

Tadh OG_5095

metacells

TrH2 OG_8660 TrH2_TrispH2_003010-RA Homeodomains.HG2.67:like:ALX1/ALX3/ALX4/ARX

metacells

Hhon OG_8660 Hhon_g03913.t1 Homeodomains.HG2.67:like:ALX1/ALX3/ALX4/ARX

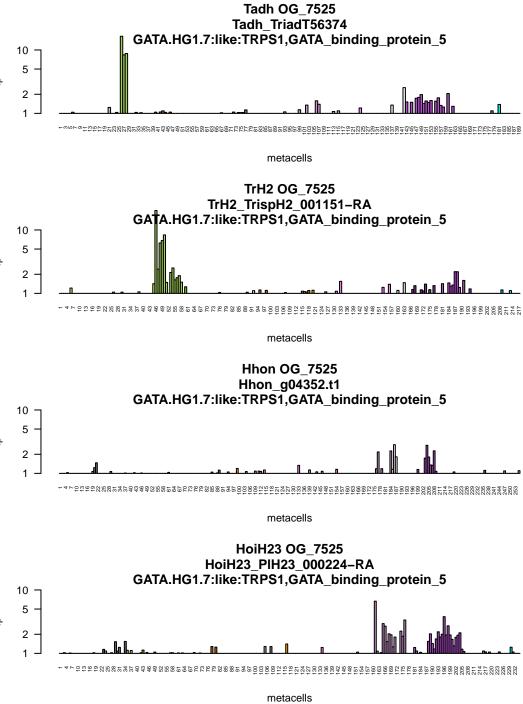
metacells

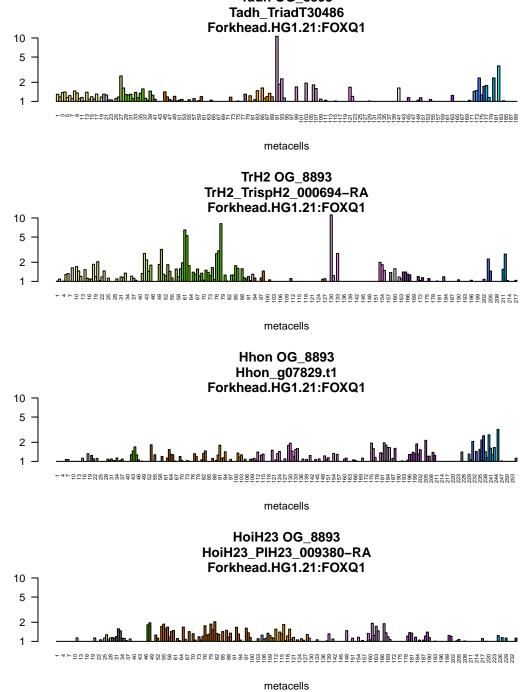
10

HoiH23 OG_8660
HoiH23_PIH23_004901-RA
Homeodomains.HG2.67:like:ALX1/ALX3/ALX4/ARX

metacells

Tadh OG_5906 Tadh_TriadT63874 IRF.HG1.7:IRF1/IRF2 10 metacells TrH2 OG_5906 TrH2_TrispH2_001644-RA IRF.HG1.7:IRF1/IRF2 10 metacells **Hhon OG_5906** Hhon_g01017.t1 IRF.HG1.7:IRF1/IRF2 metacells HoiH23 OG_5906 HoiH23_PIH23_010547-RA IRF.HG1.7:IRF1/IRF2 10 metacells





Tadh OG_8893

Tadh OG_9005 Tadh_TriadT52275 Fc_fragment_of_lgG_binding_protein,Myb.HG1.19:SNAPC4 10 metacells TrH2 OG_9005 TrH2_TrispH2_001042-RA Fc_fragment_of_IgG_binding_protein,Myb.HG1.19:SNAPC4 10 $\begin{smallmatrix} 1&4&5&5&5&5&6\\ 1&4&5&5&5&6\\ 1&4&5&5&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6&6\\ 1&4&5&6&6&6$ metacells Hhon OG_9005 Hhon_g02879.t1 Fc_fragment_of_lgG_binding_protein,Myb.HG1.19:SNAPC4 metacells HoiH23 OG_9005 HoiH23_PIH23_008293-RA $Fc_fragment_of_IgG_binding_protein, Myb. HG1.19: SNAPC4$ 10 metacells