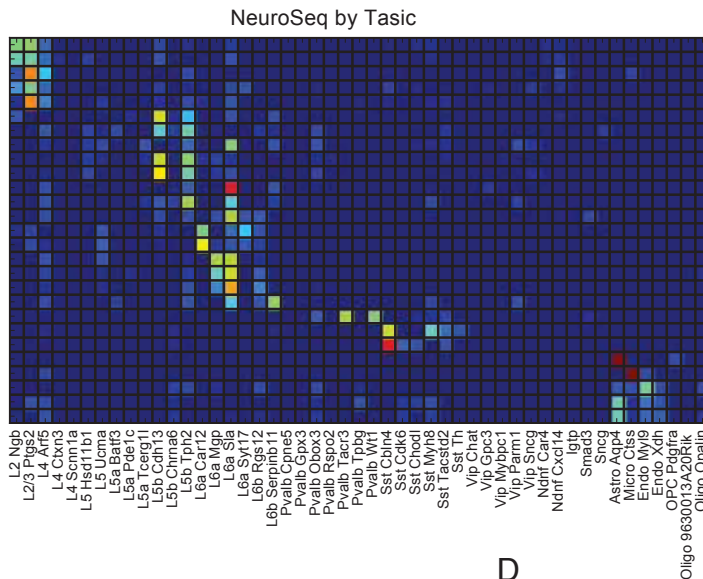
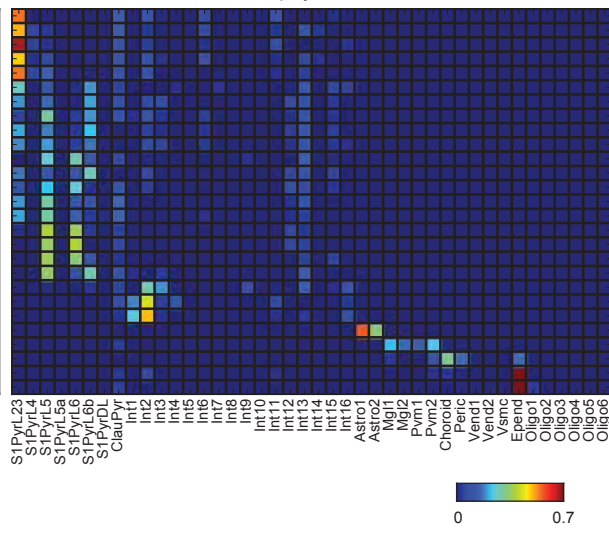


A

AL.L23_glu_P157
 ORBm.L23_glu_P157
 SSp.L23_glu_S2.inj
 AL.L23_glu_Grp
 MO.L23_glu_Grp
 ECT.L5_glu_Sla.a
 SSP.L5_glu_P084
 SSp.L5_glu_GPr2.a
 SSp.L5_glu_BPN.inj
 SSP.L5_glu_IRt.inj
 RSP.V.L6_glu_P162
 ViSP.L6_glu_P078
 ECT.L6_glu_P038
 ViSP.L6_glu_S6L
 SSP.L6_glu_S6L
 SSP.L6_glu_P162
 SSp.L6_glu_P139
 ViSP.L6a_glu_Nisr.b
 SSp.L6b_glu_GPr2.b
 SSp.L6b_GABA_Pvalb
 ISO.L5_GABA_Xv8
 ISO.L5_GABA_GIN
 ASTR0iso_GFAP
 MOP.L5_Cx3cx1
 Choroid Plexus
 Ependyma_Pvalb
 Ependyma_TCLU

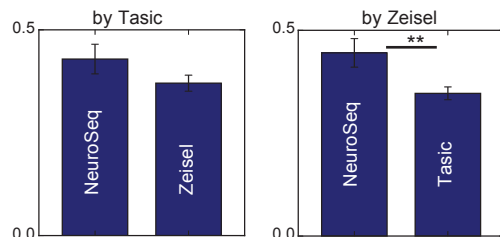


NeuroSeq by Zeisel



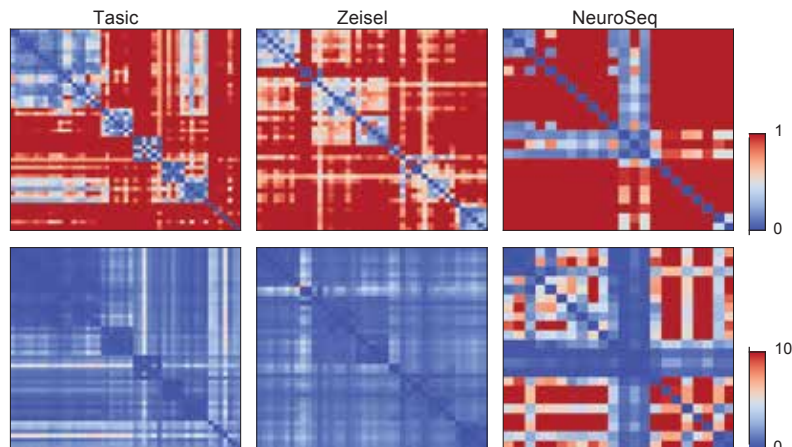
B

Mean Purity Scores

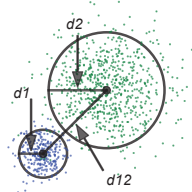


D

Separability of clusters



C



$$separability = \frac{d_{12}}{d_1 + d_2}$$

$$d1, d2 = 3 * \sigma$$