



For further information on how to interpret these results or to get a copy of the MEME software please access <http://meme.nbcr.net>.

If you use TOMTOM in your research, please cite the following paper:

Shobhit Gupta, JA Stamatoyannopoulos, Timothy Bailey and William Stafford Noble, "Quantifying similarity between motifs", *Genome Biology*, 8(2):R24, 2007. [\[full text\]](#)

[QUERY MOTIFS](#) | [TARGET DATABASES](#) | [MATCHES](#) | [SETTINGS](#) | [PROGRAM INFORMATION](#)

QUERY MOTIFS

[Next Top](#)

Database	Name	Alt. Name	Preview	Matches	
motif group5	motif_10_2	1-NNKCAGTCGN		3	UP00243 UP00247
motif group5	motif_Train1_2	1-NNKCAGTYDN		5	MA0100 UP00092
motif group5	motif_Train2_2	1-DTCAGTCG		4	UP00081 UP00092
motif group5	motif_Train3_22	23-GACGCAGCAG		16	MA0591 MA0501
motif group5	motif_Train4_25	16-CACGTG		44	MA0409 MA0004
motif group5	motif_Train5_23	22-AATRAGCAAAAC		11	MA0326 MA0010
motif group5	motif_Train6_2	1-NRTCAGTCGH		3	UP00080 UP00080
motif group5	motif_Train7_2	1-TCAGTCVB		2	UP00081 UP00092
motif group5	motif_Train8_1	1-NNKCAGTYGN		1	MA0127

motif
group5

motif_Train9_2

1-KTCAGTCG



0

TARGET DATABASES

[Previous](#) [Next](#) [Top](#)

Database	Used	Matched
uniprobe mouse	386	20
JASPAR CORE 2014	593	63

MATCHES TO MOTIF_Train5_23 (22-AATRAGCAAAAC)

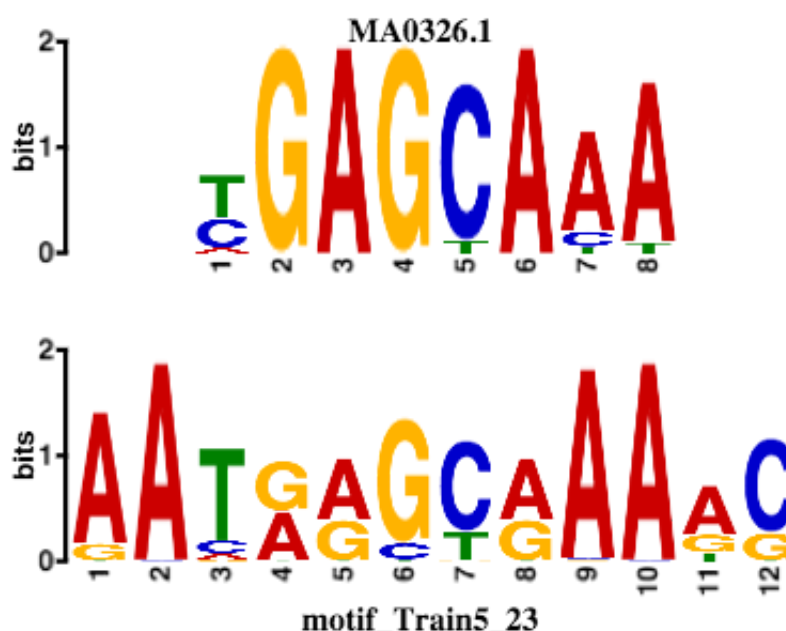
[Previous](#) [Next](#) [Top](#)

Summary

Name	MA0326.1 (MAC1)
Database	JASPAR CORE 2014
<i>p</i> -value	1.19e-04
<i>E</i> -value	1.17e-01
<i>q</i> -value	1.61e-01
Overlap	8
Offset	-2
Orientation	Reverse Complement

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Alignment

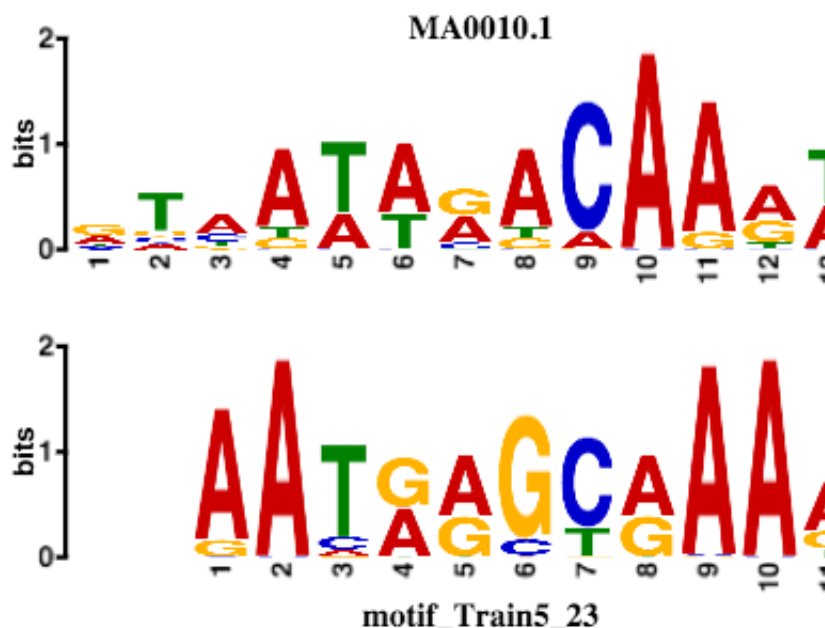


Summary

Name	MA0010.1 (br_Z1)
Database	JASPAR CORE 2014
<i>p</i> -value	1.64e-04
<i>E</i> -value	1.61e-01
<i>q</i> -value	1.61e-01
Overlap	12
Offset	2
Orientation	Normal

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Alignment



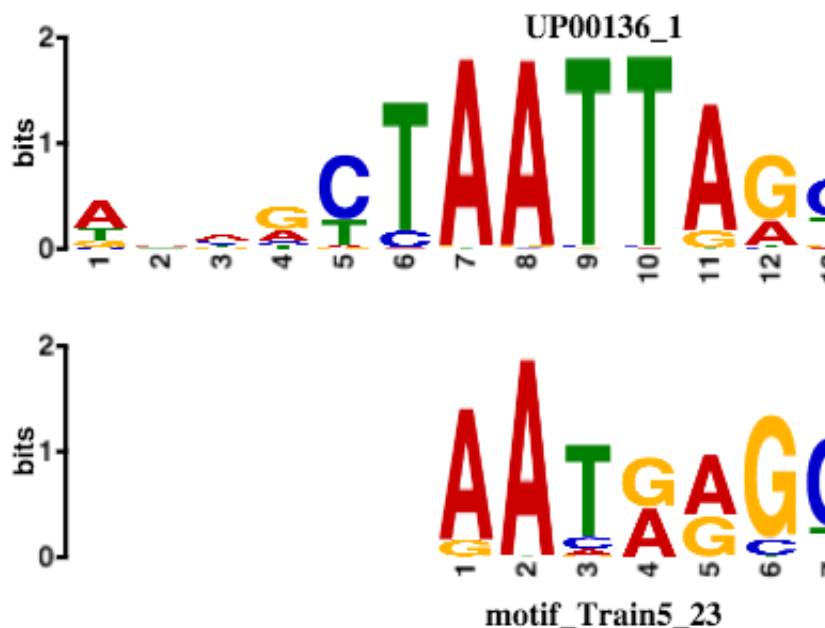
Summary

Name [UP00136_1](#)
[\(Prrx2 3072.1\)](#)
Database uniprobe mouse
p-value 2.09e-03
E-value 2.04e+00
q-value 1.00e+00

Overlap 11
Offset 6
Orientation Normal

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Alignment



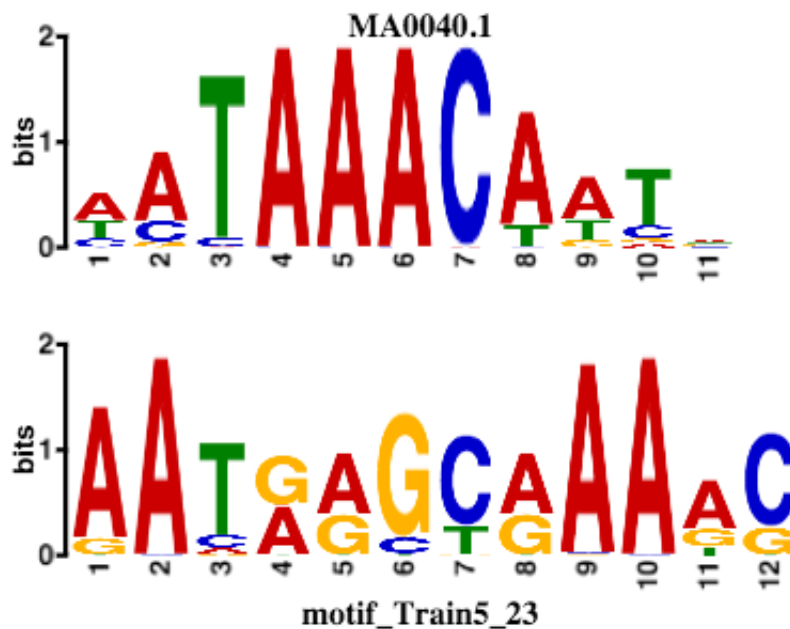
Summary

Name [MA0040.1](#)
[\(Foxq1\)](#)
Database JASPAR CORE
 2014
p-value 3.55e-03
E-value 3.48e+00
q-value 1.00e+00

Overlap 11
Offset 0
Orientation Reverse
 Complement

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Alignment



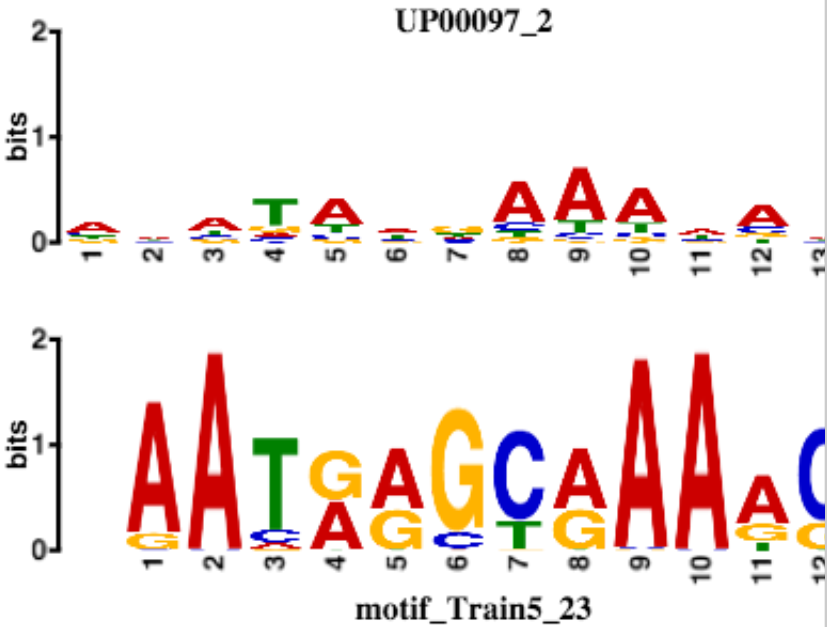
Summary

Name [UP00097_2](#)
[\(Mtf1_secondary\)](#)
Database uniprobe mouse
p-value 4.68e-03
E-value 4.58e+00
q-value 1.00e+00

Overlap 12
Offset 1
Orientation Normal

Alignment

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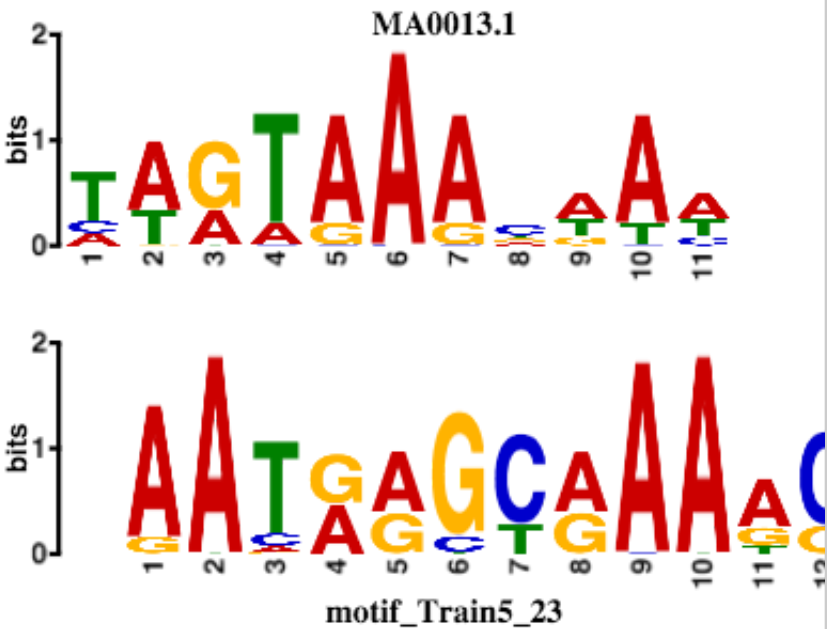


Summary

Name	MA0013.1 (br_Z4)
Database	JASPAR CORE 2014
p-value	5.29e-03
E-value	5.18e+00
q-value	1.00e+00
Overlap	10
Offset	1
Orientation	Normal

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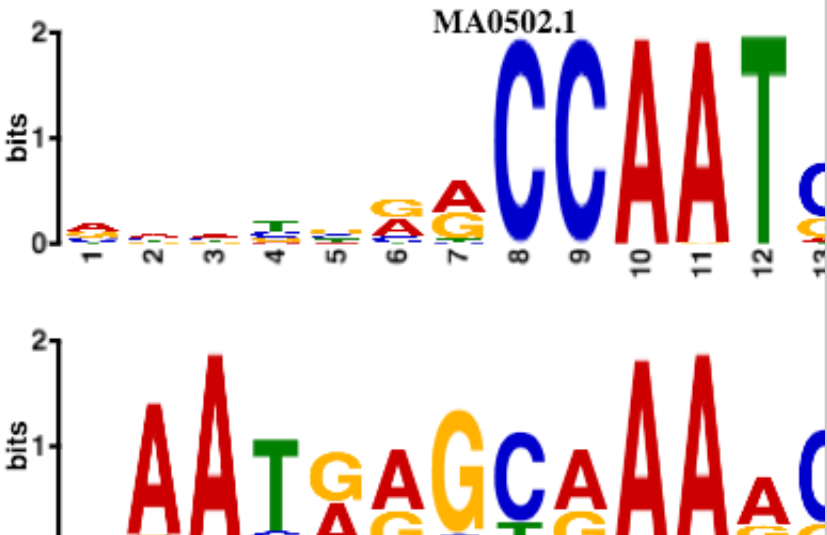
Alignment



Summary

Name	MA0502.1 (NFYB)
Database	JASPAR CORE 2014
p-value	6.27e-03
E-value	6.13e+00
q-value	1.00e+00
Overlap	12
Offset	1
Orientation	Normal

Alignment



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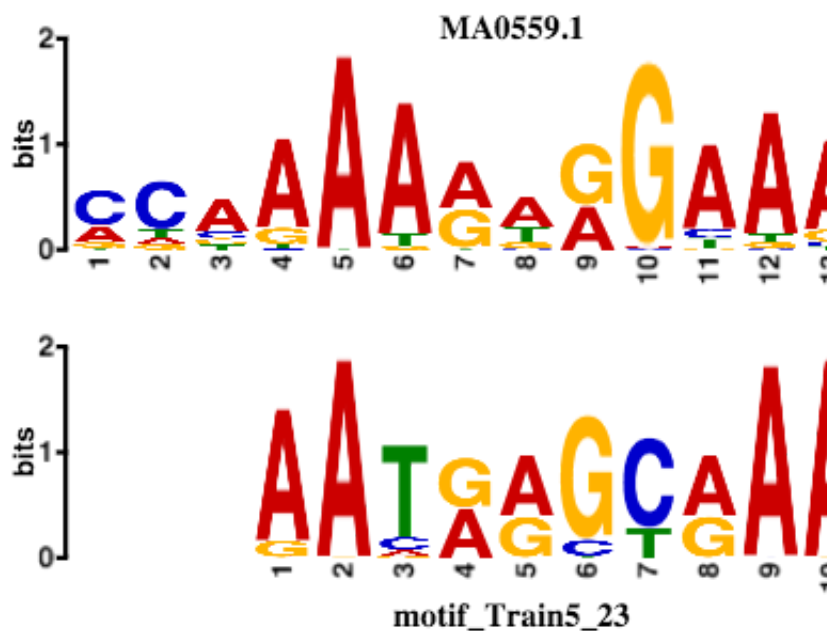

Summary

Name	MA0559.1 (PI)
Database	JASPAR CORE 2014
p-value	6.41e-03
E-value	6.28e+00
q-value	1.00e+00

Overlap	11
Offset	3
Orientation	Normal

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Alignment



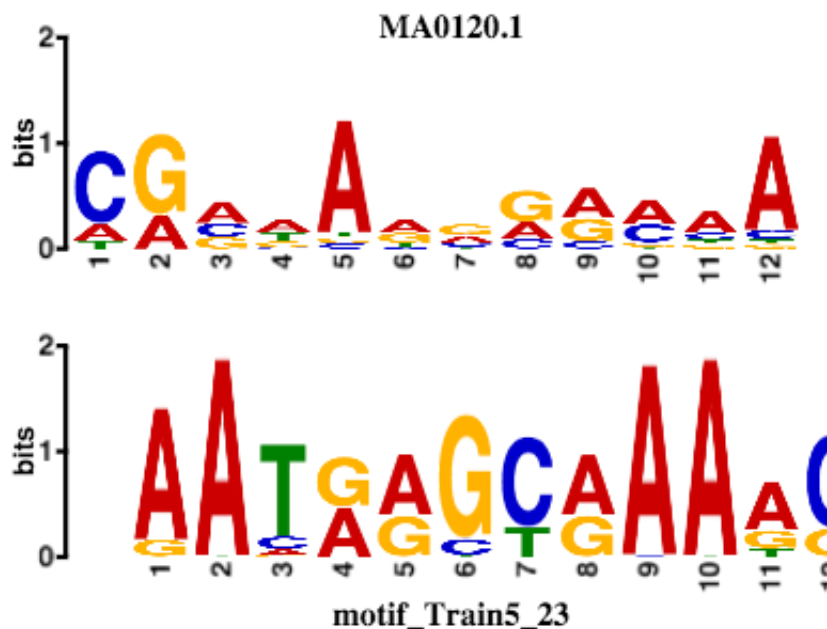
Summary

Name	MA0120.1 (id1)
Database	JASPAR CORE 2014
p-value	7.20e-03
E-value	7.05e+00
q-value	1.00e+00

Overlap	11
Offset	1
Orientation	Reverse Complement

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Alignment



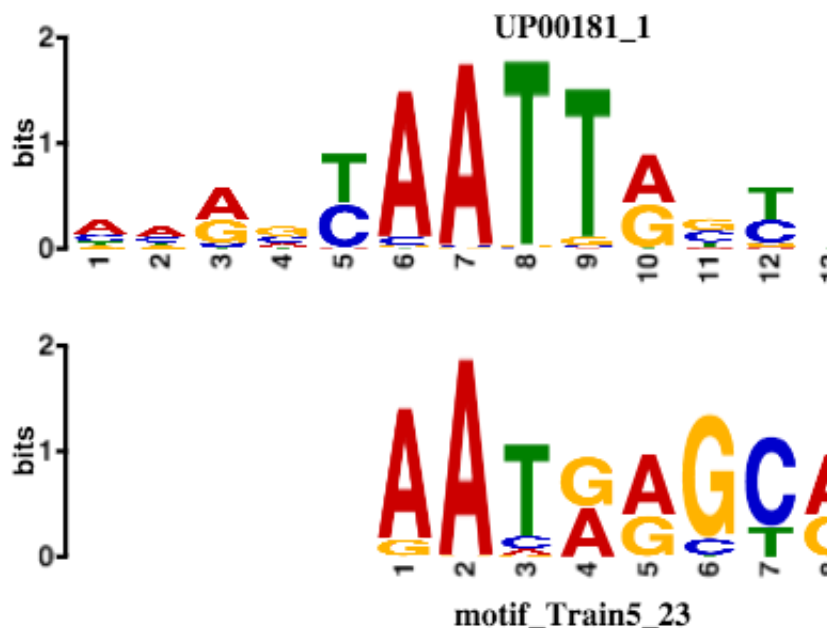
Summary

Alignment

Name [UP00181_1](#)
([Barx1_2877.1](#))
Database uniprobe mouse
p-value 7.55e-03
E-value 7.39e+00
q-value 1.00e+00

Overlap 11
Offset 5
Orientation Normal

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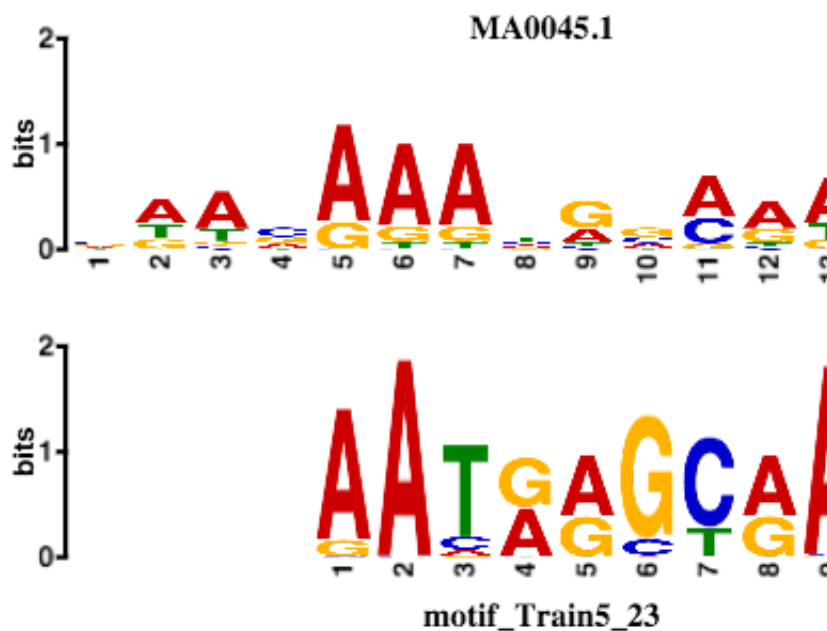
Summary

Alignment

Name [MA0045.1 \(HMG-I/Y\)](#)
Database JASPAR CORE
2014
p-value 8.54e-03
E-value 8.36e+00
q-value 1.00e+00

Overlap 12
Offset 4
Orientation Normal

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MATCHES TO MOTIF_Train6_2 (1-NRTCAGTCGH)

[Previous](#) [Next](#) [Top](#)

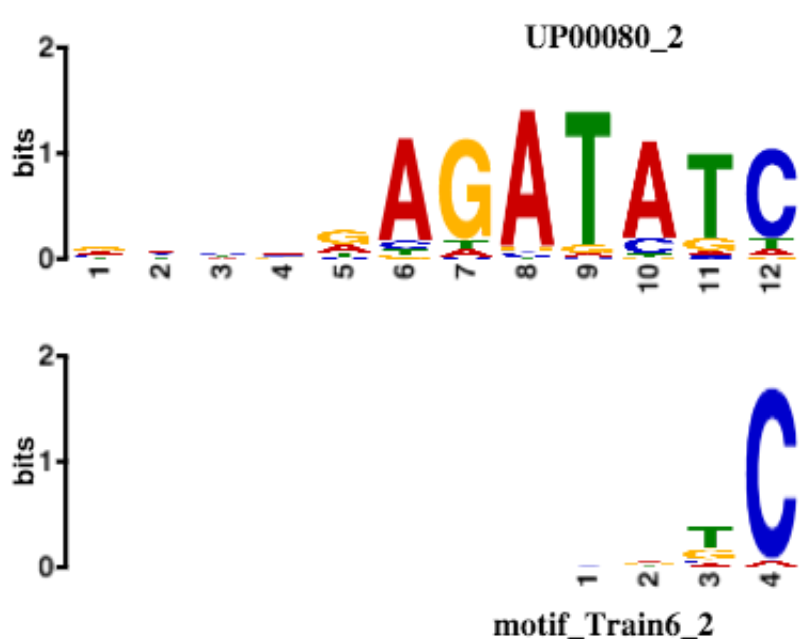
Summary

Alignment

Name [UP00080_2](#)
([Gata5_secondary](#))
Database uniprobe mouse

p-value	1.57e-03
E-value	1.54e+00
q-value	1.00e+00
Overlap	9
Offset	8
Orientation	Normal

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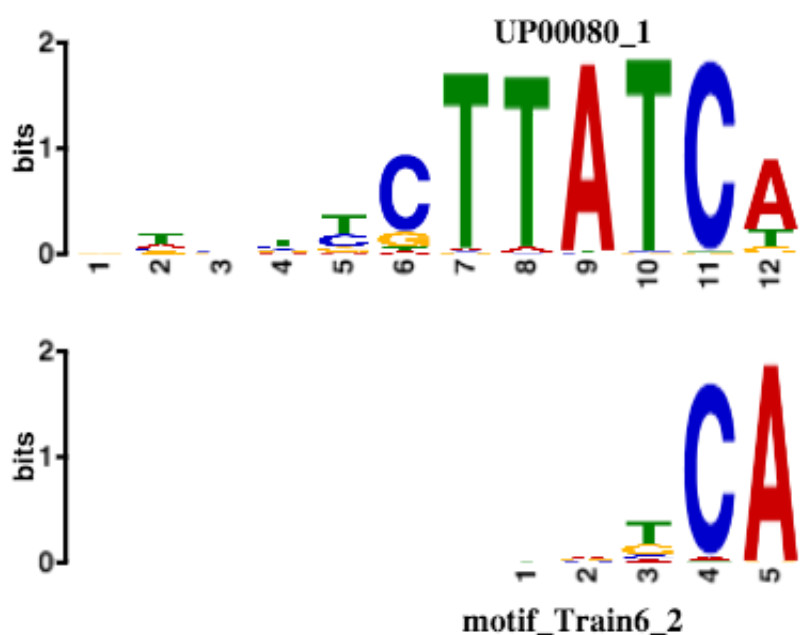


Summary

Name	UP00080_1 (Gata5_primary)
Database	uniprobe mouse
p-value	6.36e-03
E-value	6.23e+00
q-value	1.00e+00
Overlap	10
Offset	7
Orientation	Reverse Complement

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Alignment

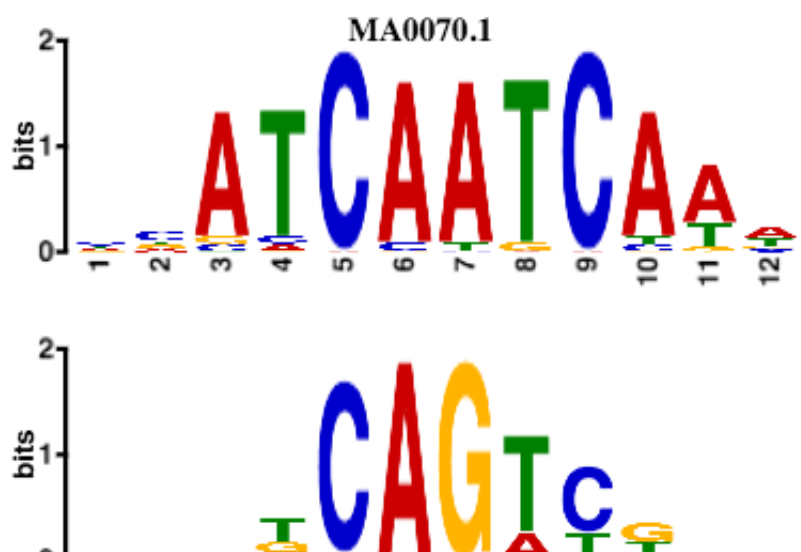


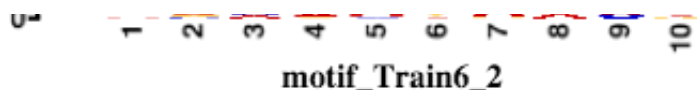
Summary

Name	MA0070.1 (PBX1)
Database	JASPAR CORE 2014
p-value	1.02e-02
E-value	9.97e+00
q-value	1.00e+00
Overlap	10
Offset	1
Orientation	Normal

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Alignment





MATCHES TO MOTIF_Train1_2 (1-NNKCAGTYDN)

[Previous](#) [Next](#) [Top](#)

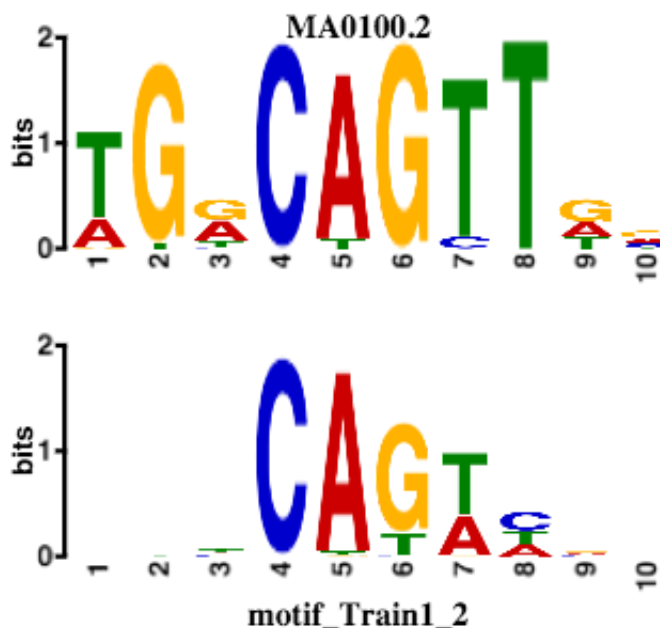
Summary

Name [MA0100.2 \(Myb\)](#)
Database JASPAR CORE 2014
p-value 3.31e-04
E-value 3.24e-01
q-value 3.47e-01

Overlap 10
Offset 0
Orientation Reverse Complement

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Alignment



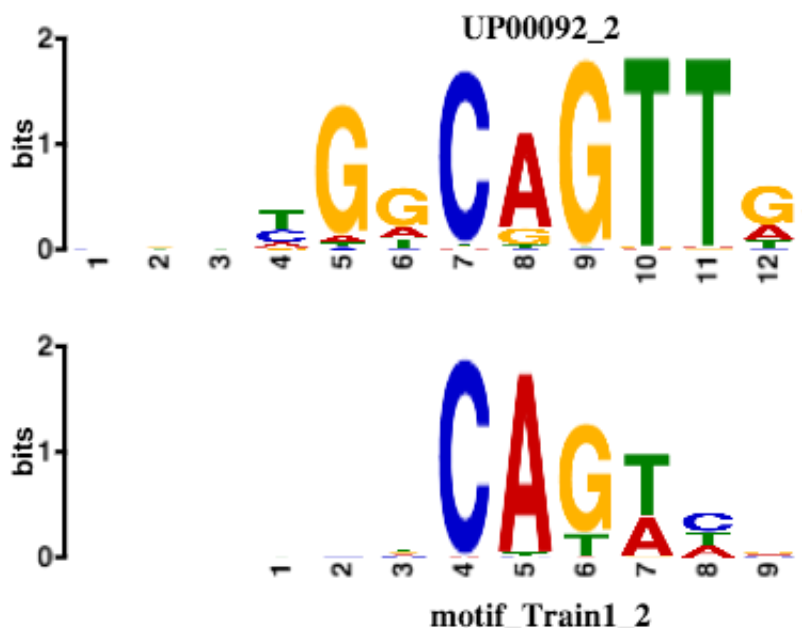
Summary

Name [UP00092_2 \(Myb_secondary\)](#)
Database uniprobe mouse
p-value 3.54e-04
E-value 3.47e-01
q-value 3.47e-01

Overlap 10
Offset 3
Orientation Reverse Complement

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Alignment



Summary

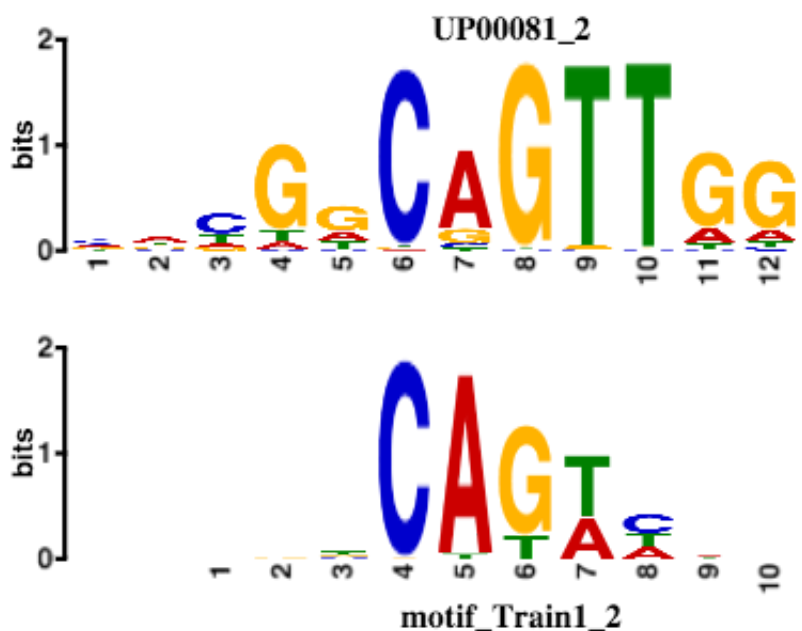
Name [UP00081_2 \(Mybl1_secondary\)](#)
Database uniprobe mouse
p-value 1.33e-03

Alignment

E-value 1.30e+00
q-value 6.82e-01

Overlap 10
Offset 2
Orientation Reverse
 Complement

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Summary

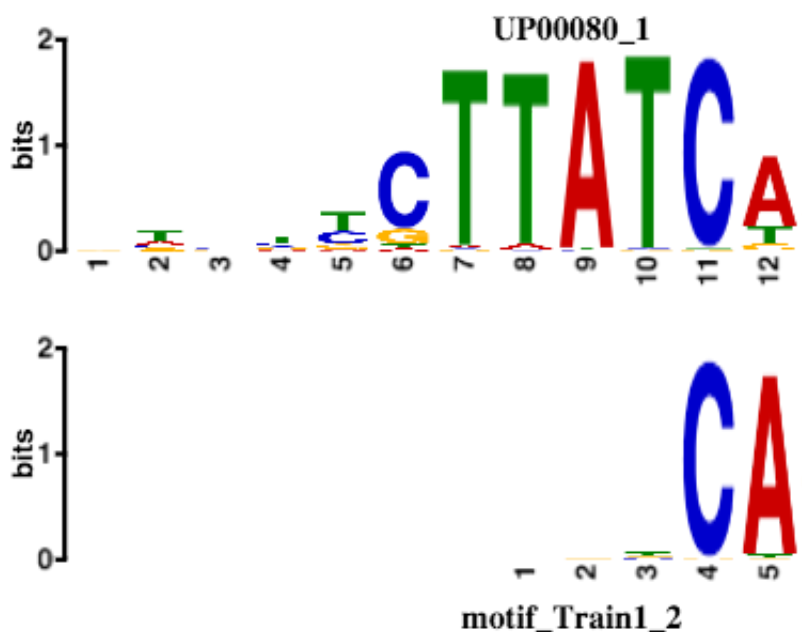
Name [UP00080_1](#)
 (Gata5_primary)
Database uniprobe mouse

p-value 1.39e-03
E-value 1.36e+00
q-value 6.82e-01

Overlap 10
Offset 7
Orientation Reverse
 Complement

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Alignment



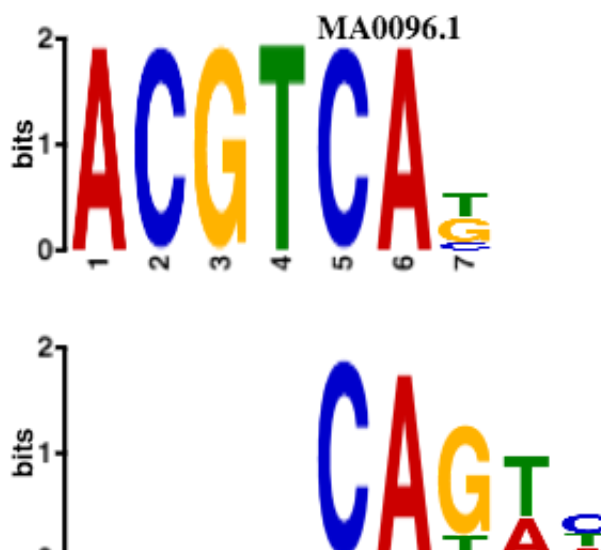
Summary

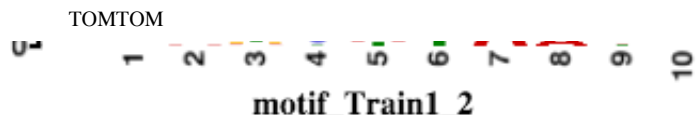
Name [MA0096.1](#)
 (bZIP910)
Database JASPAR CORE
 2014

p-value 1.01e-02
E-value 9.88e+00
q-value 1.00e+00

Overlap 6
Offset 1
Orientation Reverse
 Complement

Alignment



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MATCHES TO MOTIF_Train2_2 (1-DTCAGTCG)

[Previous](#) [Next](#) [Top](#)

Summary

Name [UP00081_2](#)
(Mybl1 secondary)

Database uniprobe mouse

p-value 1.04e-03

E-value 1.01e+00

q-value 1.00e+00

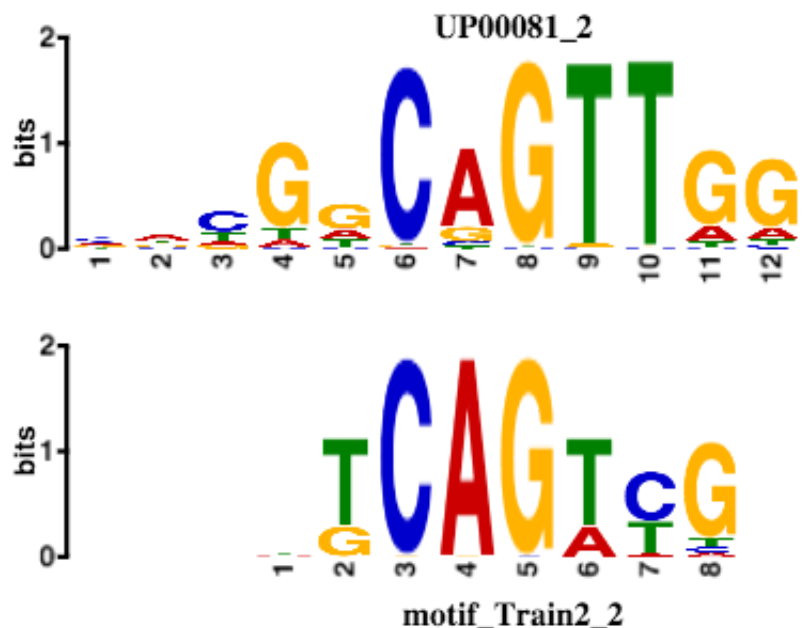
Overlap 8

Offset 3

Orientation Reverse
Complement

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Alignment



Summary

Name [UP00092_2](#)
(Myb secondary)

Database uniprobe mouse

p-value 1.57e-03

E-value 1.54e+00

q-value 1.00e+00

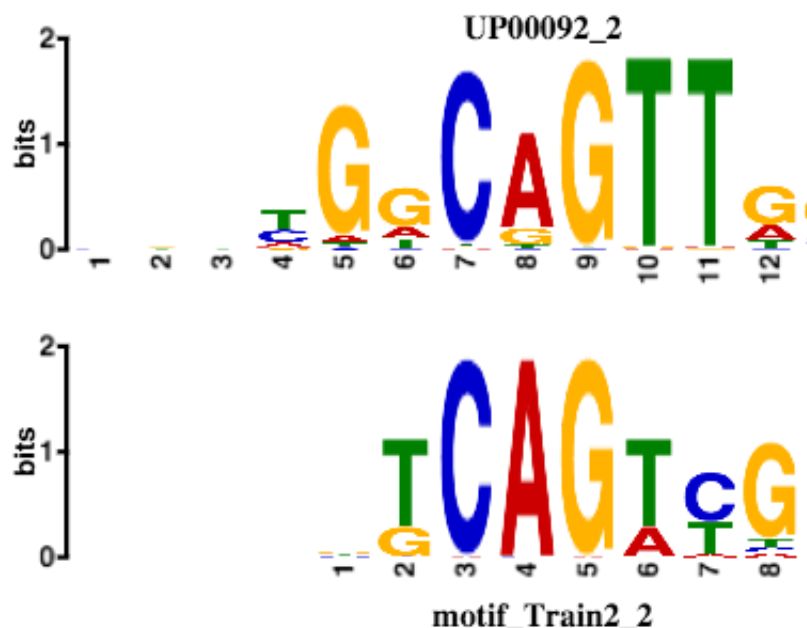
Overlap 8

Offset 4

Orientation Reverse
Complement

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Alignment



Summary

Name [MA0100.2 \(Myb\)](#)

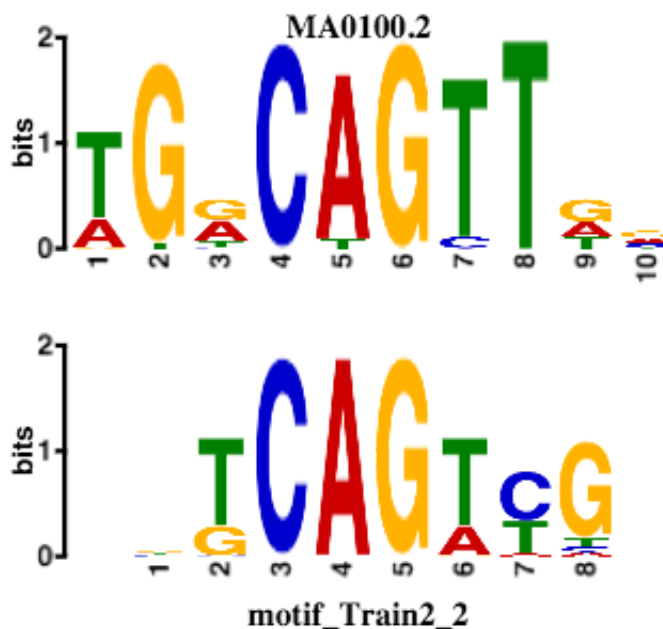
Database JASPAR CORE
2014

p-value 1.84e-03

Alignment

E-value 1.80e+00
q-value 1.00e+00
Overlap 8
Offset 1
Orientation Reverse
 Complement

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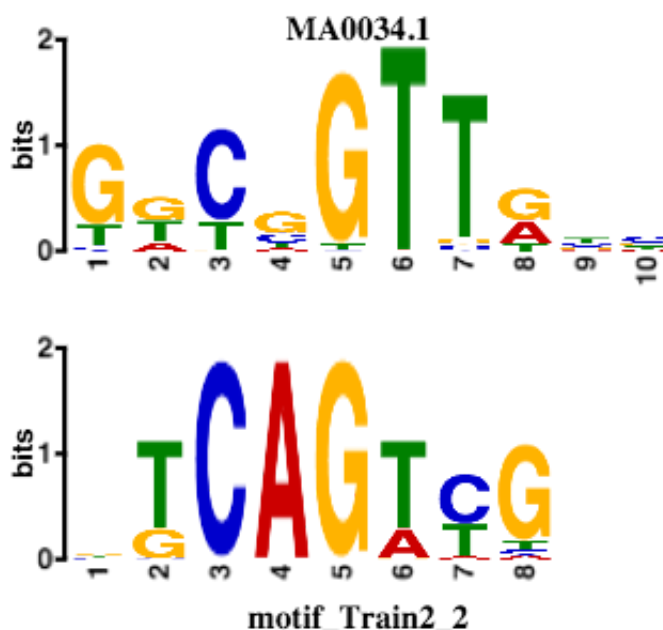


Summary

Name [MA0034.1 \(Gamyb\)](#)
Database JASPAR CORE 2014
p-value 8.19e-03
E-value 8.02e+00
q-value 1.00e+00
Overlap 8
Offset 0
Orientation Reverse
 Complement

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Alignment



MATCHES TO MOTIF_Train7_2 (1-TCAGTCVB)

[Previous](#) [Next](#) [Top](#)

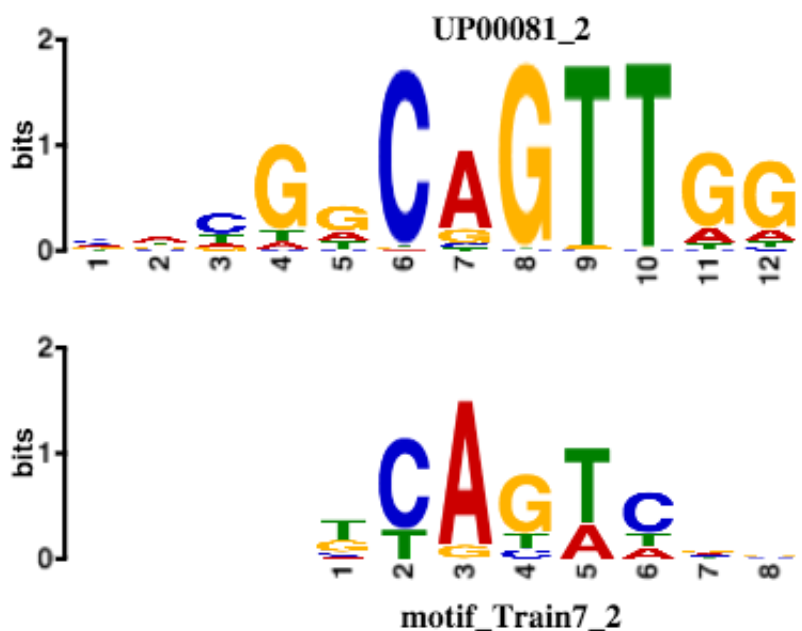
Summary

Name [UP00081_2 \(Mybl1_secondary\)](#)
Database uniprobe mouse
p-value 5.38e-03
E-value 5.27e+00
q-value 1.00e+00
Overlap 8

Alignment

Offset 4
Orientation Reverse
 Complement

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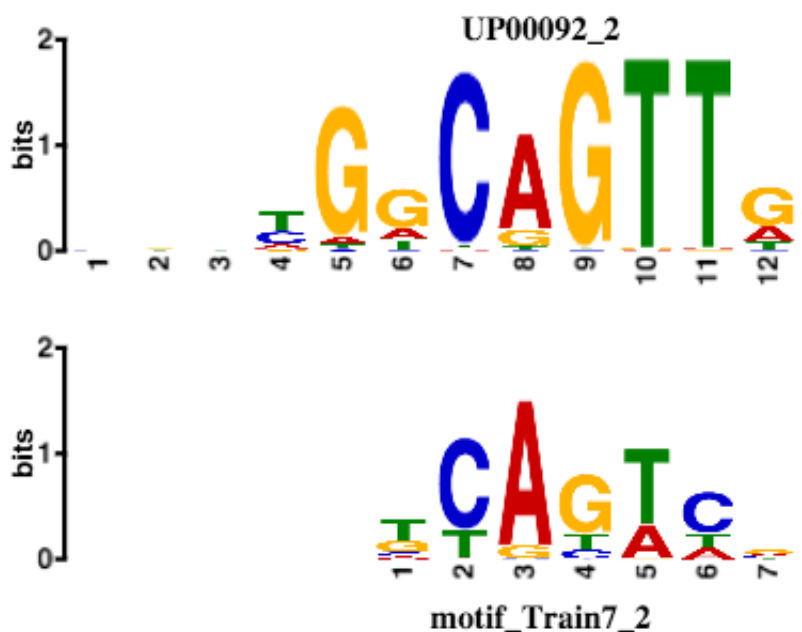
Summary

Name [UP00092_2](#)
 (Myb_secondary)
Database uniprobe mouse
p-value 8.39e-03
E-value 8.22e+00
q-value 1.00e+00

Overlap 8
Offset 5
Orientation Reverse
 Complement

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Alignment



MATCHES TO MOTIF_Train3_22 (23-GACGCAGCAG)

[Previous](#) [Next](#) [Top](#)

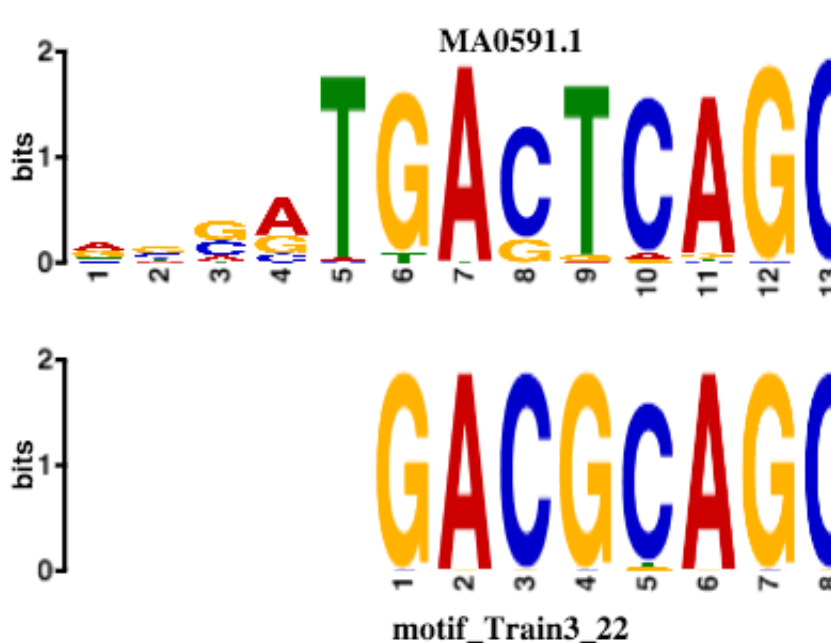
Summary

Name [MA0591.1](#)
 (Bach1::Mafk)
Database JASPAR CORE
 2014
p-value 5.86e-05
E-value 5.73e-02
q-value 1.14e-01

Alignment

Overlap 10
Offset 5
Orientation Normal

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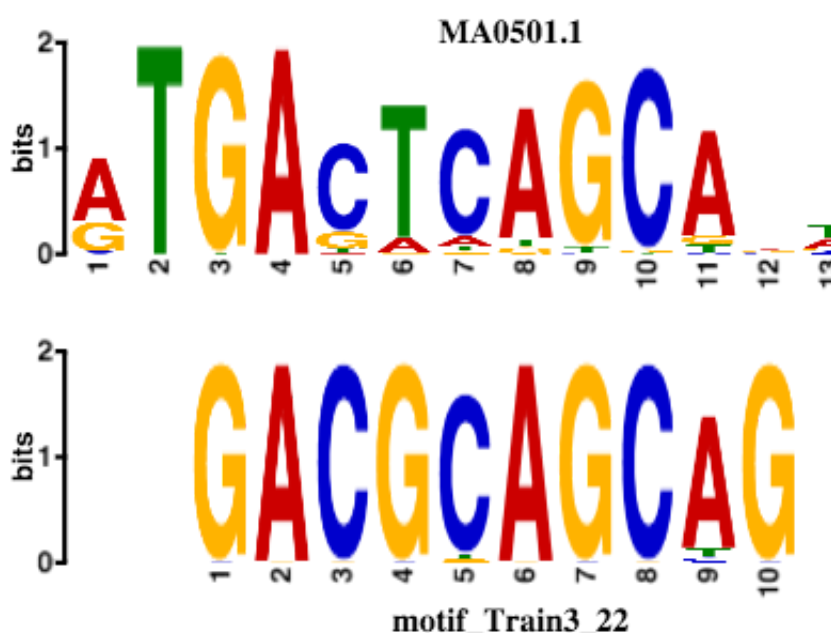
Summary

Name [MA0501.1](#)
[\(NFE2::MAF\)](#)
Database JASPAR CORE
 2014
p-value 1.76e-04
E-value 1.72e-01
q-value 1.72e-01

Overlap 10
Offset 2
Orientation Normal

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Alignment



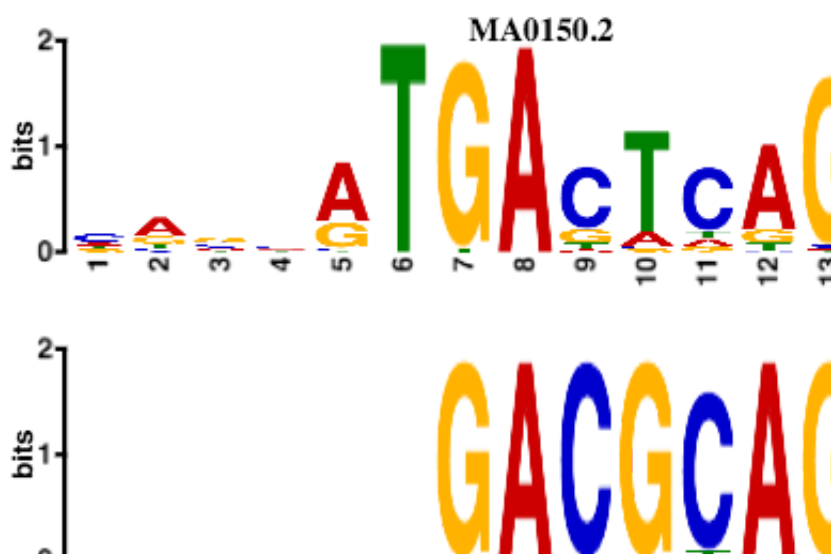
Summary

Name [MA0150.2](#)
[\(Nfe2l2\)](#)
Database JASPAR CORE
 2014
p-value 4.97e-04
E-value 4.86e-01
q-value 2.59e-01

Overlap 9
Offset 6
Orientation Normal

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Alignment



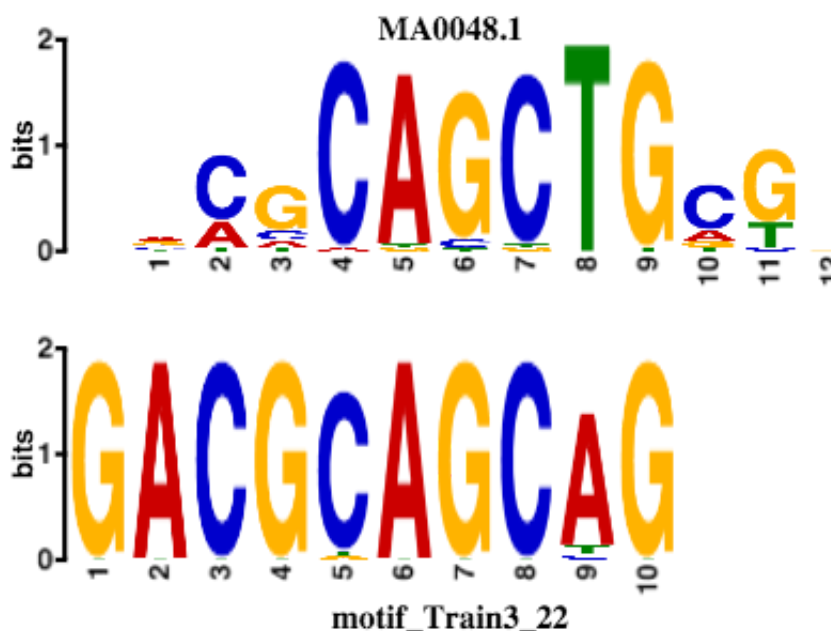
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1 2 3 4 5 6 7
motif_Train3_22

Summary

Alignment

Name	MA0048.1 (NHLH1)
Database	JASPAR CORE 2014
p-value	5.31e-04
E-value	5.19e-01
q-value	2.59e-01
Overlap	9
Offset	-1
Orientation	Reverse Complement

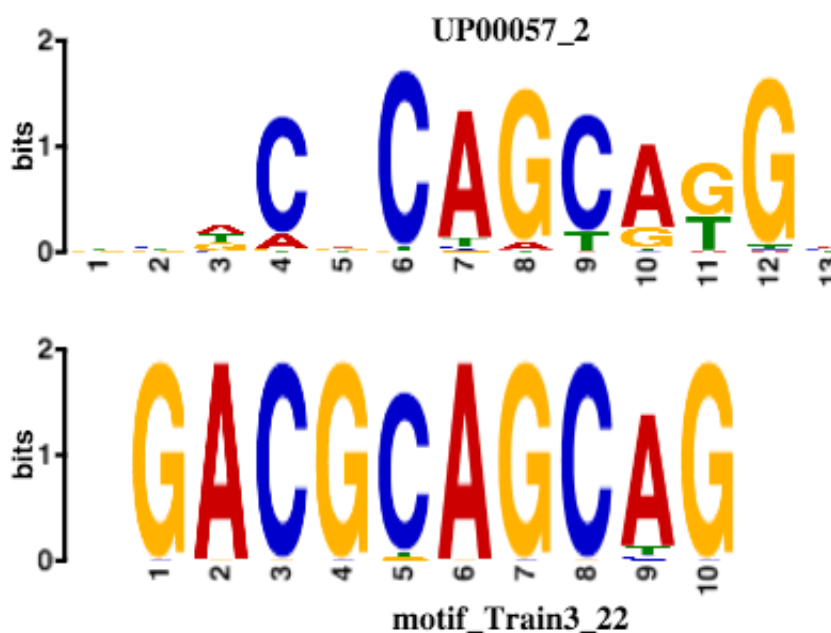
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Summary

Alignment

Name	UP00057_2 (Zic2 secondary)
Database	uniprobe mouse
p-value	1.53e-03
E-value	1.49e+00
q-value	5.72e-01

Overlap	10
Offset	1
Orientation	Normal

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Summary

Alignment

Name	UP00102_2 (Zic1 secondary)
Database	uninrobe mouse

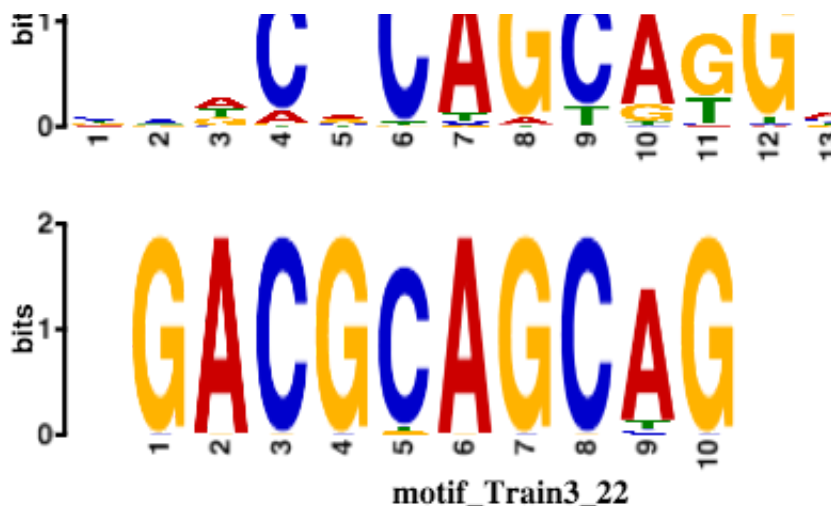


Database uniprobe mouse

p-value 2.07e-03
E-value 2.03e+00
q-value 5.72e-01

Overlap 10
Offset 1
Orientation Normal

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Summary

Alignment

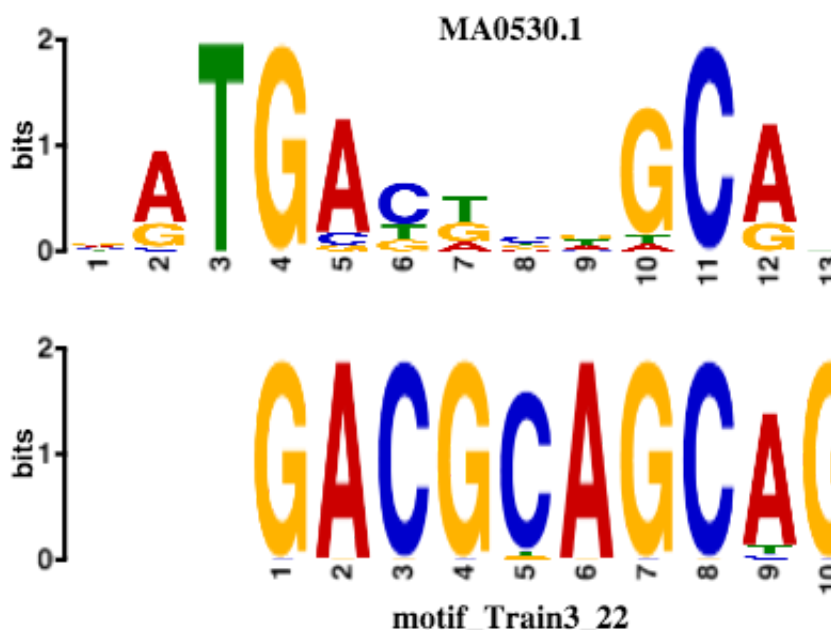
Name [MA0530.1](#)
[\(CNC::maf-S\)](#)

Database JASPAR CORE
 2014

p-value 2.39e-03
E-value 2.34e+00
q-value 5.72e-01

Overlap 10
Offset 3
Orientation Normal

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Summary

Alignment

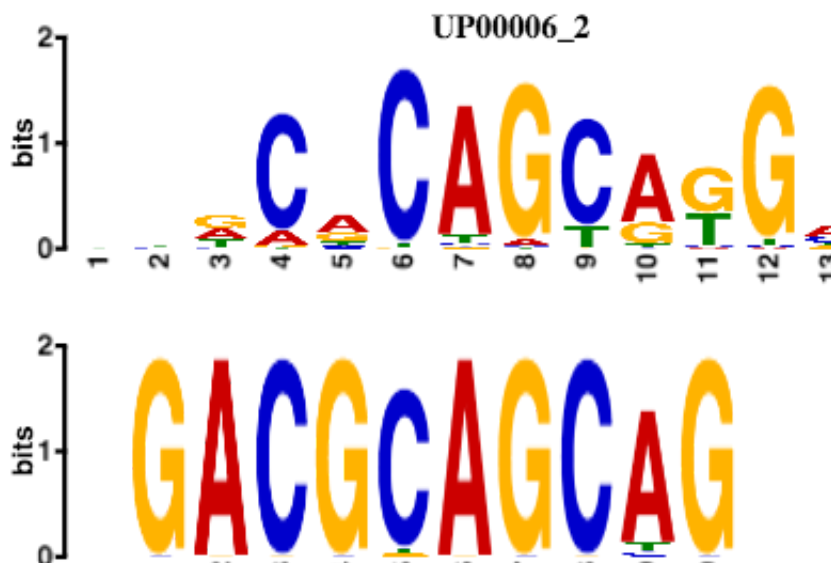
Name [UP00006_2](#)
[\(Zic3_secondary\)](#)

Database uniprobe mouse

p-value 2.63e-03
E-value 2.58e+00
q-value 5.72e-01

Overlap 10
Offset 1
Orientation Normal

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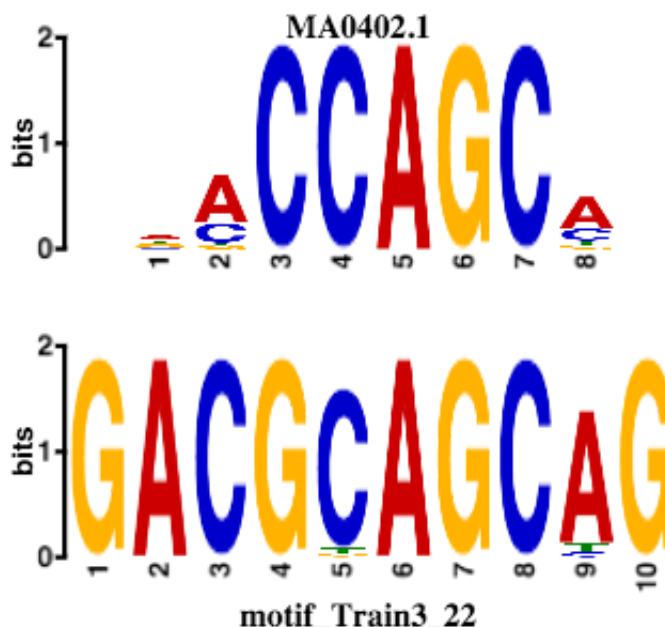
motif_Train3_22

Summary

Name	MA0402.1 (SWI5)
Database	JASPAR CORE 2014
p-value	3.12e-03
E-value	3.05e+00
q-value	5.82e-01
Overlap	8
Offset	-1
Orientation	Reverse Complement

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Alignment

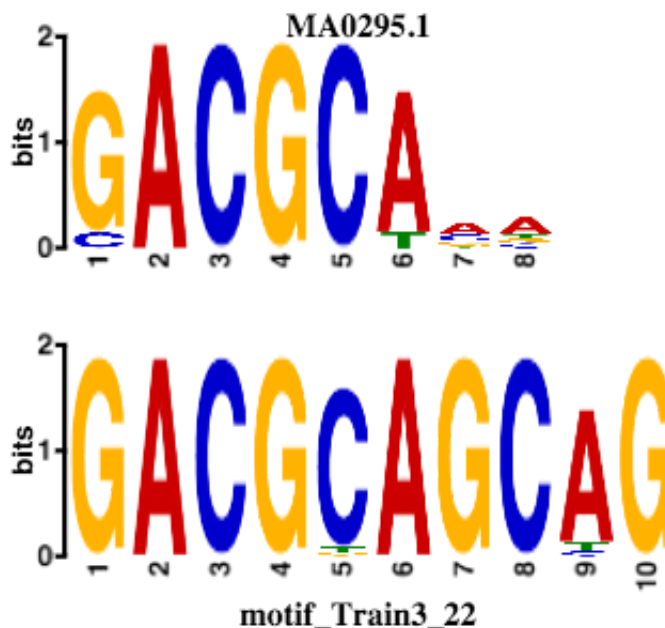


Summary

Name	MA0295.1 (FHL1)
Database	JASPAR CORE 2014
p-value	3.28e-03
E-value	3.21e+00
q-value	5.82e-01
Overlap	8
Offset	0
Orientation	Normal

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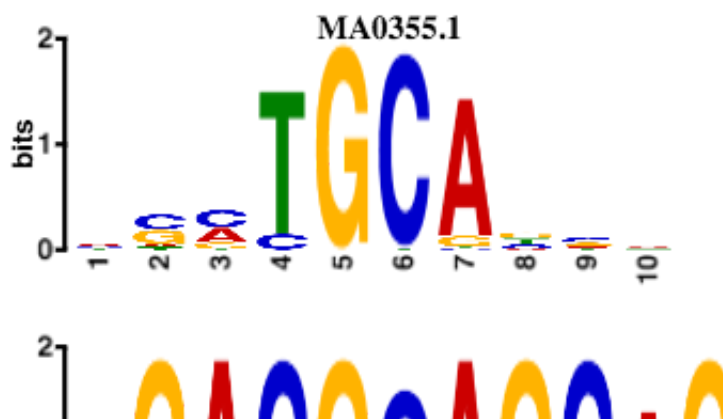
Alignment



Summary

Name	MA0355.1 (PHD1)
Database	JASPAR CORE 2014
p-value	3.62e-03
E-value	3.54e+00
q-value	5.89e-01
Overlap	9

Alignment



Offset 1
Orientation Normal
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Summary

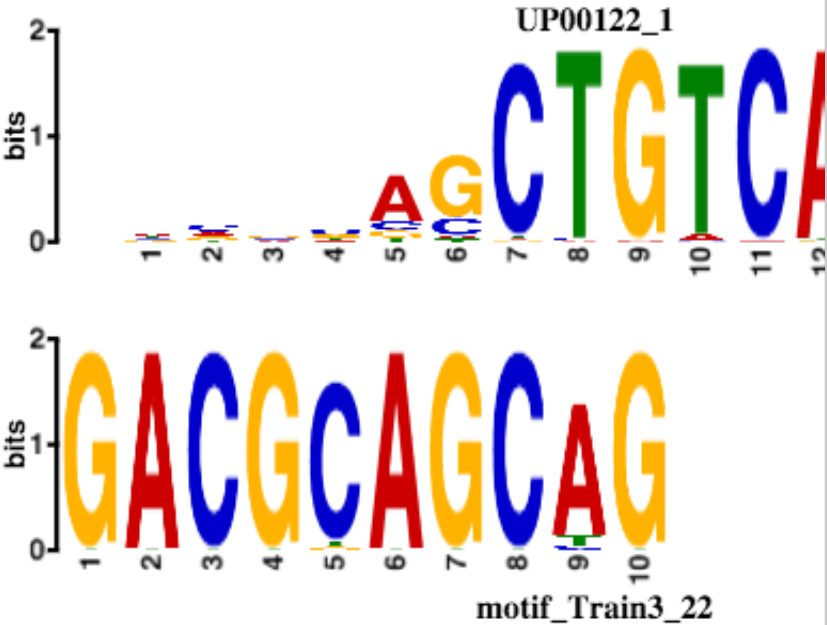
Alignment

Name [UP00122_1 \(Tqif1_2342.2\)](#)
Database uniprobe mouse

p-value 3.95e-03
E-value 3.86e+00
q-value 5.94e-01

Overlap 9
Offset -1
Orientation Reverse Complement

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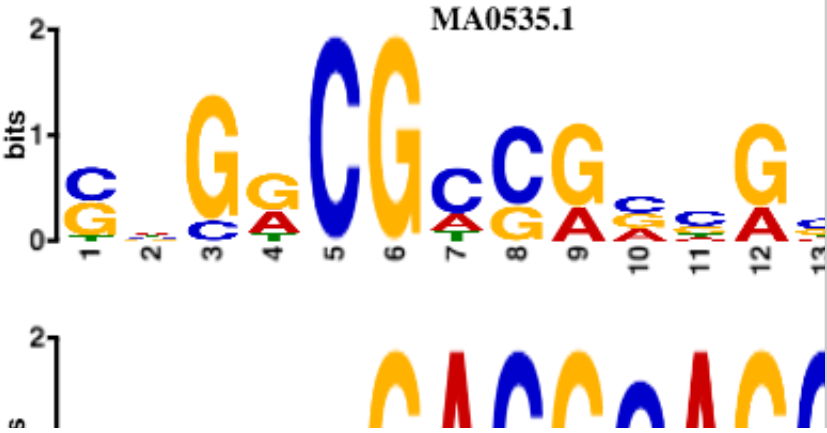
Summary

Alignment

Name [MA0535.1 \(Mad\)](#)
Database JASPAR CORE 2014

p-value 6.46e-03
E-value 6.33e+00
q-value 8.25e-01

Overlap 10
Offset 5



OrientationNormal

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Summary

Alignment

Name

[MA0495.1 \(MAFF\)](#)

Database

JASPAR CORE 2014

p-value

6.60e-03

E-value

6.46e+00

q-value

8.25e-01

Overlap

10

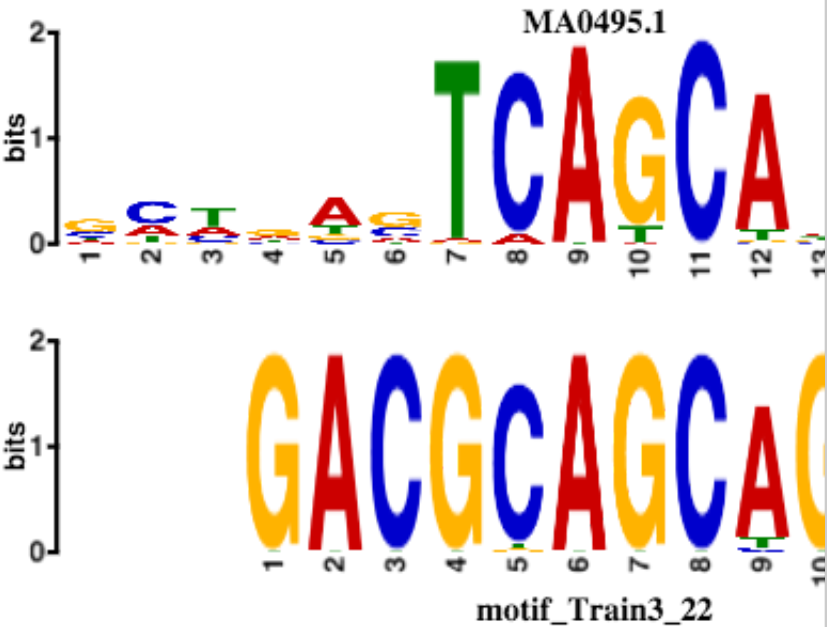
Offset

3

Orientation

Normal

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Summary

Alignment

Name

[MA0496.1 \(MAFK\)](#)

Database

JASPAR CORE 2014

p-value

6.76e-03

E-value

6.61e+00

q-value

8.25e-01

Overlap

10

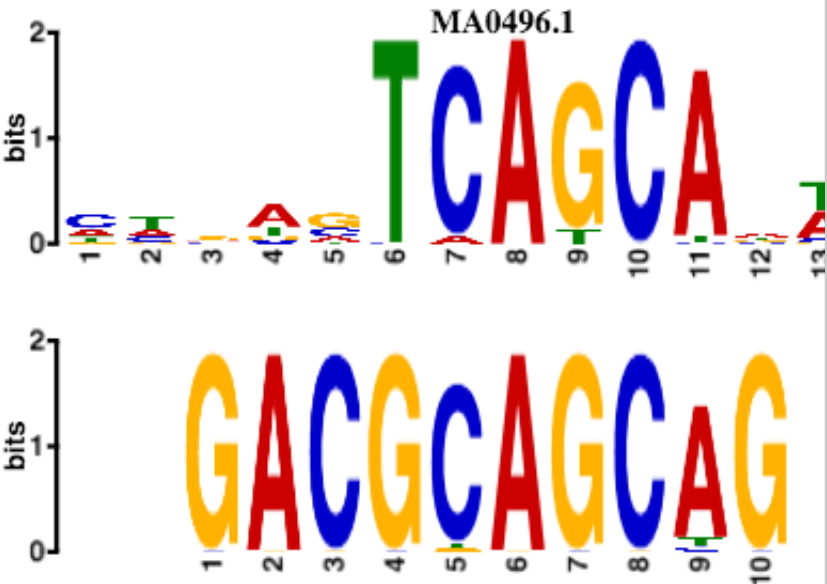
Offset

2

Orientation

Normal

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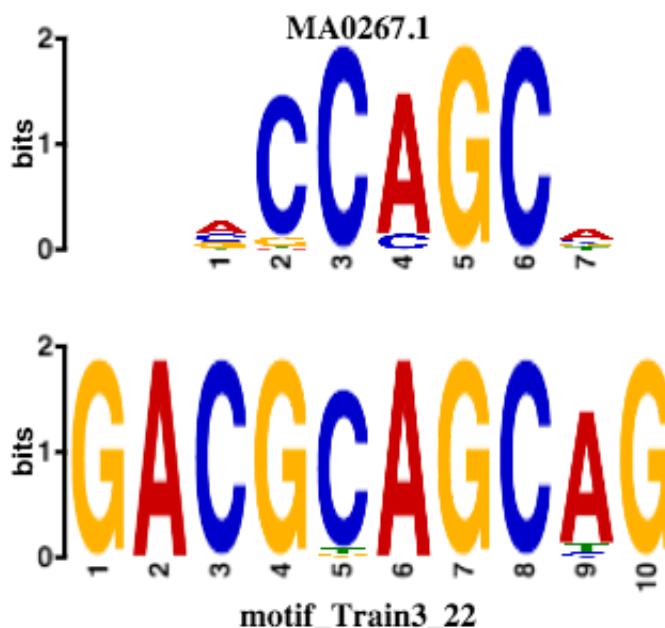


motif_Train3_22

Summary

Alignment

Name	MA0267.1 (ACE2)
Database	JASPAR CORE 2014
p-value	8.55e-03
E-value	8.37e+00
q-value	9.83e-01
Overlap	7
Offset	-2
Orientation	Normal
Show logo download options	



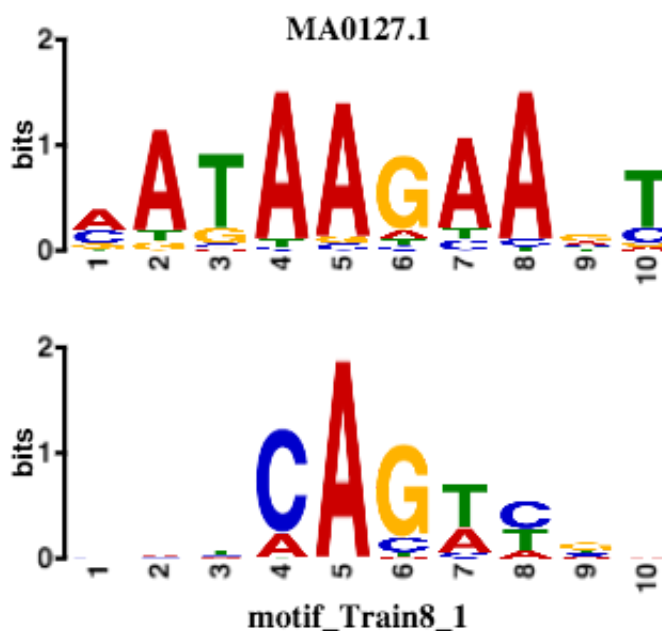
MATCHES TO MOTIF_Train8_1 (1-NNKCAGTYGN)

[Previous](#) [Next](#) [Top](#)

Summary

Alignment

Name	MA0127.1 (PEND)
Database	JASPAR CORE 2014
p-value	4.95e-03
E-value	4.84e+00
q-value	1.00e+00
Overlap	10
Offset	0
Orientation	Reverse Complement
Show logo download options	



MATCHES TO MOTIF_10_2 (1-NNKCAGTCGN)

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Summary

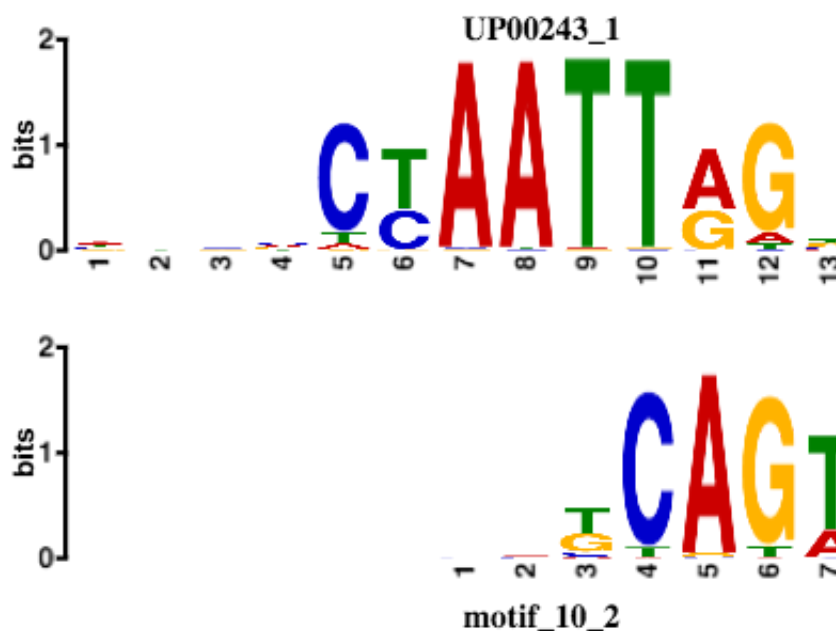
Alignment

Name	UP00243_1 (Isx_3445.1)
Database	uniprobe mouse

p-value 5.10e-03
E-value 5.00e+00
q-value 1.00e+00

Overlap 10
Offset 6
Orientation Normal

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Summary

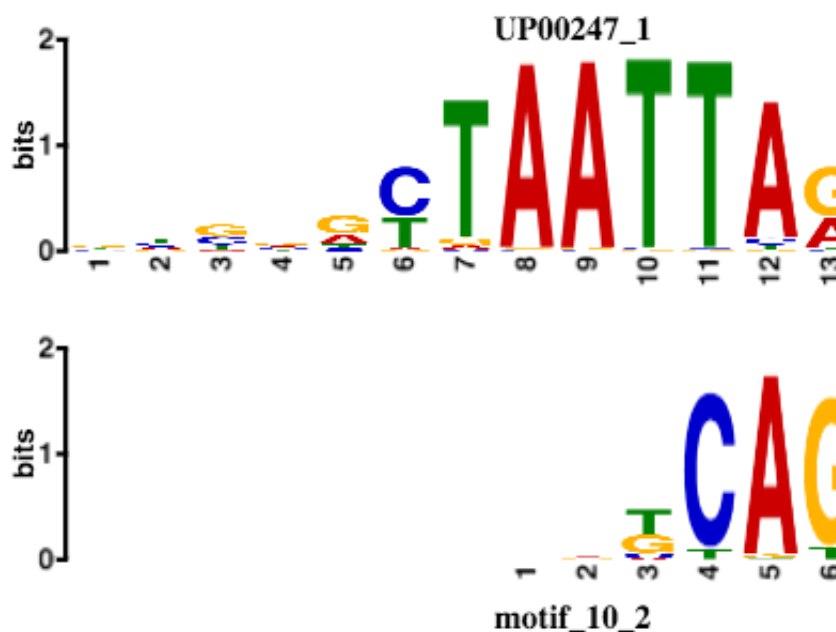
Name [UP00247_1](#)
[\(Pax4_3989.2\)](#)
Database uniprobe mouse

p-value 5.64e-03
E-value 5.53e+00
q-value 1.00e+00

Overlap 10
Offset 7
Orientation Reverse
 Complement

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Alignment



Summary

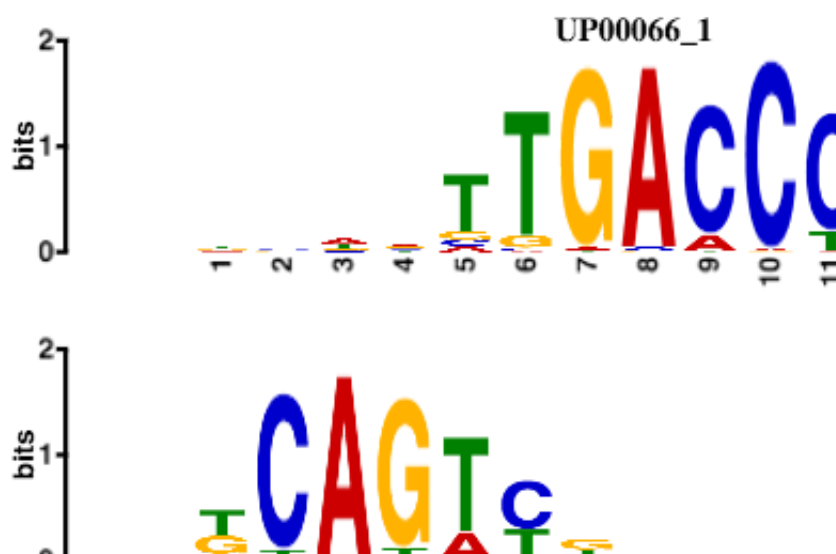
Name [UP00066_1](#)
[\(Hnf4a_primary\)](#)
Database uniprobe mouse

p-value 7.12e-03
E-value 6.97e+00
q-value 1.00e+00

Overlap 8
Offset -2
Orientation Reverse
 Complement

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Alignment





MATCHES TO MOTIF_Train4_25 (16-CACGTG)

[Previous](#) [Next](#) [Top](#)

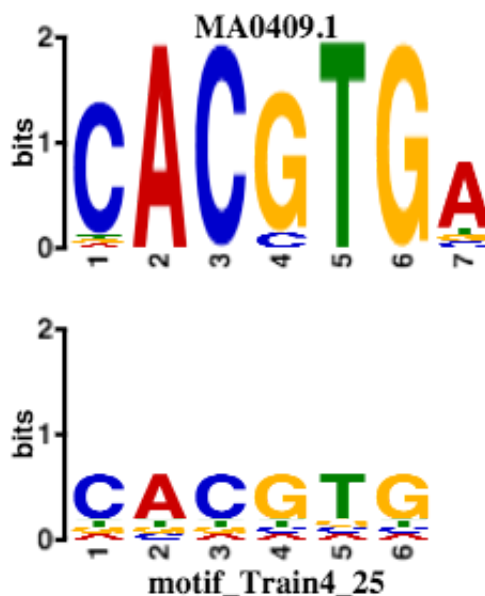
Summary

Name [MA0409.1 \(TYE7\)](#)
Database JASPAR CORE 2014
p-value 3.87e-06
E-value 3.79e-03
q-value 9.14e-04

Overlap 6
Offset 0
Orientation Normal

[Show logo download options](#)

Alignment



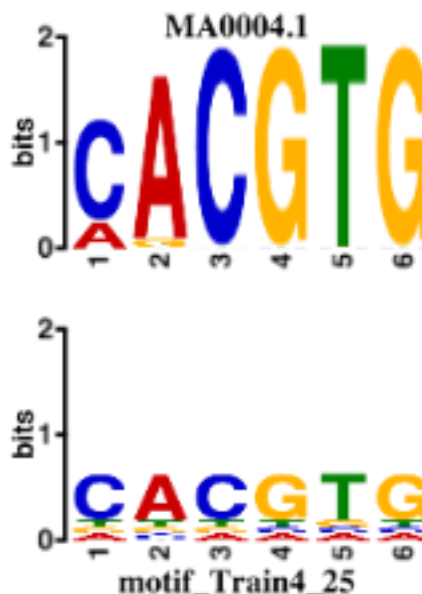
Summary

Name [MA0004.1 \(Arnt\)](#)
Database JASPAR CORE 2014
p-value 5.07e-06
E-value 4.97e-03
q-value 9.14e-04

Overlap 6
Offset 0
Orientation Normal

[Show logo download options](#)

Alignment



Summary

Name [MA0104.3 \(Mycn\)](#)
Database JASPAR CORE 2014
p-value 5.80e-06
E-value 5.68e-03

Alignment

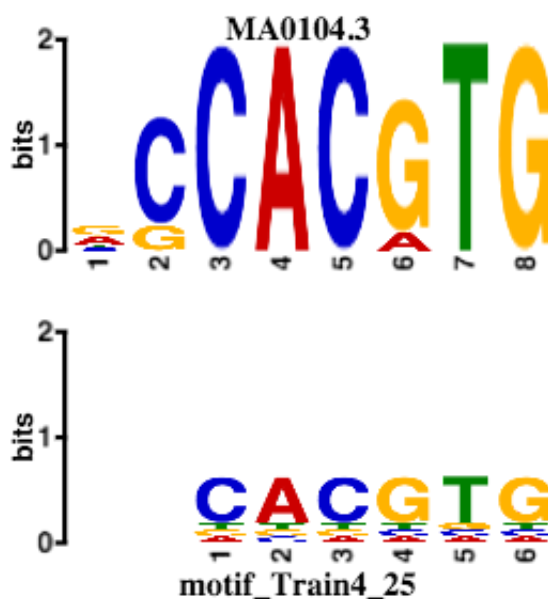
q-value 9.14e-04

Overlap 6

Offset 2

Orientation Normal

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Summary

Name [MA0281.1 \(CBF1\)](#)

Database JASPAR CORE 2014

p-value 5.80e-06

E-value 5.68e-03

q-value 9.14e-04

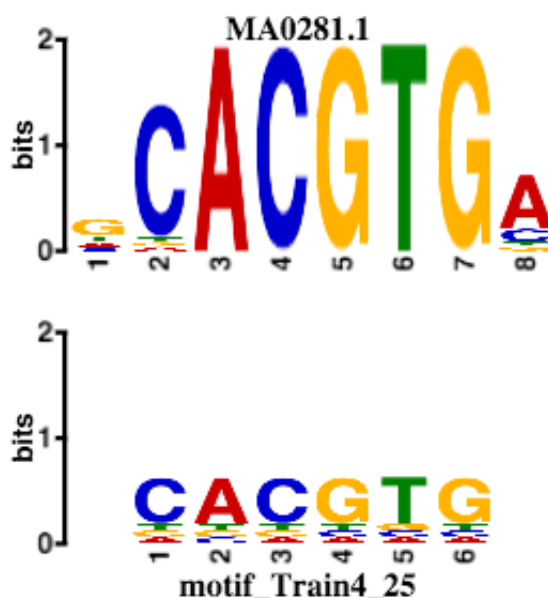
Overlap 6

Offset 1

Orientation Normal

[Show logo download options](#)

Alignment



Summary

Name [MA0357.1 \(PHO4\)](#)

Database JASPAR CORE 2014

p-value 5.80e-06

E-value 5.68e-03

q-value 9.14e-04

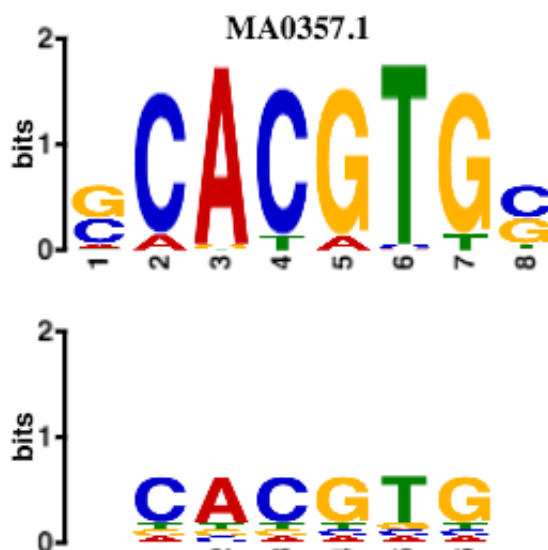
Overlap 6

Offset 1

Orientation Normal

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Alignment



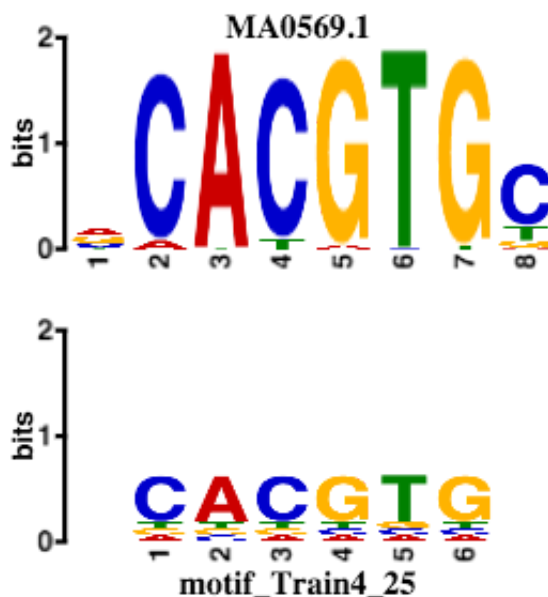
motif_Train4_25

Summary

Name	MA0569.1 (MYC4)
Database	JASPAR CORE 2014
p-value	5.80e-06
E-value	5.68e-03
q-value	9.14e-04
Overlap	6
Offset	1
Orientation	Normal

[Show logo download options](#)

Alignment

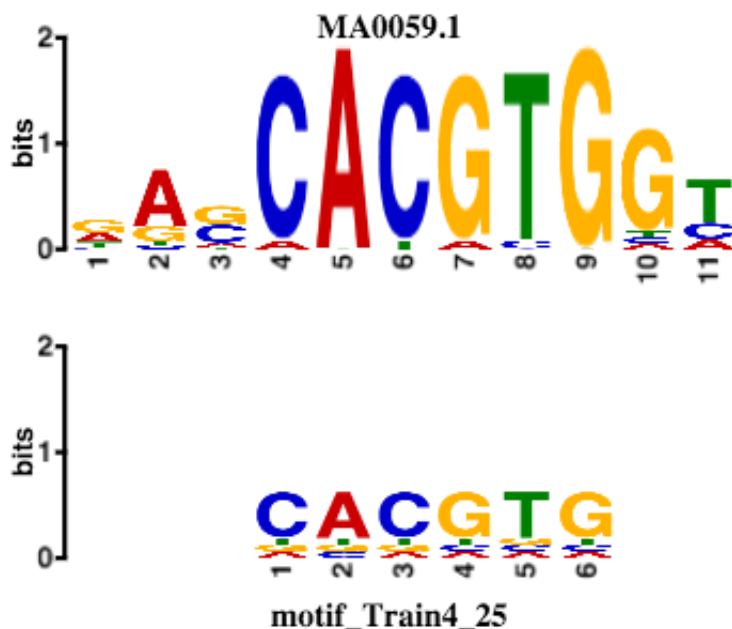


Summary

Name	MA0059.1 (MYC::MAX)
Database	JASPAR CORE 2014
p-value	1.16e-05
E-value	1.14e-02
q-value	1.37e-03
Overlap	6
Offset	3
Orientation	Normal

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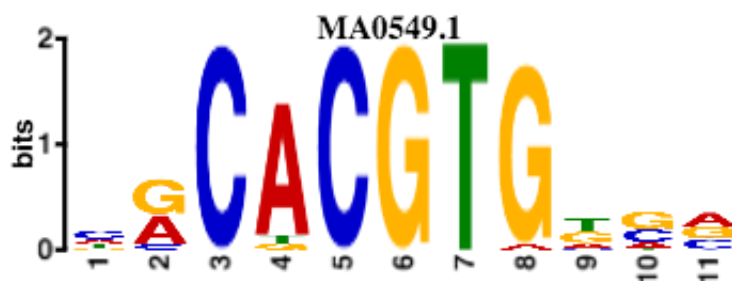
Alignment



Summary

Name	MA0549.1 (BES1)
Database	JASPAR CORE 2014
p-value	1.16e-05
E-value	1.14e-02
q-value	1.37e-03

Alignment



Overlap 6
Offset 2
Orientation Normal

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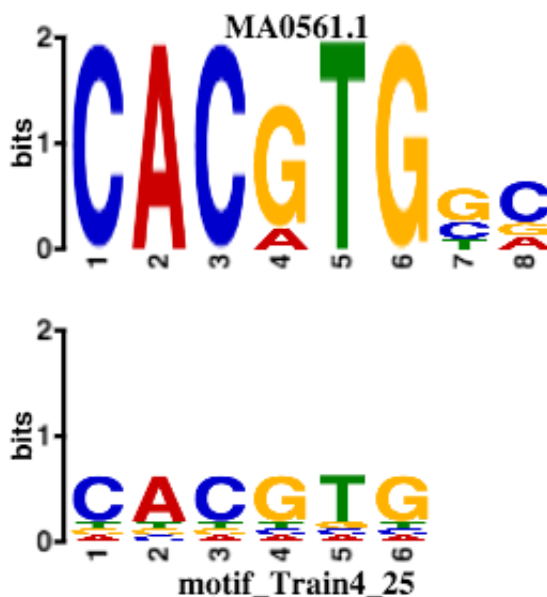
Summary

Name [MA0561.1 \(PIF4\)](#)
Database JASPAR CORE 2014
p-value 1.52e-05
E-value 1.49e-02
q-value 1.37e-03

Overlap 6
Offset 0
Orientation Normal

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Alignment



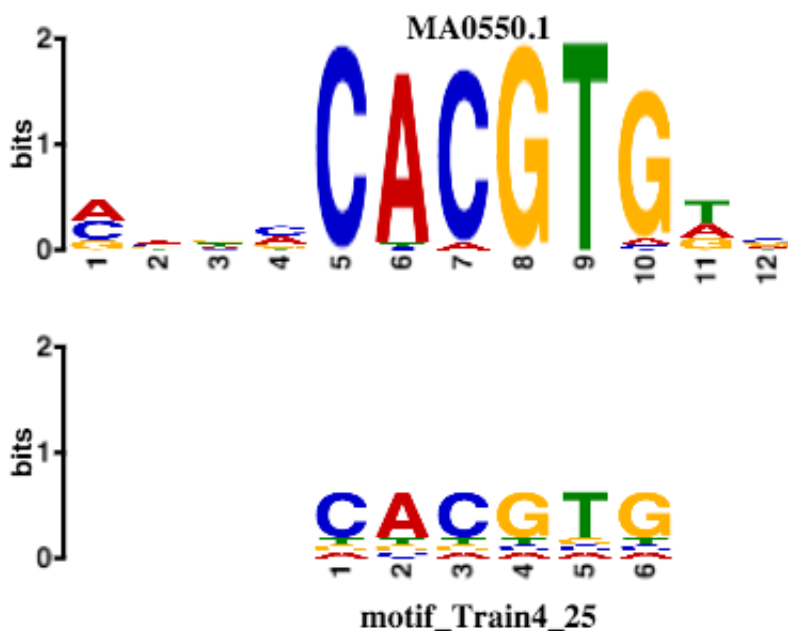
Summary

Name [MA0550.1 \(BZR1\)](#)
Database JASPAR CORE 2014
p-value 1.74e-05
E-value 1.70e-02
q-value 1.37e-03

Overlap 6
Offset 4
Orientation Normal

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Alignment



Summary

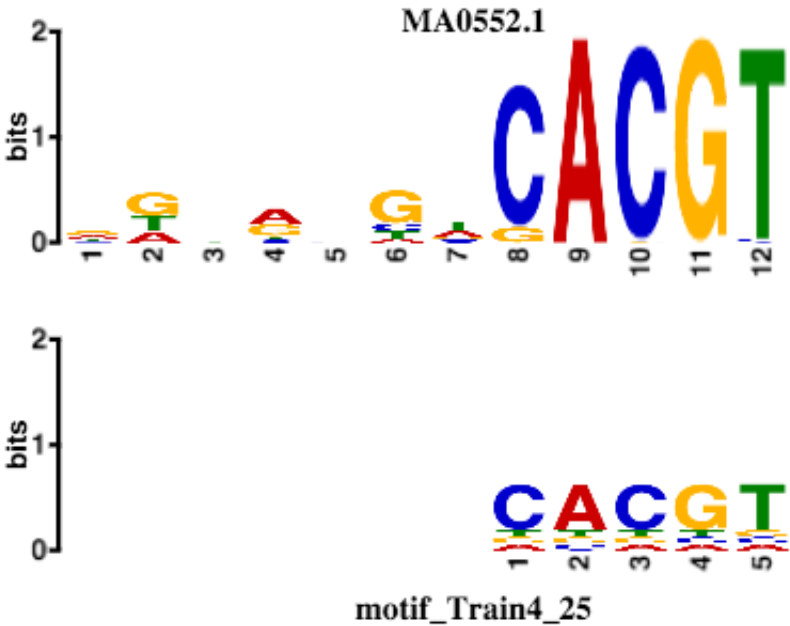
Name [MA0552.1 \(PIL5\)](#)
Database JASPAR CORE 2014

Alignment

p-value 1.74e-05
E-value 1.70e-02
q-value 1.37e-03

Overlap 6
Offset 7
Orientation Normal

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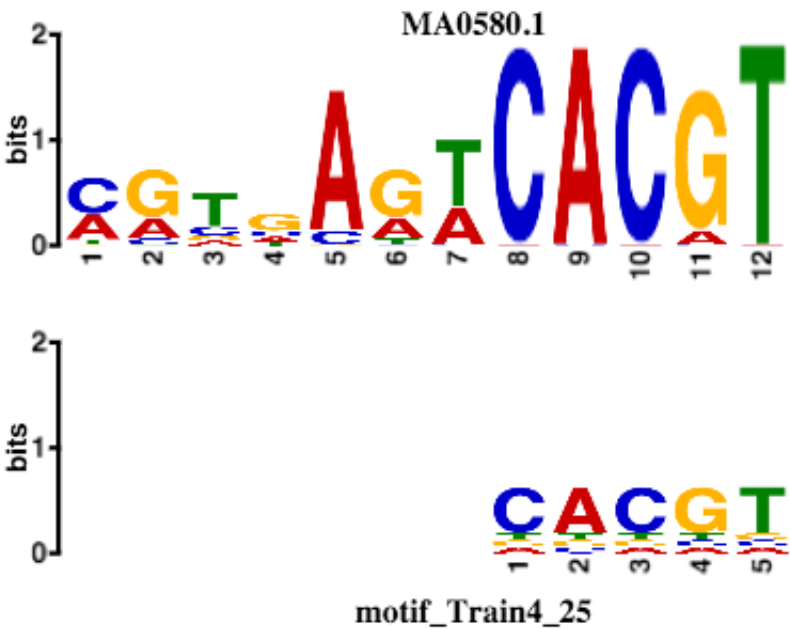
Summary

Name [MA0580.1 \(DYT1\)](#)
Database JASPAR CORE 2014
p-value 1.74e-05
E-value 1.70e-02
q-value 1.37e-03

Overlap 6
Offset 7
Orientation Normal

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Alignment



Summary

Name [UP00060_1 \(Max_primary\)](#)
Database uniprobe mouse

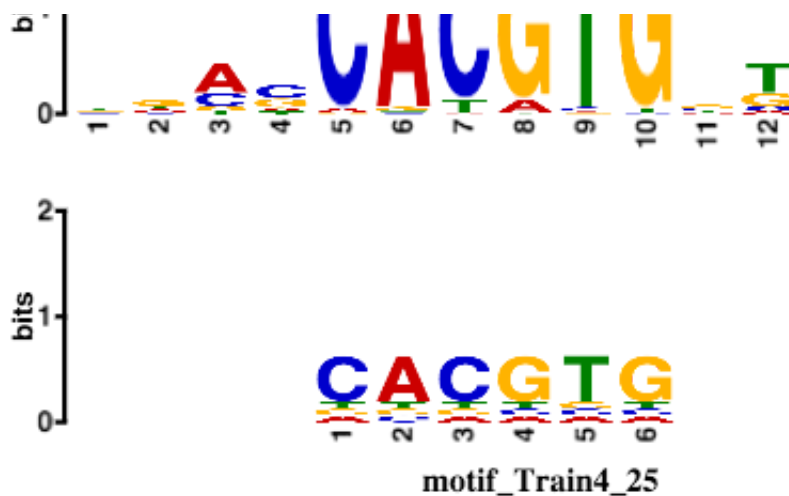
Alignment



p-value 2.13e-05
E-value 2.08e-02
q-value 1.44e-03

Overlap 6
Offset 4
Orientation Normal

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Summary

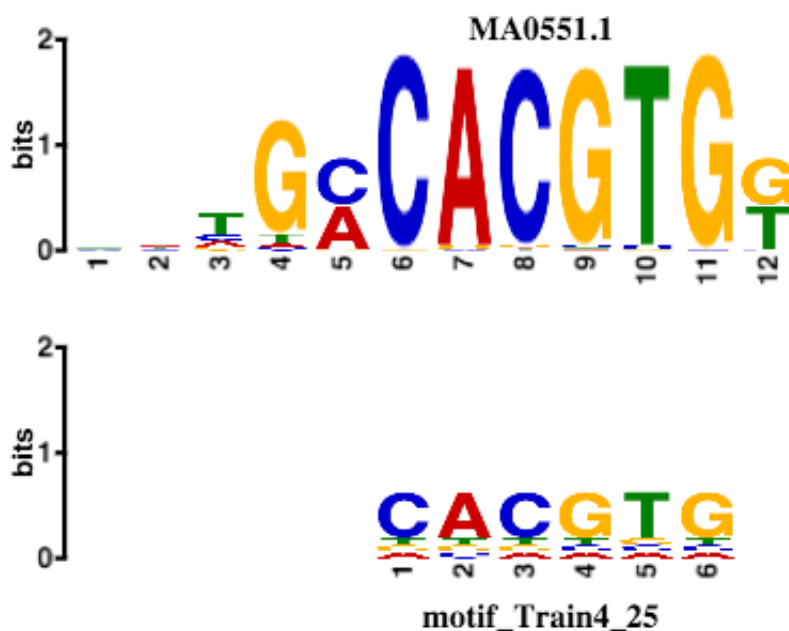
Alignment

Name [MA0551.1 \(HY5\)](#)
Database JASPAR CORE 2014

p-value 2.13e-05
E-value 2.08e-02
q-value 1.44e-03

Overlap 6
Offset 5
Orientation Normal

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Summary

Alignment

Name [MA0566.1 \(MYC2\)](#)
Database JASPAR CORE 2014

p-value 2.65e-05
E-value 2.59e-02
q-value 1.56e-03



Overlap 6
Offset 1
Orientation Normal

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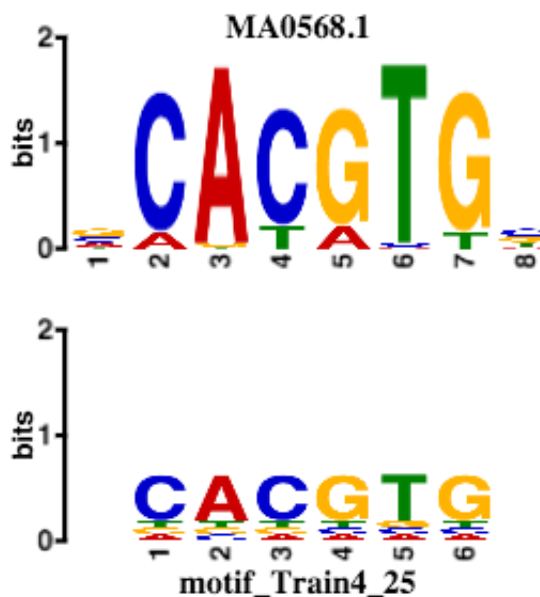
Summary

Name [MA0568.1 \(MYC3\)](#)
Database JASPAR CORE 2014
p-value 2.65e-05
E-value 2.59e-02
q-value 1.56e-03

Overlap 6
Offset 1
Orientation Normal

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Alignment



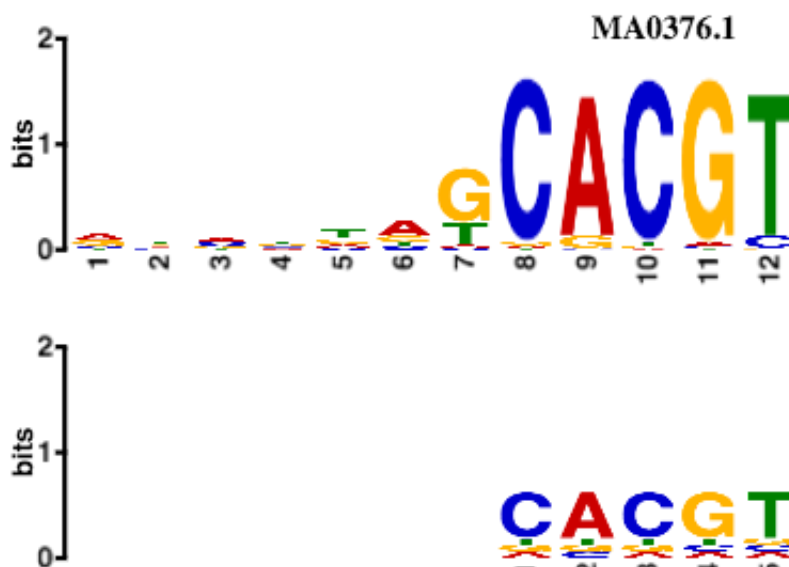
Summary

Name [MA0376.1 \(RTG3\)](#)
Database JASPAR CORE 2014
p-value 2.90e-05
E-value 2.84e-02
q-value 1.60e-03

Overlap 6
Offset 7
Orientation Normal

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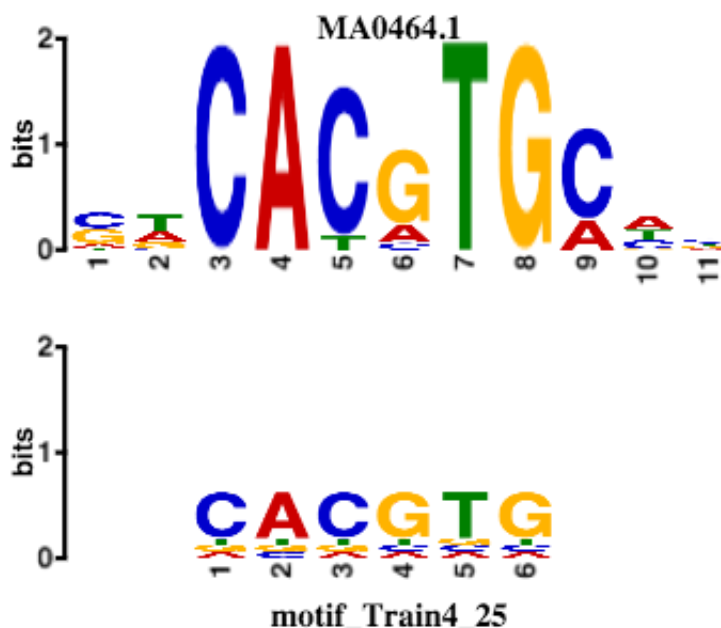
Alignment



Summary

Name	MA0464.1 (Bhlhe40)
Database	JASPAR CORE 2014
p-value	3.04e-05
E-value	2.98e-02
q-value	1.60e-03
Overlap	6
Offset	2
Orientation	Normal
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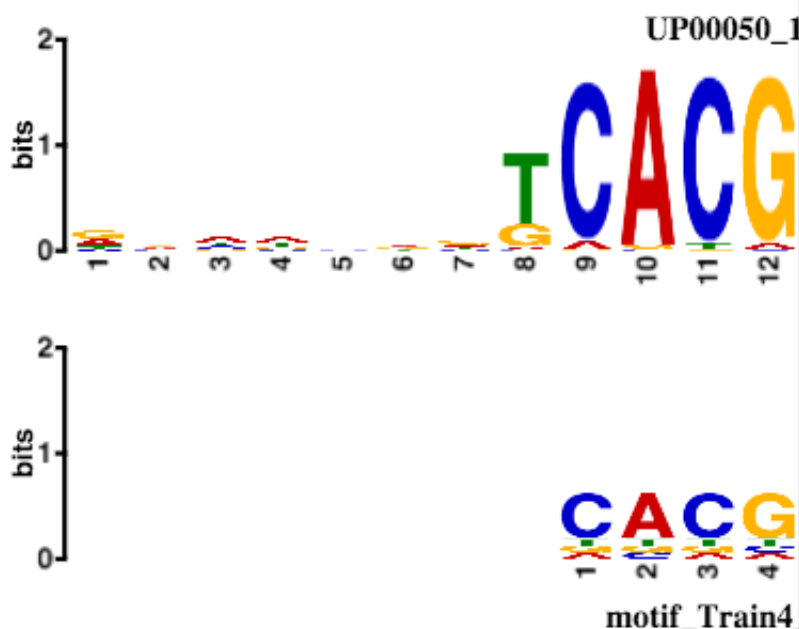
Alignment



Summary

Name	UP00050_1 (Bhlhb2_primary)
Database	uniprobe mouse
p-value	3.29e-05
E-value	3.22e-02
q-value	1.63e-03
Overlap	6
Offset	8
Orientation	Normal
	Show logo download options

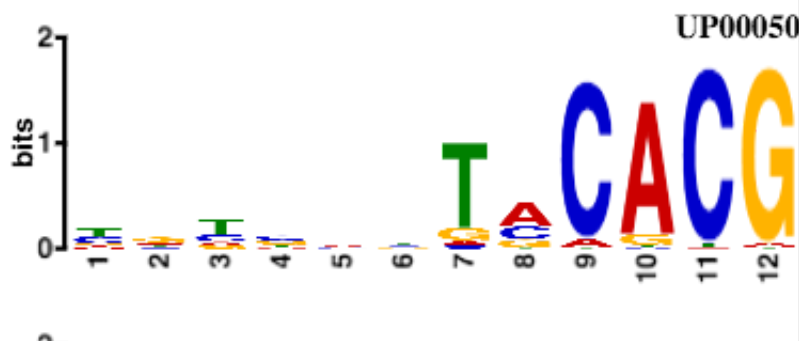
Alignment



Summary

Name	UP00050_2 (Bhlhb2_secondary)
Database	uniprobe mouse
p-value	3.48e-05
E-value	3.41e-02
q-value	1.64e-03
Overlap	6
Offset	8

Alignment



Orientation Normal[Show logo download options](#)**Summary**

Name [MA0560.1 \(PIF3\)](#)

Database JASPAR CORE 2014

p-value 4.41e-05

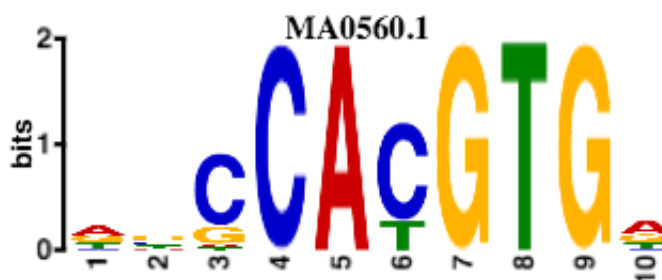
E-value 4.32e-02

q-value 1.98e-03

Overlap 6

Offset 3

Orientation Normal

[Show logo download options](#)**Alignment****Summary**

Name [MA0128.1 \(EmBP-1\)](#)

Database JASPAR CORE 2014

p-value 6.64e-05

E-value 6.50e-02

q-value 2.61e-03

Overlap 6

Offset 1

Orientation Normal

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Name [MA0310.1 \(HAC1\)](#)

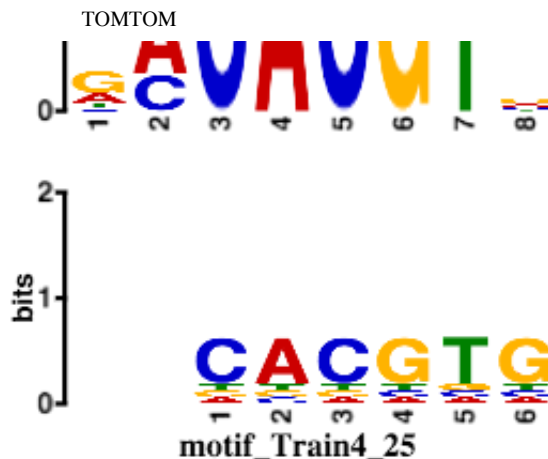
Database JASPAR CORE 2014

Alignment

p-value 6.64e-05
E-value 6.50e-02
q-value 2.61e-03

Overlap 6
Offset 2
Orientation Normal

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Summary

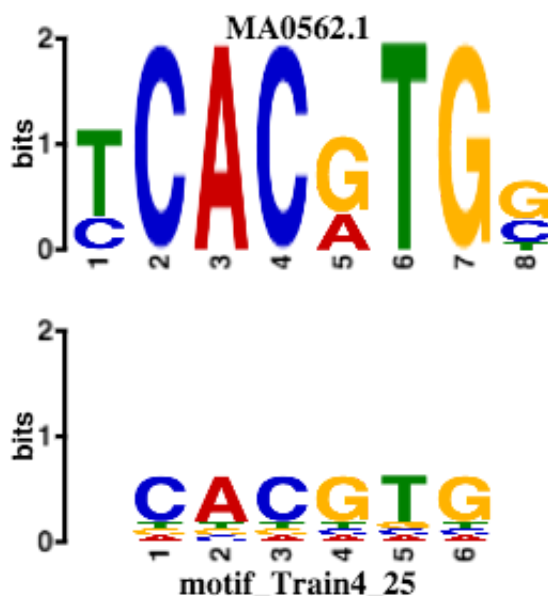
Name [MA0562.1 \(PIF5\)](#)
Database JASPAR CORE 2014

p-value 6.64e-05
E-value 6.50e-02
q-value 2.61e-03

Overlap 6
Offset 1
Orientation Normal

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Alignment



Summary

Name [MA0570.1 \(ABF1\)](#)
Database JASPAR CORE 2014

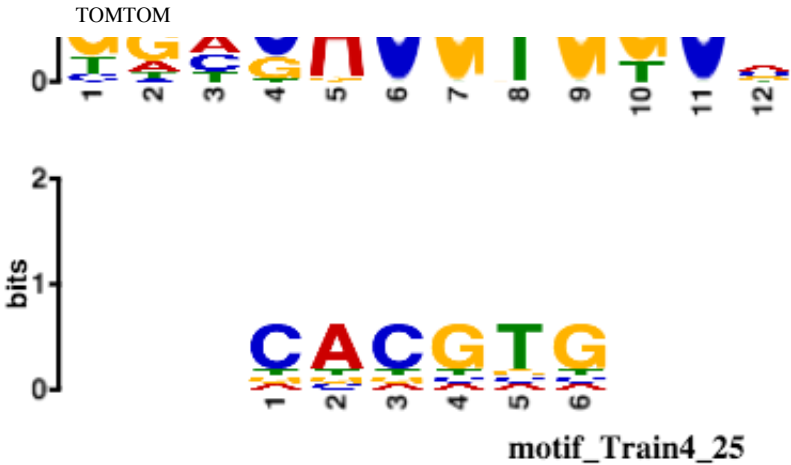
p-value 7.10e-05
E-value 6.95e-02
q-value 2.68e-03

Overlap 6
Offset 3
Orientation Normal

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Alignment





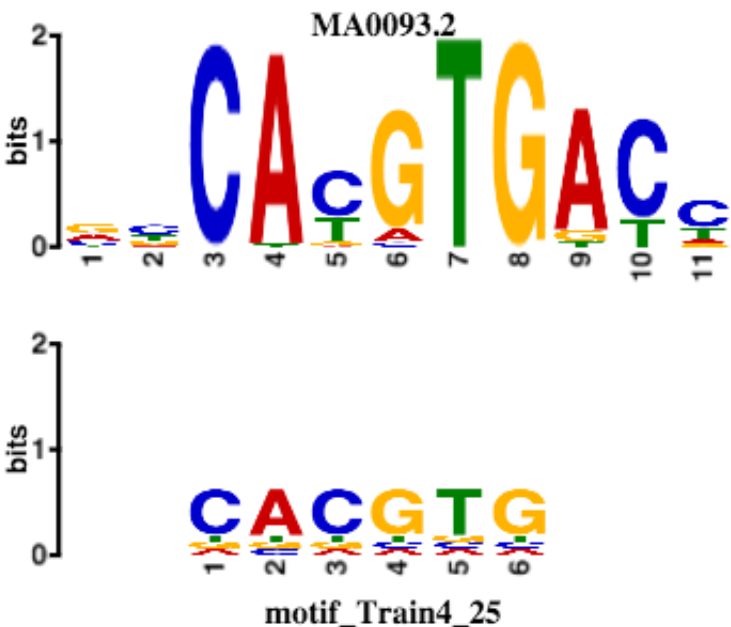
Summary

Name	MA0093.2 (USF1)
Database	JASPAR CORE 2014
<i>p</i> -value	1.63e-04
<i>E</i> -value	1.60e-01
<i>q</i> -value	5.93e-03

Overlap	6
Offset	2
Orientation	Normal

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Alignment



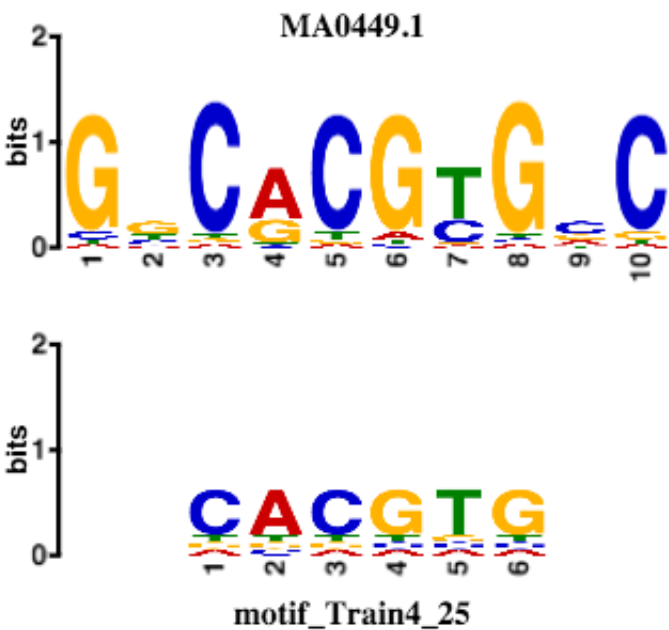
Summary

Name	MA0449.1 (h)
Database	JASPAR CORE 2014
<i>p</i> -value	1.92e-04
<i>E</i> -value	1.88e-01
<i>q</i> -value	6.70e-03

Overlap	6
Offset	2
Orientation	Normal

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Alignment



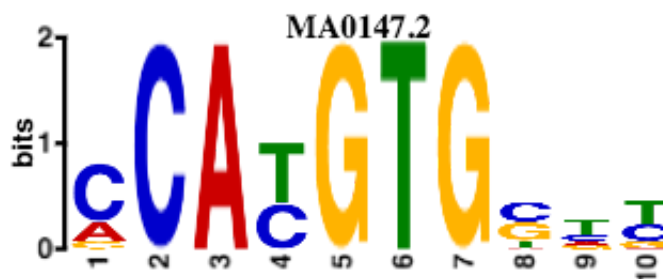
Summary

Name	MA0147.2 (Myc)
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Alignment

Name	MA0147.2 (HIF)
Database	JASPAR CORE 2014
<i>p</i>-value	9.13e-04
<i>E</i>-value	8.94e-01
<i>q</i>-value	3.08e-02
Overlap	6
Offset	1
Orientation	Normal

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Summary

Alignment

Name	MA0259.1 (HIF1A::ARNT)
Database	JASPAR CORE 2014
<i>p</i>-value	1.03e-03
<i>E</i>-value	1.01e+00
<i>q</i>-value	3.35e-02
Overlap	6
Offset	1
Orientation	Normal

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motif_Train4_25

Summary

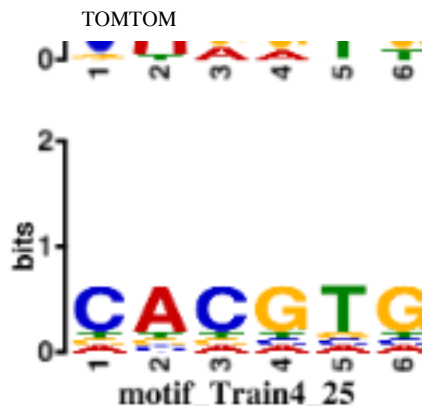
Alignment

Name	MA0086.1 (sna)
Database	JASPAR CORE 2014
<i>p</i>-value	1.35e-03
<i>E</i>-value	1.32e+00



E-value 1.52e+00
q-value 4.26e-02
Overlap 6
Offset 0
Orientation Normal

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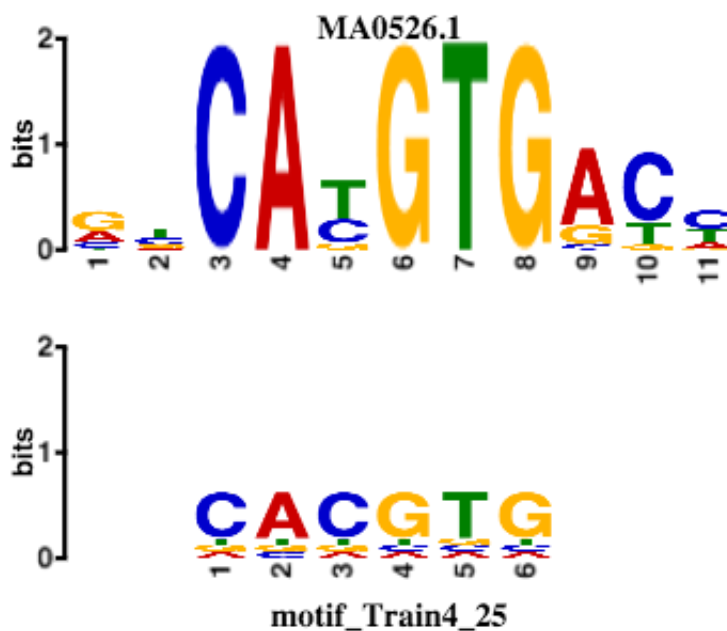


Summary

Name [MA0526.1 \(USF2\)](#)
Database JASPAR CORE 2014
p-value 1.57e-03
E-value 1.53e+00
q-value 4.77e-02
Overlap 6
Offset 2
Orientation Normal

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Alignment

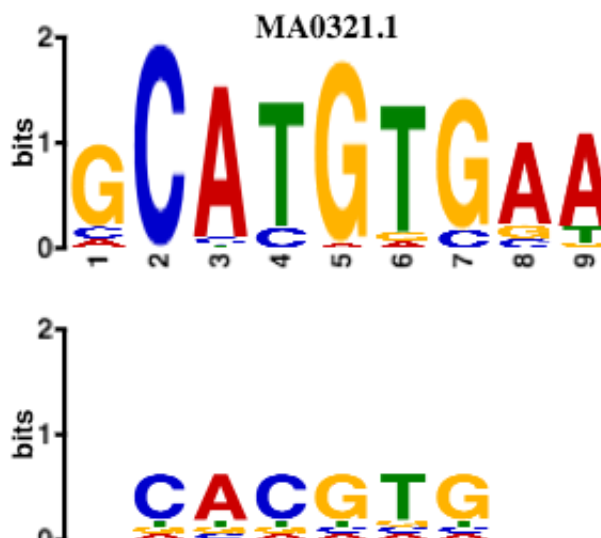


Summary

Name [MA0321.1 \(INO2\)](#)
Database JASPAR CORE 2014
p-value 2.65e-03
E-value 2.60e+00
q-value 7.83e-02
Overlap 6
Offset 1
Orientation Normal

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Alignment



motif_Train4_25

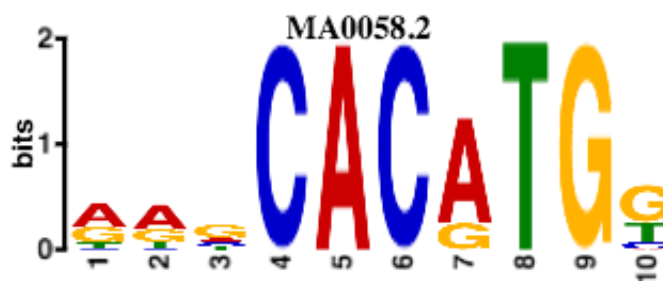
Summary

Name [MA0058.2 \(MAX\)](#)
Database JASPAR CORE 2014
p-value 2.85e-03
E-value 2.79e+00
q-value 8.16e-02

Overlap 6
Offset 3
Orientation Normal

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Alignment



motif_Train4_25

Summary

Name [UP00048_1](#)
Database uniprobe mouse
p-value 3.20e-03
E-value 3.14e+00
q-value 8.90e-02

Overlap 6
Offset 10
Orientation Normal

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Alignment

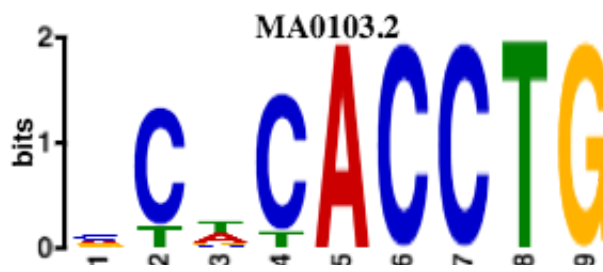


motif_Train4_25

Summary

Name [MA0103.2 \(ZEB1\)](#)
Database JASPAR CORE 2014
p-value 4.01e-03
E-value 3.93e+00
q-value 1.00e-01

Alignment



q-value 1.08e-01

Overlap 6

Offset 3

Orientation Normal

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Summary

Name [MA0322.1 \(INO4\)](#)

Database JASPAR CORE 2014

p-value 4.44e-03

E-value 4.35e+00

q-value 1.16e-01

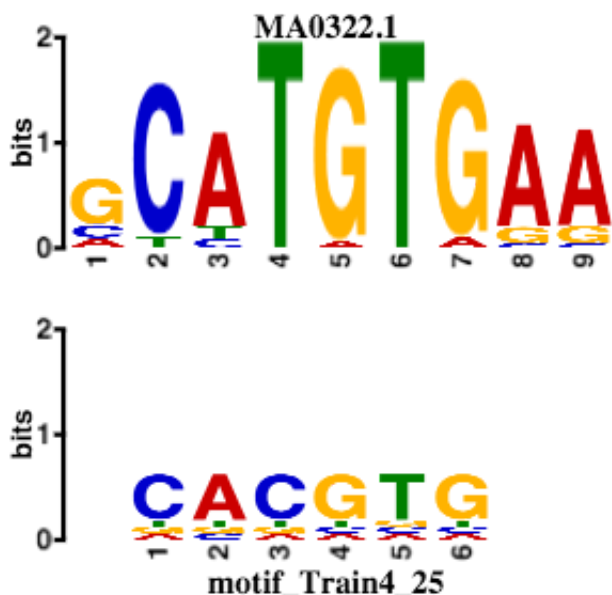
Overlap 6

Offset 1

Orientation Normal

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Alignment



Summary

Name [UP00060_2 \(Max_secondary\)](#)

Database uniprobe mouse

p-value 5.67e-03

E-value 5.55e+00

q-value 1.45e-01

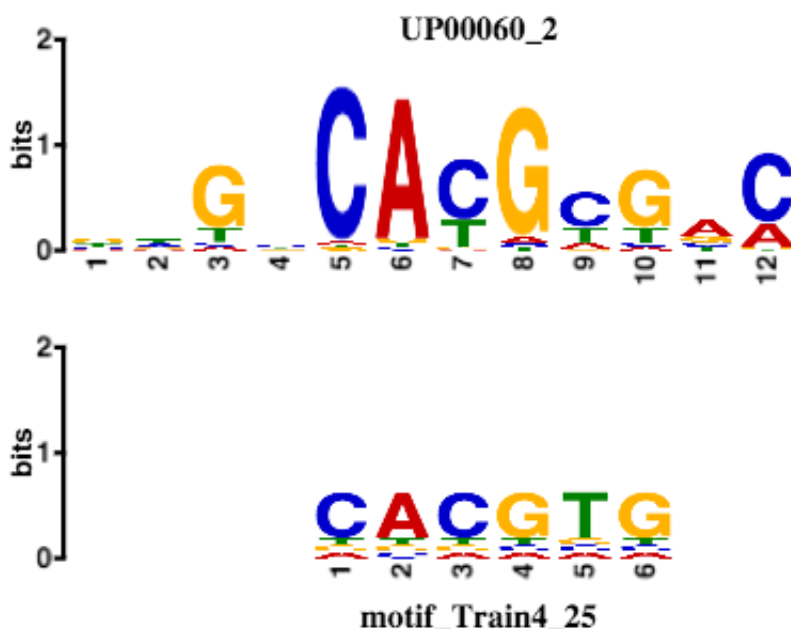
Overlap 6

Offset 4

Orientation Normal

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Alignment



Summary

Name [MA0557.1 \(FHY3\)](#)

Database JASPAR CORE 2014

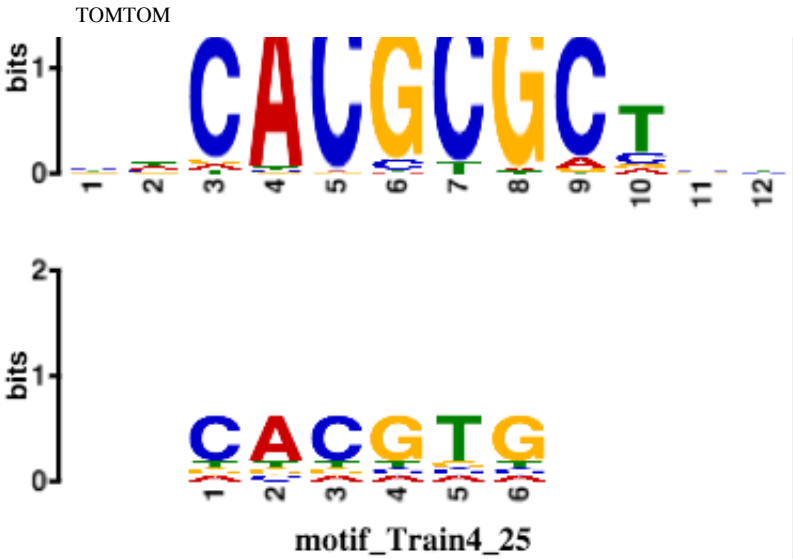
Alignment



p-value 6.05e-03
E-value 5.92e+00
q-value 1.50e-01

Overlap 6
Offset 2
Orientation Normal

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Summary

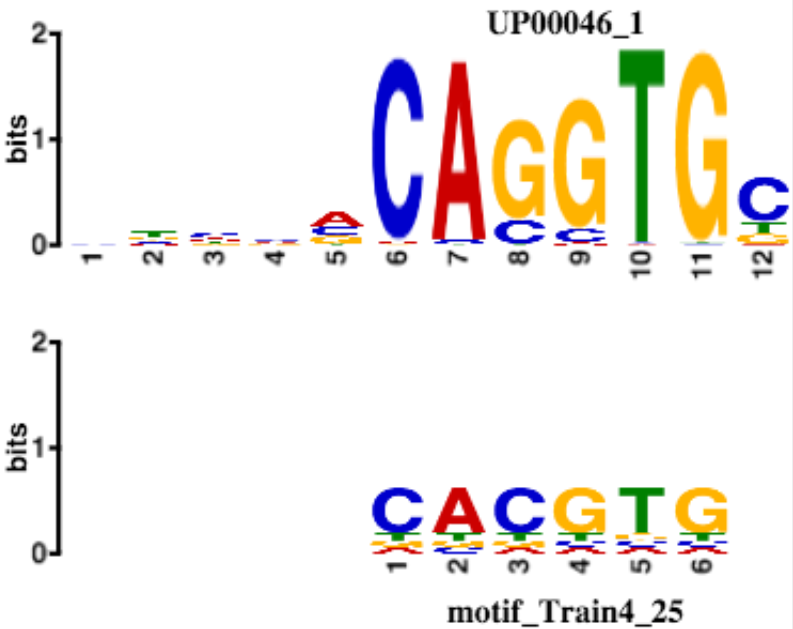
Name [UP00046_1](#)
[\(Tcfe2a_primary\)](#)
Database uniprobe mouse

p-value 6.83e-03
E-value 6.68e+00
q-value 1.65e-01

Overlap 6
Offset 5
Orientation Normal

[Show logo download options](#)

Alignment



Summary

Name [MA0097.1](#)
[\(bZIP911\)](#)
Database JASPAR CORE 2014

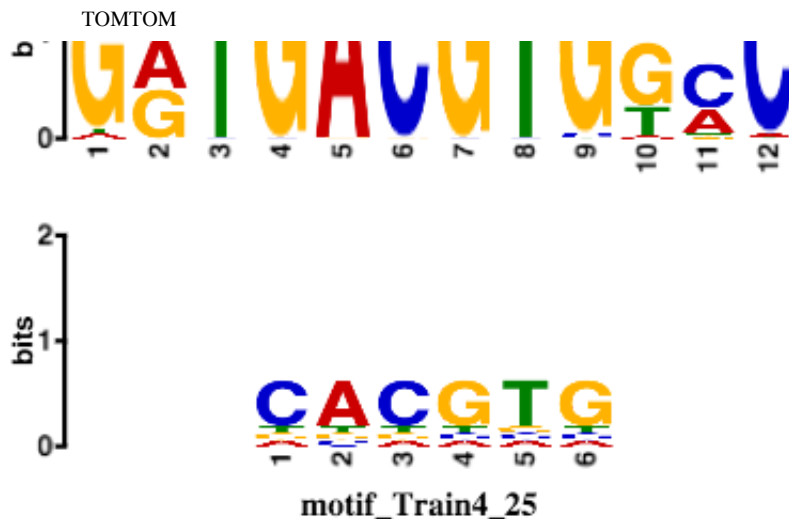
Alignment



p-value 6.98e-03
E-value 6.83e+00
q-value 1.65e-01

Overlap 6
Offset 3
Orientation Normal

[Show logo download options](#)



Summary

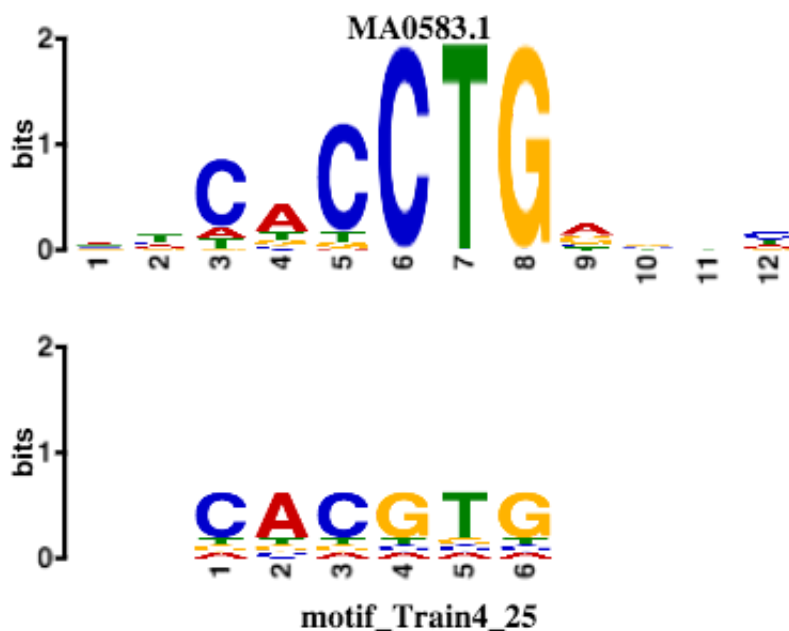
Name [MA0583.1 \(RAV1\)](#)
Database JASPAR CORE 2014

p-value 7.68e-03
E-value 7.52e+00
q-value 1.77e-01

Overlap 6
Offset 2
Orientation Normal

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Alignment



Summary

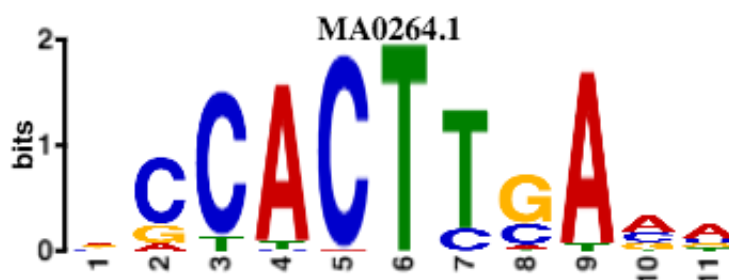
Name [MA0264.1 \(ceh-22\)](#)
Database JASPAR CORE 2014

p-value 8.73e-03
E-value 8.55e+00
q-value 1.96e-01

Overlap 6
Offset 2
Orientation Normal

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Alignment

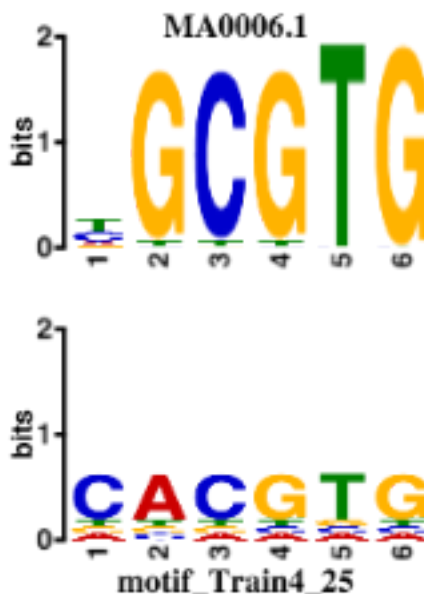




Summary

Name	MA0006.1 (Arnt::Ahr)
Database	JASPAR CORE 2014
<i>p</i>-value	9.19e-03
<i>E</i>-value	9.00e+00
<i>q</i>-value	2.02e-01
Overlap	6
Offset	0
Orientation	Normal
Show logo download options	

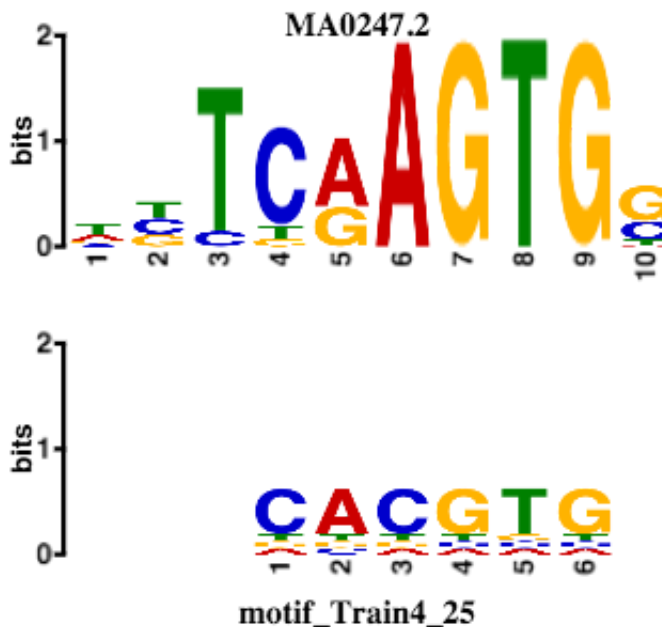
Alignment



Summary

Name	MA0247.2 (tin)
Database	JASPAR CORE 2014
<i>p</i>-value	9.43e-03
<i>E</i>-value	9.23e+00
<i>q</i>-value	2.02e-01
Overlap	6
Offset	3
Orientation	Normal
Show logo download options	

Alignment



SETTINGS

[Previous](#) [Next](#) [Top](#)

Background

Name	Bg.				Bg.	Name
Adenine	0.250	A	~	T	0.250	Thymine
Cytosine	0.250	C	~	G	0.250	Guanine

Other Settings

Strand Handling

Motifs may be reverse complemented before comparison to find a better match.

Distance Measure

Pearson correlation coefficient

Match Threshold

Matches must have a E-value of 10 or smaller.

[Previous Top](#)

TOMTOM version

4.11.1 (Release date: Fri Jan 15 12:51:59 2016 -0800)

Reference

Shobhit Gupta, JA Stamatoyannopolous, Timothy Bailey and William Stafford Noble, "Quantifying similarity between motifs", *Genome Biology*, 8(2):R24, 2007.

Command line

```
tomtom -no-ssc -oc . -verbosity 1 -min-overlap 5 -dist pearson -evaluate -thresh 10.0 motif_group5.txt
```

Result calculation took 5.729 seconds

