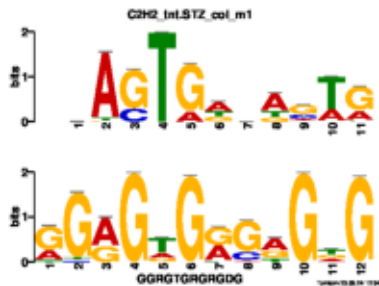
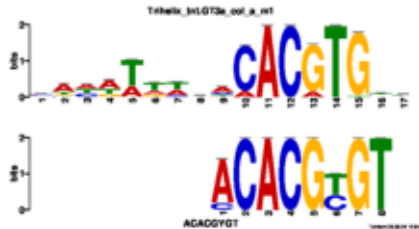


Nr: 1)
 Target ID: C2H2_tnt.STZ_col_m1
 p-value: 3.82294e-05
 E-value: 0.037656
 q-value: 0.0752293



Nr: 1)
 Target ID: Trihelix_tnt.GT3a_col_a_m1
 p-value: 2.75873e-05
 E-value: 0.0271735
 q-value: 0.0262807

ABA_JA_up



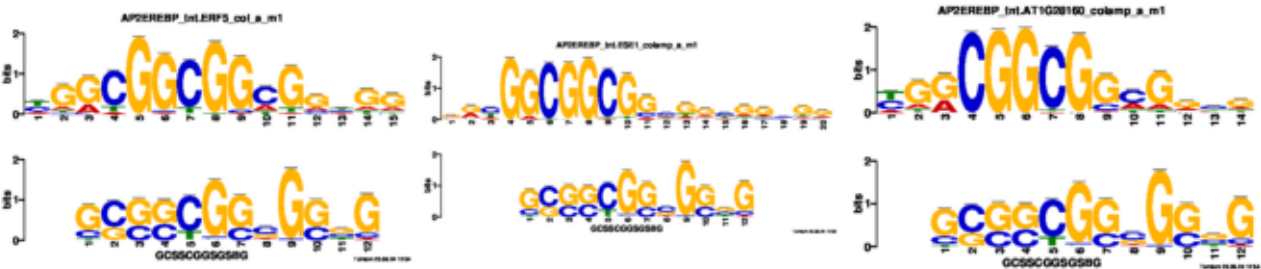
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
AP2EREBP_tnt.ERF5_col_a_m1
4.93562e-07
0.000486159
0.000263962

2)
AP2EREBP_tnt.ESE1_colamp_a_m1
5.16633e-07
0.000508883
0.000263962

3)
AP2EREBP_tnt.AT1G28160_colamp_a_m1
5.58583e-07
0.000550204
0.000263962

ABA_down



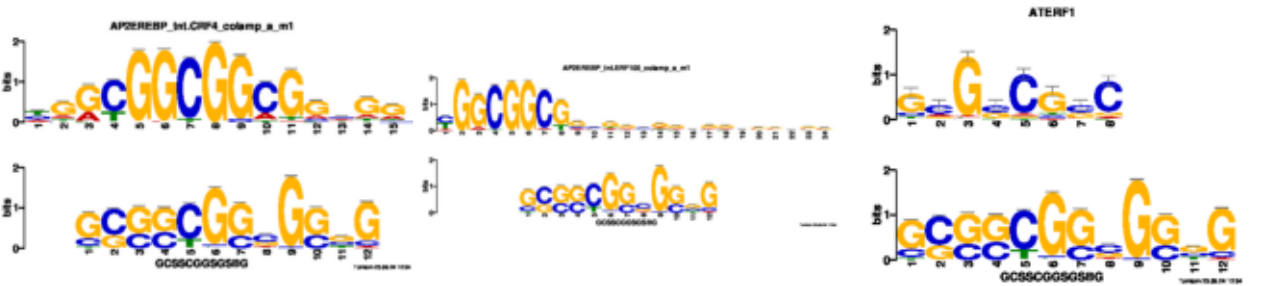
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
AP2EREBP_tnt.CRF4_colamp_a_m1
8.65817e-07
0.00085283
0.000263962

5)
AP2EREBP_tnt.ERF105_colamp_a_m1
9.61714e-07
0.000947288
0.000263962

6)
ATERF1
9.63705e-07
0.000949249
0.000263962

ABA_down



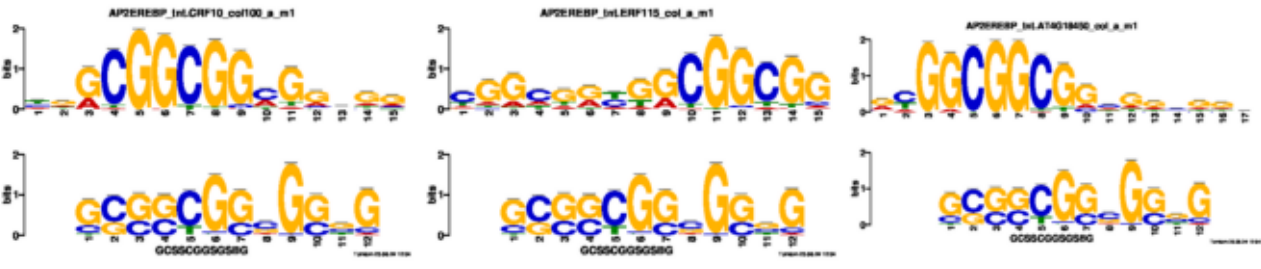
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
AP2EREBP_tnt.CRF10_col100_a_m1
1.0112e-06
0.00099603
0.000263962

8)
AP2EREBP_tnt.ERF115_col_a_m1
1.37096e-06
0.0013504
0.00028146

9)
AP2EREBP_tnt.AT4G18450_col_a_m1
1.50454e-06
0.00148197
0.00028146

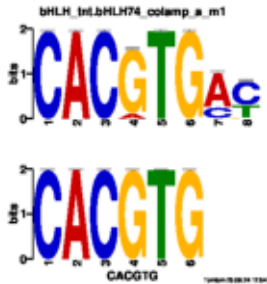
ABA_down



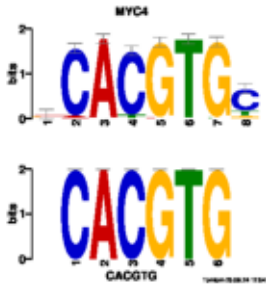
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
bHLH_tnt.bHLH74_colamp_a_m1
9.25646e-06
0.00911762
0.00170073

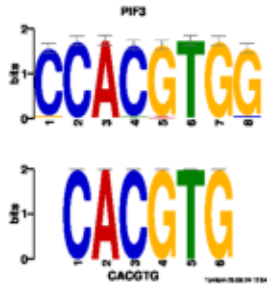
ABA_up



2)
MYC4
9.25646e-06
0.00911762
0.00170073



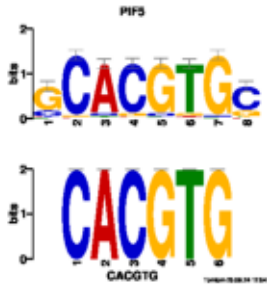
3)
PIF3
9.25646e-06
0.00911762
0.00170073



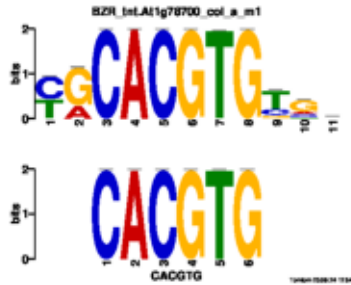
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
PIF5
9.25646e-06
0.00911762
0.00170073

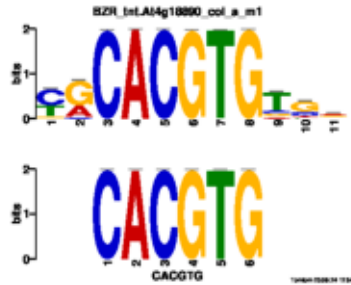
ABA_up



5)
BZR_tnt.At1g78700_col_a_m1
1.85128e-05
0.0182351
0.00170073



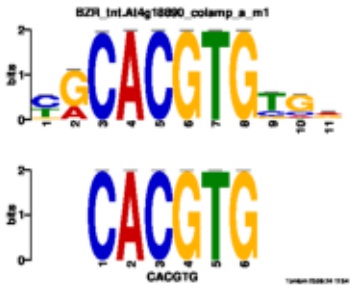
6)
BZR_tnt.At4g18890_col_a_m1
1.85128e-05
0.0182351
0.00170073



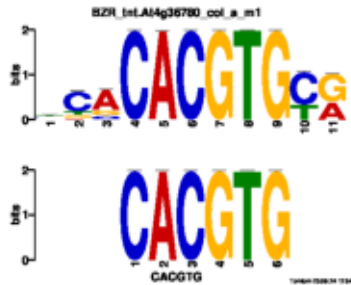
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
BZR_tnt.At4g18890_colamp_a_m1
1.85128e-05
0.0182351
0.00170073

ABA_up



8)
BZR_tnt.At4g36780_col_a_m1
1.85128e-05
0.0182351
0.00170073

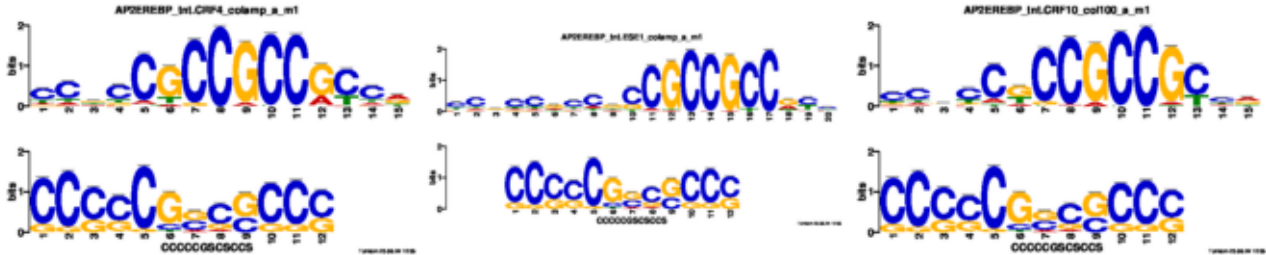


9)
BZR_tnt.At4g36780_colamp_a_m1
2.77691e-05
0.0273526
0.00170073



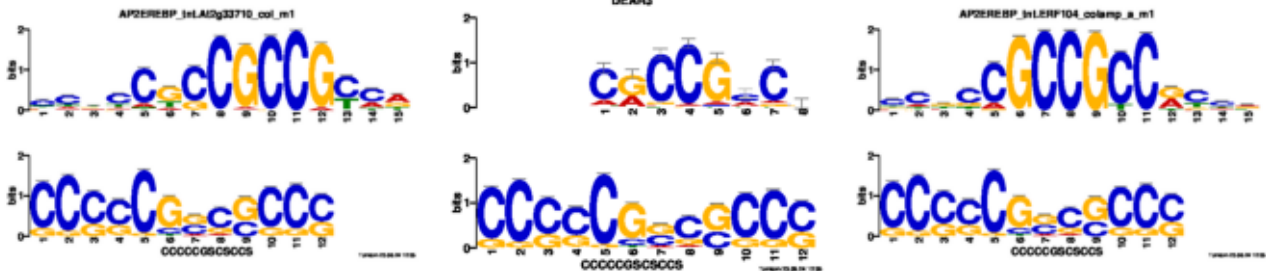
Nr:	1)	2)	3)
Target ID:	AP2EREBP_tnt.CRF4_colamp_a_m1	AP2EREBP_tnt.ESE1_colamp_a_m1	AP2EREBP_tnt.CRF10_col100_a_m1
p-value:	9.4122e-06	1.01657e-05	1.13851e-05
E-value:	0.00927102	0.0100132	0.0112143
q-value:	0.0024157	0.0024157	0.0024157

JA_down



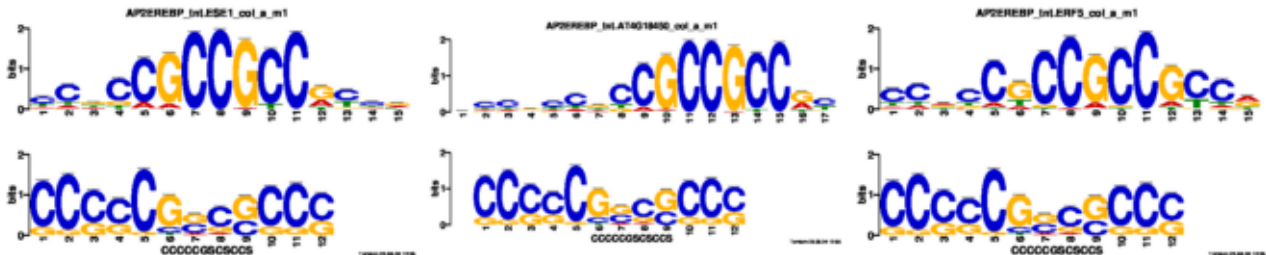
Nr:	4)	5)	6)
Target ID:	AP2EREBP_tnt.At2g33710_col_m1	DEAR3	AP2EREBP_tnt.ERF104_colamp_a_m1
p-value:	1.21305e-05	1.36207e-05	1.37713e-05
E-value:	0.0119485	0.0134164	0.0135647
q-value:	0.0024157	0.0024157	0.0024157

JA_down



Nr:	7)	8)	9)
Target ID:	AP2EREBP_tnt.ESE1_col_a_m1	AP2EREBP_tnt.AT4G18450_col_a_m1	AP2EREBP_tnt.ERF5_col_a_m1
p-value:	1.37713e-05	1.47552e-05	1.56339e-05
E-value:	0.0135647	0.0145339	0.0153994
q-value:	0.0024157	0.0024157	0.0024157

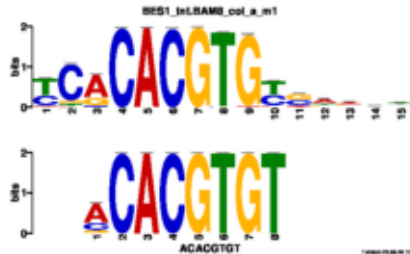
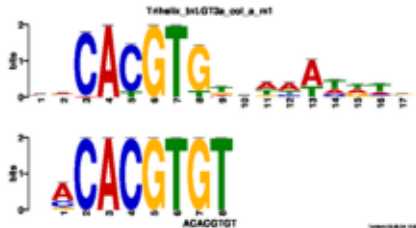
JA_down



Nr: 1)
 Target ID: Trihelix_tnt.GT3a_col_a_m1
 p-value: 4.47836e-06
 E-value: 0.00441119
 q-value: 0.00638493

2)
 BES1_tnt.BAM8_col_a_m1
 3.56239e-05
 0.0350896
 0.0137908

JA_up



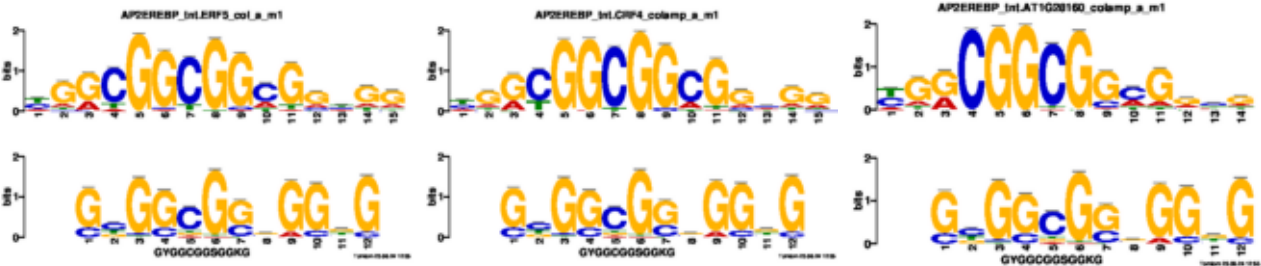
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
AP2EREBP_tnt.ERF5_col_a_m1
2.7148e-06
0.00267408
0.00160525

2)
AP2EREBP_tnt.CRF4_colamp_a_m1
5.65816e-06
0.00557329
0.00160525

3)
AP2EREBP_tnt.AT1G28160_colamp_a_m1
6.77058e-06
0.00666902
0.00160525

SA_JA_down



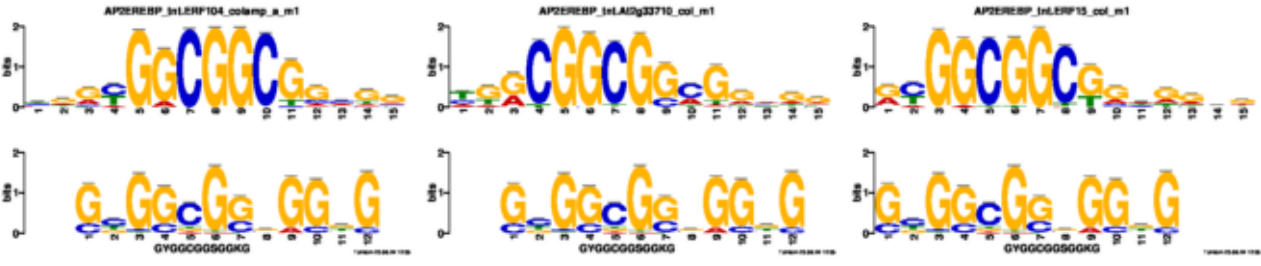
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
AP2EREBP_tnt.ERF104_colamp_a_m1
8.31431e-06
0.0081896
0.00160525

5)
AP2EREBP_tnt.At2g33710_col_m1
8.31431e-06
0.0081896
0.00160525

6)
AP2EREBP_tnt.ERF15_col_m1
8.85732e-06
0.00872446
0.00160525

SA_JA_down



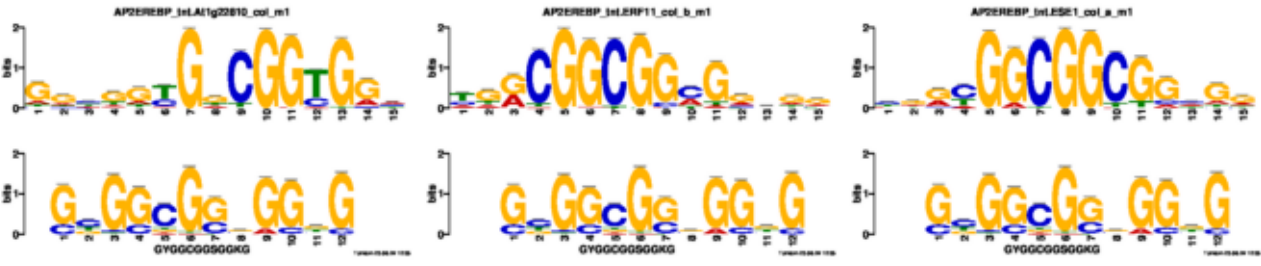
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
AP2EREBP_tnt.At1g22810_col_m1
9.43363e-06
0.00929212
0.00160525

8)
AP2EREBP_tnt.ERF11_col_b_m1
9.43363e-06
0.00929212
0.00160525

9)
AP2EREBP_tnt.ESE1_col_a_m1
1.06943e-05
0.0105339
0.00160525

SA_JA_down



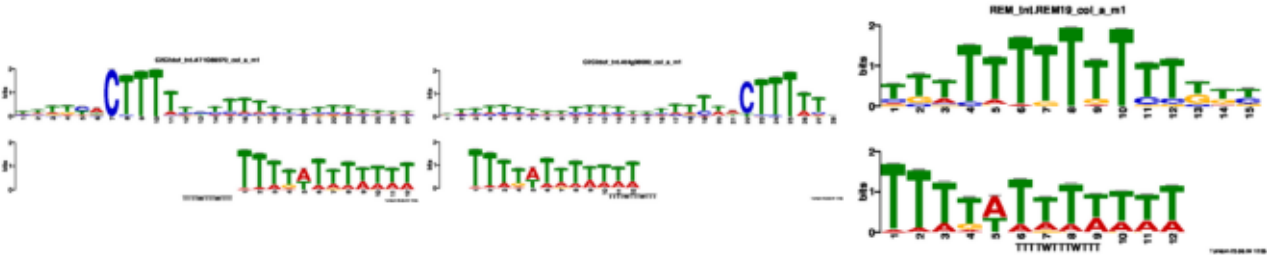
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
C2C2dof_tnt.AT1G69570_col_a_m1
2.78707e-06
0.00274527
0.00180231

2)
C2C2dof_tnt.At4g38000_col_a_m1
2.96127e-06
0.00291685
0.00180231

3)
REM_tnt.REM19_col_a_m1
2.99187e-06
0.002947
0.00180231

SA_JA_up



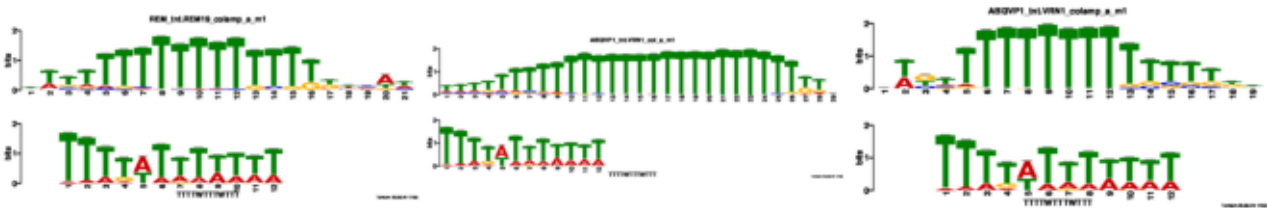
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
REM_tnt.REM19_colamp_a_m1
3.79537e-06
0.00373844
0.00180231

5)
ABI3VP1_tnt.VRN1_col_a_m1
5.9025e-06
0.00581396
0.00186052

6)
ABI3VP1_tnt.VRN1_colamp_a_m1
7.68971e-06
0.00757436
0.00186052

SA_JA_up



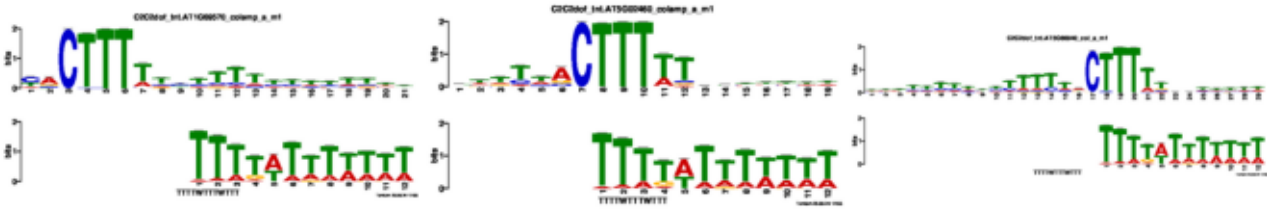
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
C2C2dof_tnt.AT1G69570_colamp_a_m1
8.48817e-06
0.00836085
0.00186052

8)
C2C2dof_tnt.AT5G02460_colamp_a_m1
8.68765e-06
0.00855734
0.00186052

9)
C2C2dof_tnt.AT5G66940_col_a_m1
9.04844e-06
0.00891272
0.00186052

SA_JA_up



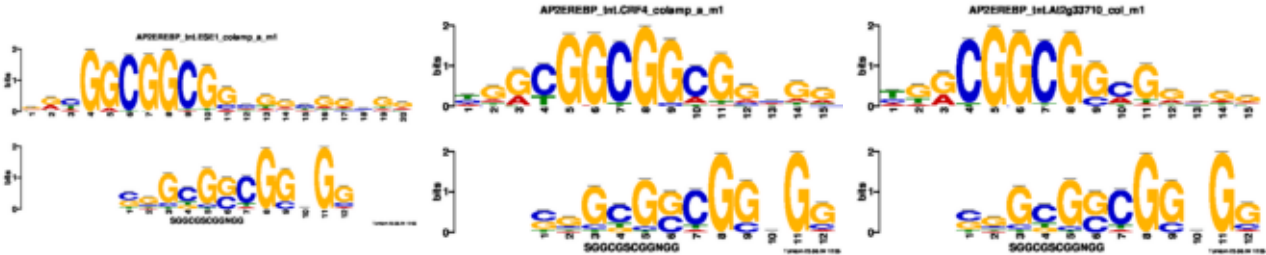
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
AP2EREBP_tnt.ESE1_colamp_a_m1
1.24676e-06
0.00122806
0.000406627

2)
AP2EREBP_tnt.CRF4_colamp_a_m1
1.35109e-06
0.00133082
0.000406627

3)
AP2EREBP_tnt.At2g33710_col_m1
1.80454e-06
0.00177747
0.000406627

SA_down



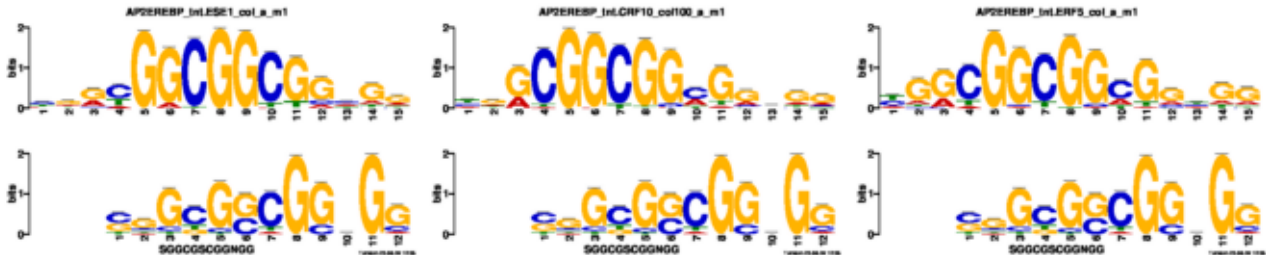
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
AP2EREBP_tnt.ESE1_col_a_m1
1.93584e-06
0.0019068
0.000406627

5)
AP2EREBP_tnt.CRF10_col100_a_m1
1.93584e-06
0.0019068
0.000406627

6)
AP2EREBP_tnt.ERF5_col_a_m1
1.93584e-06
0.0019068
0.000406627

SA_down



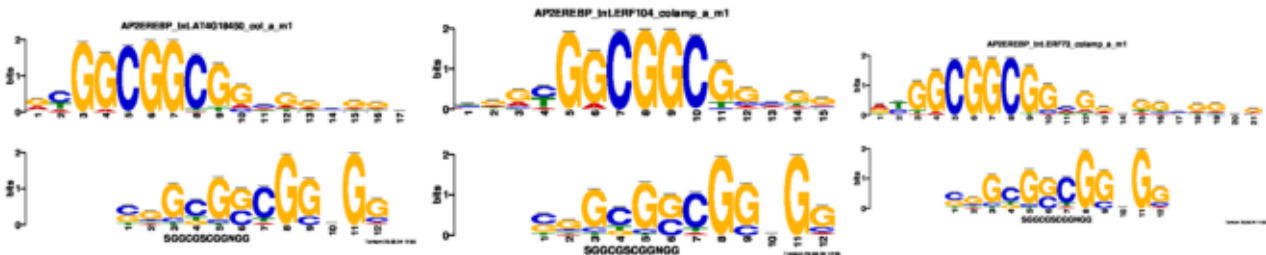
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
AP2EREBP_tnt.AT4G18450_col_a_m1
1.98472e-06
0.00195495
0.000406627

8)
AP2EREBP_tnt.ERF104_colamp_a_m1
2.07508e-06
0.00204396
0.000406627

9)
AP2EREBP_tnt.ERF73_colamp_a_m1
2.22578e-06
0.00219239
0.000406627

SA_down



Nr: 1)
 Target ID: BZR_tnt.At4g18890_col_a_m1
 p-value: 4.41241e-05
 E-value: 0.0434622
 q-value: 0.0321903

SA_up

