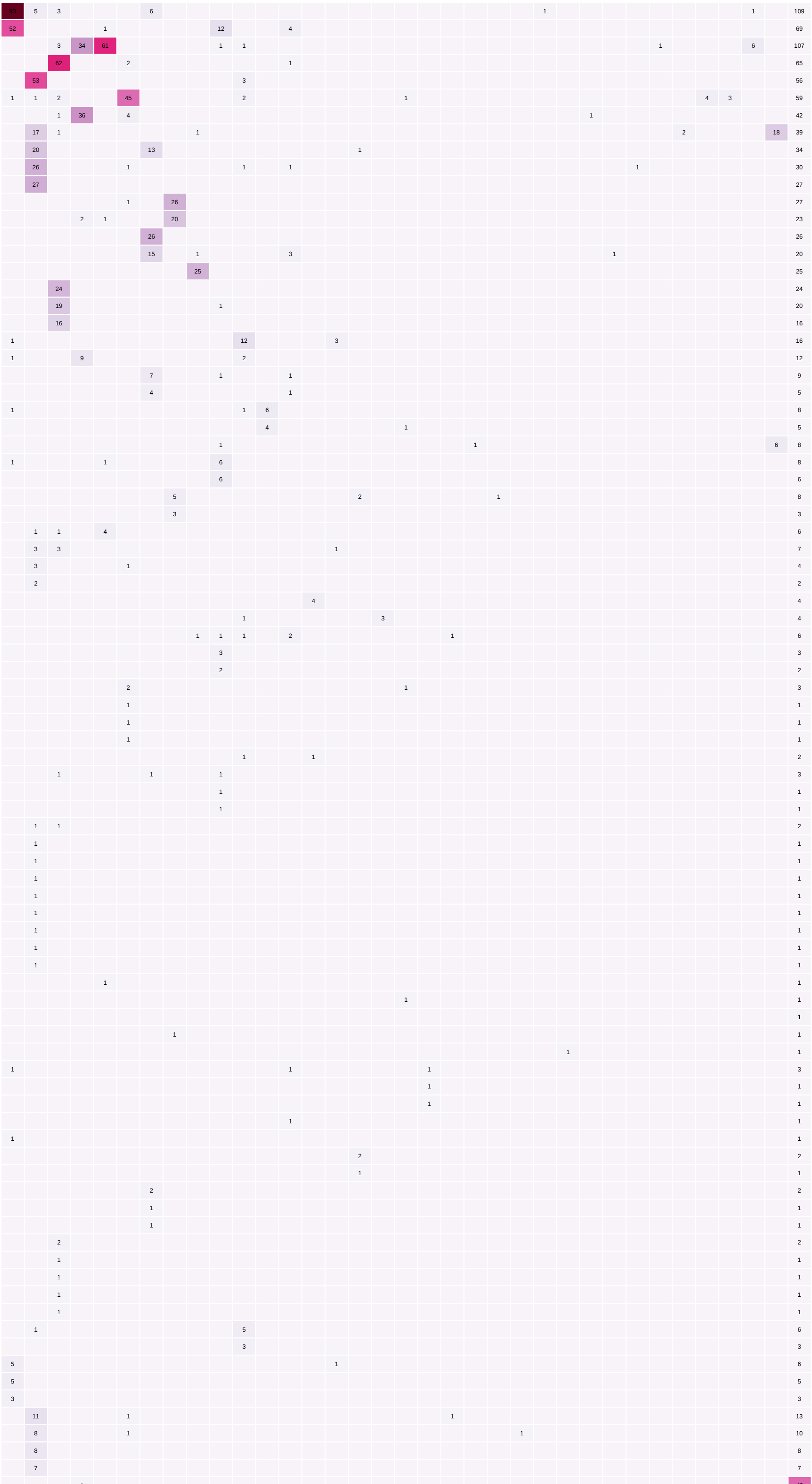
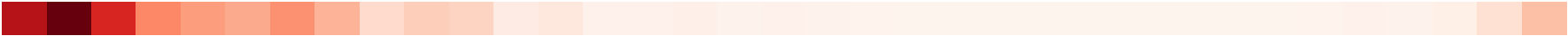


Reference groups – ARI = 0.38

RSAT matrix–clustering



- cluster_10
- cluster_23
- cluster_20
- cluster_12
- cluster_03
- cluster_06
- cluster_16
- cluster_18
- cluster_13
- cluster_01
- cluster_22
- cluster_02
- cluster_40
- cluster_49
- cluster_38
- cluster_32
- cluster_09
- cluster_47
- cluster_37
- cluster_07
- cluster_19
- cluster_17
- cluster_46
- cluster_30
- cluster_42
- cluster_14
- cluster_05
- cluster_33
- cluster_41
- cluster_48
- cluster_26
- cluster_08
- cluster_29
- cluster_52
- cluster_39
- cluster_31
- cluster_50
- cluster_56
- cluster_57
- cluster_27
- cluster_79
- cluster_63
- cluster_78
- cluster_51
- cluster_35
- cluster_71
- cluster_74
- cluster_36
- cluster_86
- cluster_83
- cluster_81
- cluster_72
- cluster_70
- cluster_66
- cluster_59
- cluster_64
- cluster_77
- cluster_73
- cluster_62
- cluster_58
- cluster_61
- cluster_43
- cluster_65
- cluster_75
- cluster_80
- cluster_85
- cluster_53
- cluster_67
- cluster_54
- cluster_69
- cluster_82
- cluster_55
- cluster_84
- cluster_76
- cluster_60
- cluster_68
- cluster_11
- cluster_45
- cluster_25
- cluster_15
- cluster_44
- cluster_34
- cluster_04
- cluster_28
- cluster_24
- cluster_21



- AP2/EREBP
- Mytophan cluster factors
- GCM domain factors
- Basic leucine zipper factors (bZIP)
- Basic helix–loop–helix factors (bHLH)
- Homeo domain factors
- Other C4 zinc finger–type factors
- C3H(C),C2HC zinc–fingers like factors
- MADS box factors
- C2H2 zinc finger factors
- B3
- C3H zinc finger factors
- CPP
- HC3 zinc ribbon factors
- Fork head/winged helix factors
- EIL
- BBR/BPC
- ARID
- RWP–RK
- RING
- S1Fa–like
- Nuclear receptors with C4 zinc fingers
- mTERF
- LEAFY
- GRAS
- CH3
- Unknown
- Basic helix–loop–helix domain factors (bHLH)
- PLATZ
- A.T hook factors
- HMG domain factors
- Heteromeric CCAAT–binding factors
- Heat shock factors
- TCP