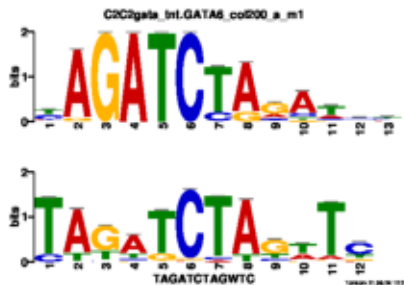
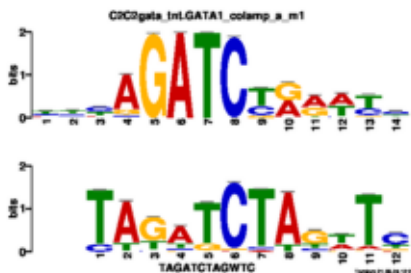


ABAJA under

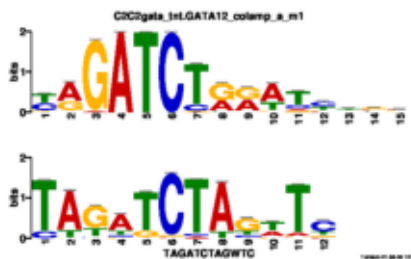


ABAJA under

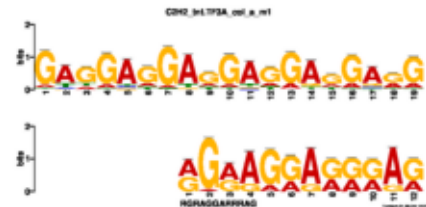


Two sequence logos are displayed. The top logo is titled "CRC2gata_int.GATA12_col_a_m1" and shows a sequence of 13 nucleotides: 1 (A), 2 (G), 3 (A), 4 (G), 5 (A), 6 (T), 7 (C), 8 (T), 9 (G), 10 (A), 11 (A), 12 (T), 13 (G). The bottom logo is titled "TAGACTAGWTC" and shows a sequence of 11 nucleotides: 1 (T), 2 (A), 3 (G), 4 (A), 5 (C), 6 (T), 7 (A), 8 (G), 9 (T), 10 (A), 11 (C). Both logos have a y-axis labeled "bits" ranging from 0 to 2.

5)
C2C2gata_tnt.GATA12_colamp_a_m1
2.95605e-05
0.0291171
0.0116222



3)
C2H2_tnt.TF3A_col_a_m1
2.72227e-05
0.0268144
0.053373

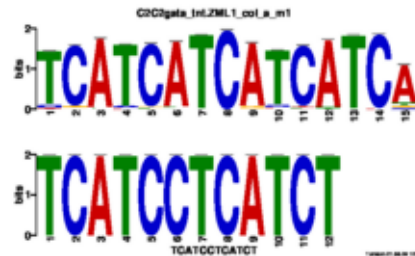
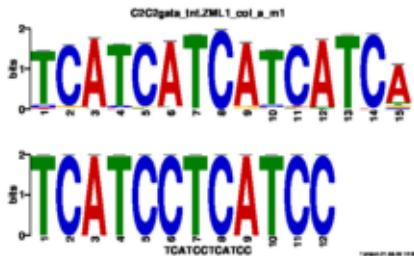


Nr: 1)
 Target ID: C2C2gata_tnt.ZML1_col_a_m1
 p-value: 3.20213e-05
 E-value: 0.0315409
 q-value: 0.0342376

2)
 MYBrelated_tnt.At2g38090_col_a_m1
 3.55131e-05
 0.0349804
 0.0342376

3)
 C2C2gata_tnt.ZML1_col_a_m1
 4.05091e-05
 0.0399015
 0.0542081

ABA_under



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

1)
C2C2dof_tnt.AT5G66940_col_a_m1
2.37584e-07
0.00023402
0.000228364

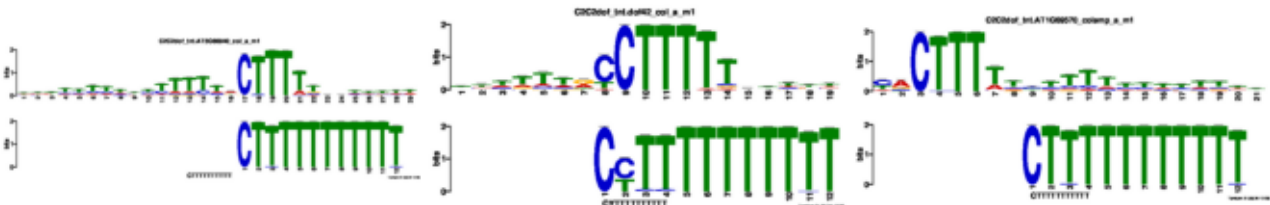
Nr:
Target ID:
p-value:
E-value:
q-value:

2)
C2C2dof_tnt.dof42_col_a_m1
2.61387e-07
0.000257466
0.000503597

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

3)
C2C2dof_tnt.AT1G69570_colamp_a_m1
4.46501e-07
0.000439803
0.000228364

JA_over



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

4)
C2C2dof_tnt.AT5G02460_colamp_a_m1
5.10351e-07
0.000502696
0.000228364

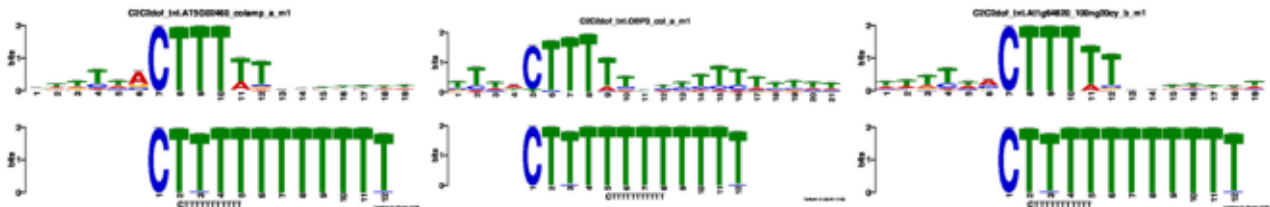
Nr:
Target ID:
p-value:
E-value:
q-value:

5)
C2C2dof_tnt.OBP3_col_a_m1
5.6882e-07
0.000560288
0.000228364

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

6)
C2C2dof_tnt.At1g64620_100ng20cy_b_m1
6.34864e-07
0.000625341
0.000228364

JA_over



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

7)
C2C2dof_tnt.AT1G69570_col_a_m1
7.14401e-07
0.000703685
0.000228364

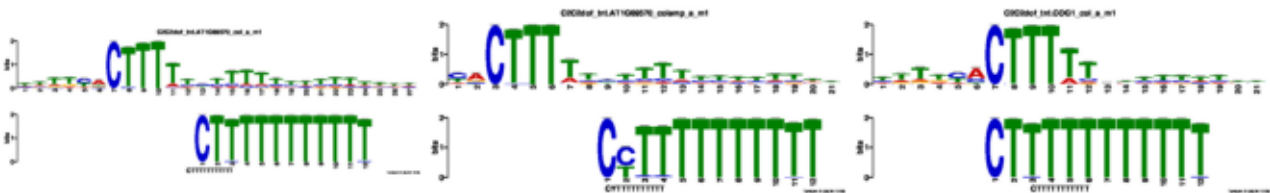
Nr:
Target ID:
p-value:
E-value:
q-value:

8)
C2C2dof_tnt.AT1G69570_colamp_a_m1
1.35485e-06
0.00133453
0.000942339

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

9)
C2C2dof_tnt.COG1_col_a_m1
1.4248e-06
0.00140343
0.000390384

JA_over



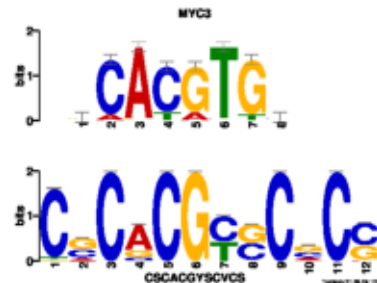
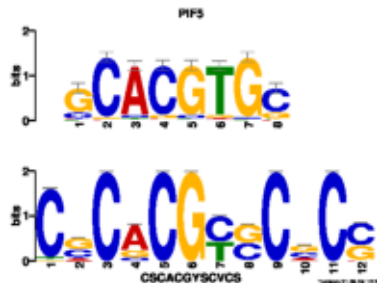
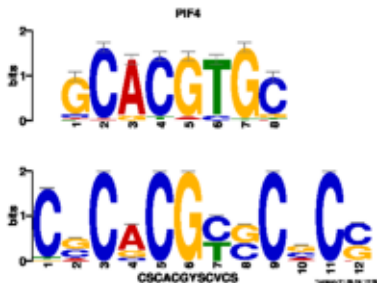
Nr:
Target ID:
p-value:
E-value:
q-value:

1)
PIF4
1.13427e-05
0.0111725
0.00629761

2)
PIF5
1.30181e-05
0.0128229
0.00629761

3)
MYC3
2.45059e-05
0.0241383
0.00662775

SAJA_under



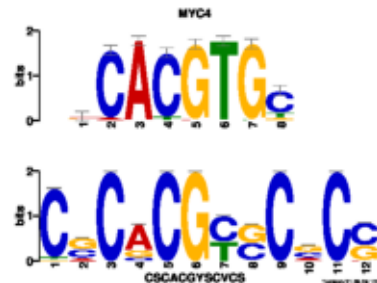
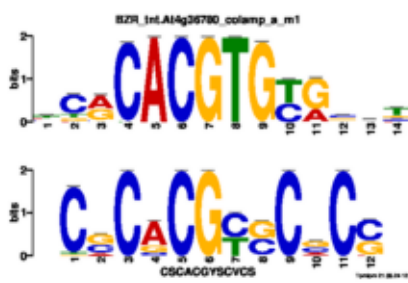
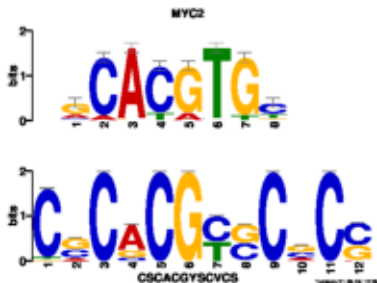
Nr:
Target ID:
p-value:
E-value:
q-value:

4)
MYC2
2.75966e-05
0.0271826
0.00662775

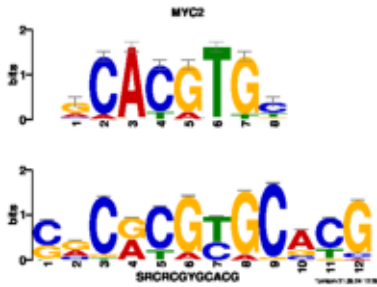
5)
BZR_tnt.At4g36780_colamp_a_m1
3.08264e-05
0.030364
0.00662775

6)
MYC4
4.30463e-05
0.0424006
0.00832956

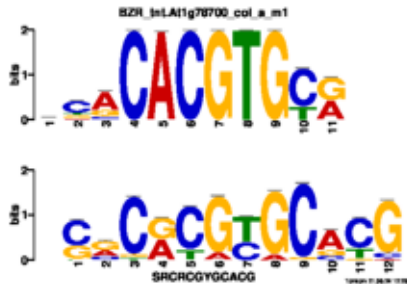
SAJA_under



SA_over



SA_over



Nr: 1)
 Target ID: BZR_tnt.At1g78700_col_a_m1
 p-value: 1.69362e-05
 E-value: 0.0166821
 q-value: 0.0146088

2)
 BZR_tnt.At4g18890_colamp_a_m1
 2.58069e-05
 0.0254198
 0.0146088

3)
 BZR_tnt.At4g36780_col_a_m1
 3.81844e-05
 0.0376116
 0.0146088

SA_under



Nr: 4)
 Target ID: BZR_tnt.At4g18890_col_a_m1
 p-value: 4.19223e-05
 E-value: 0.0412935
 q-value: 0.0146088

5)
 MYC4
 4.26386e-05
 0.041999
 0.0146088

SA_under

