



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

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Database	Name	Alt. Name	Preview	Matches	
motif group6	motif_10_3	4- TGGCAACGYTGY		9	MA0365.1 UP00084
motif group6	motif_Train1_3	5- TGGCAACGYTGC		8	MA0365.1 UP00084
motif group6	motif_Train2_3	4- TGGCAACGTTGC		12	MA0365.1 UP00084
motif group6	motif_Train4_3	4- TGGCAACGYTGH		9	MA0365.1 UP00084
motif group6	motif_Train5_3	4- TGGCAACGYTGH		9	MA0365.1 UP00084
motif group6	motif_Train6_3	4- TGGCAACGYTGY		9	MA0365.1 UP00084
motif group6	motif_Train7_3	5- TGGCAACGYTGY		9	MA0365.1 UP00084
motif group6	motif_Train8_3	5- TGGCAACGYTGY		10	MA0365.1 UP00084
motif group6	motif_Train9_3	4- TGGCAACGYTGN		9	MA0365.1 UP00084


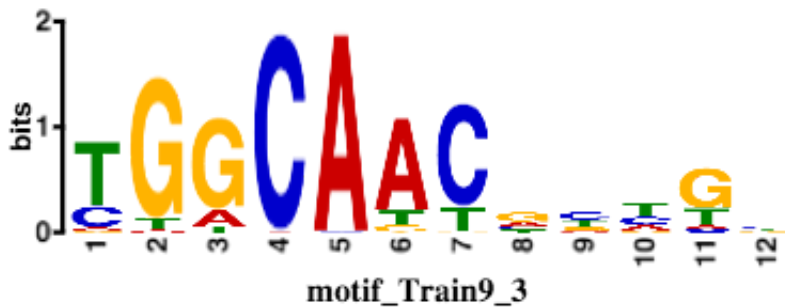
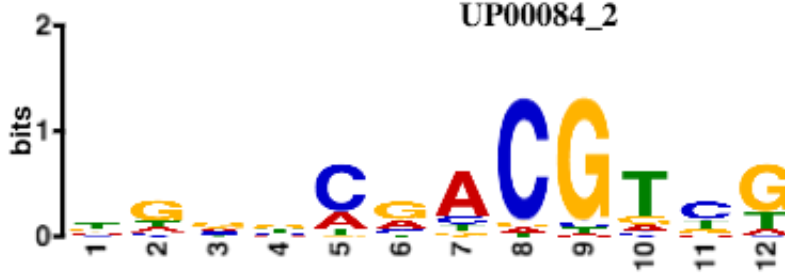

TARGET DATABASES

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Database	Used	Matched
uniprobe mouse	386	2
JASPAR CORE 2014	593	13

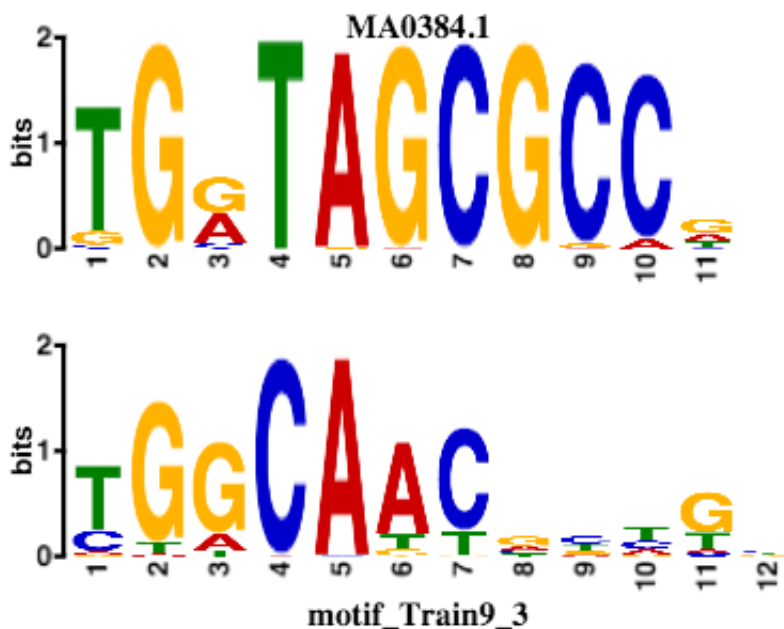
MATCHES TO MOTIF_Train9_3 (4-TGGCAACGYTGN)

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Summary	Alignment
<p>Name MA0365.1 (RFX1)</p> <p>Database JASPAR CORE 2014</p> <p>p-value 2.29e-04</p> <p>E-value 2.24e-01</p> <p>q-value 4.49e-01</p> <p>Overlap 8</p> <p>Offset 0</p> <p>Orientation Reverse Complement</p> <p>Show logo download options</p>	 <p>MA0365.1</p>  <p>motif_Train9_3</p>
<p>Name UP00084_2 (Gmeb1_secondary)</p> <p>Database uniprobe mouse</p> <p>p-value 8.31e-04</p> <p>E-value 8.14e-01</p> <p>q-value 8.14e-01</p> <p>Overlap 12</p> <p>Offset 1</p> <p>Orientation Normal</p> <p>Show logo download options</p>	 <p>UP00084_2</p>  <p>motif_Train9_3</p>
<p>Name MA0384.1 (SNT2)</p> <p>Database JASPAR CORE 2014</p>	

<i>p</i>-value	2.25e-03
<i>E</i>-value	2.20e+00
<i>q</i>-value	1.00e+00
Overlap	11
Offset	0
Orientation	Normal

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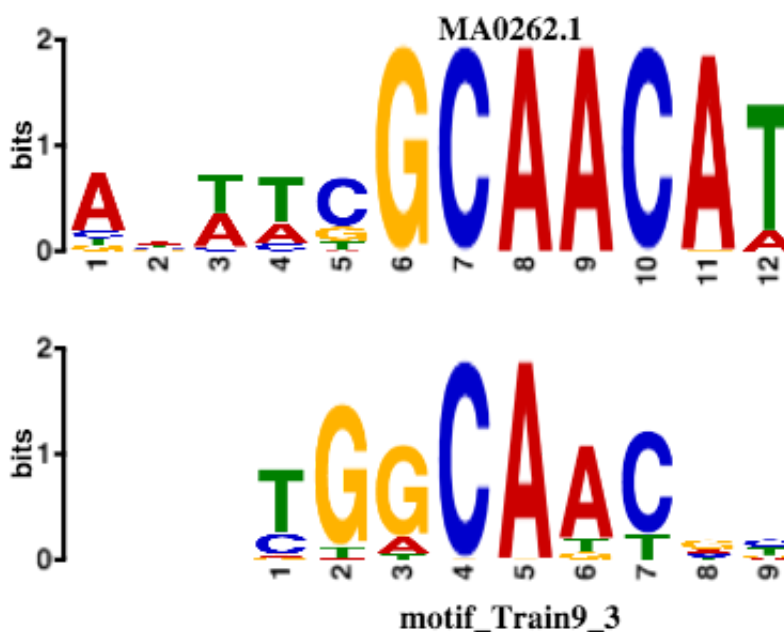


Summary

Name	MA0262.1 (mab-3)
Database	JASPAR CORE 2014
<i>p</i>-value	2.60e-03
<i>E</i>-value	2.55e+00
<i>q</i>-value	1.00e+00
Overlap	10
Offset	3
Orientation	Reverse Complement

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Alignment

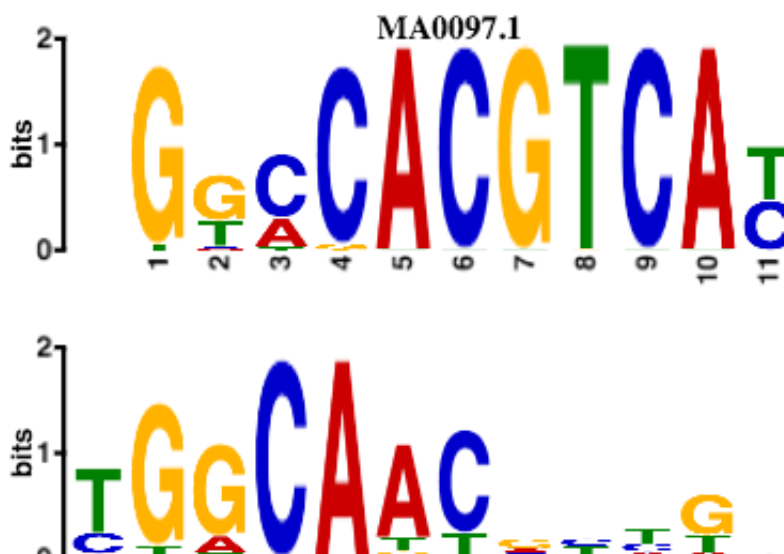


Summary

Name	MA0097.1 (bZIP911)
Database	JASPAR CORE 2014
<i>p</i>-value	4.09e-03
<i>E</i>-value	4.00e+00
<i>q</i>-value	1.00e+00
Overlap	11
Offset	-1
Orientation	Reverse Complement

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Alignment



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TOMTOM

1 2 3 4 5 6 7 8 9 10 11 12

motif_Train9_3

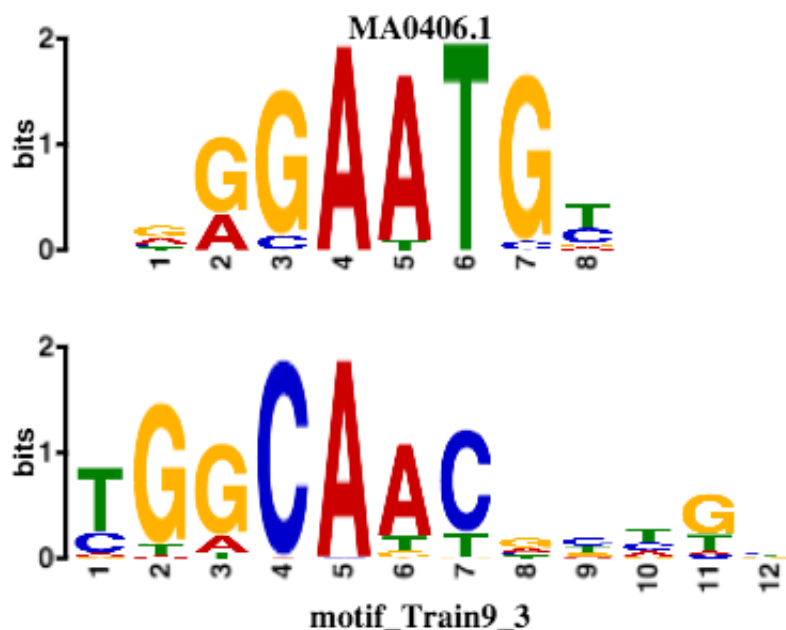
Summary

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014
p-value 4.53e-03
E-value 4.44e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse Complement

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Alignment



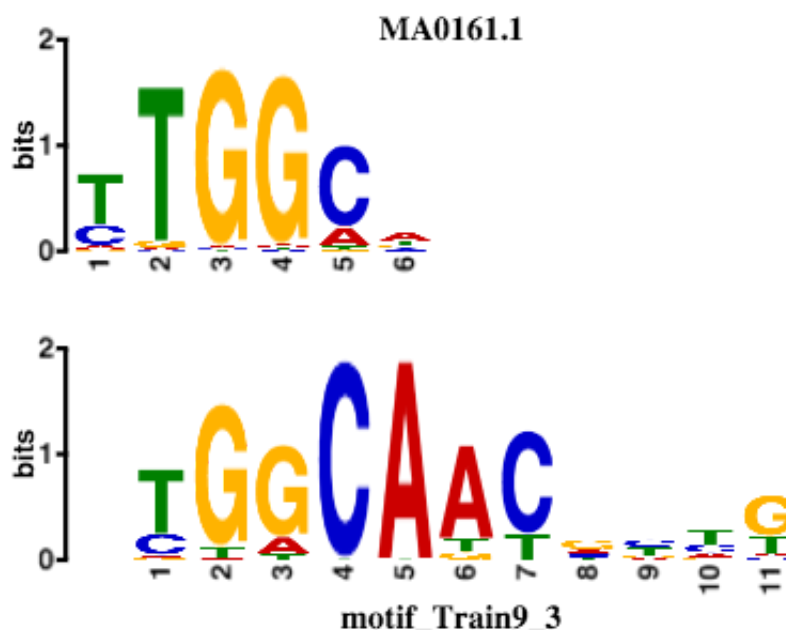
Summary

Name [MA0161.1 \(NFIC\)](#)
Database JASPAR CORE 2014
p-value 6.16e-03
E-value 6.03e+00
q-value 1.00e+00

Overlap 5
Offset 1
Orientation Normal

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Alignment



Summary

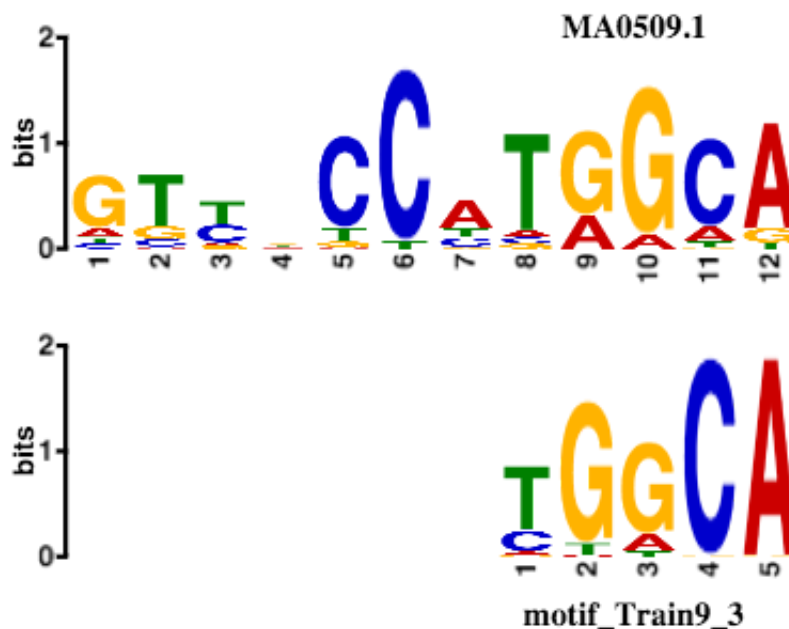
Name [MA0509.1 \(Rfx1\)](#)
Database JASPAR CORE 2014
p-value 7.10e-03
E-value 6.95e+00
q-value 1.00e+00

Overlap 7
Offset 7

Alignment

Offset /
Orientation Normal

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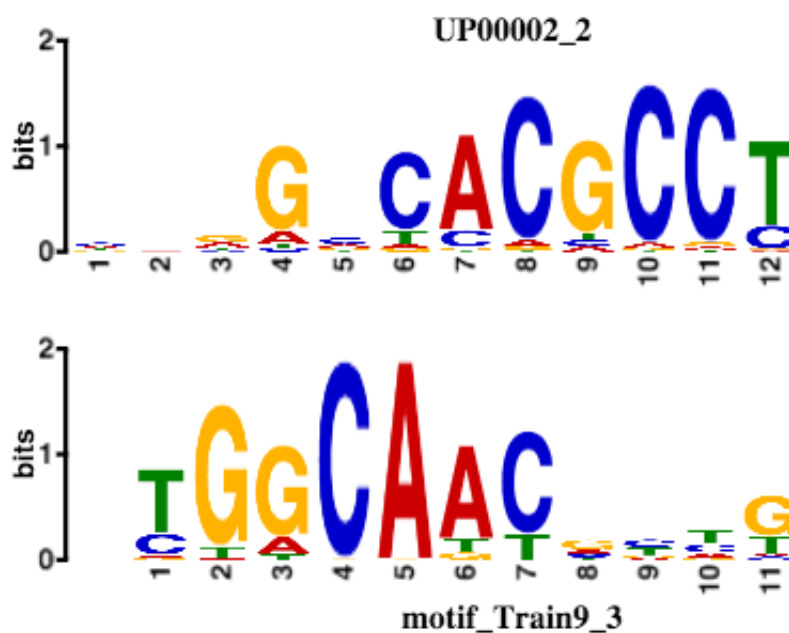
Summary

Name [UP00002_2 \(Sp4_secondary\)](#)
Database uniprobe mouse
p-value 9.69e-03
E-value 9.49e+00
q-value 1.00e+00

Overlap 12
Offset 1
Orientation Reverse Complement

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Alignment



MATCHES TO MOTIF_Train5_3 (4-TGGCAACGYTGH)

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Summary

Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014
p-value 1.96e-04

Alignment

E-value 1.92e-01
q-value 3.84e-01
Overlap 8
Offset 0
Orientation Reverse Complement

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Summary

Name [UP00084_2 \(Gmeb1_secondary\)](#)
Database uniprobe mouse
p-value 1.06e-03
E-value 1.04e+00
q-value 8.66e-01

Overlap 12
Offset 1
Orientation Normal

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Alignment



Summary

Name [MA0384.1 \(SNT2\)](#)
Database JASPAR CORE 2014
p-value 1.67e-03
E-value 1.64e+00
q-value 8.66e-01

Overlap 11
Offset 0
Orientation Normal

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Alignment

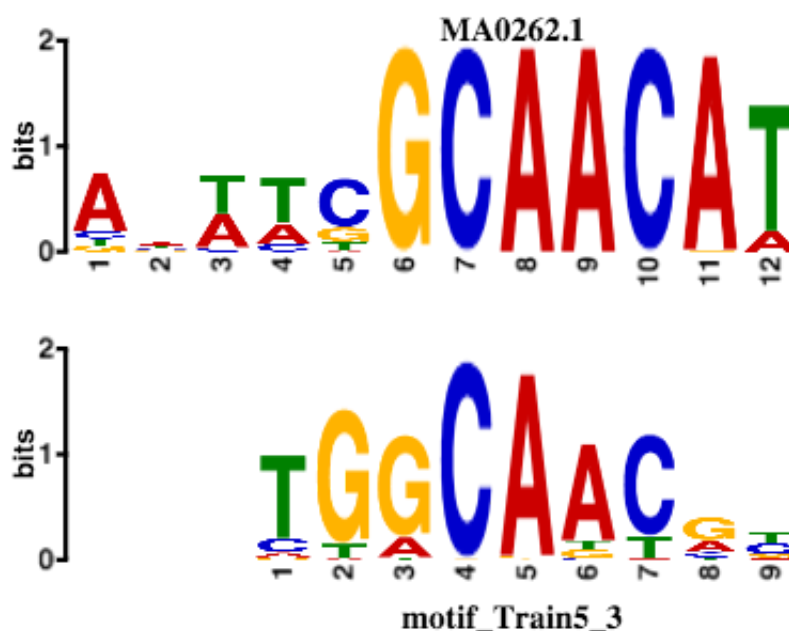


motif_Train5_3

Summary

Name	MA0262.1 (mab-3)
Database	JASPAR CORE 2014
p-value	1.77e-03
E-value	1.73e+00
q-value	8.66e-01
Overlap	10
Offset	3
Orientation	Reverse Complement
	Show logo download options

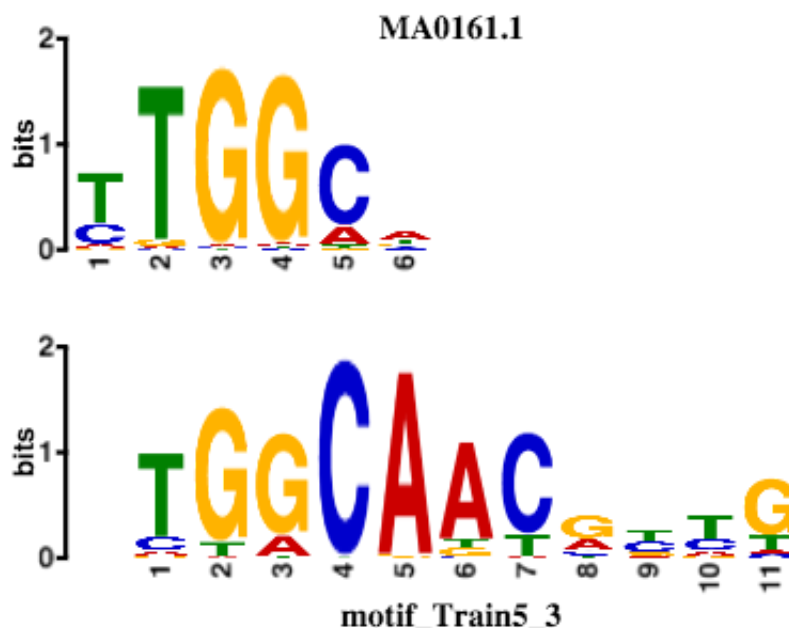
Alignment



Summary

Name	MA0161.1 (NFIC)
Database	JASPAR CORE 2014
p-value	4.35e-03
E-value	4.26e+00
q-value	1.00e+00
Overlap	5
Offset	1
Orientation	Normal
	Show logo download options

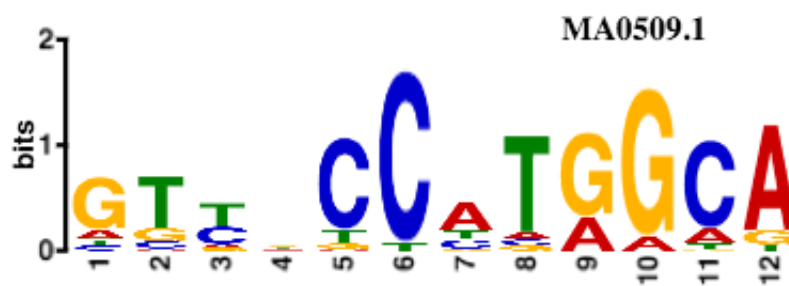
Alignment



Summary

Name	MA0509.1 (Rfx1)
Database	JASPAR CORE 2014
p-value	6.33e-03
E-value	6.20e+00
q-value	1.00e+00

Alignment



Overlap 7
Offset 7
Orientation Normal

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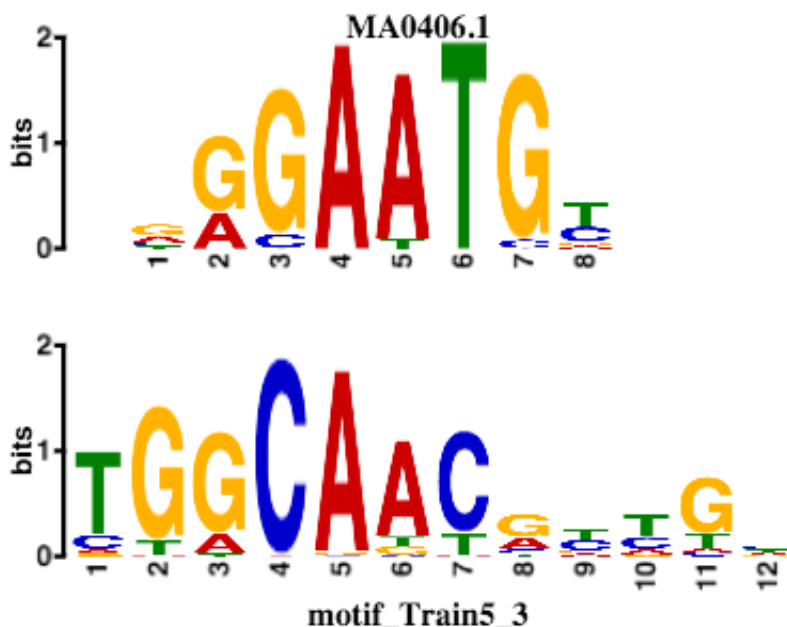
Summary

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014
p-value 6.91e-03
E-value 6.76e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse Complement

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Alignment



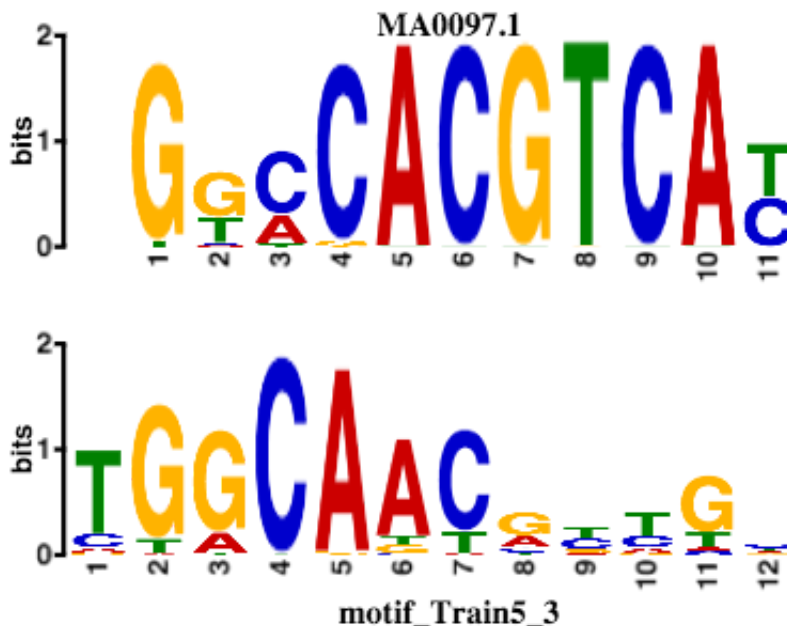
Summary

Name [MA0097.1 \(bZIP911\)](#)
Database JASPAR CORE 2014
p-value 7.13e-03
E-value 6.98e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Reverse Complement

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Alignment



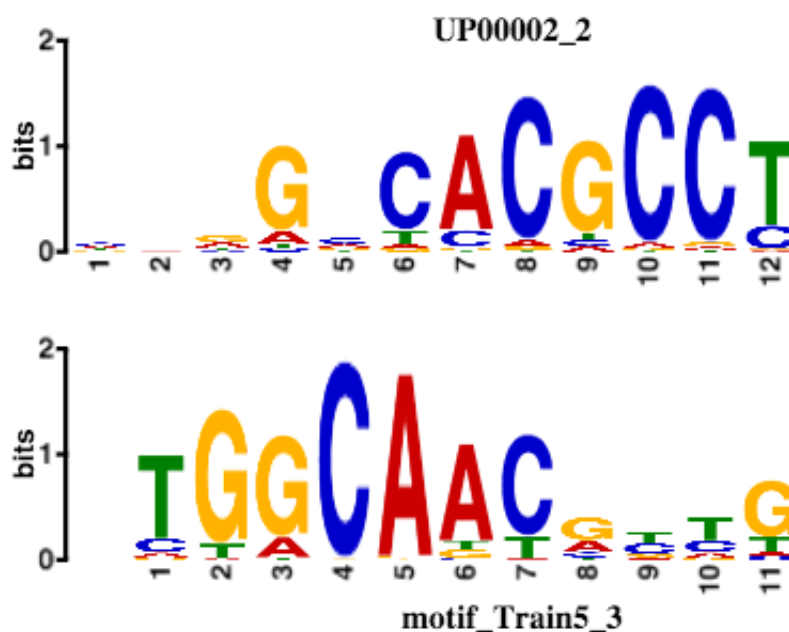
Summary

Name [UP00002_2 \(Sp4_secondary\)](#)
Database uniprobe mouse

Alignment

<i>p</i>-value	9.22e-03
<i>E</i>-value	9.03e+00
<i>q</i>-value	1.00e+00
Overlap	12
Offset	1
Orientation	Reverse
	Complement

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MATCHES TO MOTIF_TRAIN6_3 (4-TGGCAACGYTGY)

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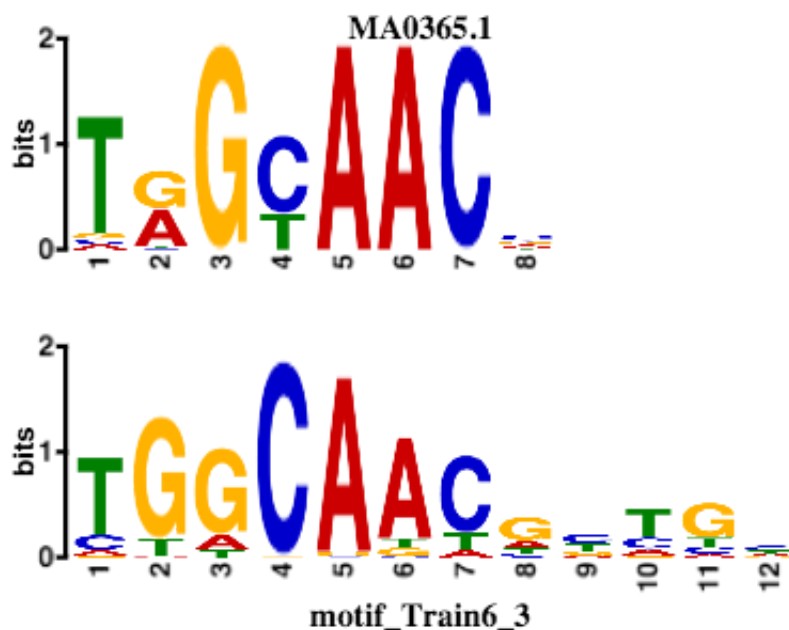
Summary

Name	MA0365.1 (RFX1)
Database	JASPAR CORE 2014
<i>p</i>-value	2.67e-04
<i>E</i>-value	2.61e-01
<i>q</i>-value	5.23e-01

Overlap	8
Offset	0
Orientation	Reverse
	Complement

[Show logo download options](#)

Alignment

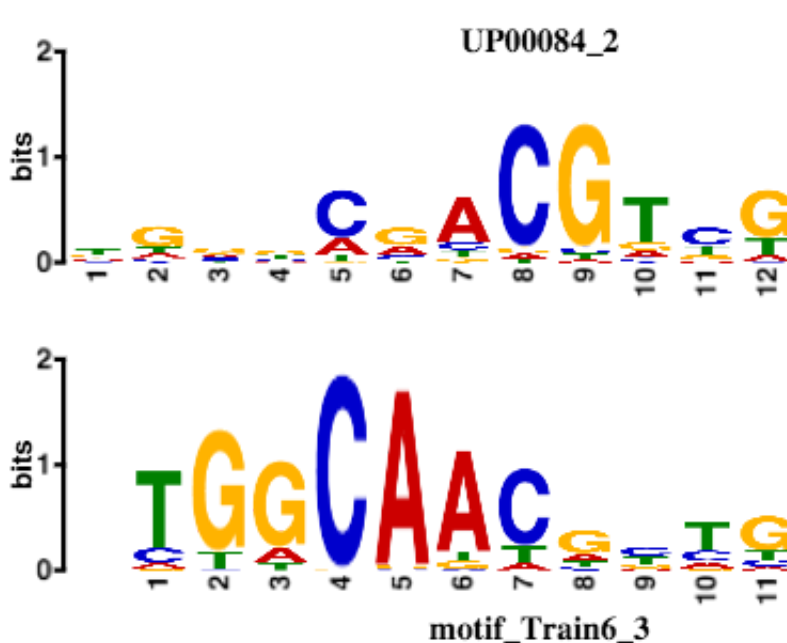


Summary

Alignment

[UP00084_2](#)

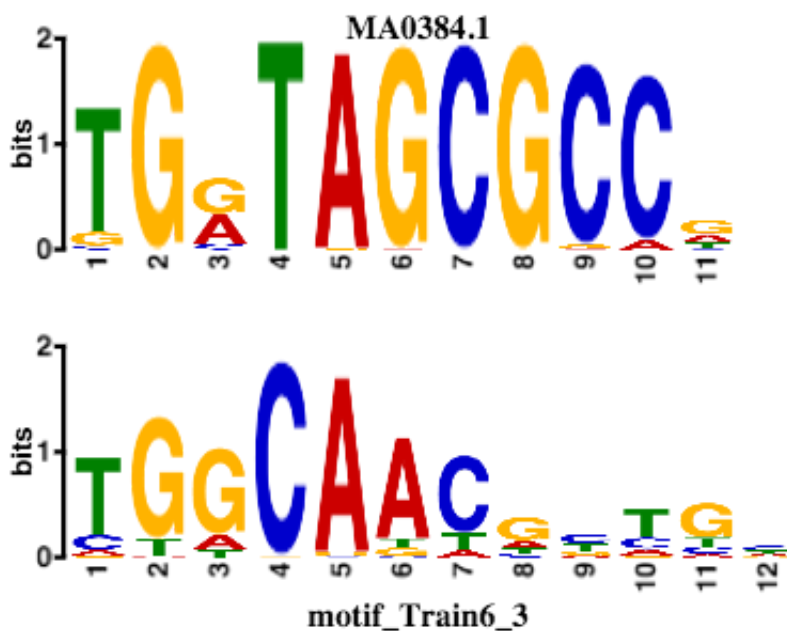
Name	(Gmeb1_secondary)
Database	uniprobe mouse
p-value	1.13e-03
E-value	1.10e+00
q-value	1.00e+00
Overlap	12
Offset	1
Orientation	Normal
	Show logo download options



Summary

Name	MA0384.1 (SNT2)
Database	JASPAR CORE 2014
p-value	2.17e-03
E-value	2.12e+00
q-value	1.00e+00
Overlap	11
Offset	0
Orientation	Normal
	Show logo download options

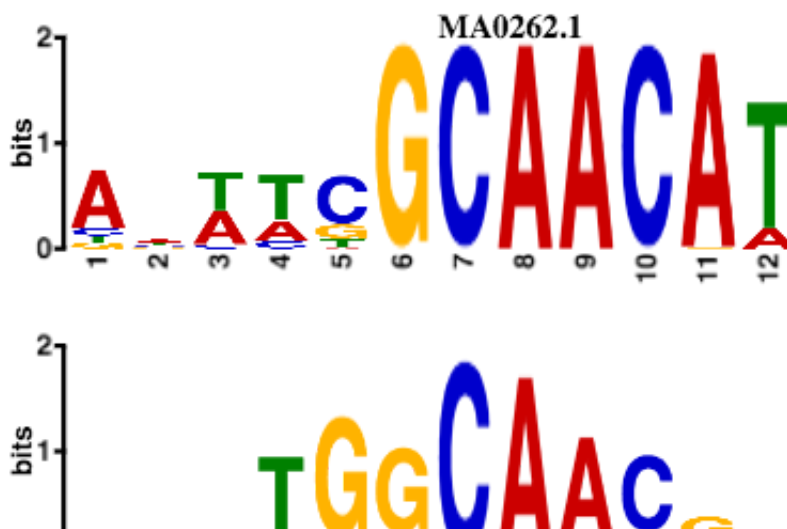
Alignment



Summary

Name	MA0262.1 (mab-3)
Database	JASPAR CORE 2014
p-value	3.69e-03
E-value	3.61e+00
q-value	1.00e+00
Overlap	10
Offset	3
Orientation	Reverse Complement
	Show logo download options

Alignment





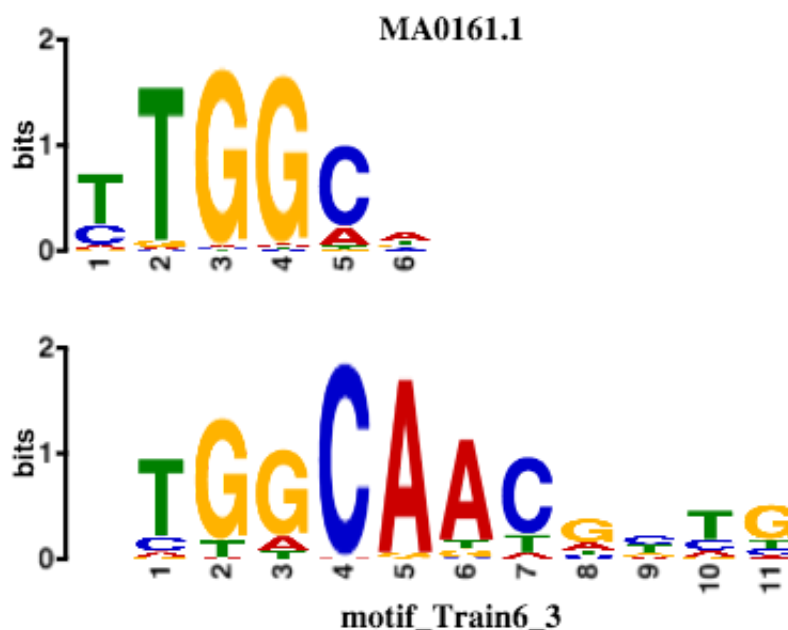
Summary

Name [MA0161.1 \(NFIC\)](#)
Database JASPAR CORE 2014
p-value 5.83e-03
E-value 5.71e+00
q-value 1.00e+00

Overlap 5
Offset 1
Orientation Normal

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Alignment



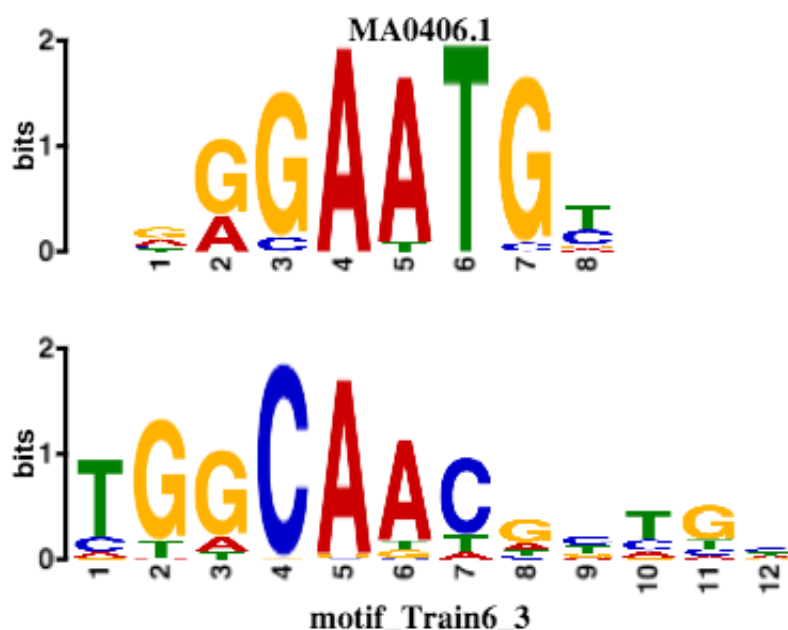
Summary

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014
p-value 7.28e-03
E-value 7.13e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse Complement

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Alignment



Summary

Name [MA0097.1 \(bZIP911\)](#)
Database JASPAR CORE 2014
p-value 7.51e-03
E-value 7.35e+00
q-value 1.00e+00

Overlap 11

Alignment

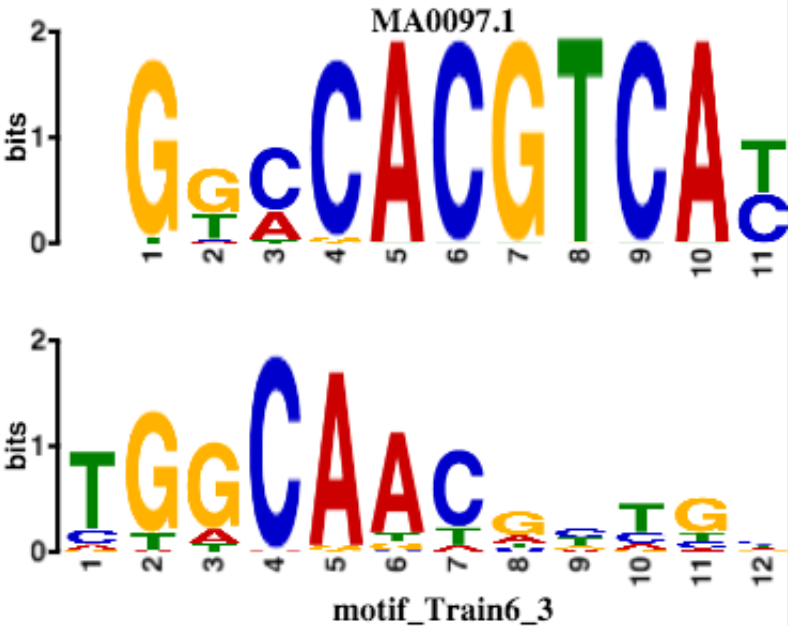
Overlap11

Offset-1

OrientationReverse

Complement

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Summary

NameMA0509.1 (Rfx1)

DatabaseJASPAR CORE 2014

p-value9.18e-03

E-value8.99e+00

q-value1.00e+00

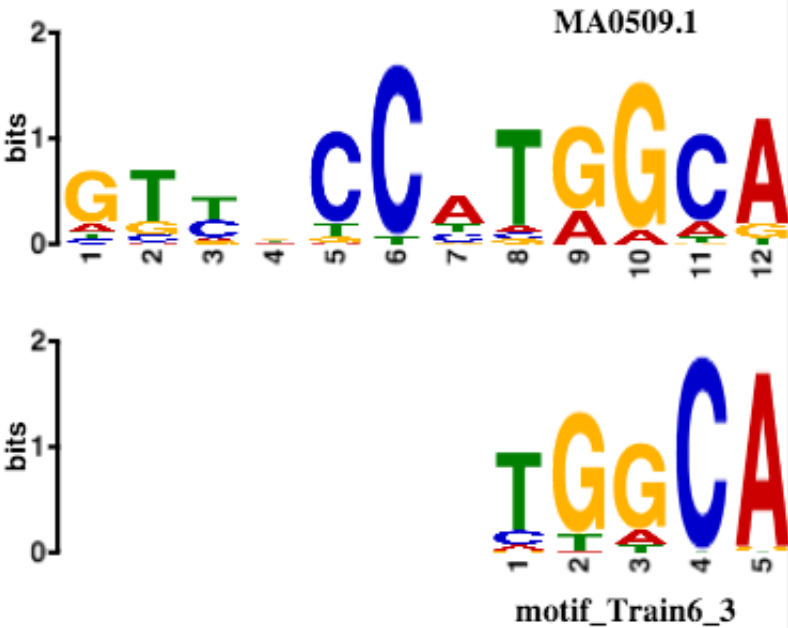
Overlap7

Offset7

OrientationNormal

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Alignment



Summary

NameUP00002_2 (Sp4 secondary)

Databaseuniprobe mouse

p-value9.94e-03

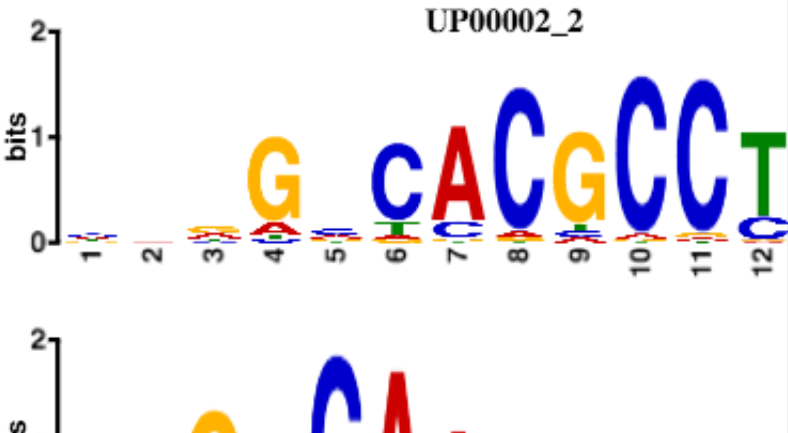
E-value9.73e+00

q-value1.00e+00

Overlap12

Offset1

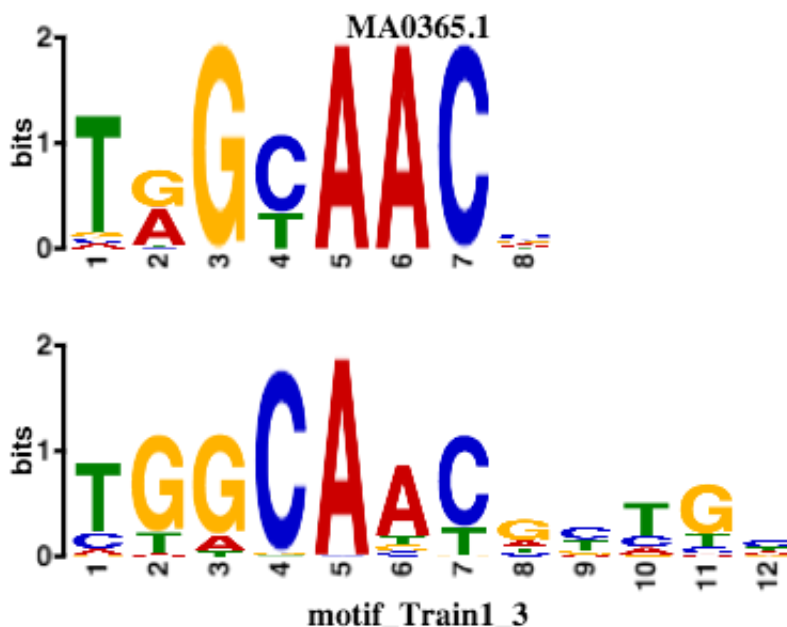
Alignment



OrientationReverse
Complement[Show logo download options](#)**MATCHES TO MOTIF_Train1_3 (5-TGGCAACGYTGC)**[Previous](#) [Next](#) [Top](#)**Summary**

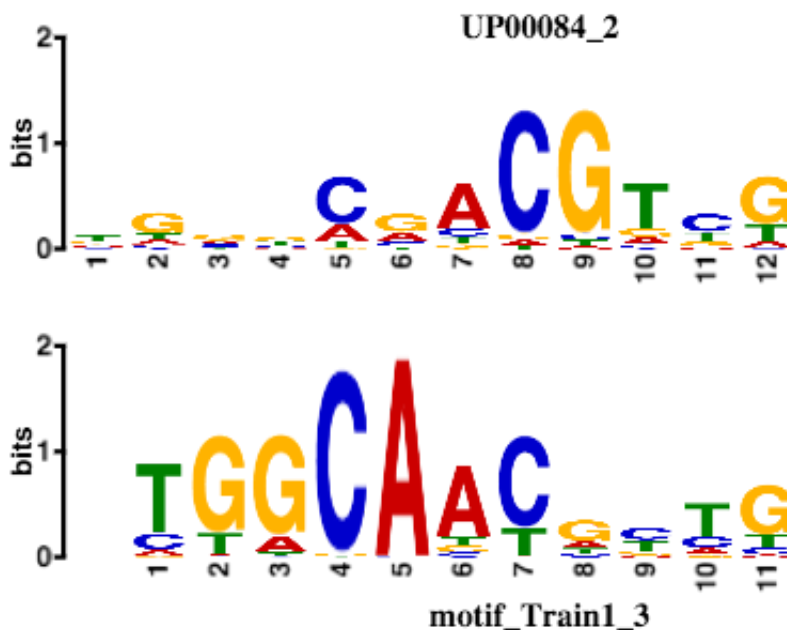
Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014
p-value 4.70e-04
E-value 4.60e-01
q-value 9.20e-01

Overlap 8
Offset 0
Orientation Reverse
 Complement

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

Name [UP00084_2](#)
[\(Gmeb1_secondary\)](#)
Database uniprobe mouse
p-value 2.77e-03
E-value 2.71e+00
q-value 1.00e+00

Overlap 12
Offset 1
Orientation Normal

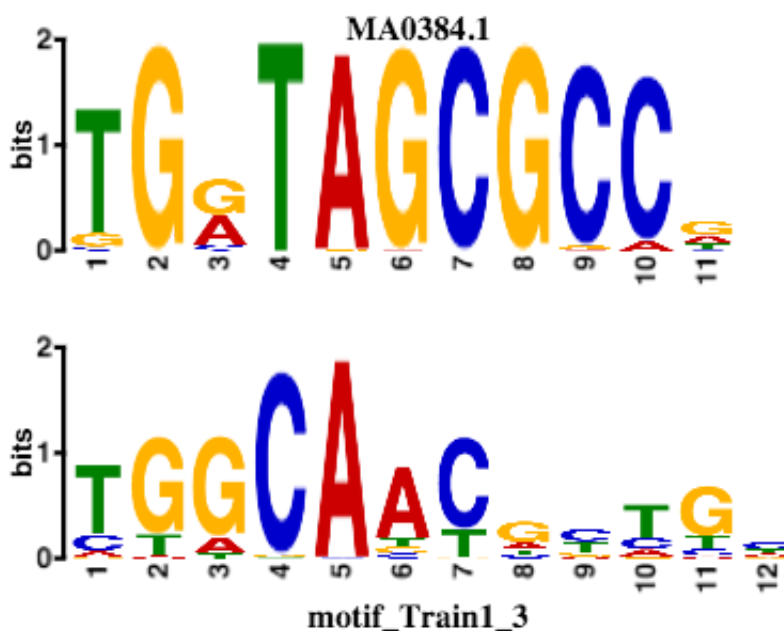
[Show logo download options](#)**Alignment****Summary**

Name [MA0384.1 \(SNT2\)](#)
Database JASPAR CORE 2014

Alignment

p-value	2.99e-03
E-value	2.92e+00
q-value	1.00e+00
Overlap	11
Offset	0
Orientation	Normal

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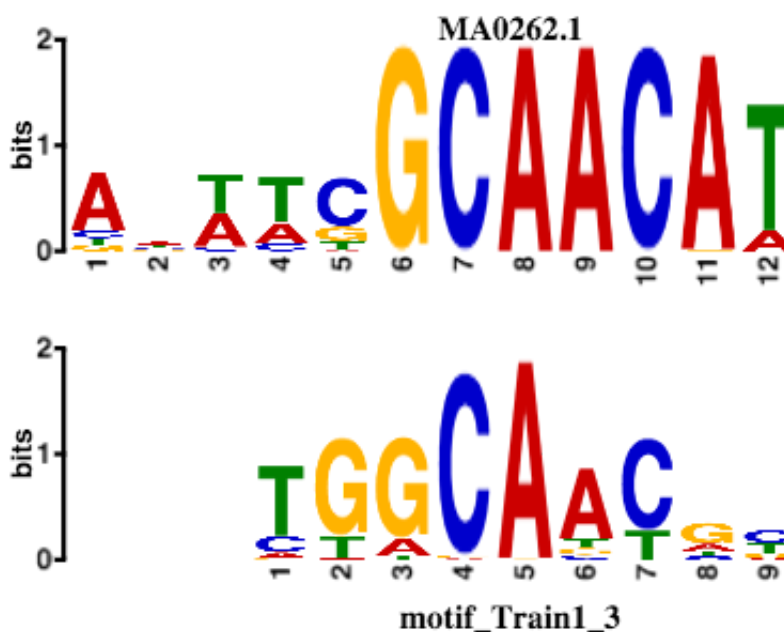
Summary

Name	MA0262.1 (mab-3)
Database	JASPAR CORE 2014
p-value	5.21e-03
E-value	5.10e+00
q-value	1.00e+00

Overlap	10
Offset	3
Orientation	Reverse Complement

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Alignment



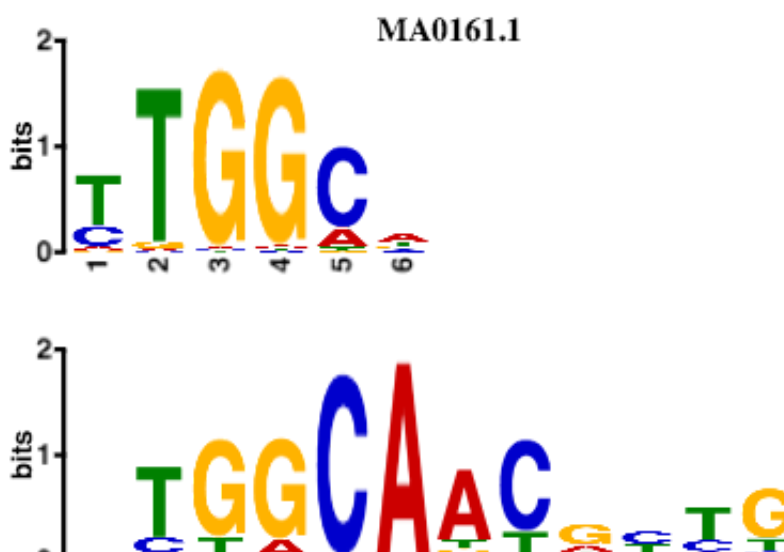
Summary

Name	MA0161.1 (NFIC)
Database	JASPAR CORE 2014
p-value	6.27e-03
E-value	6.14e+00
q-value	1.00e+00

Overlap	5
Offset	1
Orientation	Normal

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Alignment



1 2 3 4 5 6 7 8 9 10 11
motif_Train1_3

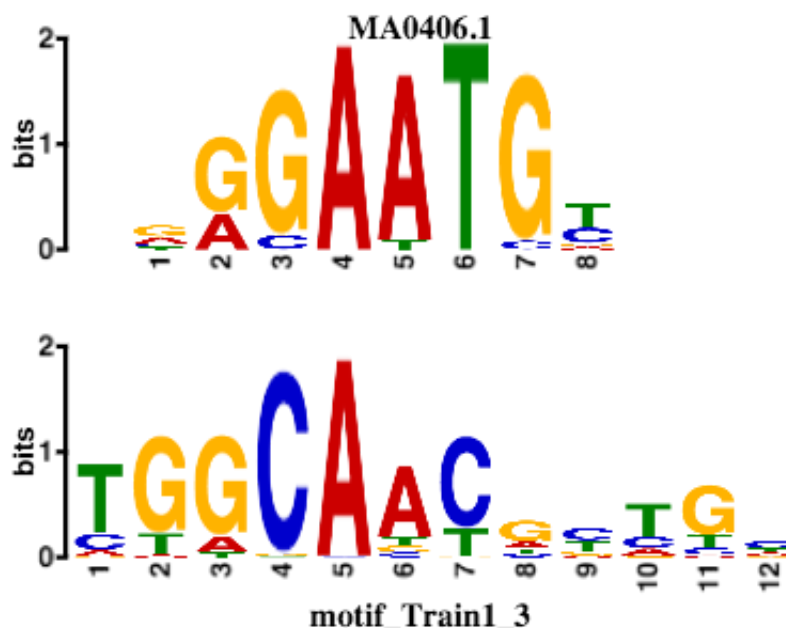
Summary

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014
p-value 6.28e-03
E-value 6.15e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse
 Complement

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Alignment



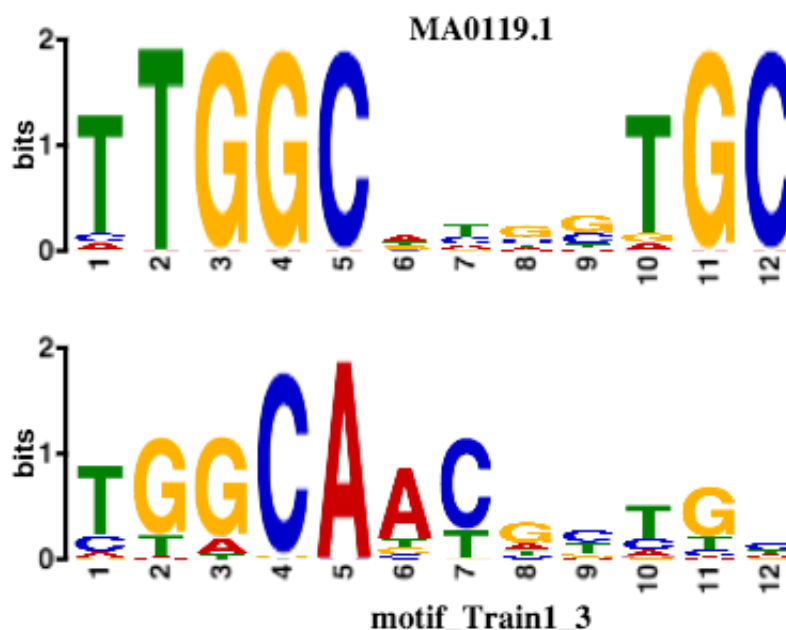
Summary

Name [MA0119.1 \(TLX1::NFIC\)](#)
Database JASPAR CORE 2014
p-value 7.97e-03
E-value 7.81e+00
q-value 1.00e+00

Overlap 12
Offset 0
Orientation Reverse
 Complement

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Alignment



Summary

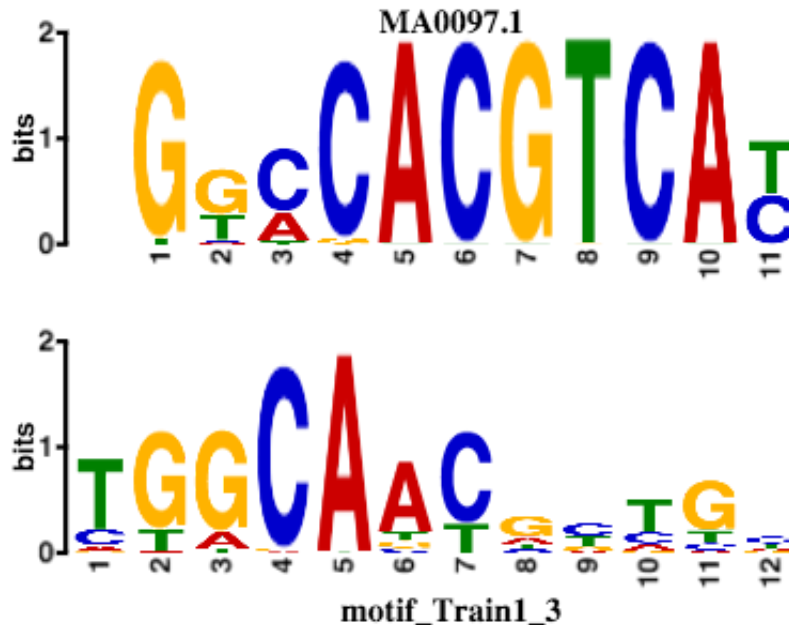
Name [MA0097.1 \(bZIP911\)](#)
Database JASPAR CORE 2014
p-value 9.93e-03
E-value 9.72e+00
q-value 1.00e+00

Overlap 11

Alignment

Offset -1
Orientation Reverse
 Complement

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MATCHES TO MOTIF_Train2_3 (4-TGGCAACGTTGC)

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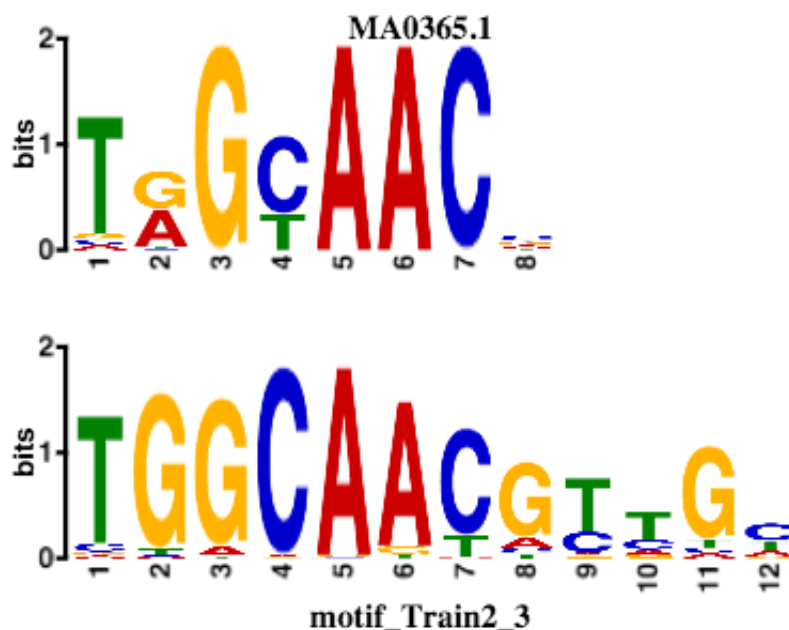
Summary

Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014
p-value 1.95e-04
E-value 1.91e-01
q-value 3.83e-01

Overlap 8
Offset 0
Orientation Reverse
 Complement

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Alignment



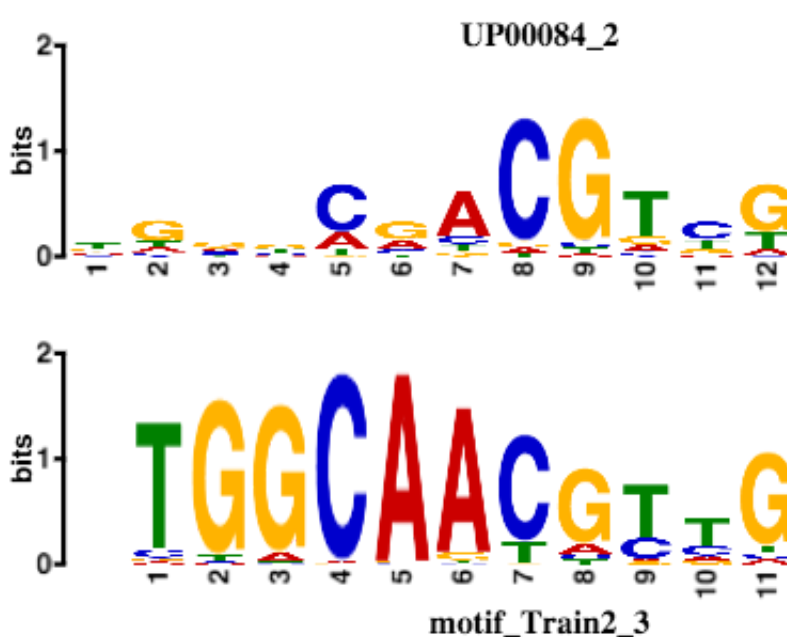
Summary

Name [UP00084_2](#)
[\(Gmeb1_secondary\)](#)
Database uniprobe mouse
p-value 7.46e-04

Alignment

E-value 7.30e-01
q-value 7.30e-01
Overlap 12
Offset 1
Orientation Normal

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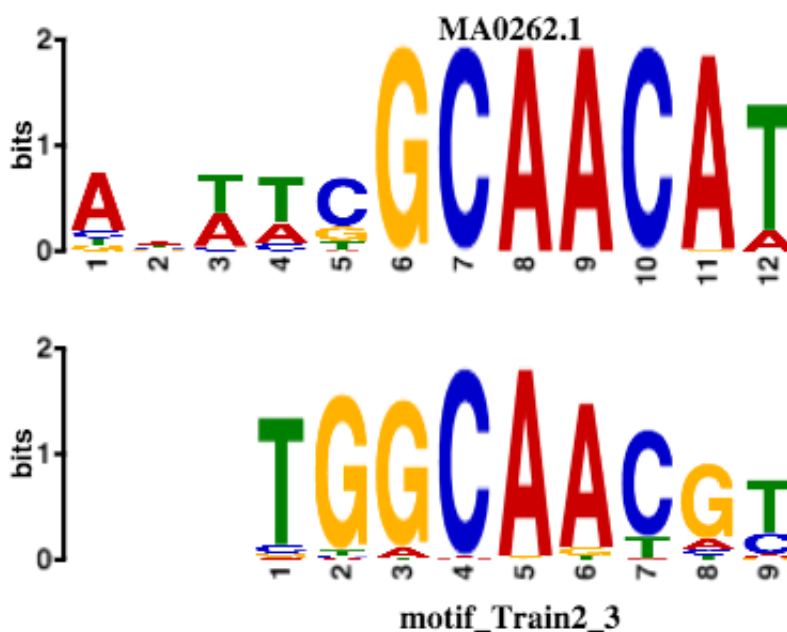
Summary

Name [MA0262.1 \(mab-3\)](#)
Database JASPAR CORE 2014
p-value 2.21e-03
E-value 2.16e+00
q-value 1.00e+00

Overlap 10
Offset 3
Orientation Reverse
 Complement

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Alignment



Summary

Name [MA0097.1 \(bZIP911\)](#)
Database JASPAR CORE 2014
p-value 3.82e-03
E-value 3.74e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Reverse
 Complement

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Alignment



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TOMTOM

1 2 3 4 5 6 7 8 9 10 11 12

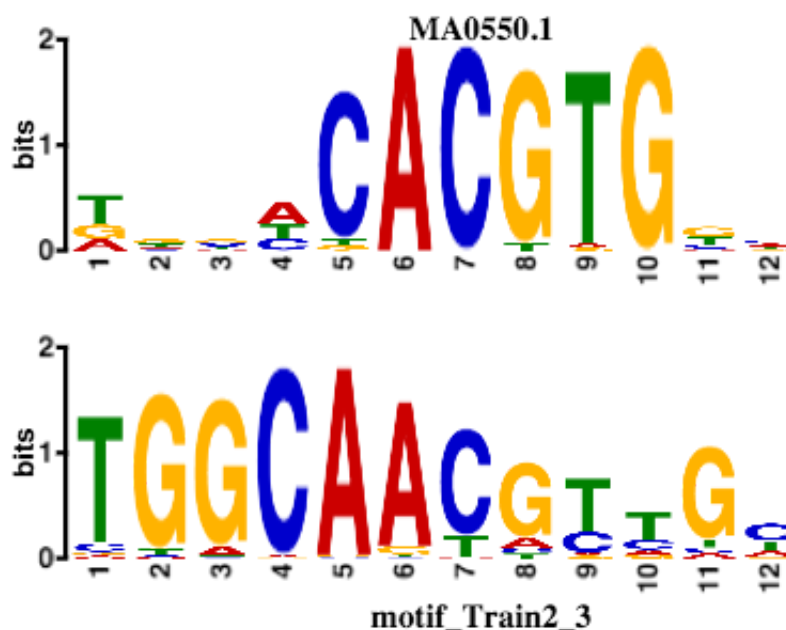
motif_Train2_3

Summary

Name	MA0550.1 (BZR1)
Database	JASPAR CORE 2014
p-value	4.31e-03
E-value	4.22e+00
q-value	1.00e+00
Overlap	12
Offset	0
Orientation	Reverse Complement

[Show logo download options](#)

Alignment

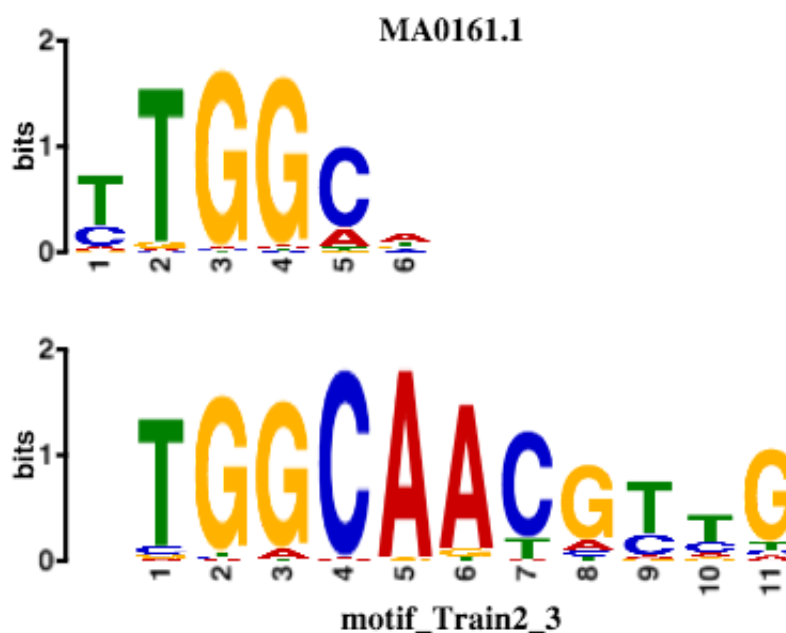


Summary

Name	MA0161.1 (NFIC)
Database	JASPAR CORE 2014
p-value	4.44e-03
E-value	4.35e+00
q-value	1.00e+00
Overlap	5
Offset	1
Orientation	Normal

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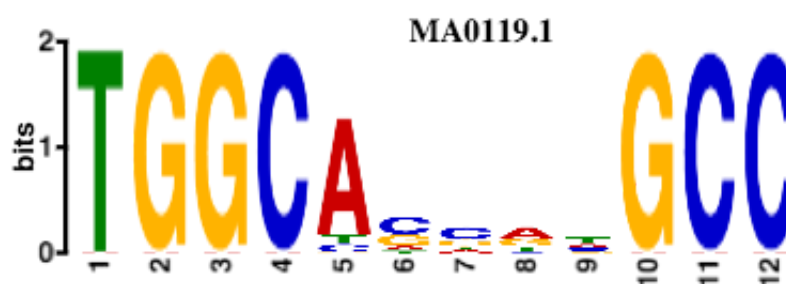
Alignment



Summary

Name	MA0119.1 (TLX1::NFIC)
Database	JASPAR CORE 2014
p-value	4.50e-03
E-value	4.40e+00
q-value	1.00e+00

Alignment



Overlap 12
Offset 0
Orientation Normal

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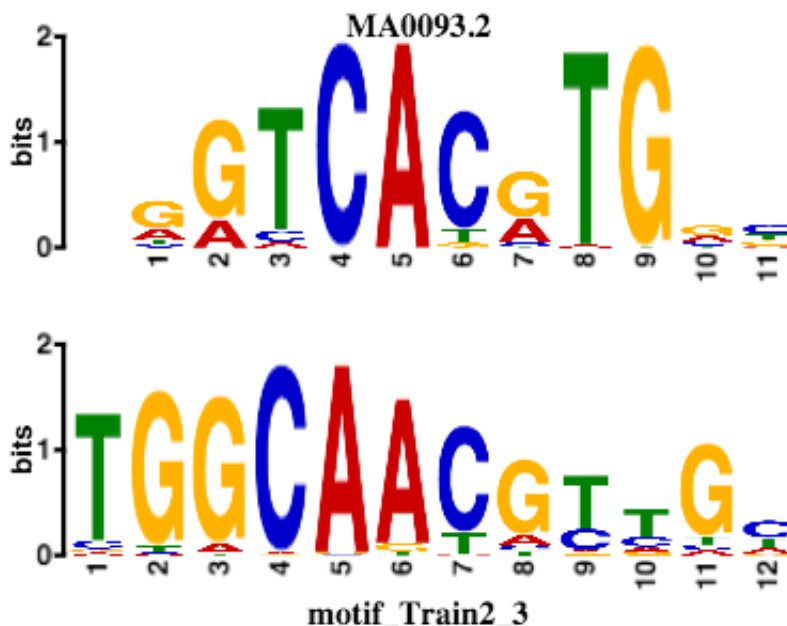
Summary

Name [MA0093.2 \(USF1\)](#)
Database JASPAR CORE 2014
p-value 5.37e-03
E-value 5.26e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Reverse Complement

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Alignment



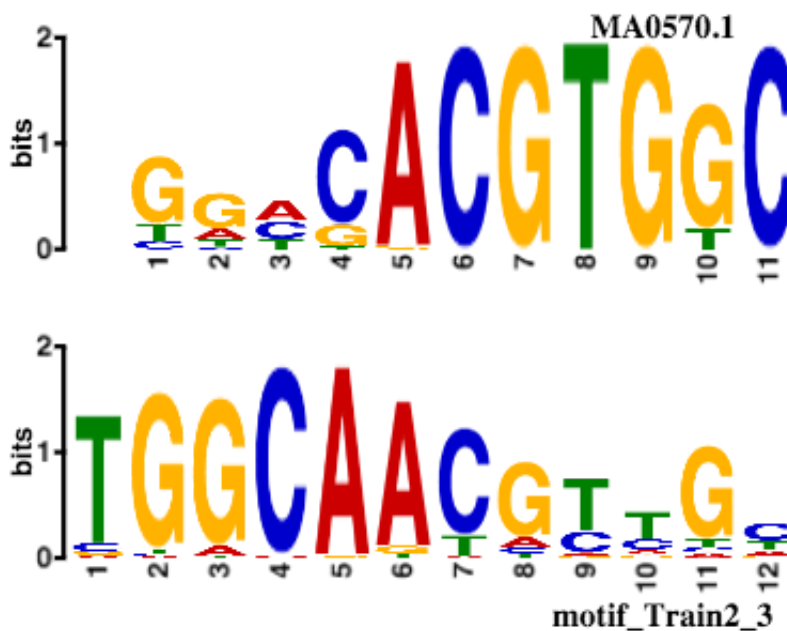
Summary

Name [MA0570.1 \(ABF1\)](#)
Database JASPAR CORE 2014
p-value 6.48e-03
E-value 6.35e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Normal

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Alignment



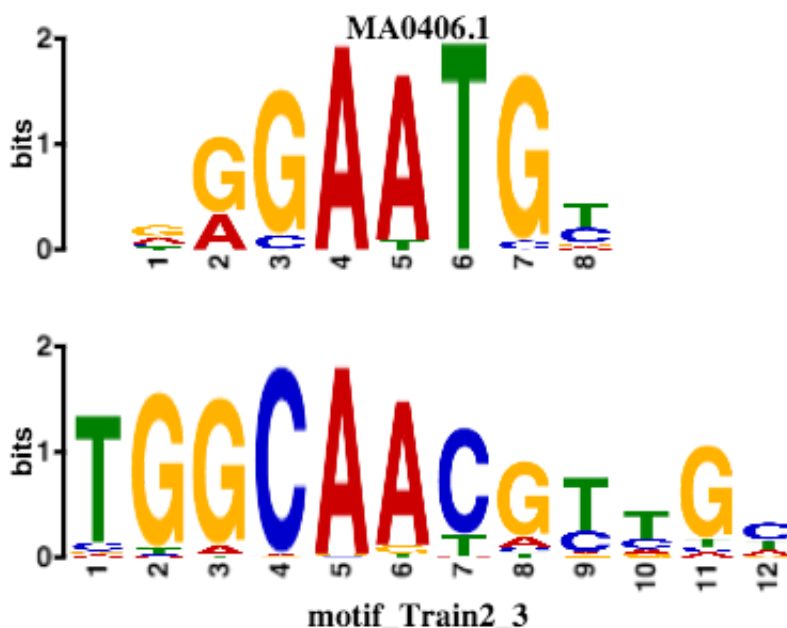
Summary

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014

Alignment

p-value	7.05e-03
E-value	6.90e+00
q-value	1.00e+00
Overlap	8
Offset	-1
Orientation	Reverse Complement

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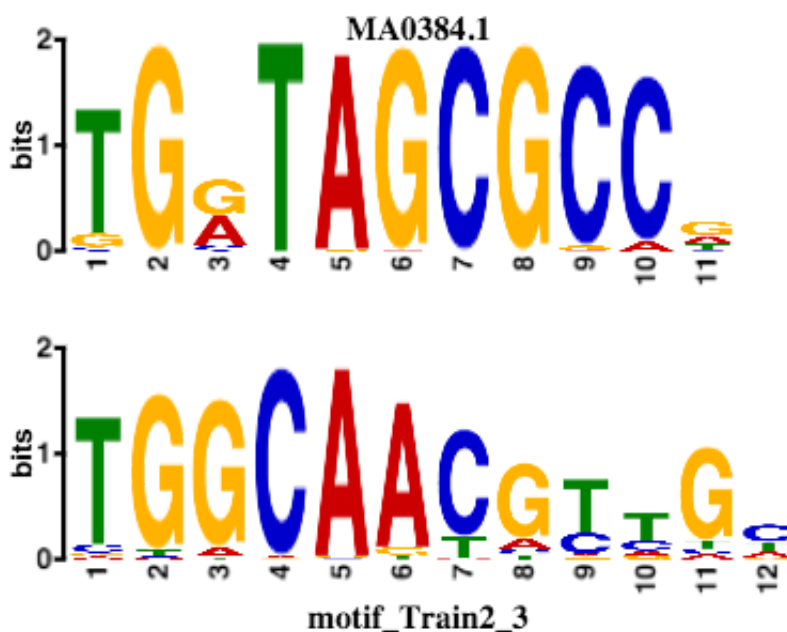
Summary

Name	MA0384.1 (SNT2)
Database	JASPAR CORE 2014
p-value	7.88e-03
E-value	7.71e+00
q-value	1.00e+00

Overlap	11
Offset	0
Orientation	Normal

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Alignment



Summary

Name	MA0567.1 (ERF1)
Database	JASPAR CORE 2014
p-value	9.59e-03

Alignment



E-value 9.39e+00
q-value 1.00e+00

Overlap 8
Offset 0
Orientation Reverse Complement

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MATCHES TO MOTIF_Train7_3 (5-TGGCAACGYTGY)

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Summary

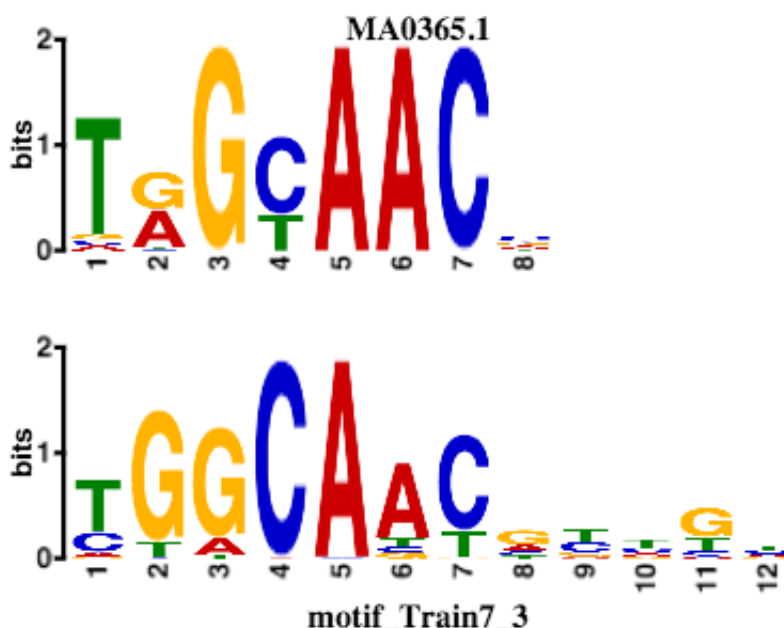
Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014

p-value 2.83e-04
E-value 2.77e-01
q-value 5.18e-01

Overlap 8
Offset 0
Orientation Reverse Complement

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Alignment



Summary

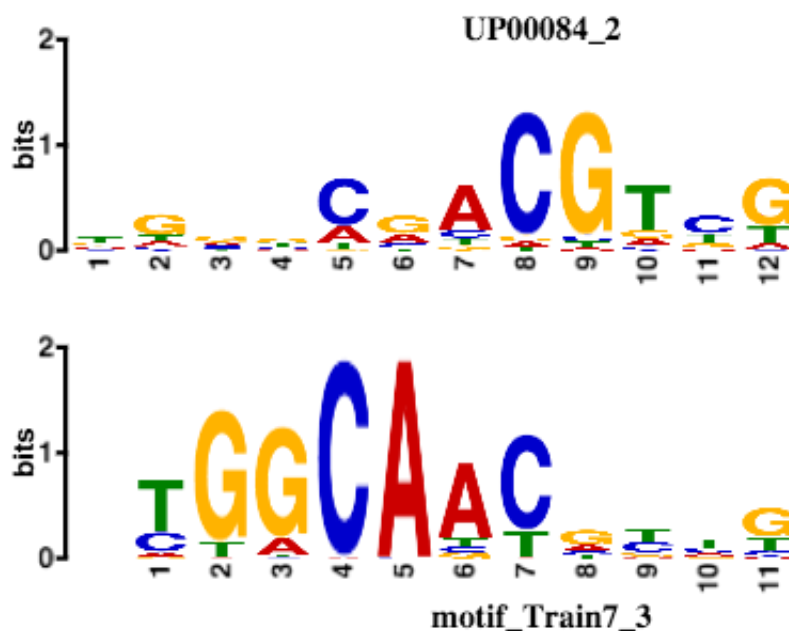
Name [UP00084_2](#)
Database [\(Gmeb1_secondary\)](#)

p-value 5.29e-04
E-value 5.18e-01
q-value 5.18e-01

Overlap 12
Offset 1
Orientation Normal

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Alignment

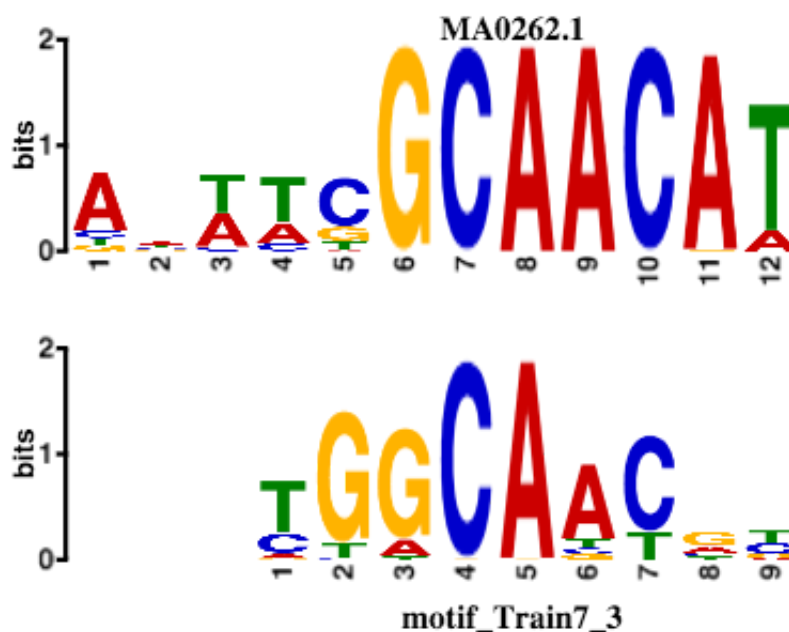


Summary

Name [MA0262.1 \(mab-3\)](#)
Database JASPAR CORE 2014
p-value 2.22e-03
E-value 2.18e+00
q-value 1.00e+00

Overlap 10
Offset 3
Orientation Reverse
 Complement
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Alignment

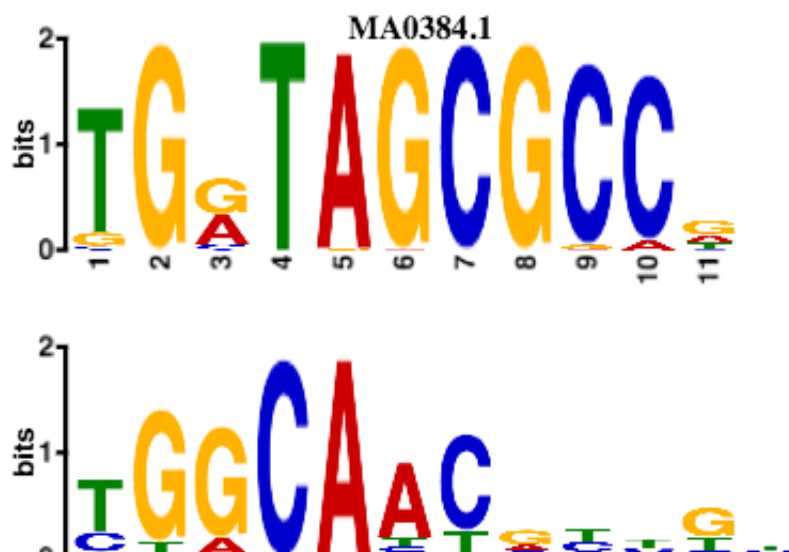


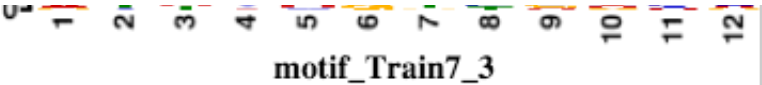
Summary

Name [MA0384.1 \(SNT2\)](#)
Database JASPAR CORE 2014
p-value 3.58e-03
E-value 3.51e+00
q-value 1.00e+00

Overlap 11
Offset 0
Orientation Normal
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Alignment

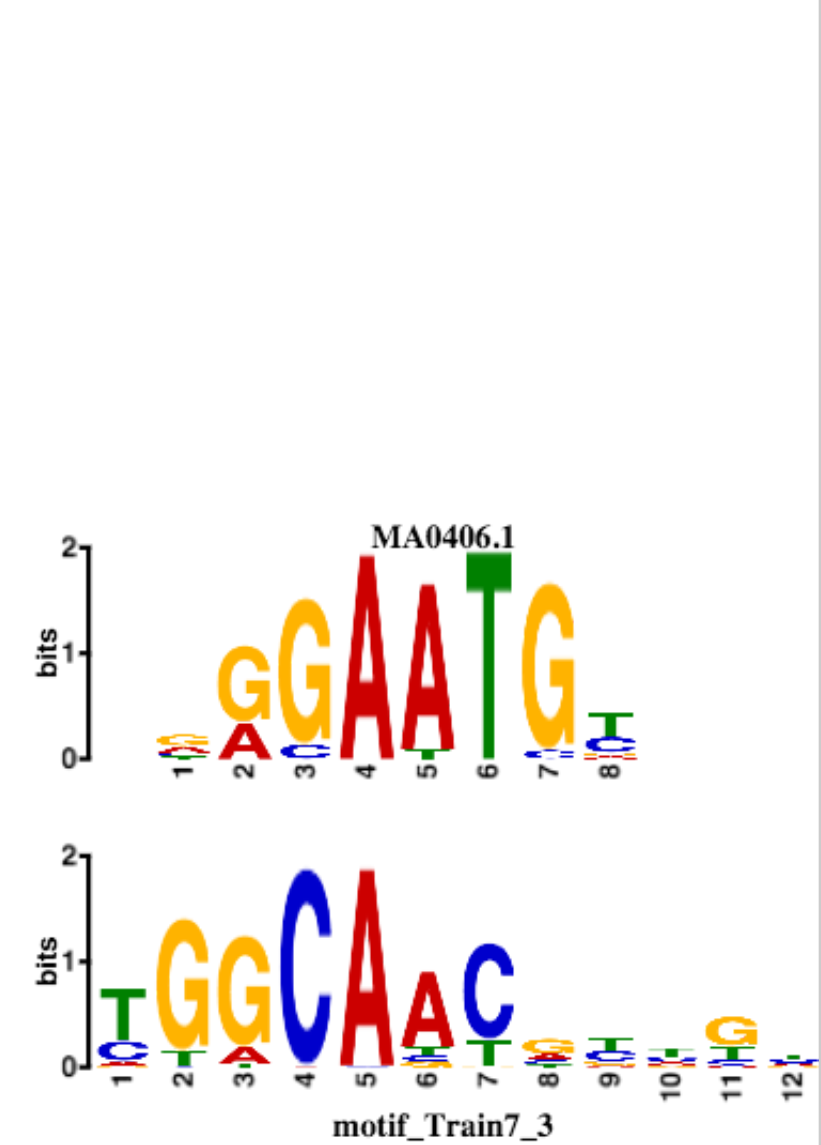




Summary

Alignment

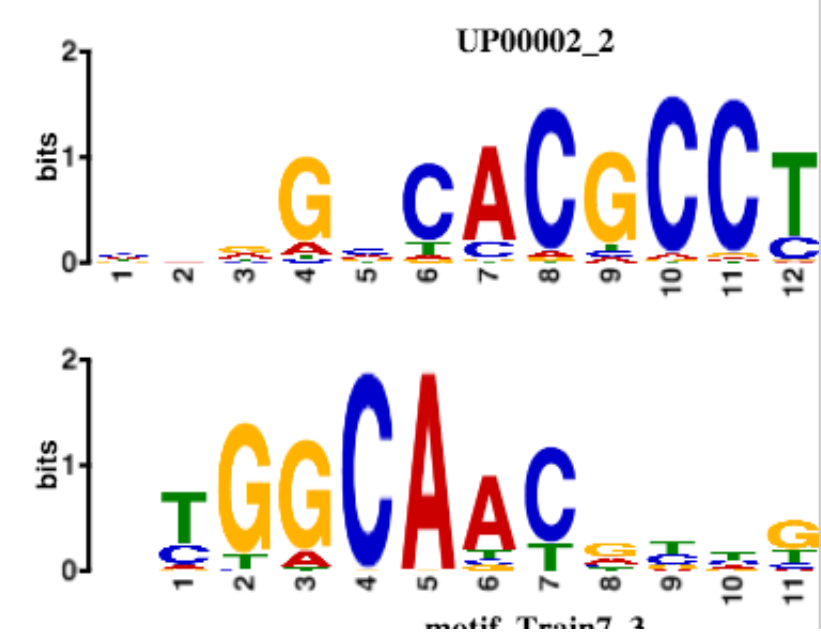
Name	MA0406.1 (TEC1)
Database	JASPAR CORE 2014
p-value	3.88e-03
E-value	3.80e+00
q-value	1.00e+00
Overlap	8
Offset	-1
Orientation	Reverse Complement
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Summary

Alignment

Name	UP00002_2 (Sp4 secondary)
Database	uniprobe mouse
p-value	5.38e-03
E-value	5.26e+00
q-value	1.00e+00
Overlap	12
Offset	1
Orientation	Reverse Complement
Show logo download options	

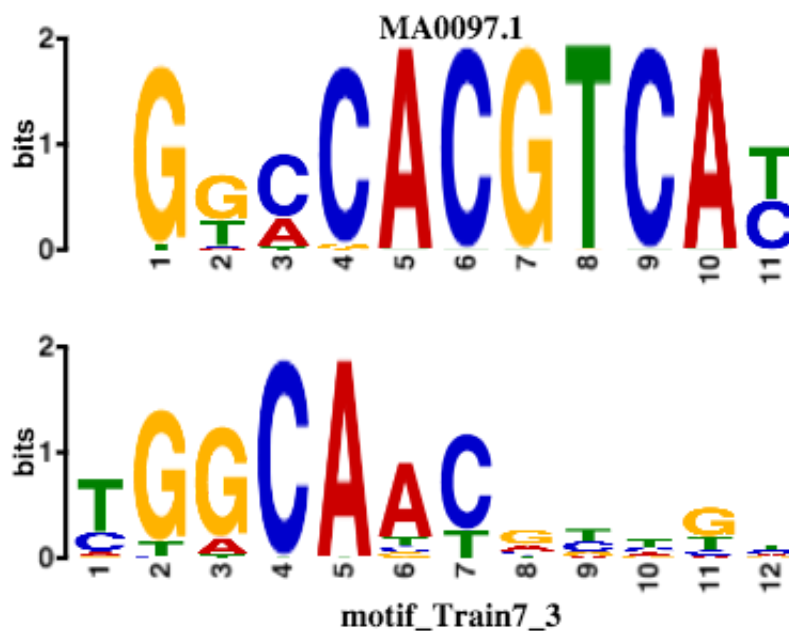


Summary

Name	MA0097.1 (bZIP911)
Database	JASPAR CORE 2014
p-value	6.44e-03
E-value	6.31e+00
q-value	1.00e+00
Overlap	11
Offset	-1
Orientation	Reverse Complement

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Alignment

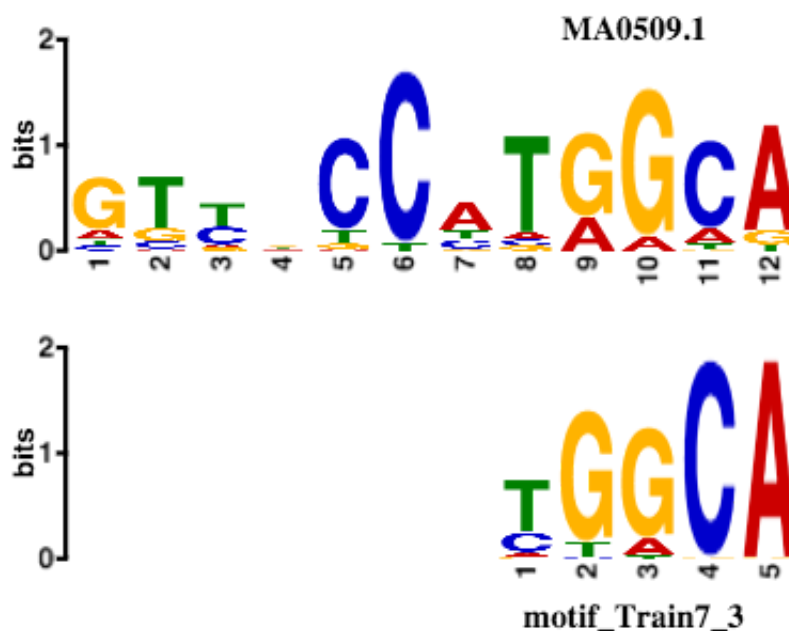


Summary

Name	MA0509.1 (Rfx1)
Database	JASPAR CORE 2014
p-value	6.49e-03
E-value	6.36e+00
q-value	1.00e+00
Overlap	7
Offset	7
Orientation	Normal

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Alignment



Summary

Name	MA0161.1 (NFIC)
Database	JASPAR CORE 2014

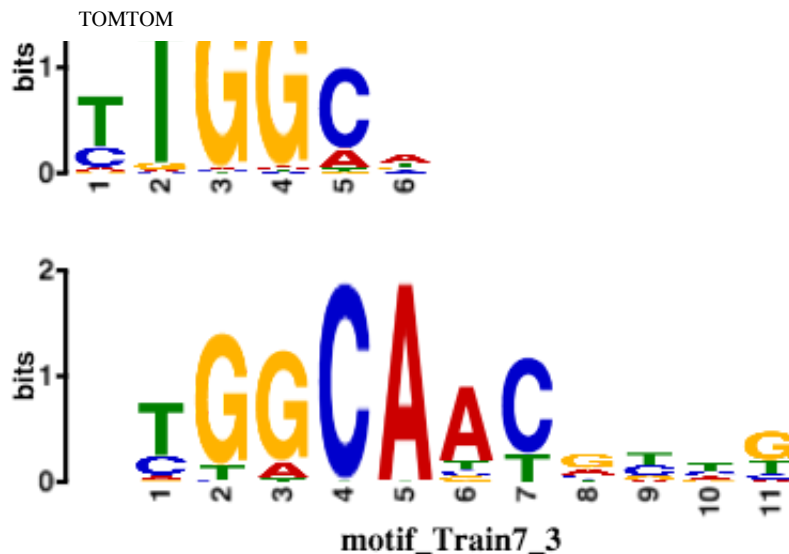
Alignment



p-value 7.82e-03
E-value 7.66e+00
q-value 1.00e+00

Overlap 5
Offset 1
Orientation Normal

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MATCHES TO MOTIF_TRAIN8_3 (5-TGGCAACGYTGY)

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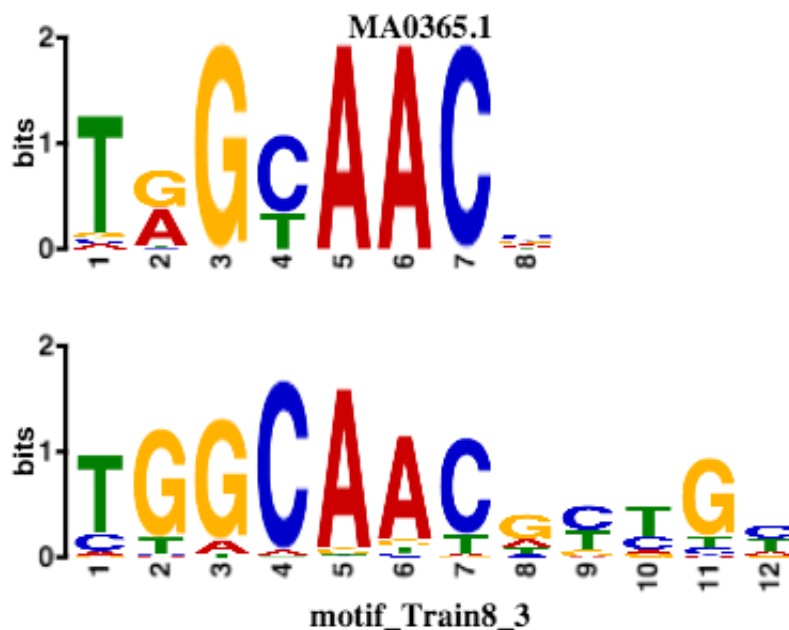
Summary

Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014
p-value 4.36e-04
E-value 4.27e-01
q-value 8.55e-01

Overlap 8
Offset 0
Orientation Reverse Complement

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Alignment



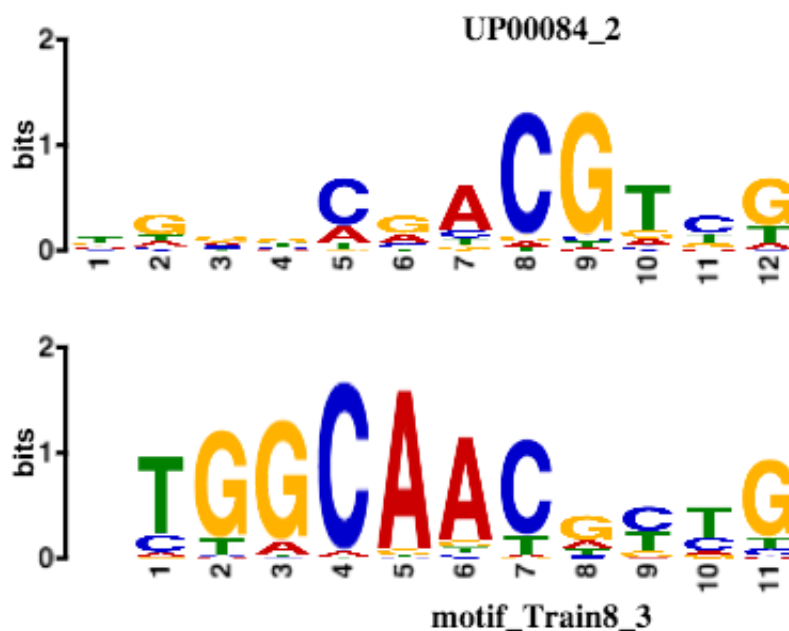
Summary

Name [UP00084_2 \(Gmeb1 secondary\)](#)
Database uniprobe mouse
p-value 1.16e-03
E-value 1.13e+00
q-value 1.00e+00

Overlap 12
Offset 1
Orientation Normal

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Alignment



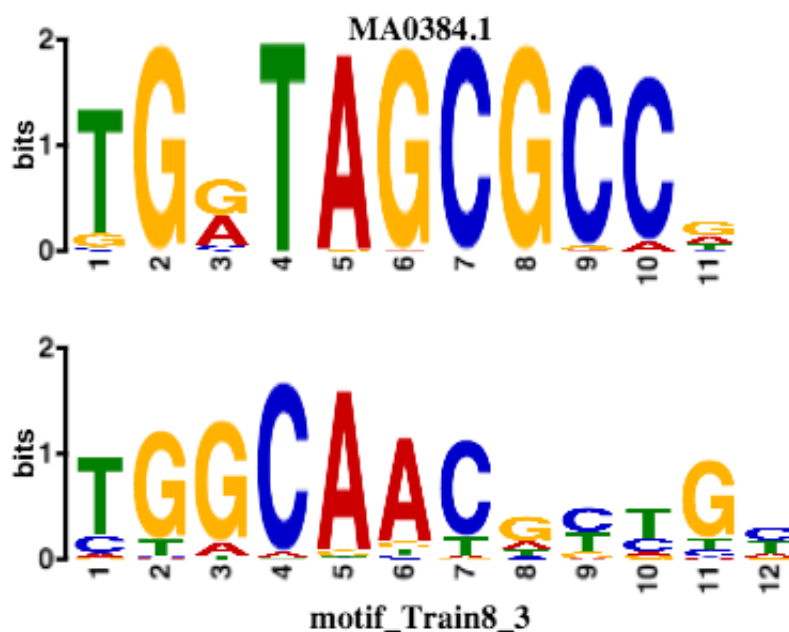
Summary

Name [MA0384.1 \(SNT2\)](#)
Database JASPAR CORE 2014
p-value 1.66e-03
E-value 1.63e+00
q-value 1.00e+00

Overlap 11
Offset 0
Orientation Normal

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Alignment



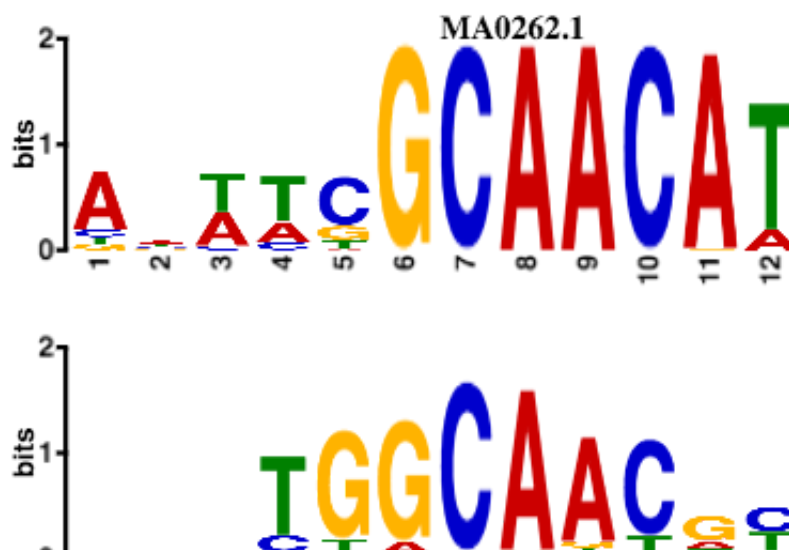
Summary

Name [MA0262.1 \(mab-3\)](#)
Database JASPAR CORE 2014
p-value 4.77e-03
E-value 4.67e+00
q-value 1.00e+00

Overlap 10
Offset 3
Orientation Reverse Complement

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Alignment



1 2 3 4 5 6 7 8 9
motif_Train8_3

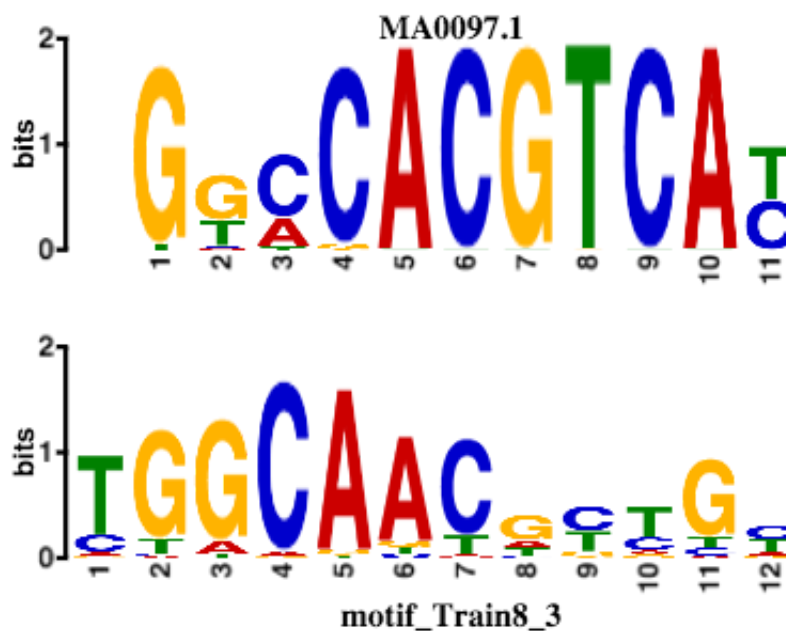
Summary

Name [MA0097.1 \(bZIP911\)](#)
Database JASPAR CORE 2014
p-value 5.93e-03
E-value 5.80e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Reverse Complement

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Alignment



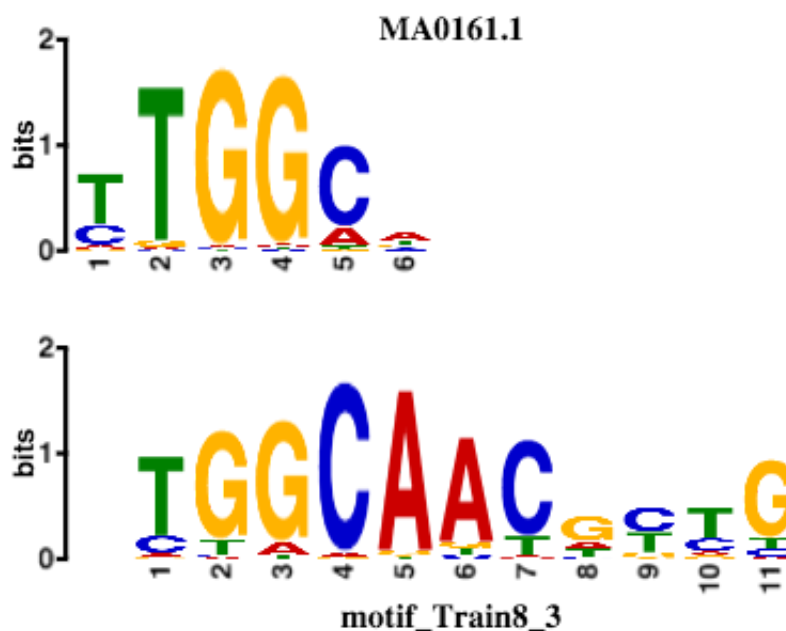
Summary

Name [MA0161.1 \(NFIC\)](#)
Database JASPAR CORE 2014
p-value 5.96e-03
E-value 5.84e+00
q-value 1.00e+00

Overlap 5
Offset 1
Orientation Normal

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Alignment



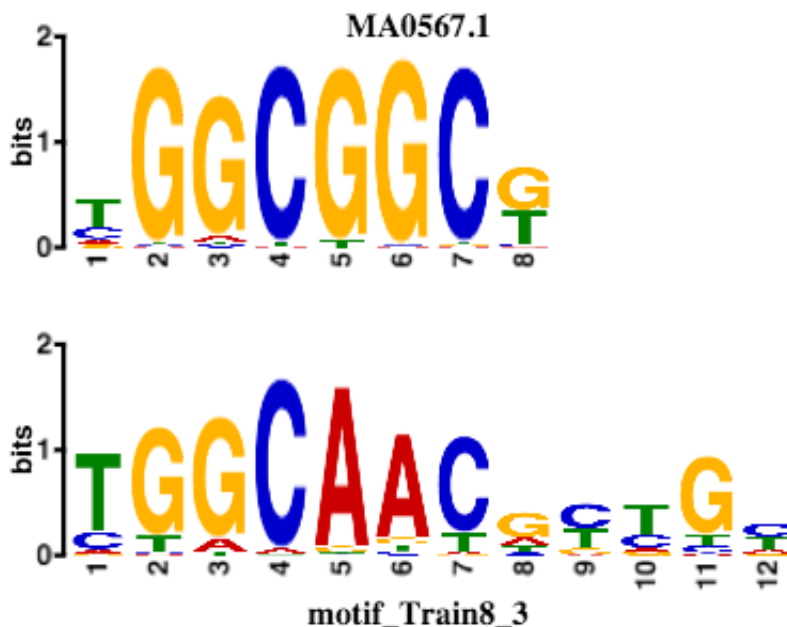
Summary

Alignment

Name [MA0567.1 \(ERF1\)](#)
Database JASPAR CORE 2014
p-value 8.42e-03
E-value 8.24e+00
q-value 1.00e+00

Overlap 8
Offset 0
Orientation Reverse
 Complement

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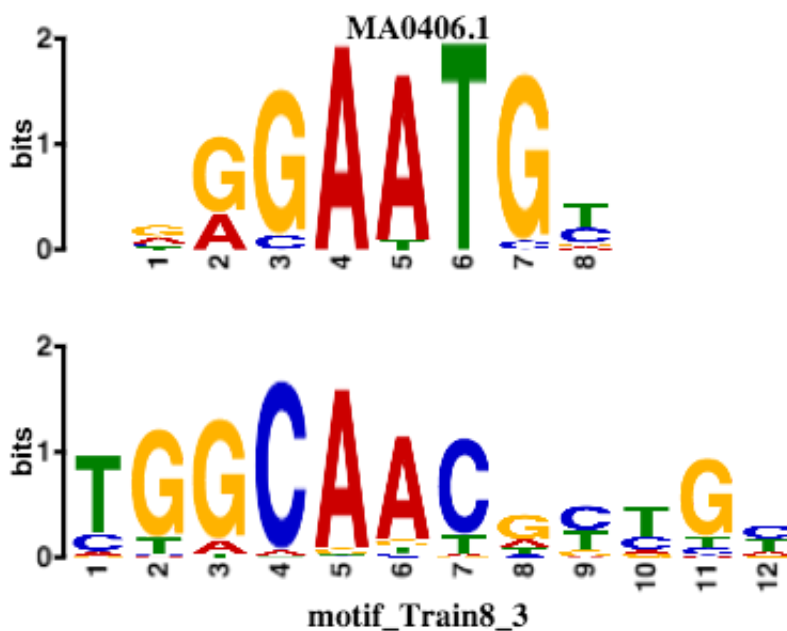
Summary

Alignment

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014
p-value 9.05e-03
E-value 8.86e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse
 Complement

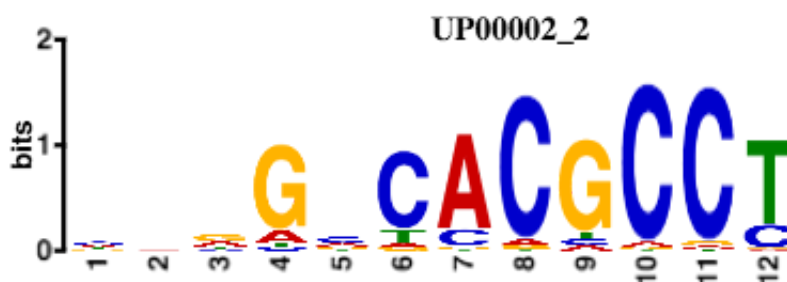
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Summary

Alignment

Name [UP00002_2](#)
Database [\(Sp4 secondary\)](#)
 uniprobe mouse
p-value 9.47e-03
E-value 9.27e+00
q-value 1.00e+00



Overlap 12
Offset 1
Orientation Reverse
 Complement

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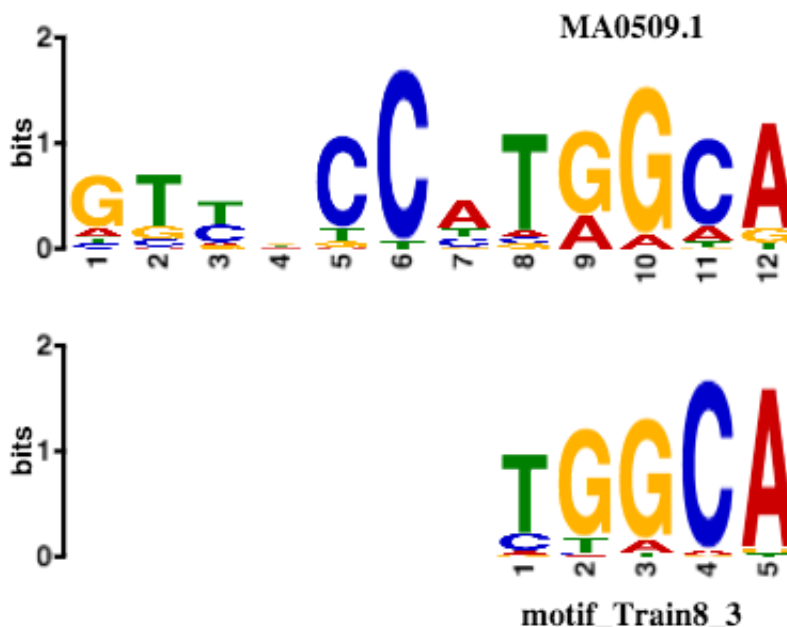
Summary

Name [MA0509.1 \(Rfx1\)](#)
Database JASPAR CORE 2014
p-value 9.81e-03
E-value 9.60e+00
q-value 1.00e+00

Overlap 7
Offset 7
Orientation Normal

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Alignment



MATCHES TO MOTIF_Train4_3 (4-TGGCAACGYTGH)

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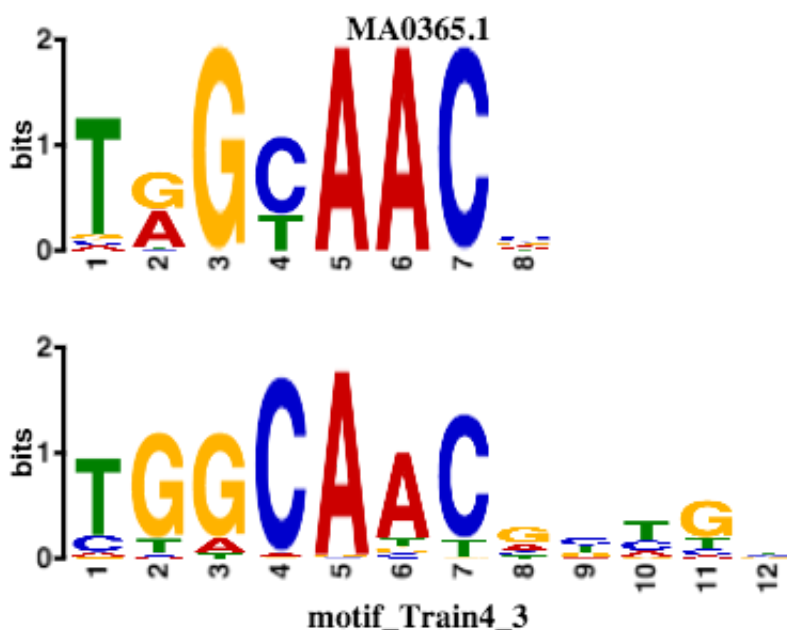
Summary

Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014
p-value 2.58e-04
E-value 2.53e-01
q-value 5.05e-01

Overlap 8
Offset 0
Orientation Reverse
 Complement

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Alignment



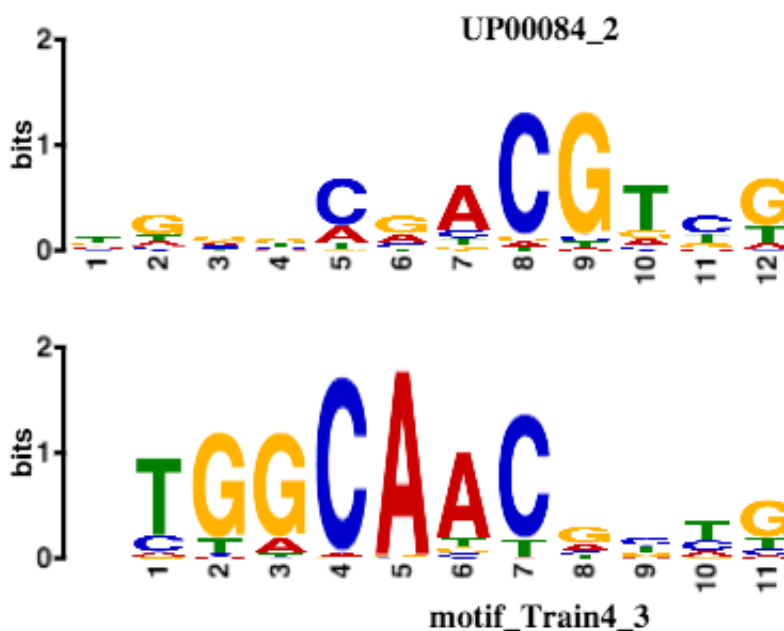
Summary

Alignment

Name [UP00084_2](#)
Database [\(Gmeh1_secondary\)](#)
Database uniprobe mouse
p-value 8.25e-04
E-value 8.07e-01
q-value 8.07e-01

Overlap 12
Offset 1
Orientation Normal

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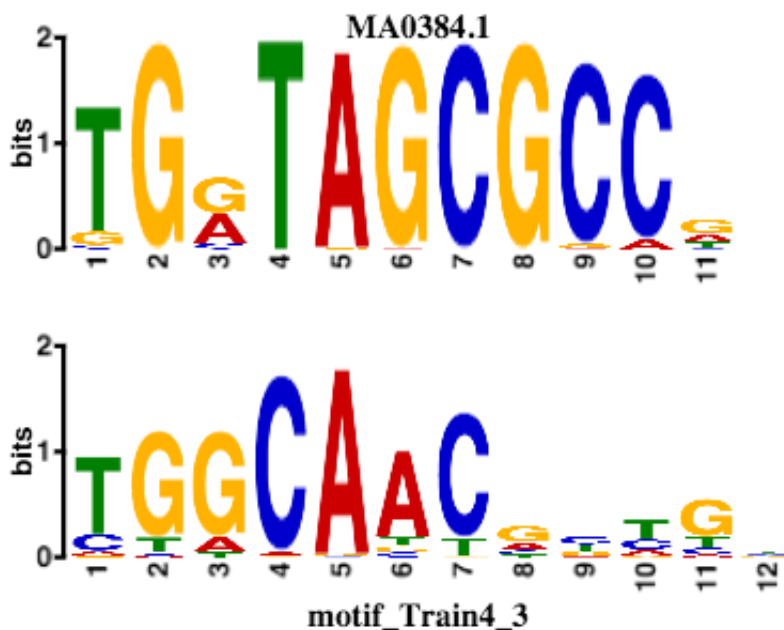
Summary

Name [MA0384.1 \(SNT2\)](#)
Database JASPAR CORE 2014
p-value 1.96e-03
E-value 1.91e+00
q-value 1.00e+00

Overlap 11
Offset 0
Orientation Normal

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Alignment



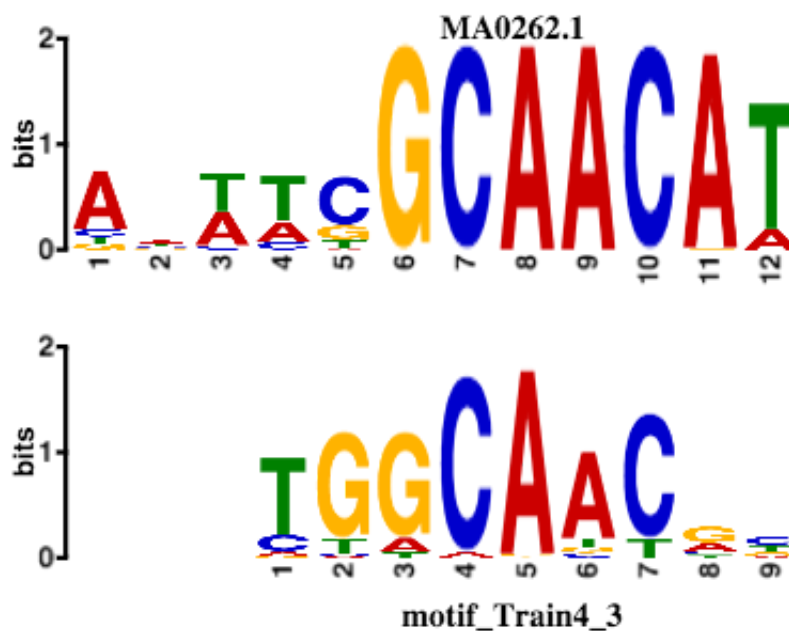
Summary

Name [MA0262.1 \(mab-3\)](#)
Database JASPAR CORE 2014
p-value 2.11e-03
E-value 2.07e+00
q-value 1.00e+00

Overlap 10
Offset 3
Orientation Reverse
 Complement

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Alignment



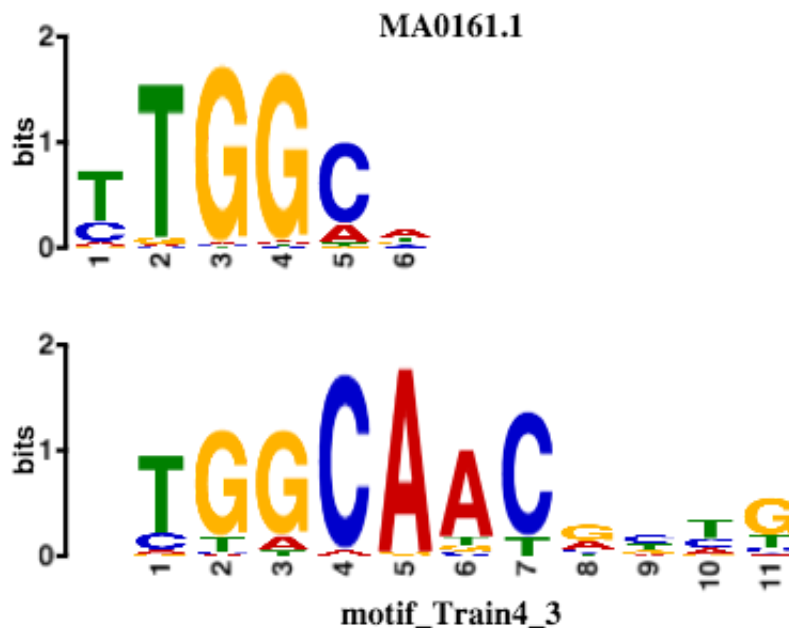
Summary

Name	MA0161.1 (NFIC)
Database	JASPAR CORE 2014
p-value	5.03e-03
E-value	4.92e+00
q-value	1.00e+00

Overlap	5
Offset	1
Orientation	Normal

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Alignment

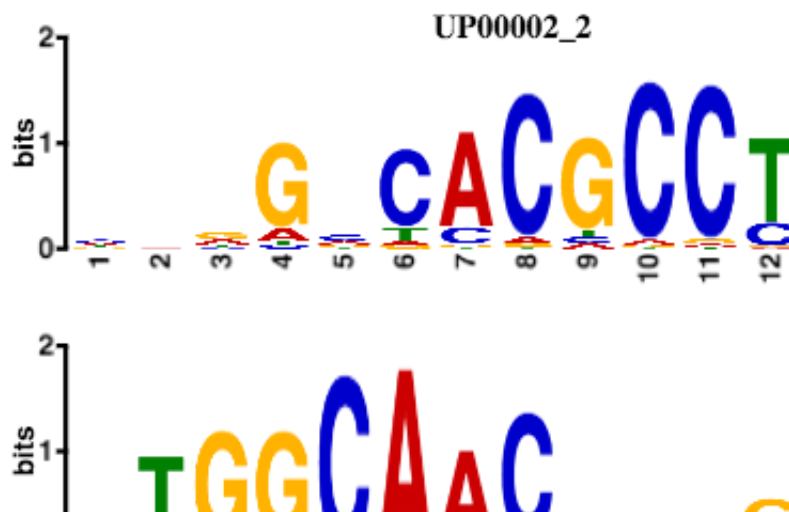


Summary

Name	UP00002_2 (Sp4_secondary)
Database	uniprobe mouse
p-value	5.87e-03
E-value	5.75e+00
q-value	1.00e+00

Overlap	12
Offset	1
Orientation	Reverse Complement

Alignment



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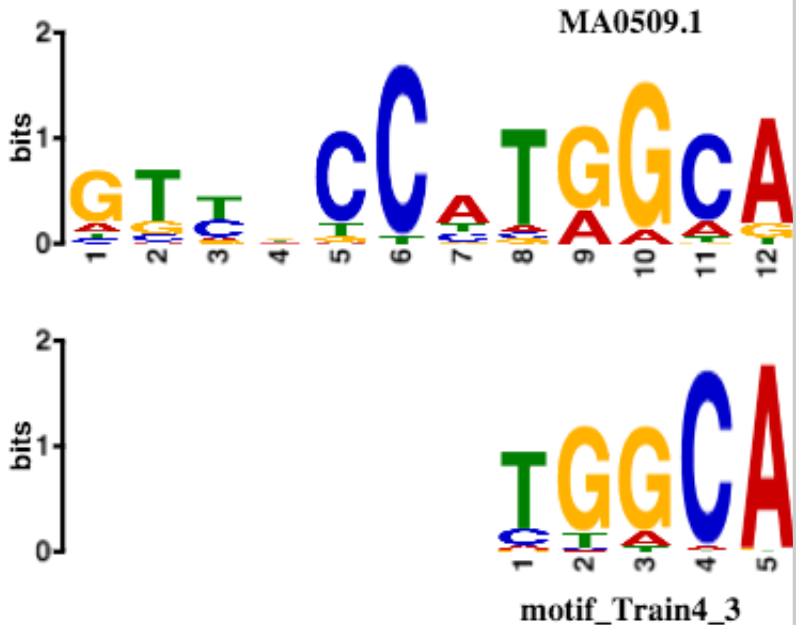
Summary

Alignment

Name	MA0509.1 (Rfx1)
Database	JASPAR CORE 2014
<i>p</i> -value	6.58e-03
<i>E</i> -value	6.44e+00
<i>q</i> -value	1.00e+00
Overlap	7
Offset	7

Orientation Normal

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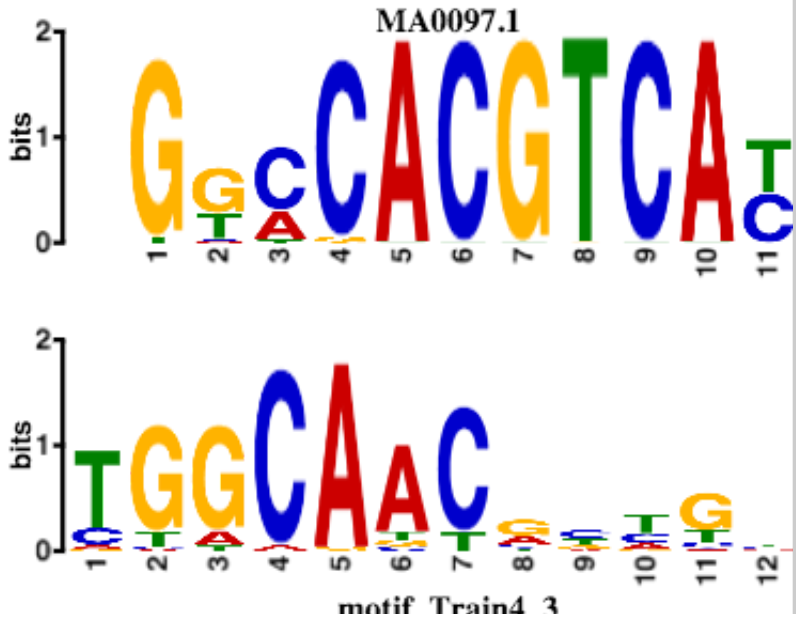


Summary

Alignment

Name	MA0097.1 (bZIP911)
Database	JASPAR CORE 2014
<i>p</i> -value	7.54e-03
<i>E</i> -value	7.38e+00
<i>q</i> -value	1.00e+00
Overlap	11
Offset	-1
Orientation	Reverse Complement

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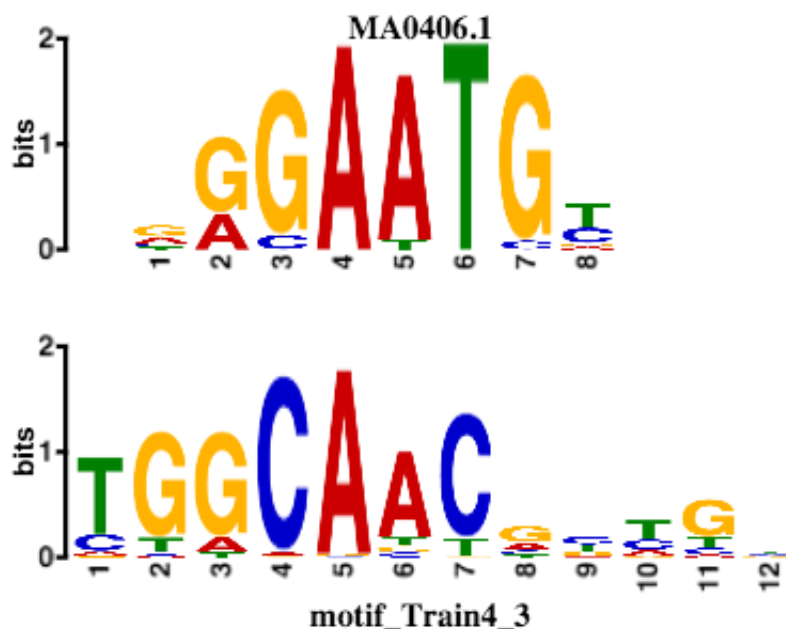
Summary

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014
p-value 9.55e-03
E-value 9.35e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse
 Complement

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Alignment



MATCHES TO MOTIF_10_3 (4-TGGCAACGYTGY)

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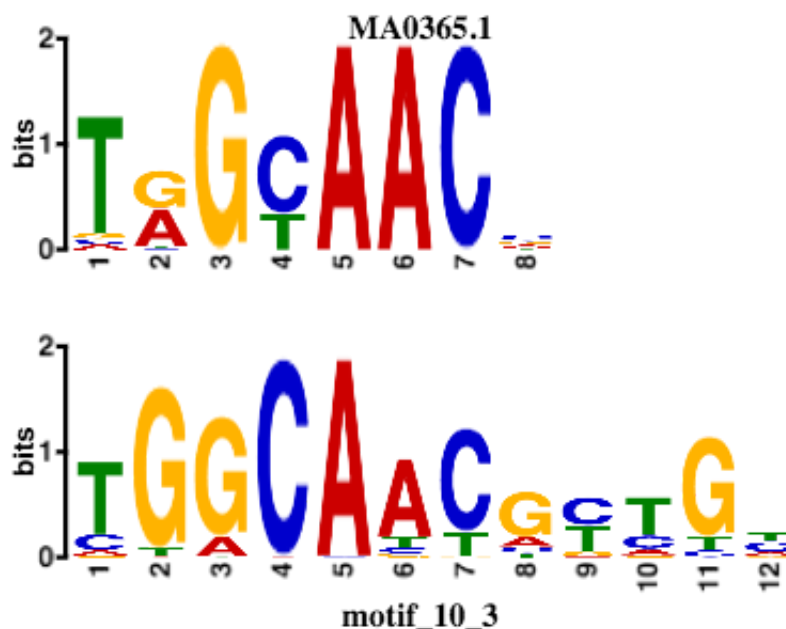
Summary

Name [MA0365.1 \(RFX1\)](#)
Database JASPAR CORE 2014
p-value 1.88e-04
E-value 1.84e-01
q-value 3.68e-01

Overlap 8
Offset 0
Orientation Reverse
 Complement

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Alignment



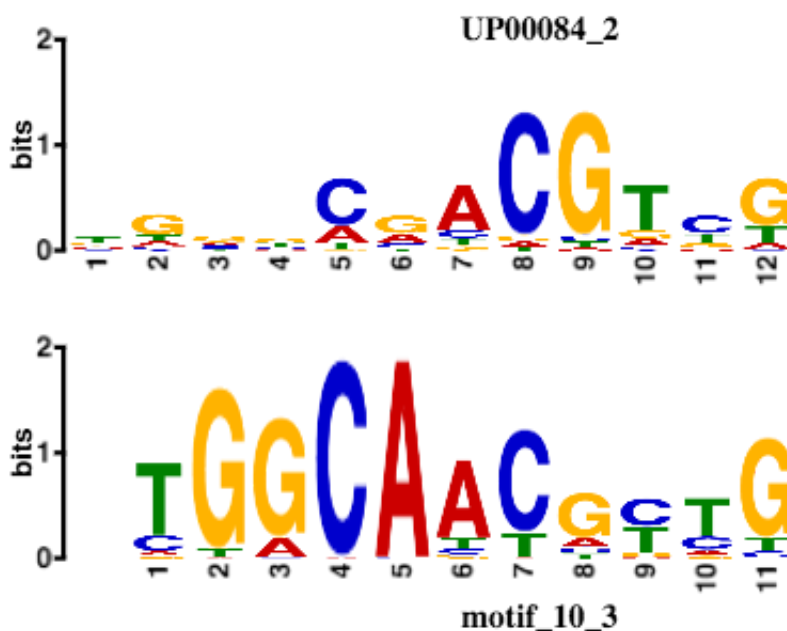
Summary

Name [UP00084_2](#)
[\(Gmeb1_secondary\)](#)
Database uniprobe mouse
p-value 7.73e-04
E-value 7.57e-01
q-value 7.57e-01

Alignment

Overlap 12
Offset 1
Orientation Normal

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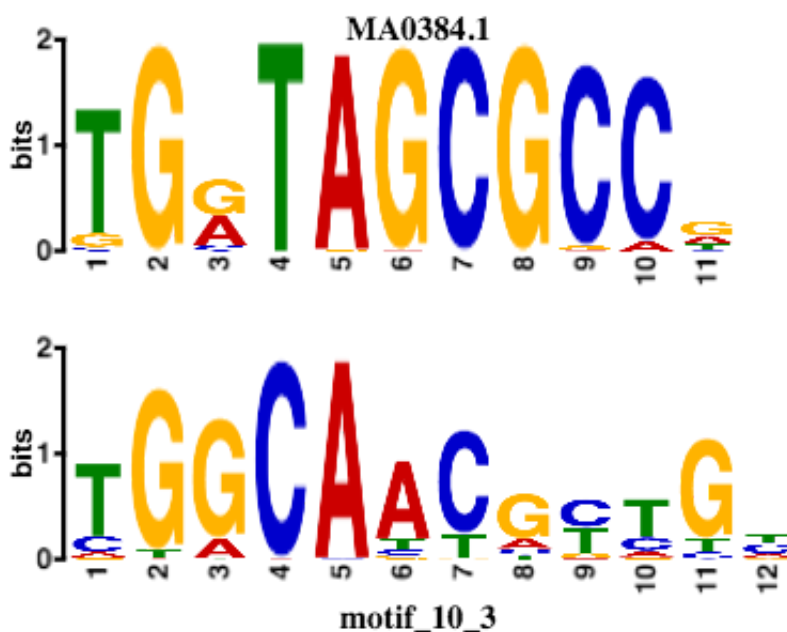
Summary

Name [MA0384.1 \(SNT2\)](#)
Database JASPAR CORE 2014
p-value 2.53e-03
E-value 2.47e+00
q-value 1.00e+00

Overlap 11
Offset 0
Orientation Normal

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Alignment



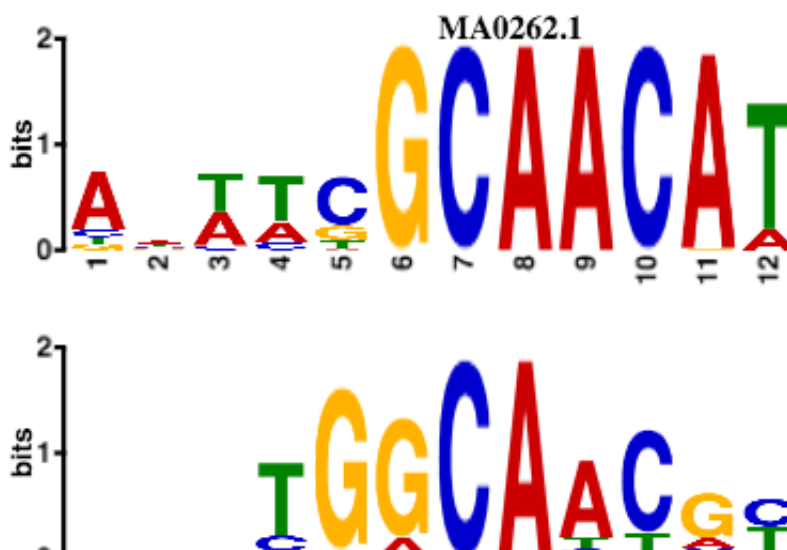
Summary

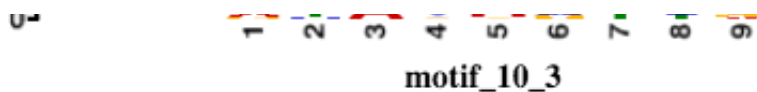
Name [MA0262.1 \(mab-3\)](#)
Database JASPAR CORE 2014
p-value 3.35e-03
E-value 3.27e+00
q-value 1.00e+00

Overlap 10
Offset 3
Orientation Reverse Complement

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Alignment





Summary

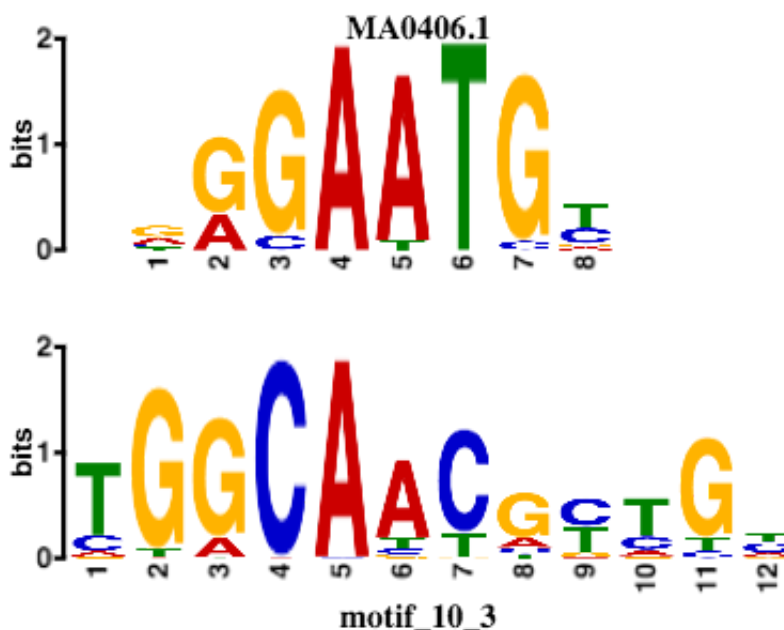
Alignment

Name [MA0406.1 \(TEC1\)](#)
Database JASPAR CORE 2014

p-value 4.84e-03
E-value 4.74e+00
q-value 1.00e+00

Overlap 8
Offset -1
Orientation Reverse Complement

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Summary

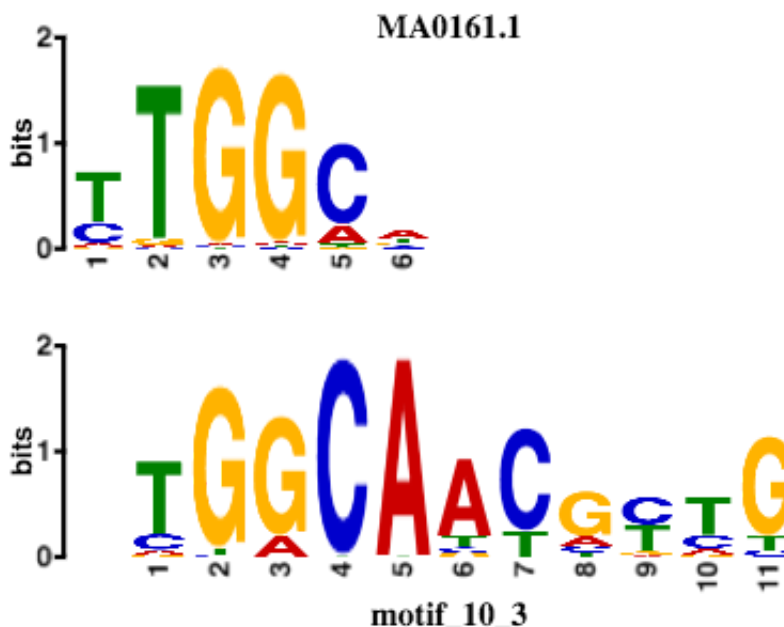
Alignment

Name [MA0161.1 \(NFIC\)](#)
Database JASPAR CORE 2014

p-value 6.05e-03
E-value 5.93e+00
q-value 1.00e+00

Overlap 5
Offset 1
Orientation Normal

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Summary

Alignment

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

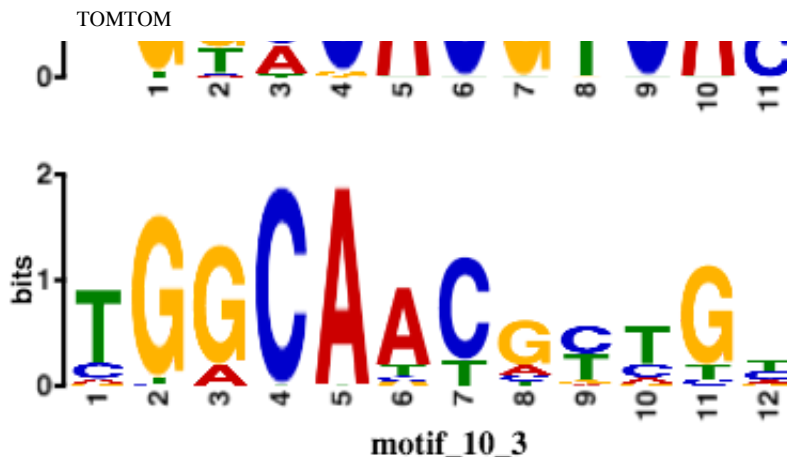
p-value 6.38e-03



p-value 6.58e-03
E-value 6.25e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Reverse Complement

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Summary

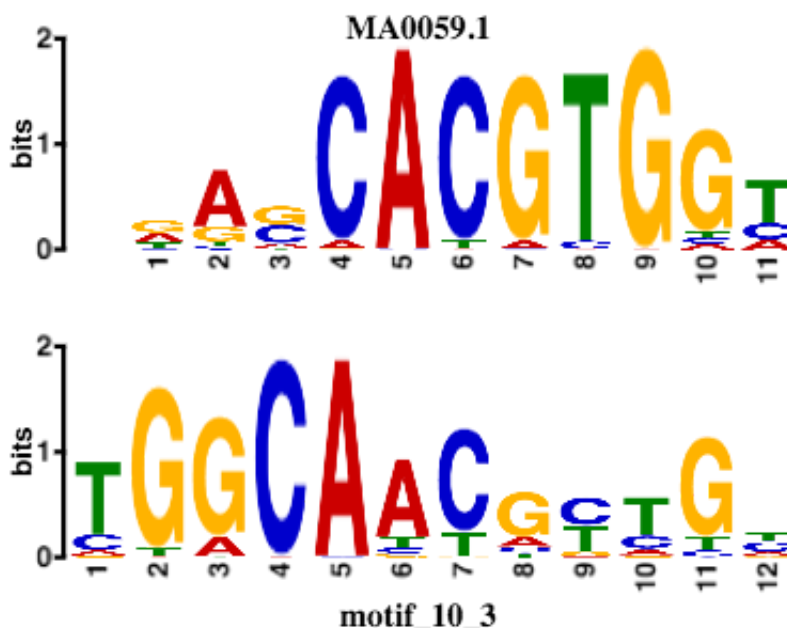
Name [MA0059.1 \(MYC::MAX\)](#)
Database JASPAR CORE 2014

p-value 1.01e-02
E-value 9.89e+00
q-value 1.00e+00

Overlap 11
Offset -1
Orientation Normal

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Alignment



Summary

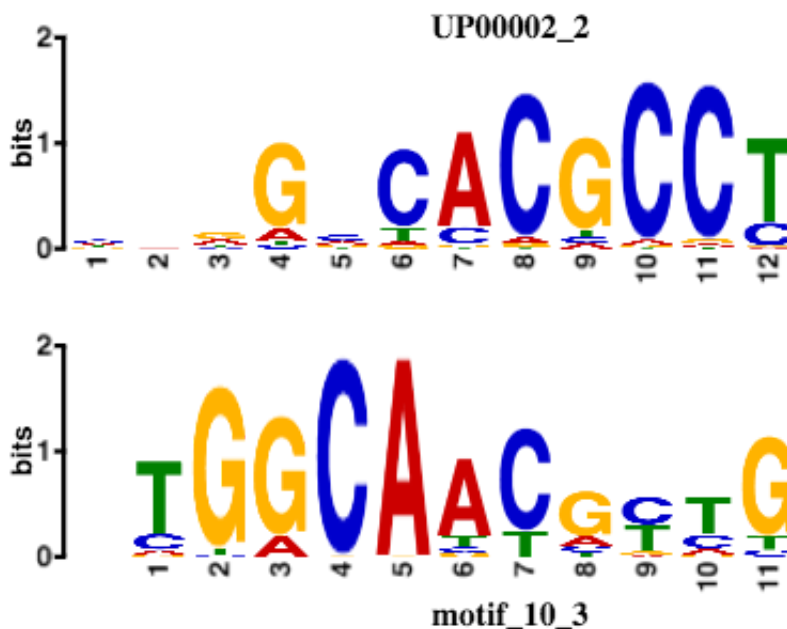
Name [UP00002_2 \(Sp4_secondary\)](#)
Database uniprobe mouse

p-value 1.02e-02
E-value 9.95e+00
q-value 1.00e+00

Overlap 12
Offset 1
Orientation Reverse Complement

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Alignment



SETTINGS

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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.

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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 -eval -thresh 10.0 motif_group6.txt
```

Result calculation took 9.742 seconds

