Reference groups − ARI = 0.42
RSAT matrix−clustering
cluster_01
cluster_07
cluster_11
cluster_03
cluster_06
cluster_10
cluster_15
cluster_04
cluster_25
cluster_17
cluster_26
cluster_05
cluster_28
cluster_35
cluster_31
cluster_43
cluster_09
cluster_14
cluster_23
cluster_18
cluster_20
cluster_21
cluster_39
cluster_19
cluster_45
cluster_46
cluster_27
cluster_65
cluster_58
cluster_61
cluster_34
cluster_64
cluster_48
cluster_59
cluster_66
cluster_62
cluster_60
cluster_57
cluster_53
cluster_52
cluster_56
cluster_51
cluster_47
cluster_50
cluster_44
cluster_63
cluster_55
cluster_49
cluster_54
cluster_24
cluster_22
cluster_38
cluster_41
cluster_42
cluster_30
cluster_29
cluster_36
cluster_33
cluster_40
cluster_12
cluster_37
cluster_32
cluster_08
cluster_13
cluster_02
cluster_16
C3H(C),C2HC zinc fingers factors
Heat shock factors
CPP
C3H zinc finger factors
Fork head/winged helix factors
ARID
EIL
RWP−RK
RING
S1Fa−like
Other all−alpha−helical DNA−binding domains
Nuclear receptors with C4 zinc fingers
LEAFY
GeBP/Storekeeper
mTERF
CH3
GRAS
PLATZ
BBR/BPC
A.T hook factors
High−mobility group (HMG) domain factors
Heteromeric CCAAT−binding factors
MADS box factors
Basic leucine zipper factors (bZIP)
Basic helix−loop−helix factors (bHLH)
AP2/EREBP