Clustering the target scatter plots according to TFs' family and Pearson correlation coefficient

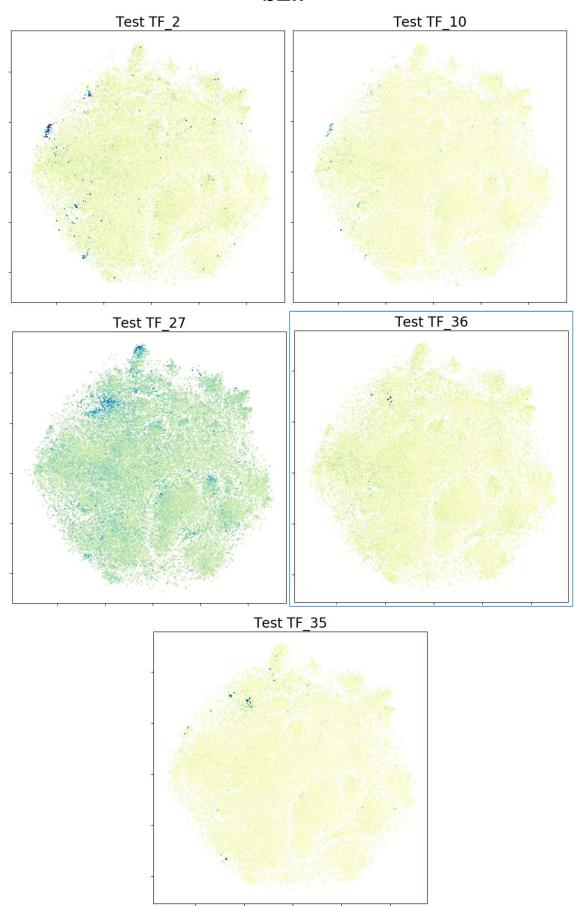
The bold text above plots indicates the family to which these TFs belong. Plots outlined by blue lines are selected to be the representative of family to constitute PBM_20. The TF name and Pearson correlation coefficient of binding intensities of paired TF are listed below plots.

All 66 TFs from the PBM data set were grouped by their families in general. In particular, TFs in the family, C2H2 ZF, were further divided into subgroups according to the Pearson correlation coefficient. Some TFs in different families with high PCC also clustered in the same group. The general cutoff of Pearson correlation coefficient is 0.4. As we can see, plots of TFs in different TF families has different patterns. In general, PBM_20 is the subset of PBM_66 without homologs.

The indicator of plots is belw:





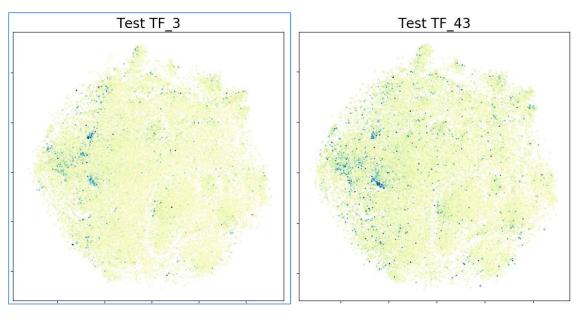


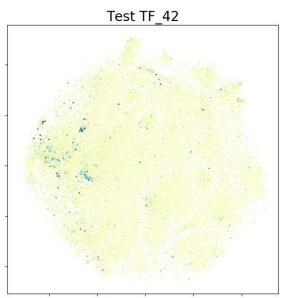
TF_2: Dbp TF_10: Nfil3 TF_27: Xbp1 TF_35: Atf3 TF_36: Atf4

TF_2, TF_35: 0.420 TF_2, TF_36: 0.506 TF_2, TF_10: 0.685 TF_2, TF_27: 0.342 TF_35, TF_36: 0.636 TF_10, TF_35: 0.462 TF_27, TF_35: 0.519 TF_10, TF_36: 0.586 TF_27, TF_36: 0.497

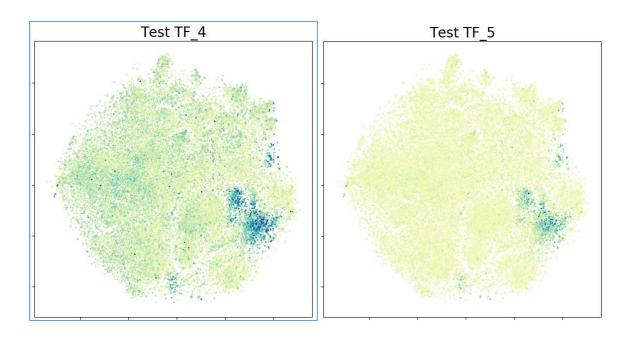
TF_10, TF_27: 0.339

Forkhead



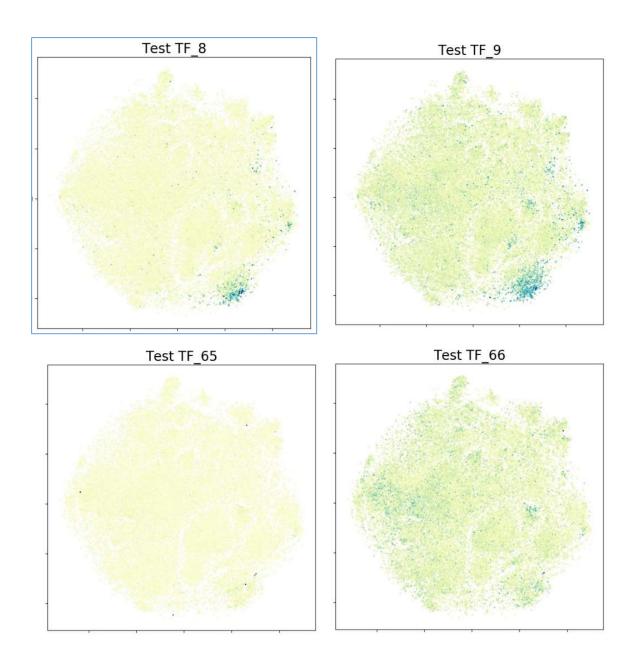


TF_3: Foxo6 TF_42: Foxc2 TF_43: Foxg1



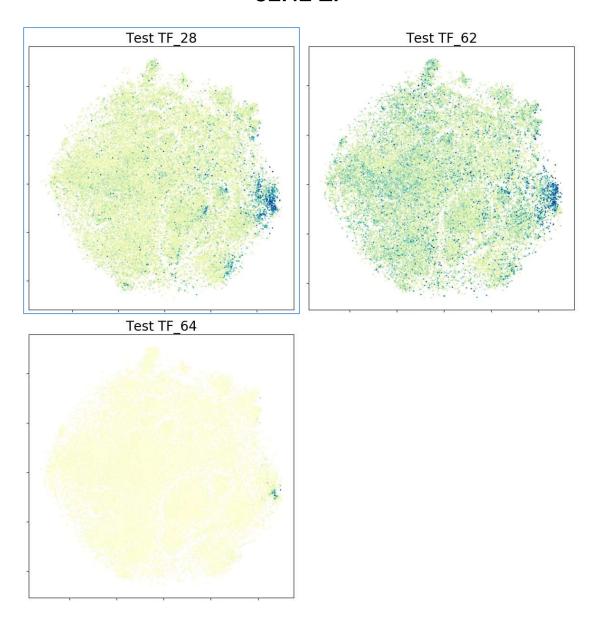
TF_4: Klf12 TF_5: Klf8

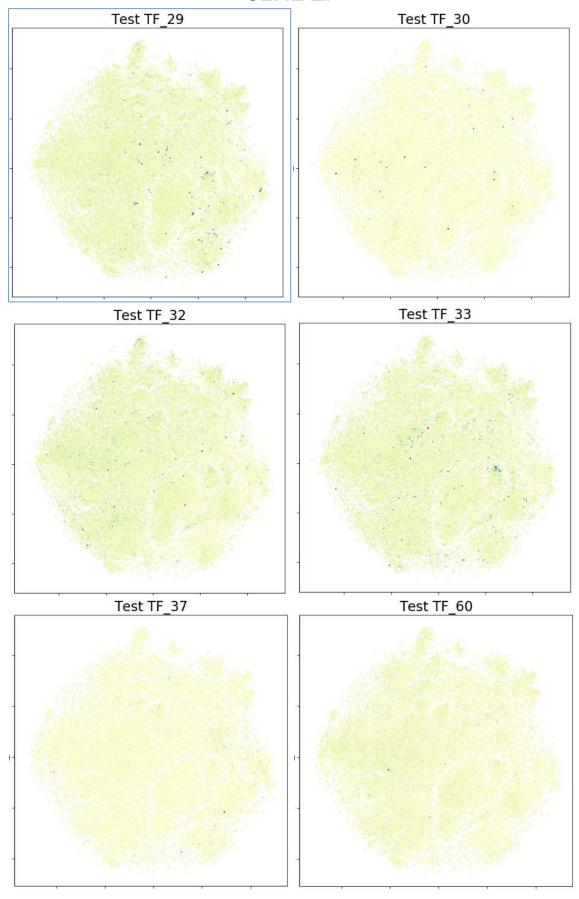
TF_4, TF_5: 0.707

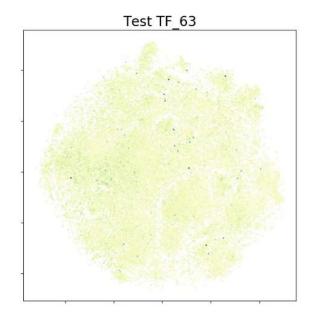


TF_8: Mzf1 TF_9: Mzf1 TF_65: Zscan10 TF_66: Zscan10

TF_8, TF_9: 0.728 TF_8, TF_65: 0.430 TF_8, TF_66: 0.480 TF_9, TF_65: 0.563 TF_9, TF_66: 0.692 TF_65, TF_66: 0.734



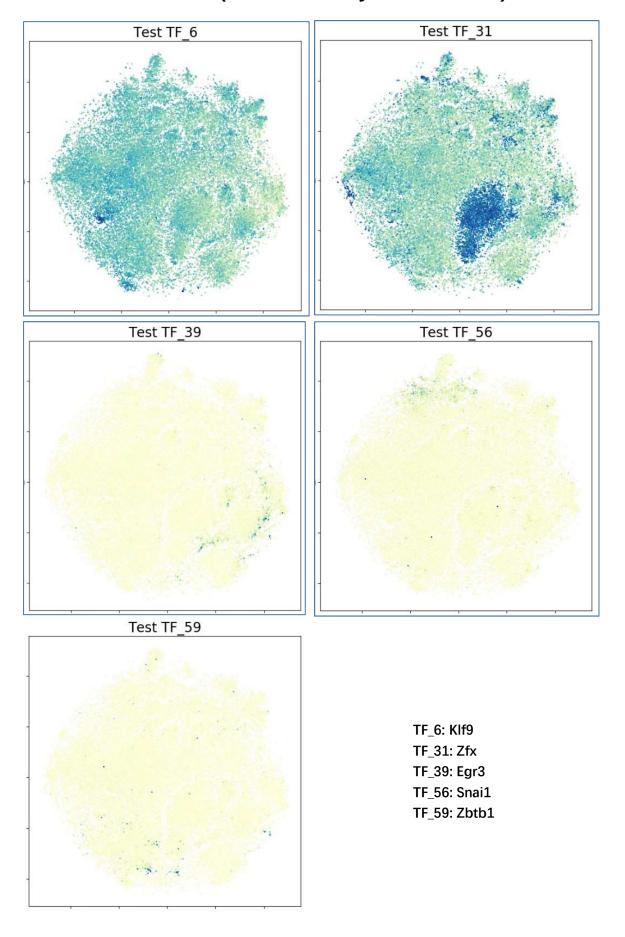




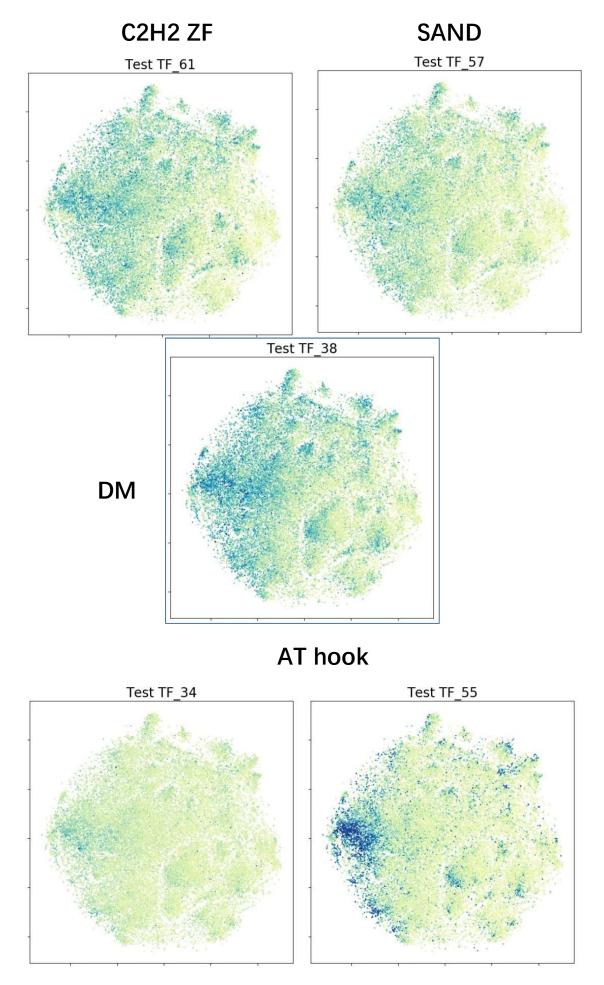
TF_29 Zfp263 TF_30 Zfp3 TF_32 Zkscan1 TF_33 Zscan10 TF_37 chr15-10391820-10399583 TF_60 Zfp300 TF_63 Zkscan5

TF_33, TF_37: 0.273	TF_33, TF_60: 0.566	TF_29, TF_33: 0.343
TF_30, TF_33: 0.467	TF_33, TF_63: 0.559	TF_32, TF_33: 0.465
TF_37, TF_60: 0.248	TF_29, TF_37: 0.170	TF_30, TF_37: 0.205
TF_37, TF_63: 0.278	TF_32, TF_37: 0.152	TF_29, TF_60: 0.283
TF_30, TF_60: 0.444	TF_60, TF_63: 0.634	TF_32, TF_60: 0.514
TF_29, TF_30: 0.222	TF_29, TF_63: 0.304	TF_29, TF_32: 0.240
TF_30, TF_63: 0.407	TF_30, TF_32: 0.371	TF_32, TF_63: 0.514

C2H2 ZF (low similarity with others)



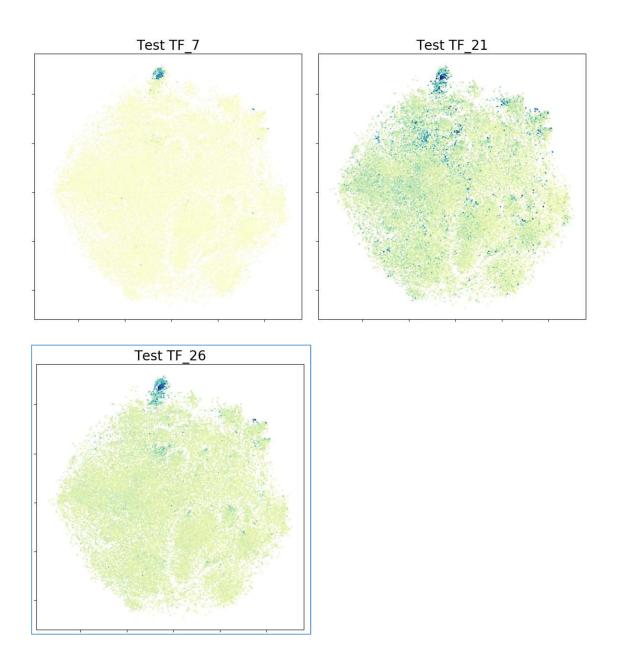
Similar but in different families



TF_34: Ahctf1 TF_55: Sdccag8

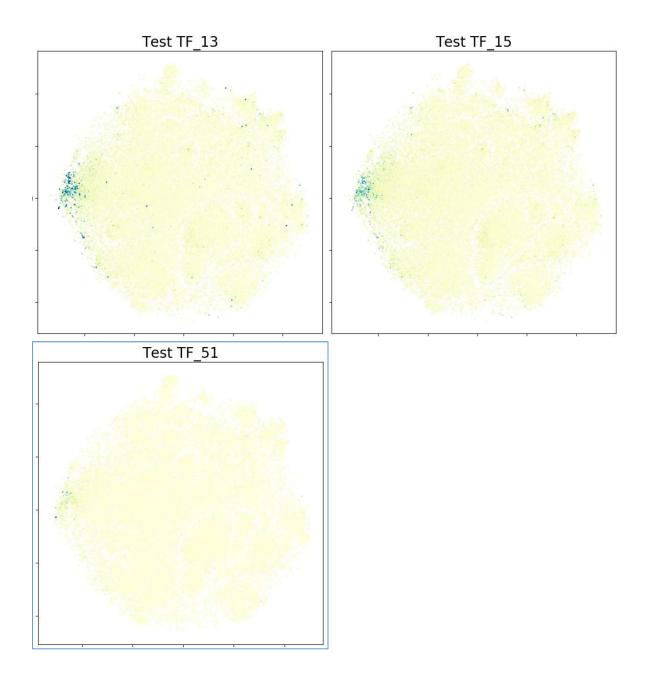
TF_57, TF_61: 0.688

bHLH

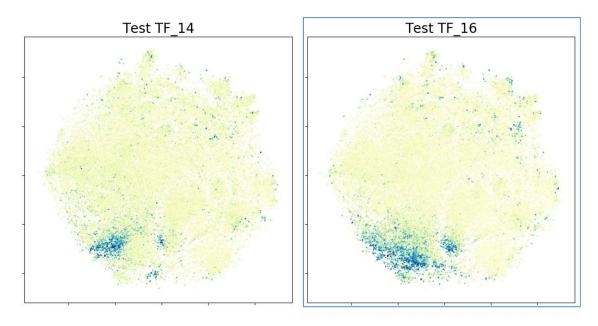


TF_7: Mlx TF_21: Srebf1 TF_26: Tfec

POU



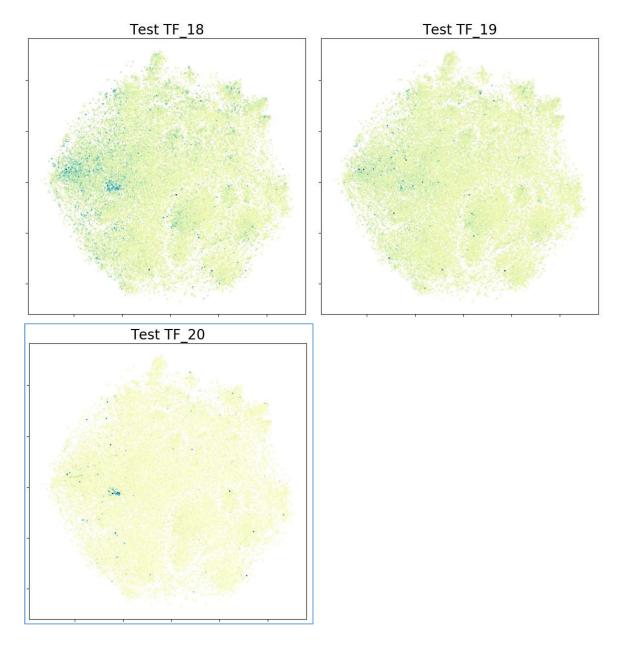
Myb/SANT



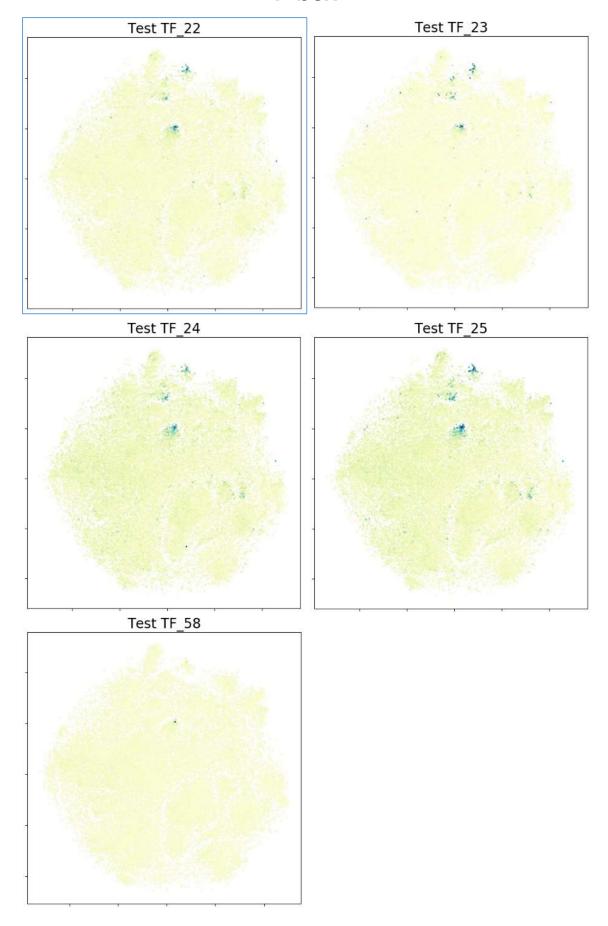
TF_14: P42pop TF_16: Prdm11

TF_14, TF_16: 0.414

Sox

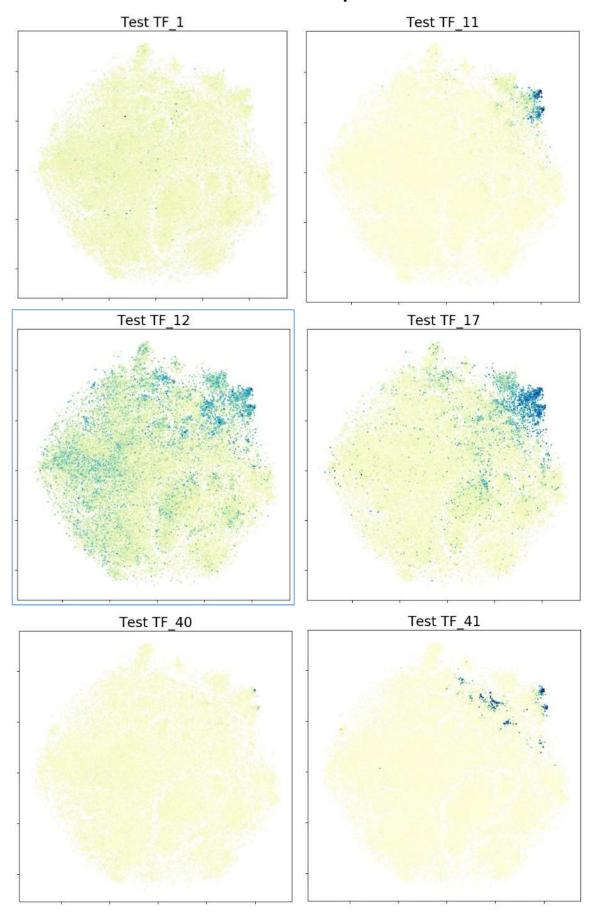


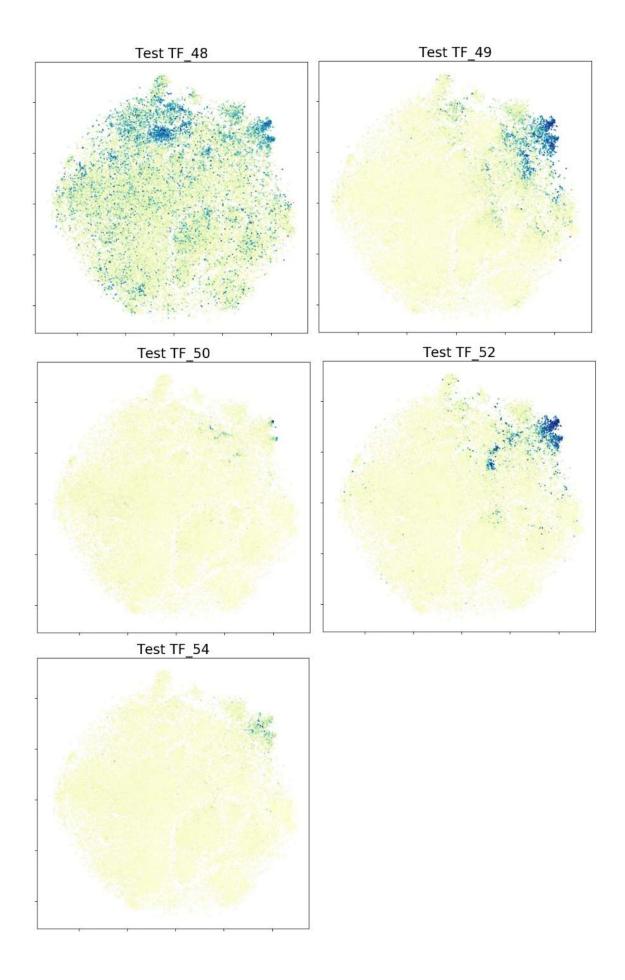
T-box



TF_22, TF_23: 0.692 TF_22, TF_24: 0.788 TF_22, TF_25: 0.800
TF_22, TF_58: 0.499 TF_23, TF_24: 0.597 TF_23, TF_25: 0.626
TF_23, TF_58: 0.366 TF_24, TF_25: 0.913 TF_24, TF_58: 0.481
TF_25, TF_58: 0.421

Nuclear receptor





TF_1: Ar	TF_11: Nr2f6	TF_12: Nr4a2
TF_17: Rorb	TF_40: Esrrb	TF_41: Esrrg
TF_48: Nr2e1	TF_49: Nr2f1	TF_50: Nr5a2

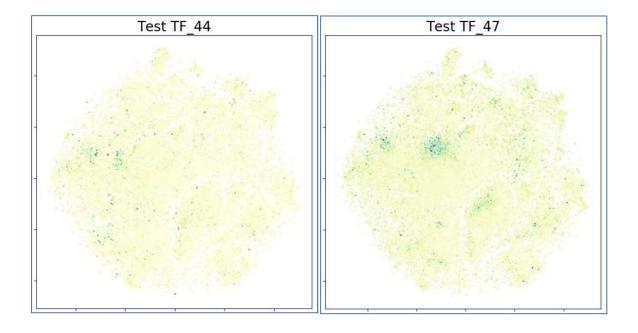
TF_52: Rarg TF_54: Rora

TF_1, TF_40: 0.457	TF_1, TF_41: 0.0545	TF_1, TF_11: 0.0203
TF_1, TF_12: 0.280	TF_1, TF_48: 0.157	TF_1, TF_17: 0.0397
TF_1, TF_49: -0.0251	TF_1, TF_50: 0.262	TF_1, TF_52: 0.136
TF_1, TF_54: 0.256	TF_40, TF_41: 0.290	TF_11, TF_40: 0.266
TF_12, TF_40: 0.482	TF_40, TF_48: 0.336	TF_17, TF_40: 0.204
TF_40, TF_49: 0.118	TF_40, TF_50: 0.606	TF_40, TF_52: 0.357
TF_40, TF_54: 0.431	TF_11, TF_41: 0.451	TF_12, TF_41: 0.285
TF_41, TF_48: 0.207	TF_17, TF_41: 0.375	TF_41, TF_49: 0.409
TF_41, TF_50: 0.508	TF_41, TF_52: 0.457	TF_41, TF_54: 0.282
TF_11, TF_12: 0.386	TF_11, TF_48: 0.280	TF_11, TF_17: 0.651
TF_11, TF_49: 0.610	TF_11, TF_50: 0.400	TF_11, TF_52: 0.720
TF_11, TF_54: 0.507	TF_12, TF_48: 0.389	TF_12, TF_17: 0.362
TF_12, TF_49: 0.337	TF_12, TF_50: 0.437	TF_12, TF_52: 0.508
TF_12, TF_54: 0.477	TF_17, TF_48: 0.251	TF_48, TF_49: 0.262
TF_48, TF_50: 0.263	TF_48, TF_52: 0.376	TF_48, TF_54: 0.309
TF_17, TF_49: 0.640	TF_17, TF_50: 0.326	TF_17, TF_52: 0.681
TF_17, TF_54: 0.750	TF_49, TF_50: 0.266	TF_49, TF_52: 0.659
TF_49, TF_54: 0.449	TF_50, TF_52: 0.433	TF_50, TF_54: 0.453
TF_52, TF_54: 0.607		

Others

GATA

Homeodomain



RFX

