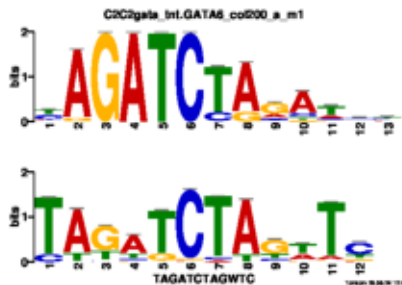
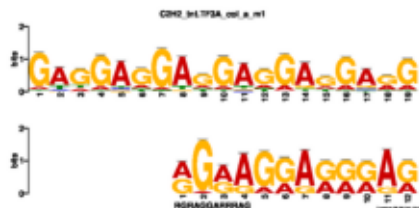


ABAJA_under



Nr: 4)
Target ID: C2H2_tnt.TF3A_col_a_m1
p-value: 2.53e-05
E-value: 2.49e-02
q-value: 4.98e-02



ABAJA_under

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

1)
C2C2gata_tnt.ZML1_col_a_m1
8.49e-07
8.36e-04
1.67e-03

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

2)
C2C2gata_tnt.ZML1_col_a_m1
9.27e-07
9.13e-04
1.82e-03

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

3)
C2C2gata_tnt.ZML2_col_a_m1
9.24e-06
9.11e-03
9.10e-03

ABA_under



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

4)
C2C2gata_tnt.ZML2_col_a_m1
1.10e-05
1.08e-02
1.08e-02

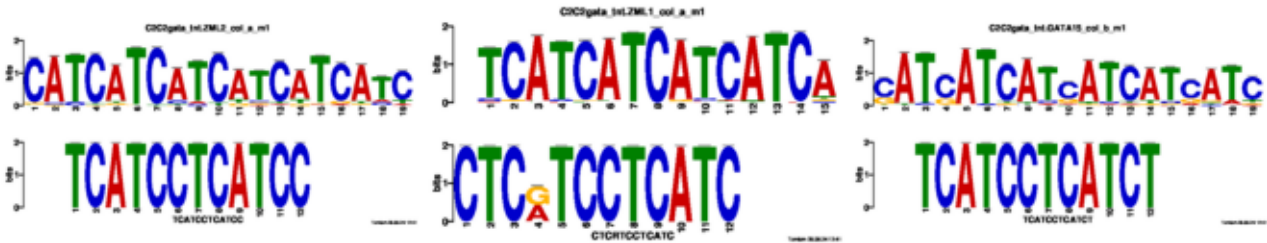
Nr:
Target ID:
p-value:
E-value:
q-value:

5)
C2C2gata_tnt.ZML1_col_a_m1
6.55e-06
6.45e-03
1.29e-02

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

6)
C2C2gata_tnt.GATA15_col_b_m1
4.86e-05
4.79e-02
3.19e-02

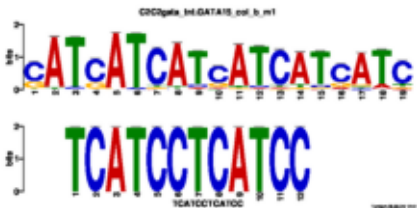
ABA_under



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

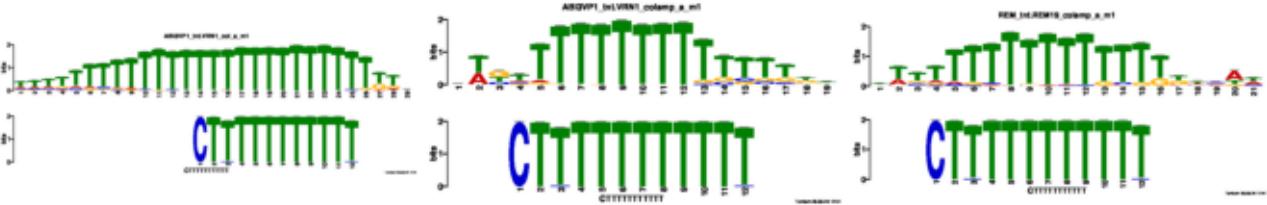
7)
C2C2gata_tnt.GATA15_col_b_m1
4.98e-05
4.91e-02
3.26e-02

ABA_under



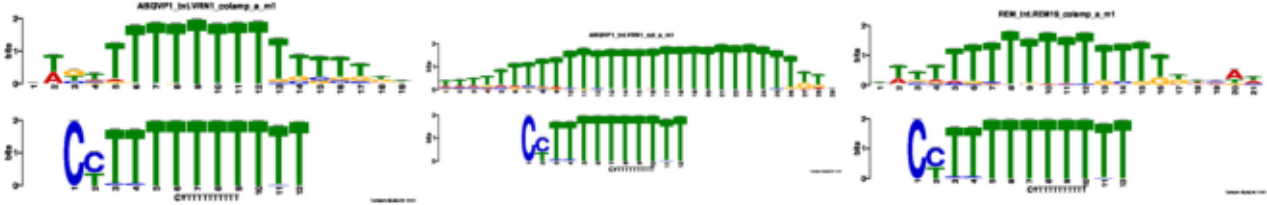
Nr:	1)	2)	3)
Target ID:	ABI3VP1_tnt.VRN1_col_a_m1	ABI3VP1_tnt.VRN1_colamp_a_m1	REM_tnt.REM19_colamp_a_m1
p-value:	1.70e-08	6.58e-08	1.64e-07
E-value:	1.67e-05	6.48e-05	1.62e-04
q-value:	3.27e-05	6.34e-05	1.06e-04

JA_over



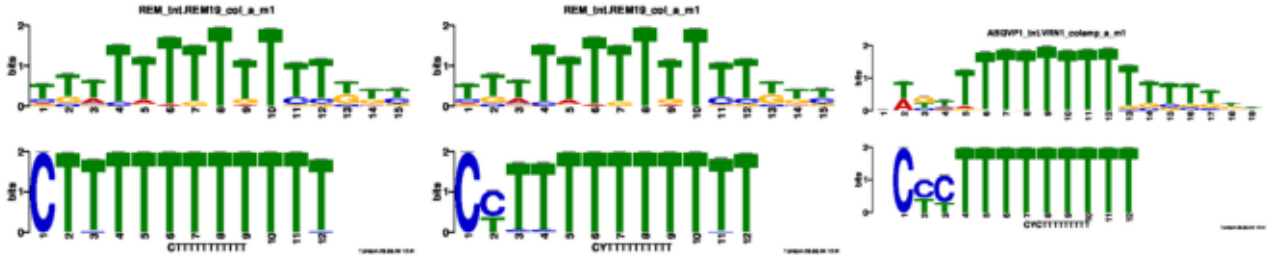
Nr:	4)	5)	6)
Target ID:	ABI3VP1_tnt.VRN1_colamp_a_m1	ABI3VP1_tnt.VRN1_col_a_m1	REM_tnt.REM19_colamp_a_m1
p-value:	6.91e-08	4.22e-07	4.00e-07
E-value:	6.81e-05	4.15e-04	3.94e-04
q-value:	1.34e-04	2.74e-04	2.74e-04

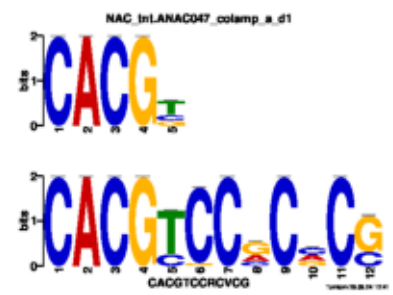
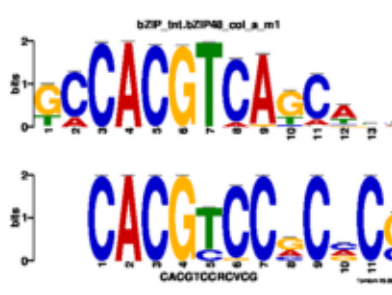



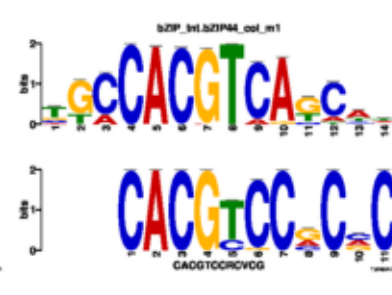

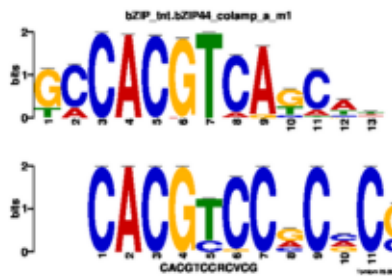
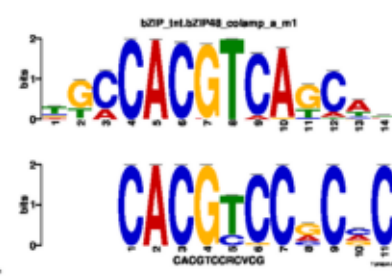
JA_over



Nr:	7)	8)	9)
Target ID:	REM_tnt.REM19_col_a_m1	REM_tnt.REM19_col_a_m1	ABI3VP1_tnt.VRN1_colamp_a_m1
p-value:	1.22e-06	1.79e-06	9.36e-07
E-value:	1.20e-03	1.76e-03	9.22e-04
q-value:	5.86e-04	8.71e-04	1.83e-03

JA_over



SAJA_over	<p>Nr: 1) Target ID: NAC_tnt.ANAC047_colamp_a_d1 p-value: 1.55e-05 E-value: 1.52e-02 q-value: 1.72e-02</p> 	<p>Nr: 2) Target ID: bZIP_tnt.bZIP48_col_a_m1 p-value: 7.31e-05 E-value: 7.20e-02 q-value: 1.72e-02</p> 	<p>Nr: 3) Target ID: bZIP_tnt.bZIP43_col_a_m1 p-value: 7.97e-05 E-value: 7.85e-02 q-value: 1.72e-02</p> 
SAJA_over	<p>Nr: 4) Target ID: bZIP_tnt.bZIP42_col_a_m1 p-value: 8.52e-05 E-value: 8.39e-02 q-value: 1.72e-02</p> 	<p>Nr: 5) Target ID: bZIP_tnt.GBF5_col_v3a_m1 p-value: 9.08e-05 E-value: 8.94e-02 q-value: 1.72e-02</p> 	<p>Nr: 6) Target ID: bZIP_tnt.bZIP44_col_m1 p-value: 9.70e-05 E-value: 9.55e-02 q-value: 1.72e-02</p> 
SAJA_over	<p>Nr: 7) Target ID: bZIP_tnt.bZIP42_colamp_a_m1 p-value: 1.01e-04 E-value: 9.94e-02 q-value: 1.72e-02</p> 	<p>Nr: 8) Target ID: bZIP_tnt.bZIP44_colamp_a_m1 p-value: 1.08e-04 E-value: 1.06e-01 q-value: 1.72e-02</p> 	<p>Nr: 9) Target ID: bZIP_tnt.bZIP48_colamp_a_m1 p-value: 1.11e-04 E-value: 1.09e-01 q-value: 1.72e-02</p> 

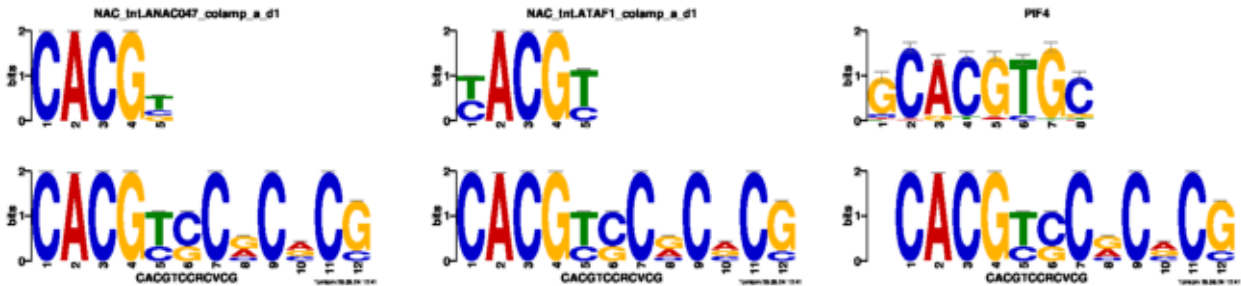
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
NAC_tnt.ANAC047_colamp_a_d1
1.96e-06
1.93e-03
3.77e-03

2)
NAC_tnt.ATAF1_colamp_a_d1
1.93e-04
1.90e-01
3.52e-02

3)
PIF4
2.00e-04
1.97e-01
3.52e-02

SAJA_under



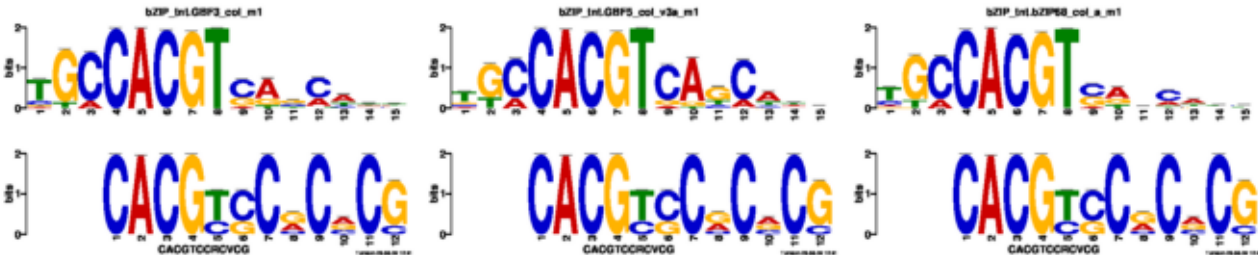
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
bZIP_tnt.GBF3_col_m1
2.15e-04
2.12e-01
3.52e-02

5)
bZIP_tnt.GBF5_col_v3a_m1
2.15e-04
2.12e-01
3.52e-02

6)
bZIP_tnt.bZIP68_col_a_m1
2.45e-04
2.41e-01
3.52e-02

SAJA_under



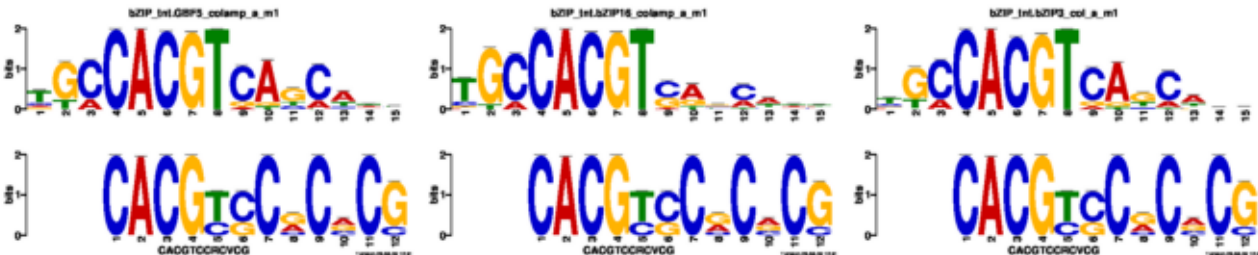
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
bZIP_tnt.GBF5_colamp_a_m1
2.45e-04
2.41e-01
3.52e-02

8)
bZIP_tnt.bZIP16_colamp_a_m1
2.45e-04
2.41e-01
3.52e-02

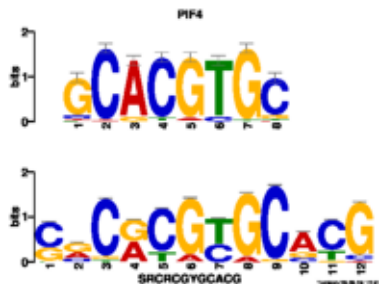
9)
bZIP_tnt.bZIP3_col_a_m1
2.45e-04
2.41e-01
3.52e-02

SAJA_under

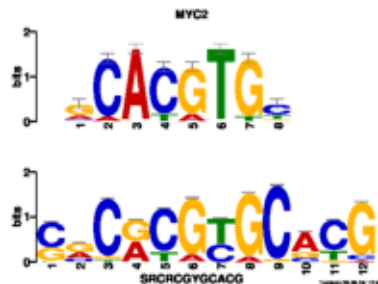


Nr: 1)
Target ID: PIF4
p-value: 4.26e-05
E-value: 4.20e-02
q-value: 2.33e-02

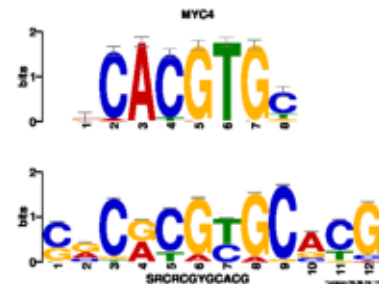
SA_over



Nr: 2)
Target ID: MYC2
p-value: 4.77e-05
E-value: 4.70e-02
q-value: 2.33e-02

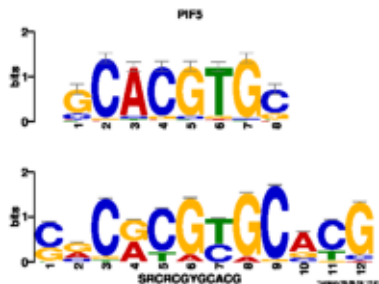


Nr: 3)
Target ID: MYC4
p-value: 8.23e-05
E-value: 8.11e-02
q-value: 2.55e-02

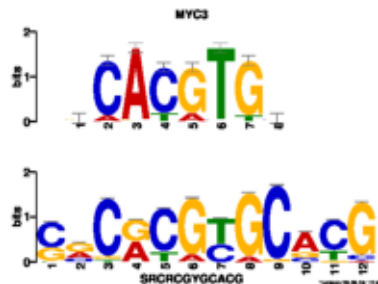


Nr: 4)
Target ID: PIF5
p-value: 9.15e-05
E-value: 9.02e-02
q-value: 2.55e-02

SA_over



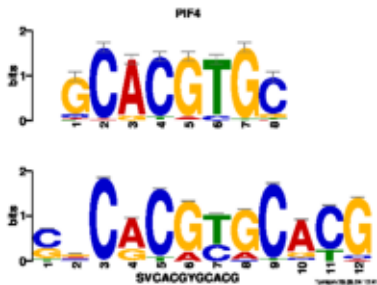
Nr: 5)
Target ID: MYC3
p-value: 1.53e-04
E-value: 1.51e-01
q-value: 3.32e-02



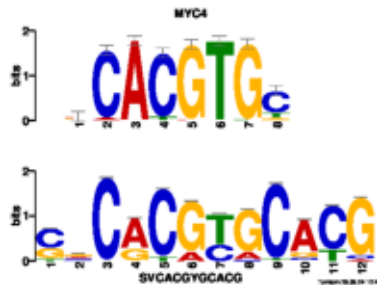
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
PIF4
1.68e-05
1.65e-02
1.18e-02

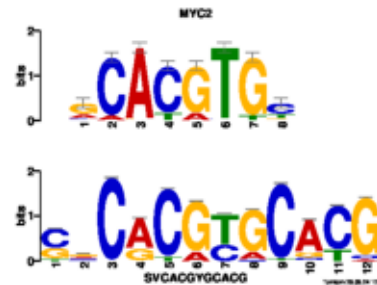
SA_under



2)
MYC4
1.88e-05
1.85e-02
1.18e-02



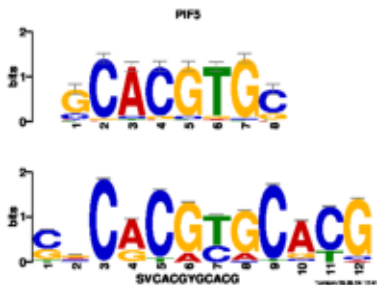
3)
MYC2
4.94e-05
4.87e-02
1.18e-02



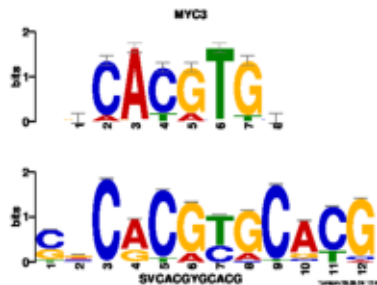
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
PIF5
4.94e-05
4.87e-02
1.18e-02

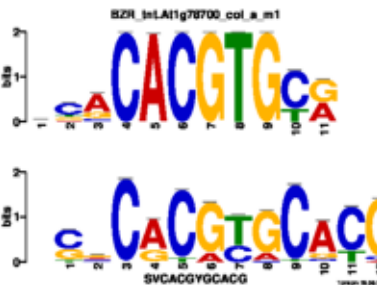
SA_under



5)
MYC3
5.48e-05
5.39e-02
1.18e-02



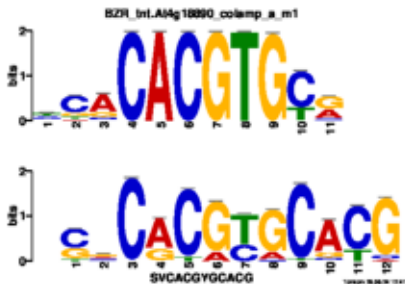
6)
BZR_tnt.At1g78700_col_a_m1
9.60e-05
9.46e-02
1.86e-02



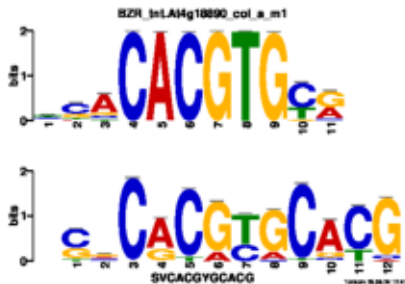
Nr:
Target ID:
p-value:
E-value:
q-value:

7)
BZR_tnt.At4g18890_colamp_a_m1
1.13e-04
1.11e-01
1.99e-02

SA_under



8)
BZR_tnt.At4g18890_col_a_m1
1.43e-04
1.40e-01
2.31e-02



9)
BZR_tnt.At4g36780_col_a_m1
2.26e-04
2.23e-01
3.38e-02

