



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 Stamatoyannopolous, 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 group7	motif_10_1	1-RTATAAAAASVS		9	<a href="#">MA0108</a> <a href="#">UP00021</a>
motif group7	motif_Train1_19	22-GTCTTCAGGGAC		5	<a href="#">MA0154</a> <a href="#">MA0364</a>
motif group7	motif_Train2_18	22-AATGGGCGAAAC		10	<a href="#">MA0502</a> <a href="#">UP00130</a>
motif group7	motif_Train3_19	22-TGAAACGAGA		5	<a href="#">MA0393</a> <a href="#">MA0085</a>
motif group7	motif_Train4_19	14-ACCATGGA		14	<a href="#">MA0138</a> <a href="#">MA0445</a>
motif group7	motif_Train5_19	19-CAGCGCACTATC		9	<a href="#">MA0291</a> <a href="#">UP00140</a>
motif group7	motif_Train6_1	1-GTATAAAARNNV		9	<a href="#">MA0108</a> <a href="#">MA0015</a>
motif group7	motif_Train7_1	1-GTATAAAAVVS		10	<a href="#">MA0108</a> <a href="#">UP0006</a>
motif group7	motif_Train8_2	1-GTATAAAARSVG		6	<a href="#">MA0108</a> <a href="#">MA0296</a>

motif  
group7

motif\_Train9\_19

20-  
GTAGGCTATGGT

6

[UP00056\\_1](#)  
[MA0511.1](#)

## TARGET DATABASES

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

Database	Used	Matched
uniprobe mouse	386	24
JASPAR CORE 2014	593	35

## MATCHES TO MOTIF\_Train9\_19 (20-GTAGGCTATGGT)

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

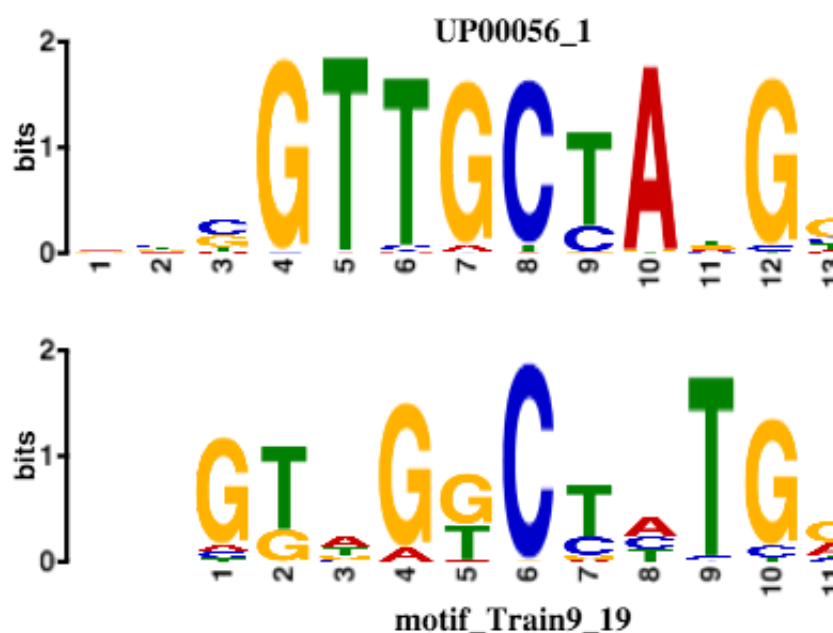
### Summary

**Name** [UP00056\\_1](#)  
[\(Rfx4\\_primary\)](#)  
**Database** uniprobe mouse  
**p-value** 2.86e-03  
**E-value** 2.80e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 2  
**Orientation** Reverse  
 Complement

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### Alignment



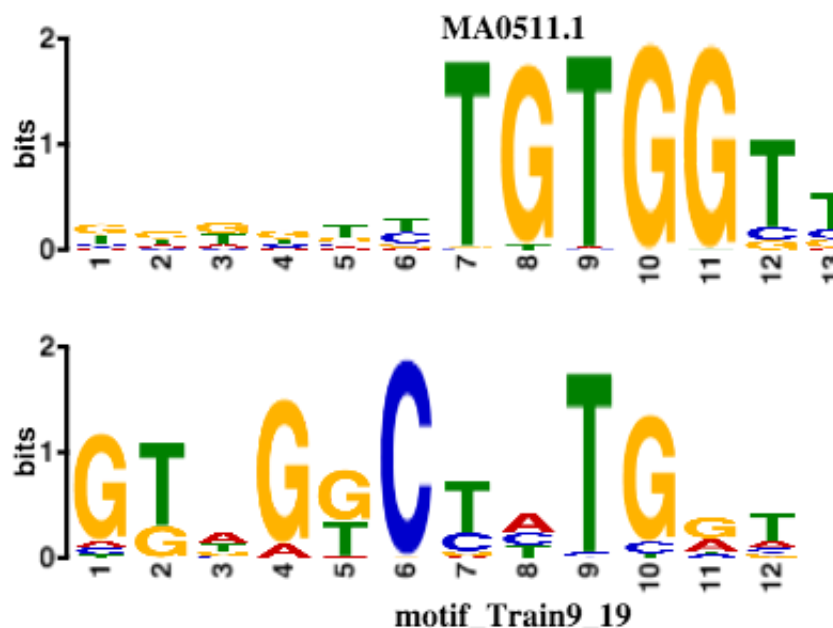
### Summary

**Name** [MA0511.1](#)  
[\(RUNX2\)](#)  
**Database** JASPAR CORE  
 2014  
**p-value** 3.48e-03  
**E-value** 3.41e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 0  
**Orientation** Normal

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### Alignment

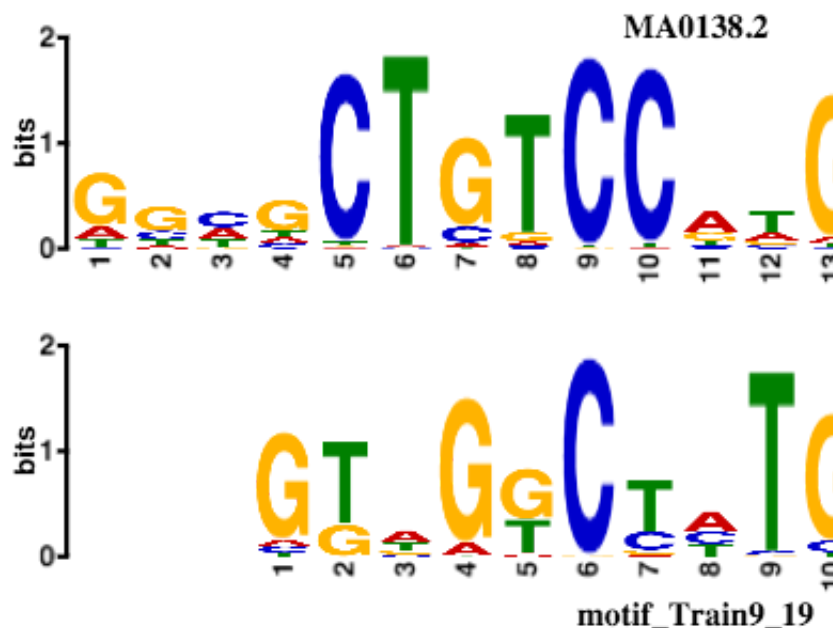


## Summary

<b>Name</b>	<a href="#">MA0138.2 (REST)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	3.57e-03
<b>E-value</b>	3.49e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	3
<b>Orientation</b>	Reverse Complement

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## Alignment

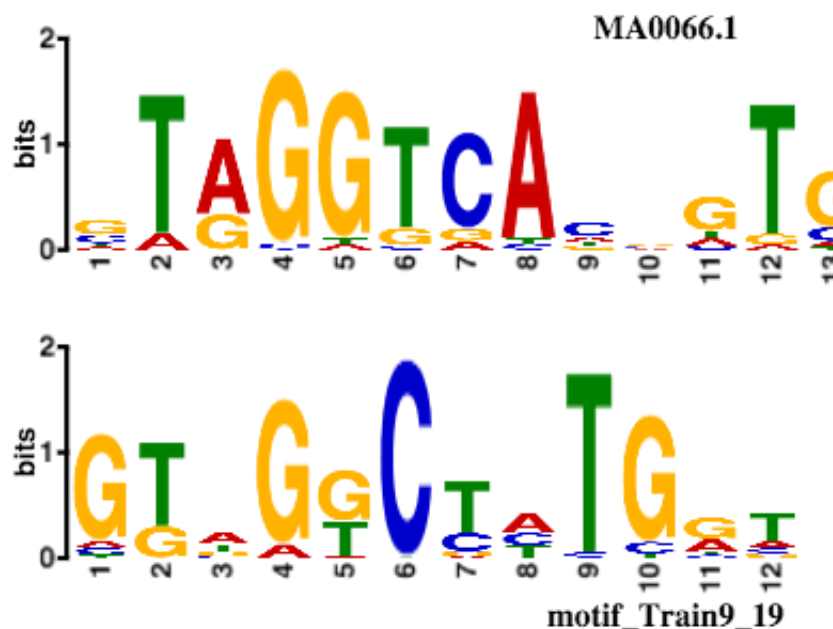


## Summary

<b>Name</b>	<a href="#">MA0066.1 (PPARG)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	5.84e-03
<b>E-value</b>	5.72e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	0
<b>Orientation</b>	Normal

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## Alignment

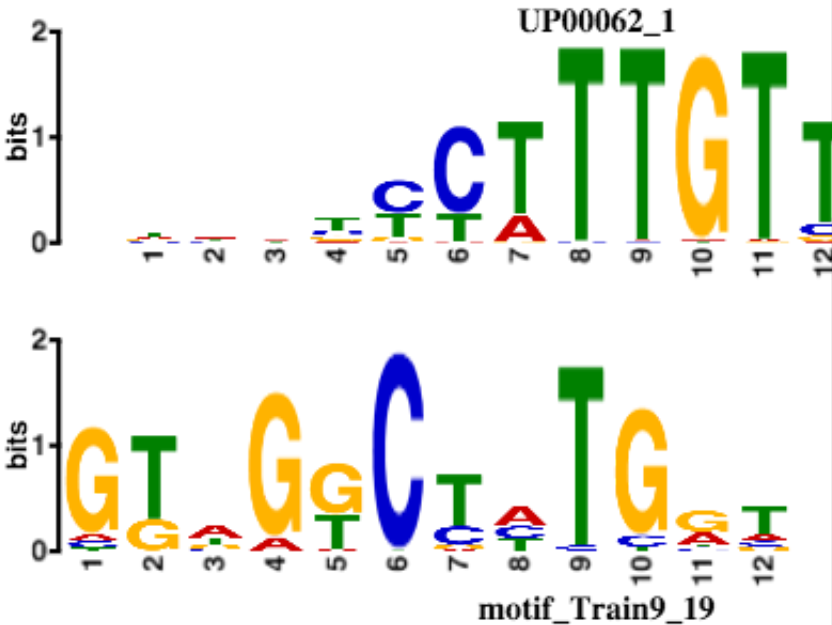


## Summary

<b>Name</b>	<a href="#">UP00062_1 (Sox4_primary)</a>
<b>Database</b>	uniprobe mouse
<b>p-value</b>	7.63e-03
<b>E-value</b>	7.47e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	11
<b>Offset</b>	-1
<b>Orientation</b>	Reverse Complement

## Alignment

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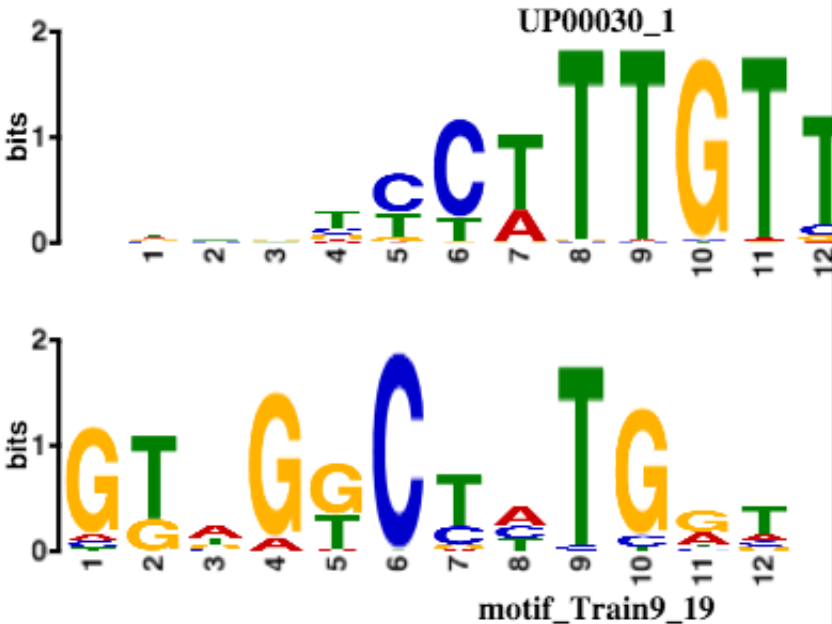


Summary

Alignment

Name	<a href="#">UP00030_1 (Sox11_primary)</a>
Database	uniprobe mouse
p-value	9.16e-03
E-value	8.97e+00
q-value	1.00e+00
Overlap	11
Offset	-1
Orientation	Reverse Complement

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MATCHES TO MOTIF\_TRAIN5\_19 (19-CAGCGCACTATC)

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

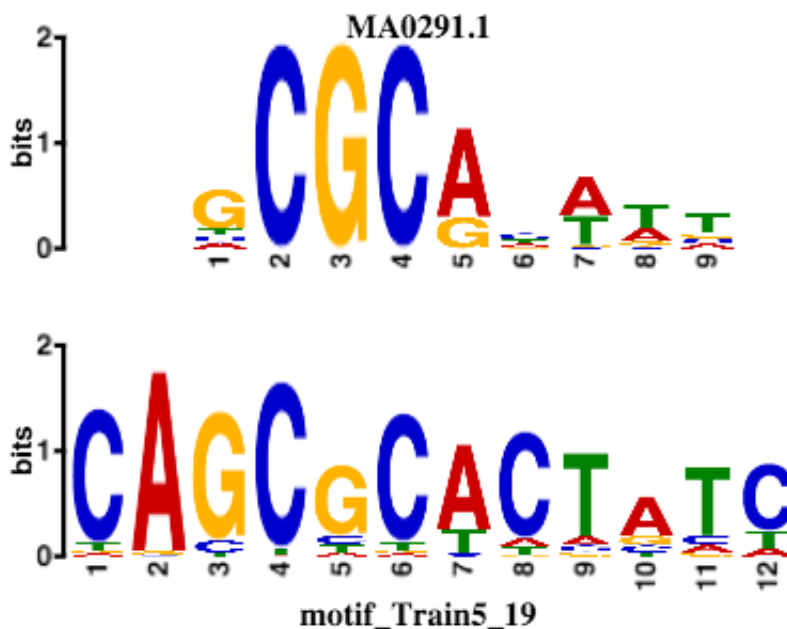
Summary

Alignment

Name	<a href="#">MA0291.1 (DAL82)</a>
Database	JASPAR CORE 2014
p-value	1.85e-04
E-value	1.81e-01
q-value	3.62e-01
Overlap	9

**Offset** -2  
**Orientation** Reverse  
 Complement

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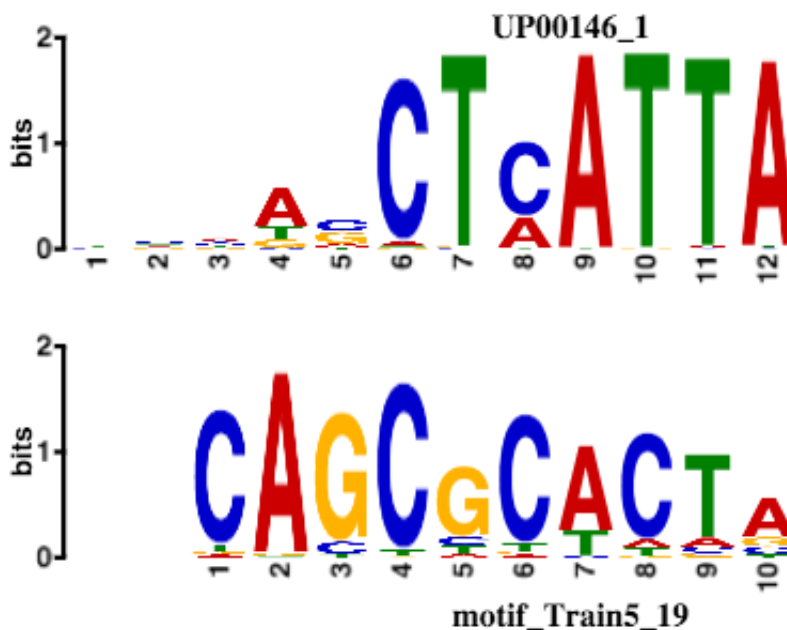
## Summary

**Name** [UP00146\\_1 \(Pou6f1 1731.2\)](#)  
**Database** uniprobe mouse  
**p-value** 1.51e-03  
**E-value** 1.48e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 2  
**Orientation** Reverse  
 Complement

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## Alignment



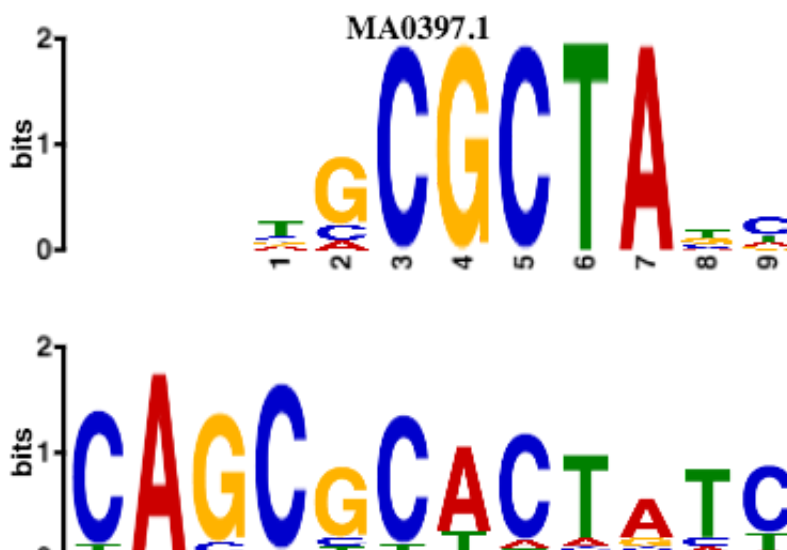
## Summary

**Name** [MA0397.1 \(STP4\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 4.99e-03  
**E-value** 4.89e+00  
**q-value** 1.00e+00

**Overlap** 9  
**Offset** -3  
**Orientation** Reverse  
 Complement

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## Alignment



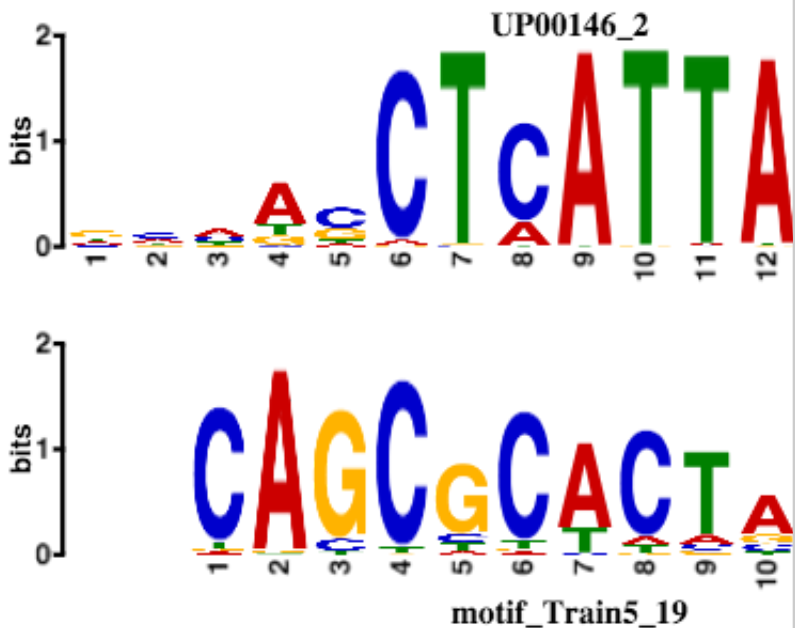


Summary

Name	<a href="#">UP00146_2</a> (Pou6f1_3733.1)
Database	uniprobe mouse
p-value	6.23e-03
E-value	6.10e+00
q-value	1.00e+00
Overlap	12
Offset	2
Orientation	Reverse Complement

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Alignment

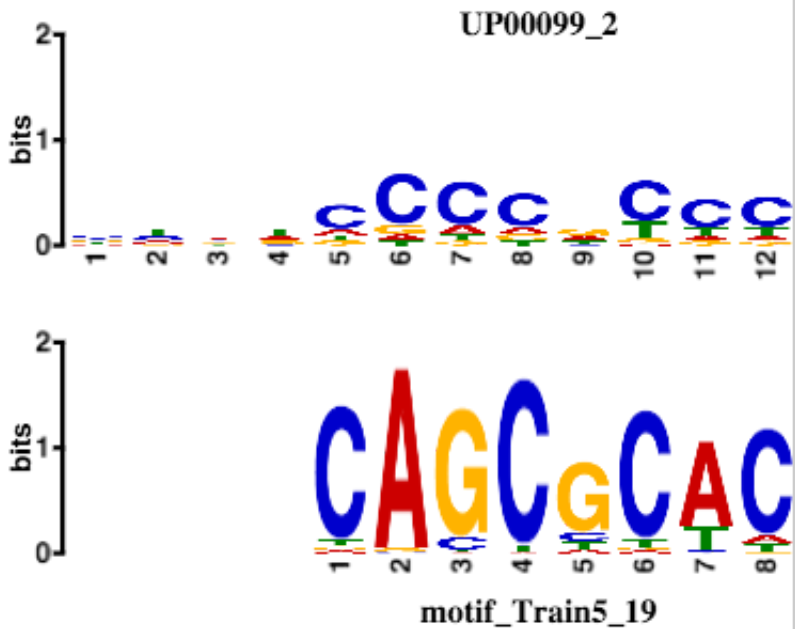


Summary

Name	<a href="#">UP00099_2</a> (Ascl2_secondary)
Database	uniprobe mouse
p-value	8.05e-03
E-value	7.88e+00
q-value	1.00e+00
Overlap	12
Offset	4
Orientation	Normal

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Alignment



Summary

Name	<a href="#">MA0327.1</a> (MATA1)
Database	JASPAR CORE 2014

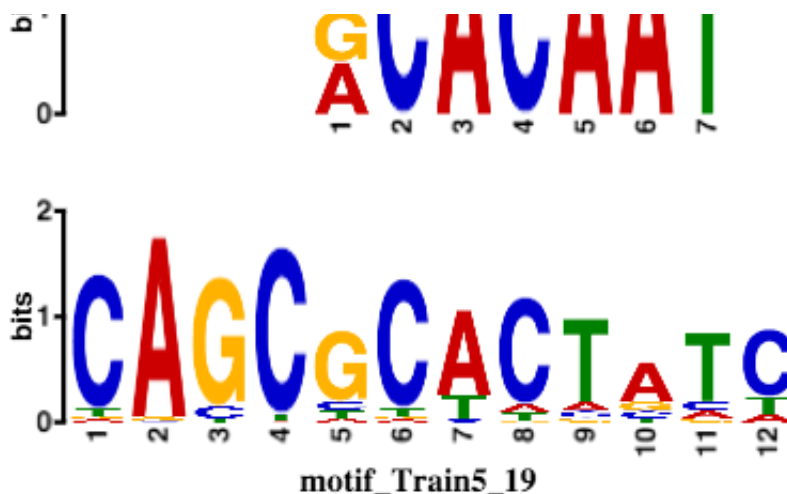
Alignment



**p-value** 8.52e-03  
**E-value** 8.34e+00  
**q-value** 1.00e+00

**Overlap** 7  
**Offset** -4  
**Orientation** Normal

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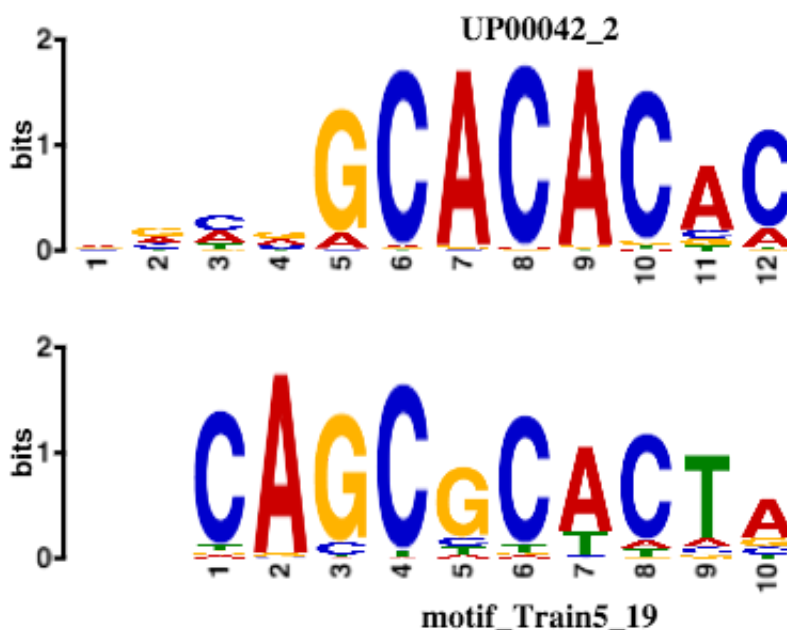


## Summary

**Name** [UP00042\\_2](#)  
[\(Gm397\\_secondary\)](#)  
**Database** uniprobe mouse  
**p-value** 8.73e-03  
**E-value** 8.54e+00  
**q-value** 1.00e+00  
**Overlap** 12  
**Offset** 2  
**Orientation** Normal

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## Alignment

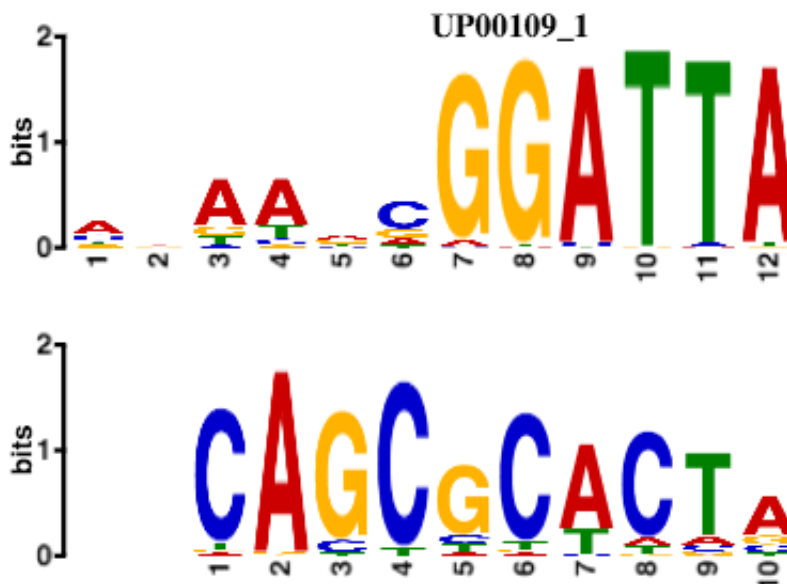


## Summary

**Name** [UP00109\\_1](#)  
[\(Obox6\\_3440.2\)](#)  
**Database** uniprobe mouse  
**p-value** 8.95e-03  
**E-value** 8.77e+00  
**q-value** 1.00e+00  
**Overlap** 12  
**Offset** 2  
**Orientation** Normal

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## Alignment





motif\_Train5\_19

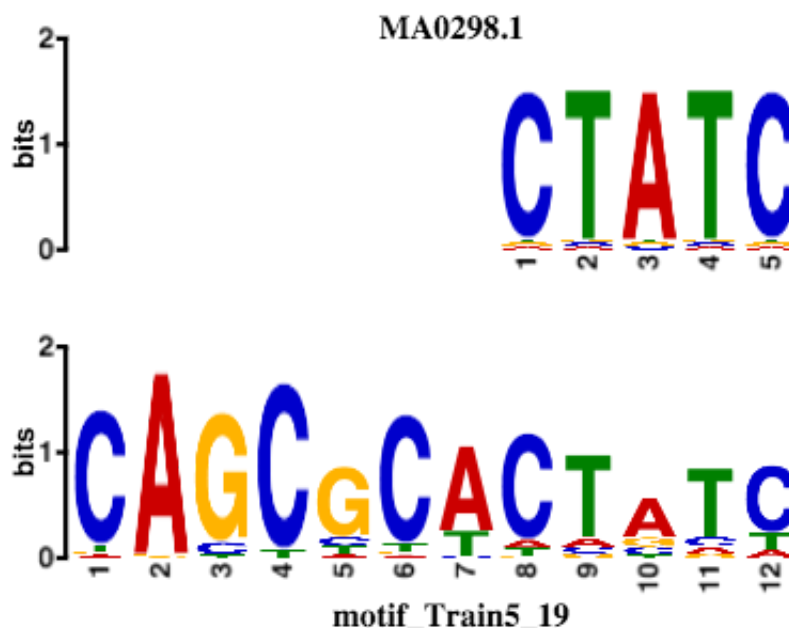
## Summary

## Alignment

**Name** [MA0298.1 \(FZF1\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 9.71e-03  
**E-value** 9.50e+00  
**q-value** 1.00e+00

**Overlap** 5  
**Offset** -7  
**Orientation** Normal

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## MATCHES TO MOTIF\_Train1\_19 (22-GTCTTCAGGGAC)

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

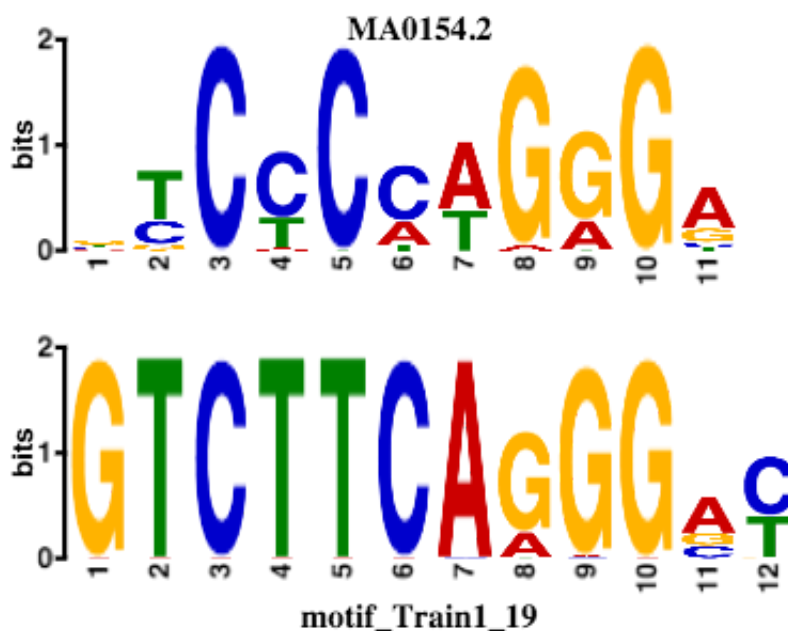
## Summary

## Alignment

**Name** [MA0154.2 \(EBF1\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 8.44e-05  
**E-value** 8.27e-02  
**q-value** 1.65e-01

**Overlap** 11  
**Offset** 0  
**Orientation** Normal

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## Summary

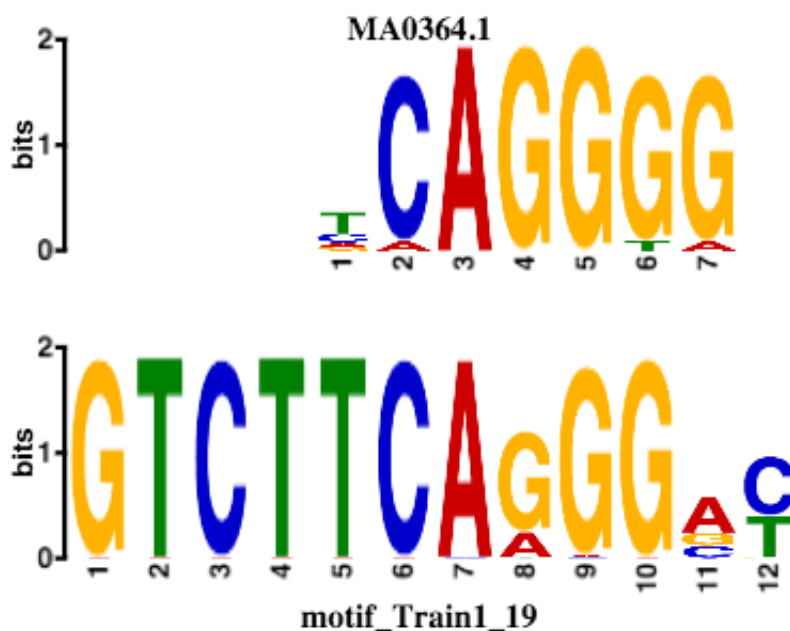
## Alignment

**Name** [MA0364.1 \(REI1\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 1.40e-03  
**E-value** 1.37e+00



**q-value** 1.00e+00  
**Overlap** 7  
**Offset** -4  
**Orientation** Reverse Complement

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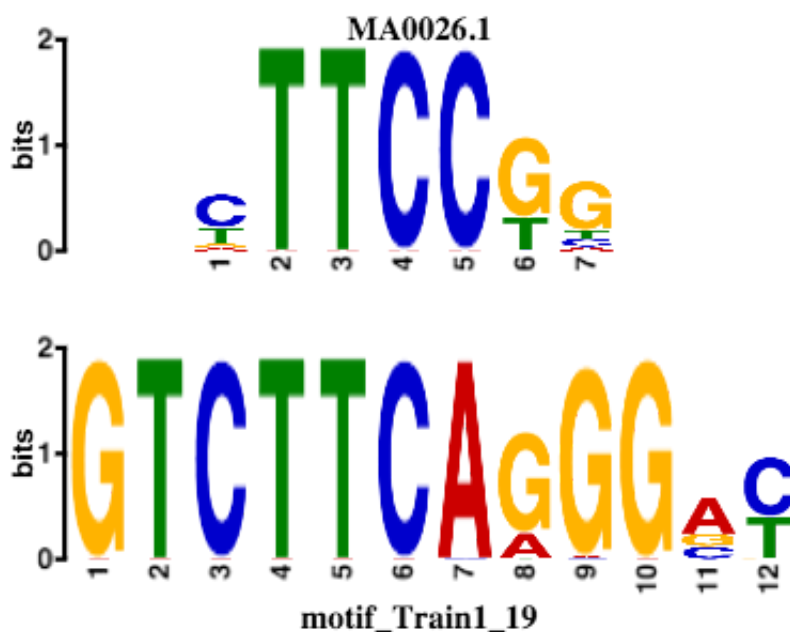
## Summary

**Name** [MA0026.1 \(Eip74EF\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 3.87e-03  
**E-value** 3.79e+00  
**q-value** 1.00e+00

**Overlap** 7  
**Offset** -2  
**Orientation** Reverse Complement

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## Alignment

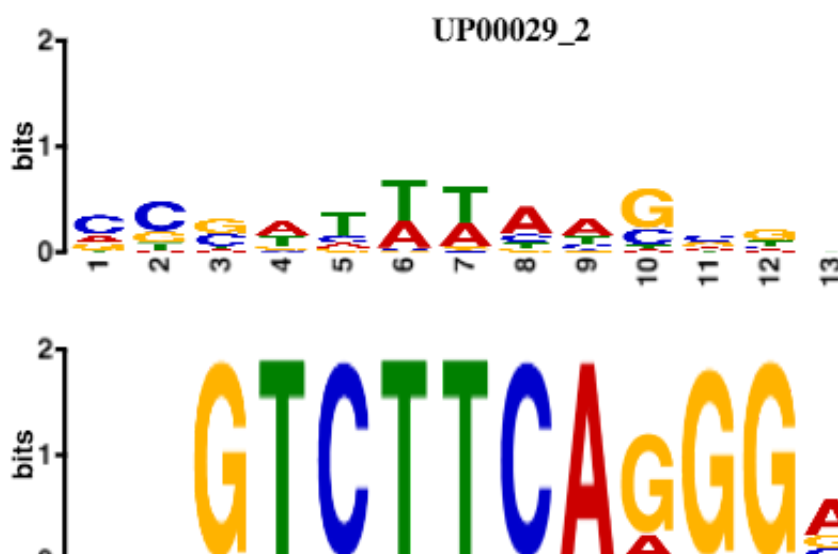


## Summary

**Name** [UP00029\\_2 \(Tbp\\_secondary\)](#)  
**Database** uniprobe mouse  
**p-value** 8.21e-03  
**E-value** 8.04e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 2  
**Orientation** Normal

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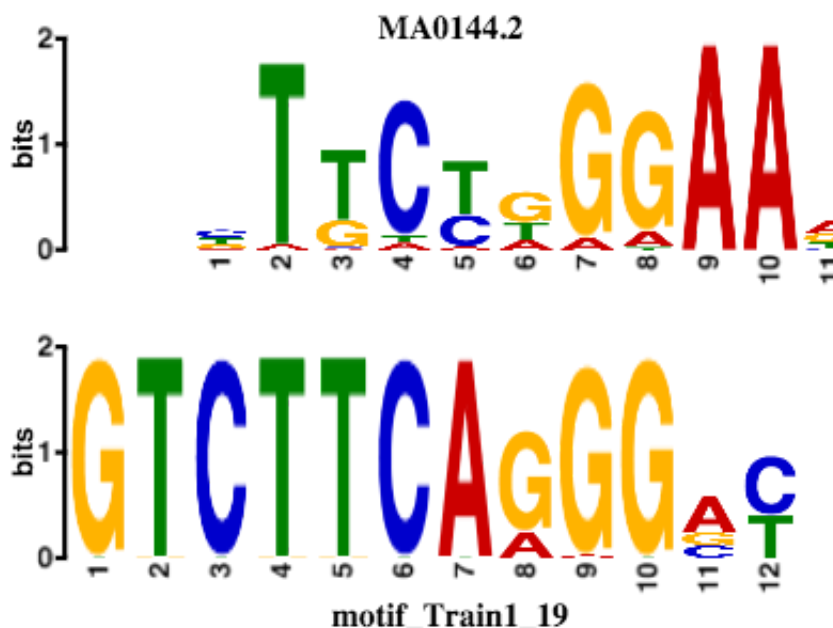
1 2 3 4 5 6 7 8 9 10 11  
motif\_Train1\_19

## Summary

## Alignment

<b>Name</b>	<a href="#">MA0144.2 (STAT3)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	9.00e-03
<b>E-value</b>	8.81e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	-2
<b>Orientation</b>	Normal

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## MATCHES TO MOTIF\_Train6\_1 (1-GTATAAAARNNV)

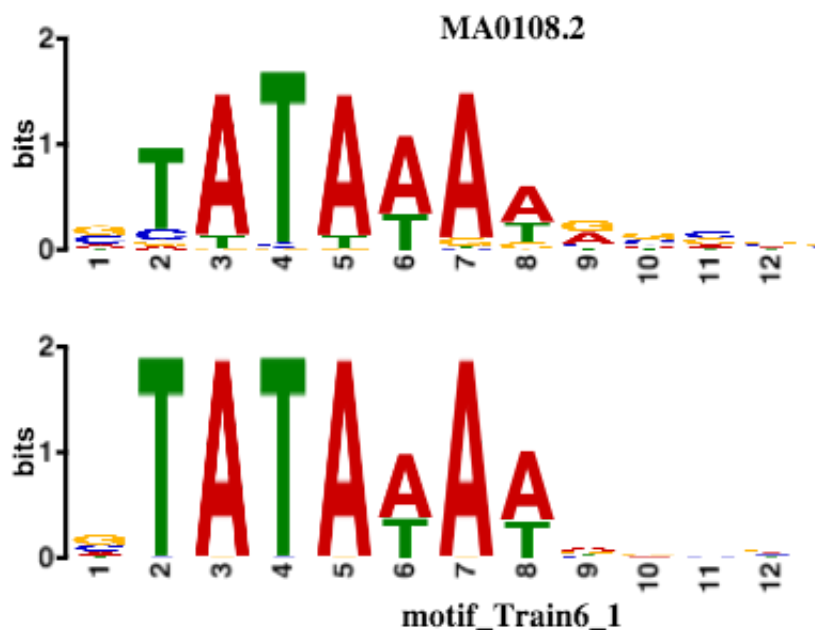
[Previous](#) [Next](#) [Top](#)

## Summary

## Alignment

<b>Name</b>	<a href="#">MA0108.2 (TBP)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	4.89e-14
<b>E-value</b>	4.79e-11
<b>q-value</b>	9.58e-11
<b>Overlap</b>	12
<b>Offset</b>	0
<b>Orientation</b>	Normal

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## Summary

## Alignment

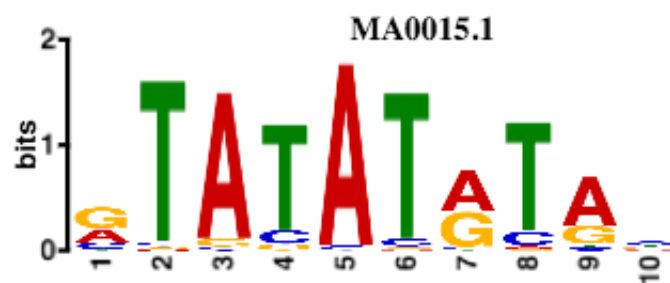
**Name** [MA0015.1](#)

**Database** [\(Cf2\\_II\)](#)  
JASPAR CORE  
2014

**p-value** 1.14e-03  
**E-value** 1.12e+00  
**q-value** 1.00e+00

**Overlap** 10  
**Offset** 0  
**Orientation** Normal

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## Summary

**Name** [UP00029\\_1](#)  
[\(Tbp\\_primary\)](#)

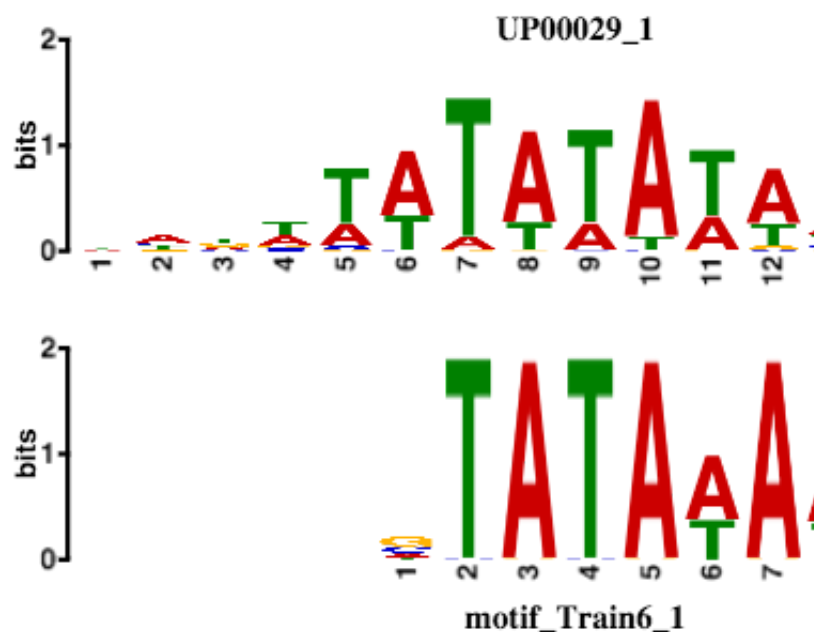
**Database** uniprobe mouse

**p-value** 4.87e-03  
**E-value** 4.77e+00  
**q-value** 1.00e+00

**Overlap** 11  
**Offset** 5  
**Orientation** Reverse  
Complement

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## Alignment



## Summary

**Name** [UP00061\\_1](#)  
[\(Foxl1\\_primary\)](#)

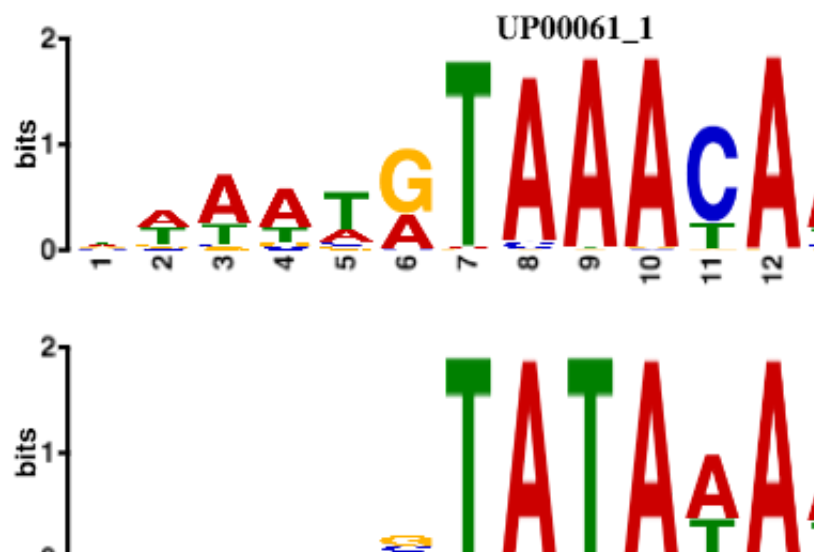
**Database** uniprobe mouse

**p-value** 4.91e-03  
**E-value** 4.81e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 5  
**Orientation** Normal

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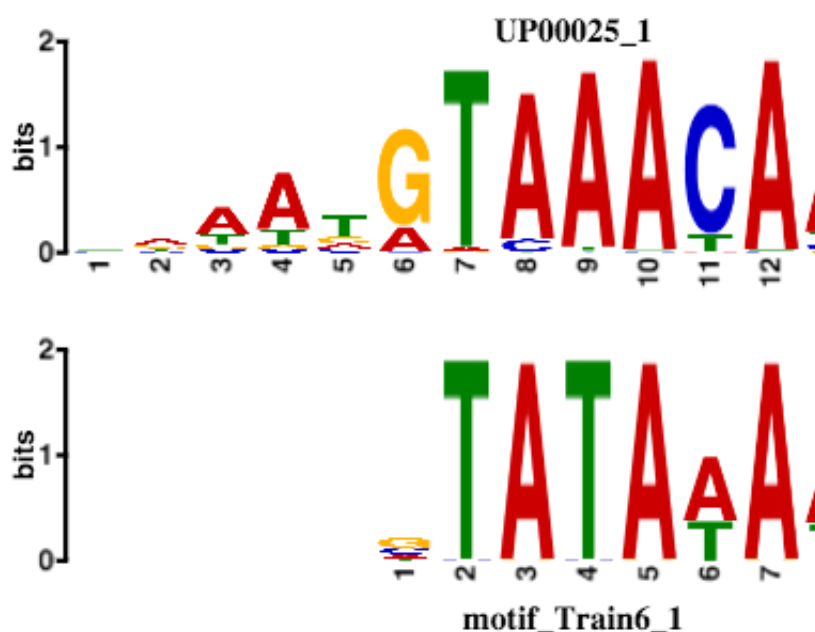
## Alignment



## Summary

<b>Name</b>	<a href="#">UP00025_1</a> (Foxk1_primary)
<b>Database</b>	uniprobe mouse
<b>p-value</b>	6.01e-03
<b>E-value</b>	5.89e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	5
<b>Orientation</b>	Normal
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## Alignment



## Summary

<b>Name</b>	<a href="#">MA0296.1 (FKH1)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	6.02e-03
<b>E-value</b>	5.90e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	6
<b>Orientation</b>	Normal
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## Alignment



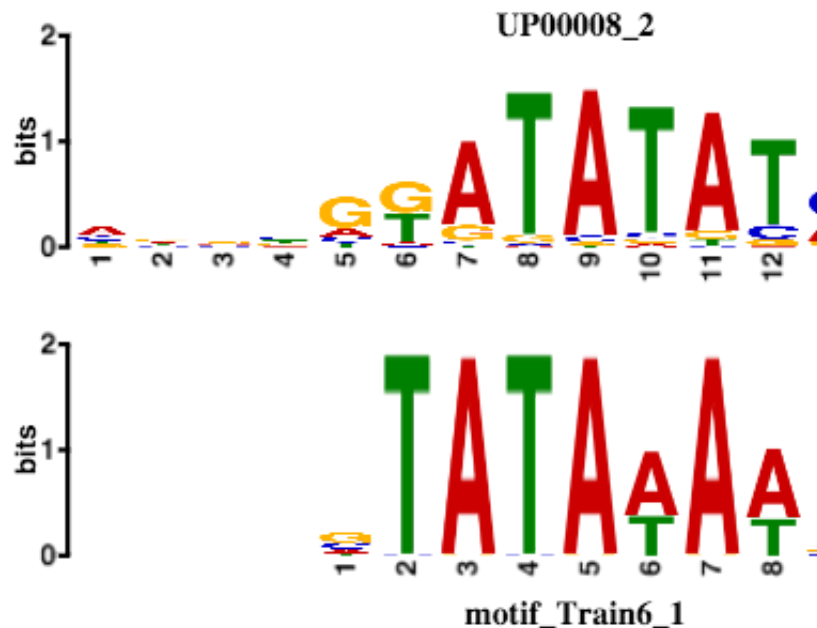
## Summary

<b>Name</b>	<a href="#">UP00008_2</a> (Six6_secondary)
<b>Database</b>	uniprobe mouse
<b>p-value</b>	6.42e-03
<b>E-value</b>	6.29e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12

## Alignment

**Offset** 4  
**Orientation** Reverse  
 Complement

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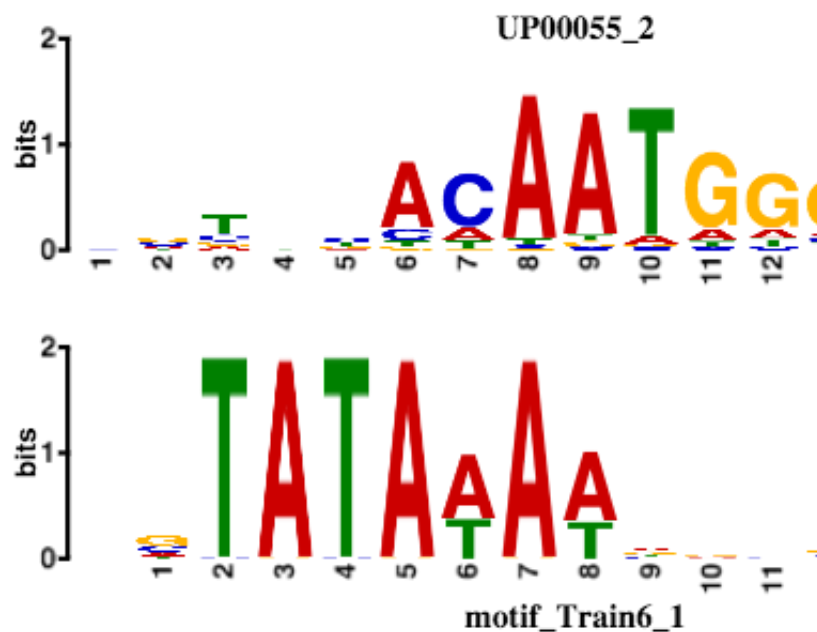
## Summary

**Name** [UP00055\\_2](#)  
 (Hbp1\_secondary)  
**Database** uniprobe mouse  
**p-value** 7.82e-03  
**E-value** 7.66e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 1  
**Orientation** Reverse  
 Complement

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## Alignment

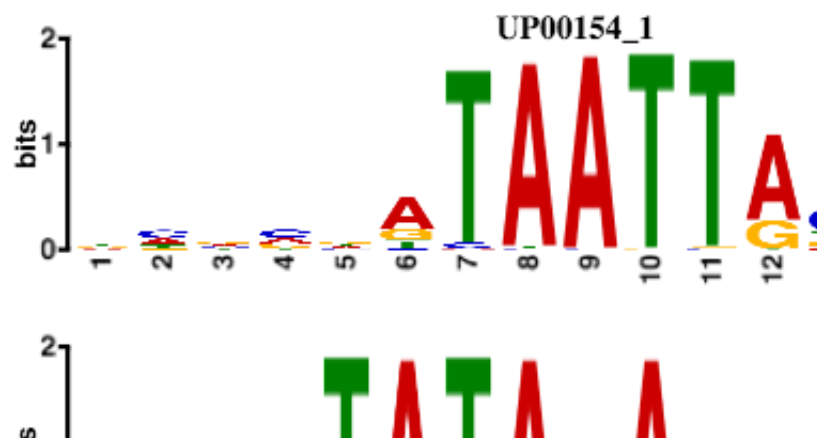


## Summary

**Name** [UP00154\\_1](#)  
 (Dlx3\_1030.1)  
**Database** uniprobe mouse  
**p-value** 9.48e-03  
**E-value** 9.28e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 3

## Alignment



**Orientation** Normal

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bits  
1  
0



## MATCHES TO MOTIF\_Train2\_18 (22-AATGGGCGAAAC)

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

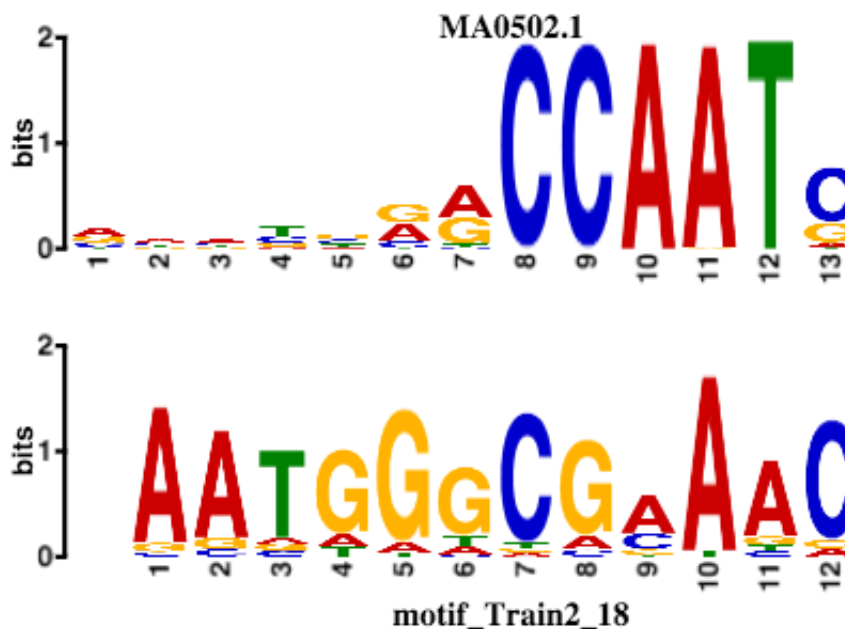
### Summary

**Name** [MA0502.1 \(NFYB\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 1.27e-03  
**E-value** 1.24e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 1  
**Orientation** Normal

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### Alignment



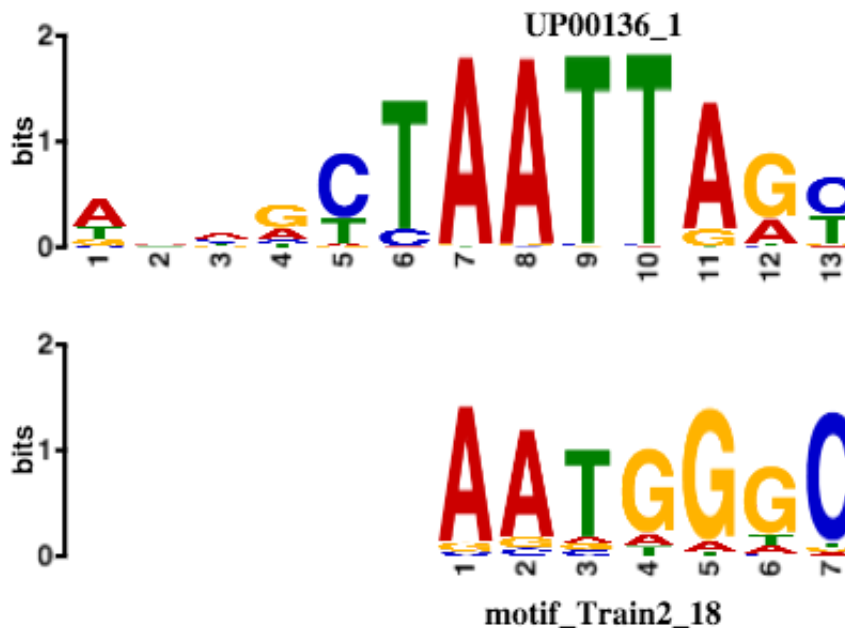
### Summary

**Name** [UP00136\\_1 \(Prrx2\\_3072.1\)](#)  
**Database** uniprobe mouse  
**p-value** 2.40e-03  
**E-value** 2.35e+00  
**q-value** 1.00e+00

**Overlap** 11  
**Offset** 6  
**Orientation** Normal

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### Alignment



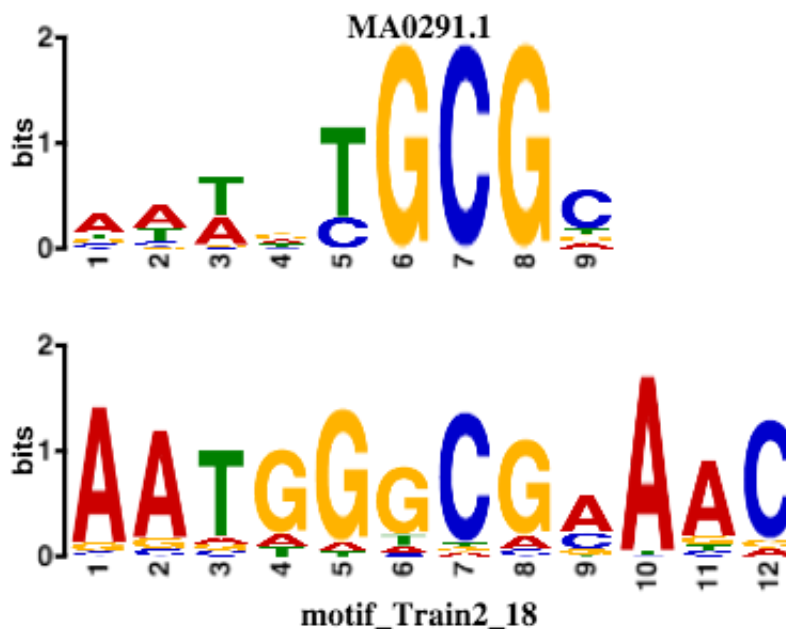
### Summary

**Name** [MA0291.1 \(DAL82\)](#)  
**Database** JASPAR CORE

### Alignment

<b>Database</b>	2014
<b><i>p</i>-value</b>	3.31e-03
<b><i>E</i>-value</b>	3.24e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	9
<b>Offset</b>	0
<b>Orientation</b>	Normal

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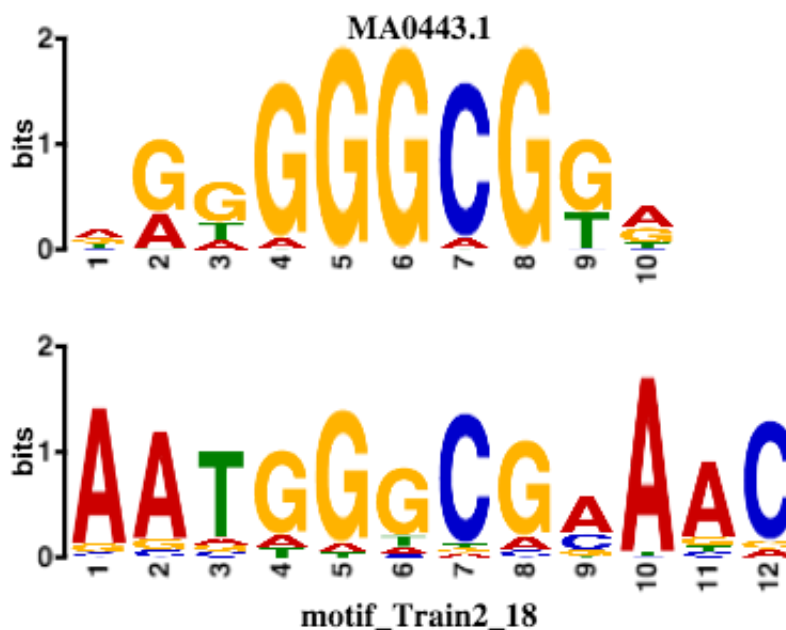


## Summary

<b>Name</b>	<a href="#">MA0443.1 (btd)</a>
<b>Database</b>	JASPAR CORE 2014
<b><i>p</i>-value</b>	3.41e-03
<b><i>E</i>-value</b>	3.34e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	0
<b>Orientation</b>	Normal

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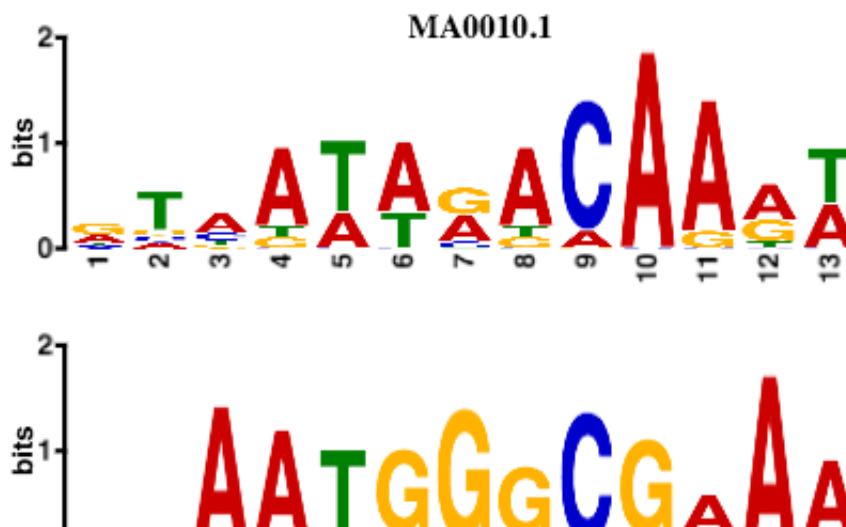
## Alignment



## Summary

<b>Name</b>	<a href="#">MA0010.1 (br_Z1)</a>
<b>Database</b>	JASPAR CORE 2014
<b><i>p</i>-value</b>	5.11e-03
<b><i>E</i>-value</b>	5.00e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	2
<b>Orientation</b>	Normal

## Alignment





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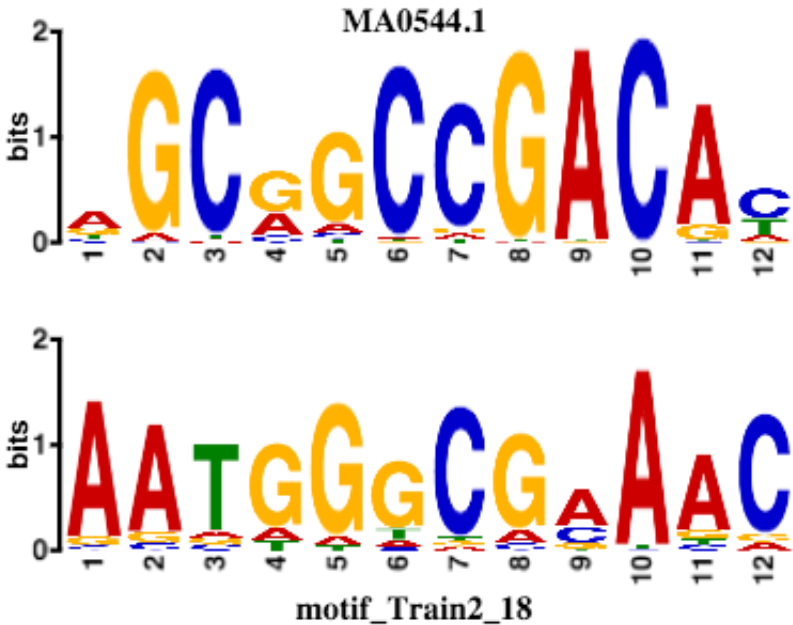


Summary

Name	<a href="#">MA0544.1 (GEI-11)</a>
Database	JASPAR CORE 2014
p-value	5.18e-03
E-value	5.07e+00
q-value	1.00e+00
Overlap	12
Offset	0
Orientation	Reverse Complement

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Alignment

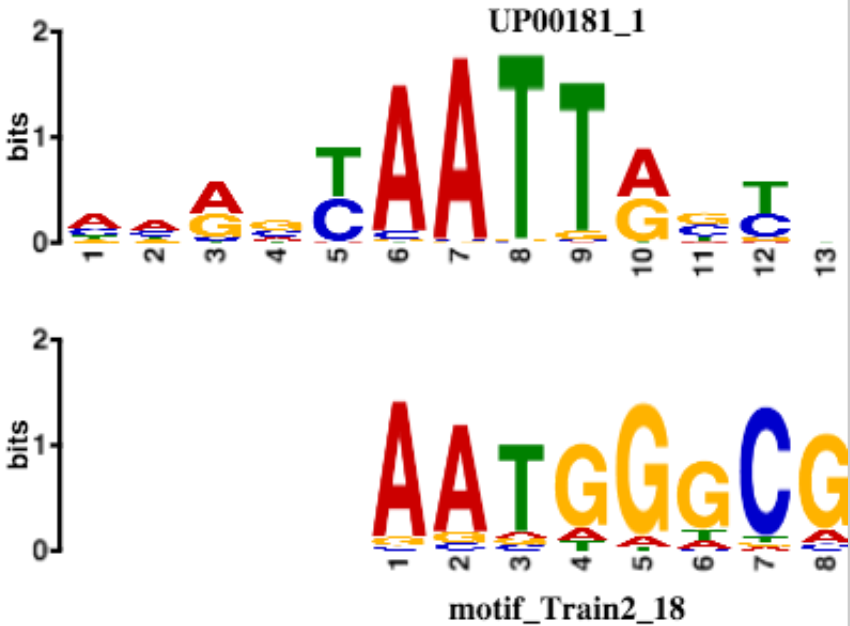


Summary

Name	<a href="#">UP00181_1 (Barx1_2877.1)</a>
Database	uniprobe mouse
p-value	7.11e-03
E-value	6.96e+00
q-value	1.00e+00
Overlap	11
Offset	5
Orientation	Normal

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Alignment



Summary

Name	<a href="#">MA0401.1 (SWI4)</a>
Database	JASPAR CORE 2014
p-value	8.81e-03
E-value	8.63e+00
q-value	1.00e+00

Alignment

Overlap

8

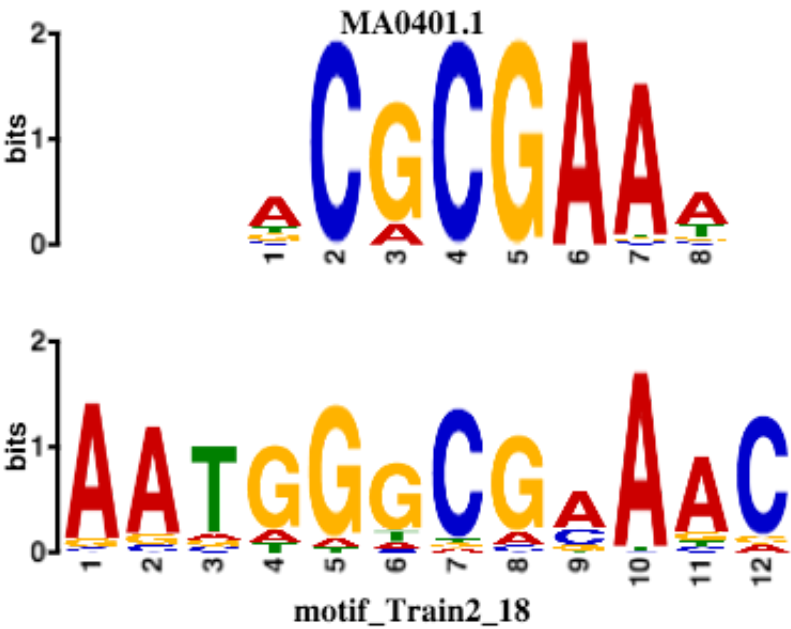
Offset

-3

Orientation

Normal

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Summary

Alignment

Name

[UP00139\\_1](#)  
(Nkx1-  
2\_3214.1)

Database

uniprobe  
mouse

p-value

9.76e-03

E-value

9.55e+00

q-value

1.00e+00

Overlap

11

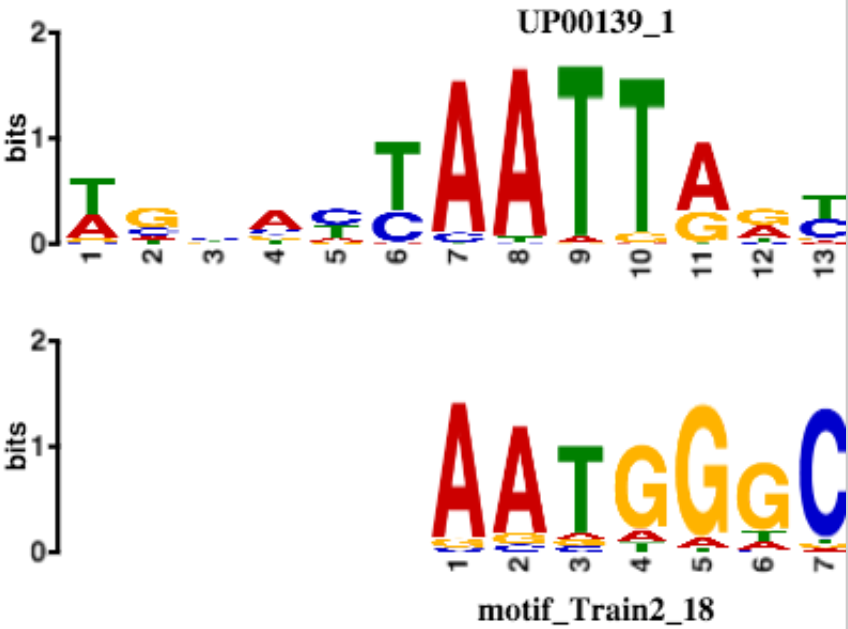
Offset

6

Orientation

Reverse  
Complement

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Summary

Alignment

Name

[UP00198\\_1](#)  
(Cphx\_3484.1)

Database

uniprobe  
mouse

p-value

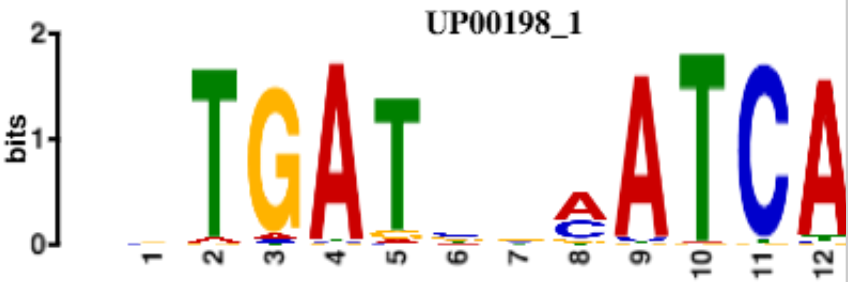
1.02e-02

E-value

9.95e+00

q-value

1.00e+00



**q-value** 1.00e+00

**Overlap** 11

**Offset** -1

**Orientation** Normal

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## MATCHES TO MOTIF\_Train7\_1 (1-GTATAAAA VVS)

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

### Summary

**Name** [MA0108.2 \(TBP\)](#)

**Database** JASPAR CORE 2014

**p-value** 8.15e-10

**E-value** 7.98e-07

**q-value** 1.60e-06

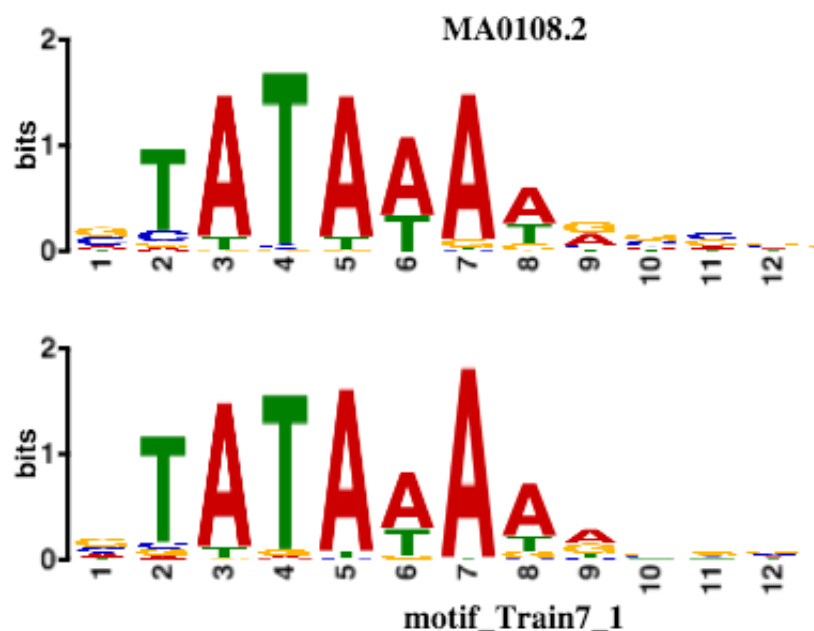
**Overlap** 12

**Offset** 0

**Orientation** Normal

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### Alignment



### Summary

**Name** [UP00061\\_1 \(Foxl1 primary\)](#)

**Database** uniprobe mouse

**p-value** 1.87e-03

**E-value** 1.83e+00

**q-value** 1.00e+00

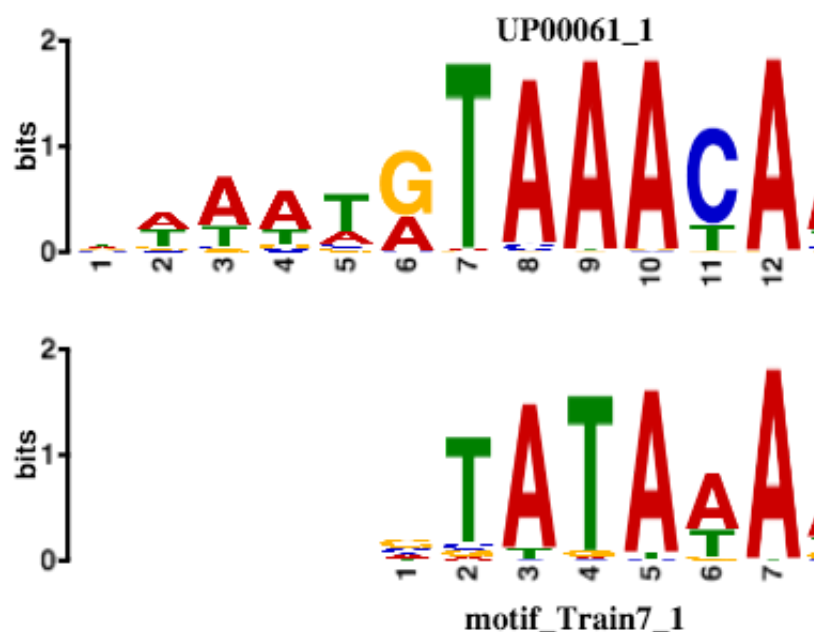
**Overlap** 12

**Offset** 5

**Orientation** Normal

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### Alignment



### Summary

### Alignment

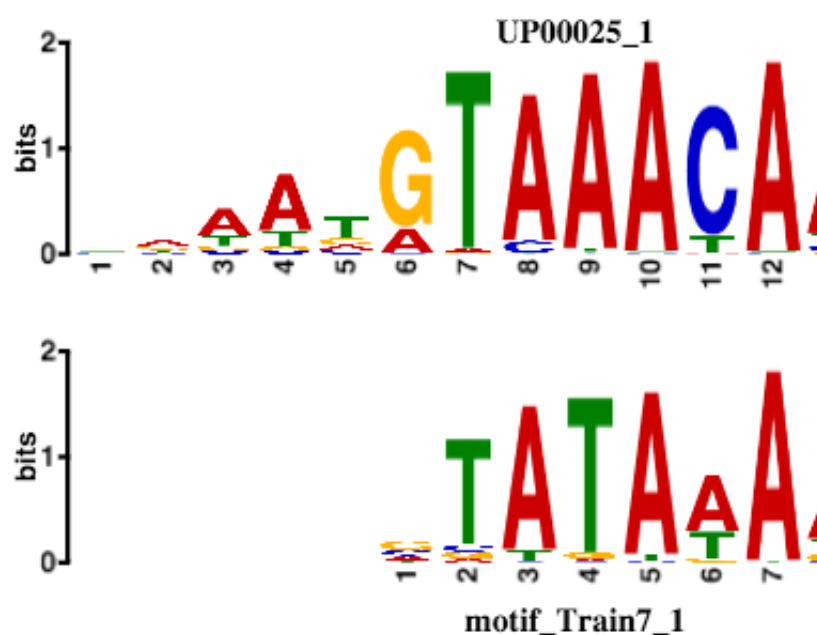
<b>Name</b>	<a href="#">MA0296.1 (FKH1)</a>
<b>Database</b>	JASPAR CORE 2014
<b><i>p</i>-value</b>	2.84e-03
<b><i>E</i>-value</b>	2.78e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	6
<b>Orientation</b>	Normal
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## Summary

<b>Name</b>	<a href="#">UP00025_1</a> <a href="#">(Foxk1_primary)</a>
<b>Database</b>	uniprobe mouse
<b><i>p</i>-value</b>	3.49e-03
<b><i>E</i>-value</b>	3.42e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	5
<b>Orientation</b>	Normal
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## Alignment

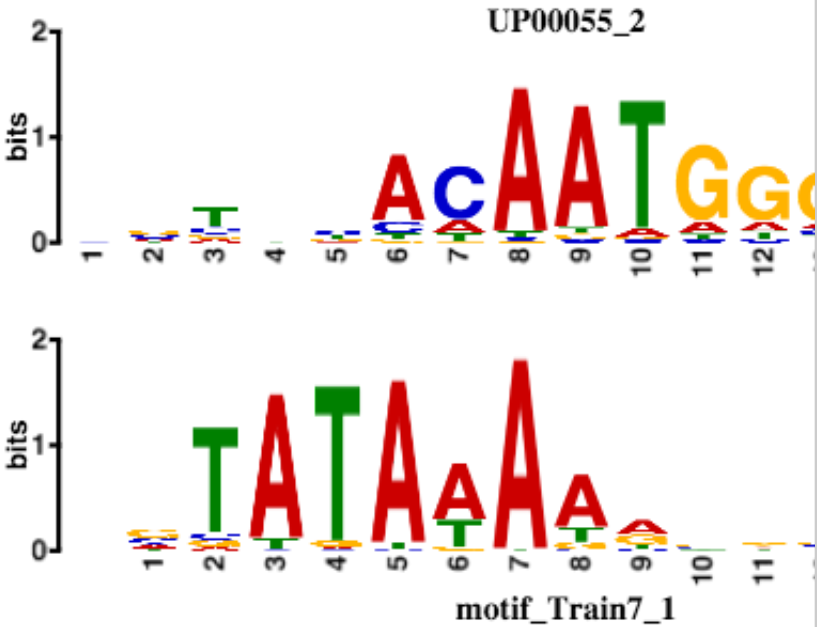


## Summary

<b>Name</b>	<a href="#">UP00055_2</a> <a href="#">(Hbp1_secondary)</a>
<b>Database</b>	uniprobe mouse
<b><i>p</i>-value</b>	3.49e-03
<b><i>E</i>-value</b>	3.42e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	1
<b>Orientation</b>	Reverse Complement
	<a href="#">Show logo download options</a>

## Alignment

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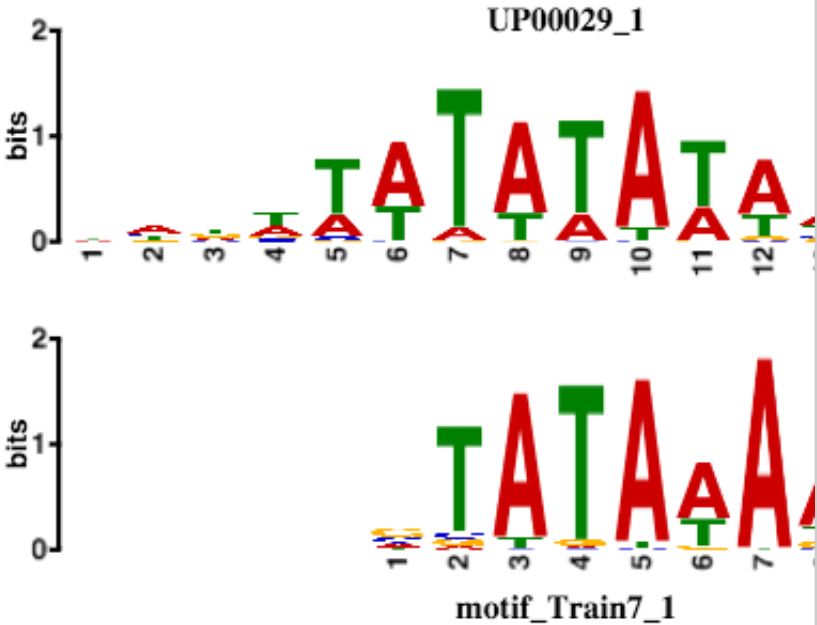


Summary

Name	<a href="#">UP00029_1</a> (Tbp_primary)
Database	uniprobe mouse
p-value	3.69e-03
E-value	3.62e+00
q-value	1.00e+00
Overlap	11
Offset	5
Orientation	Reverse Complement

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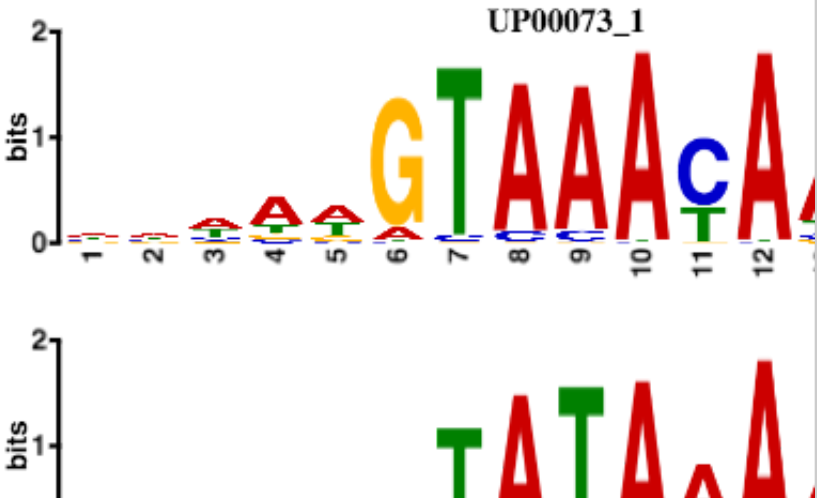
Alignment



Summary

Name	<a href="#">UP00073_1</a> (Foxa2_primary)
Database	uniprobe mouse
p-value	8.67e-03
E-value	8.49e+00
q-value	1.00e+00
Overlap	12
Offset	5
Orientation	Normal

Alignment



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0

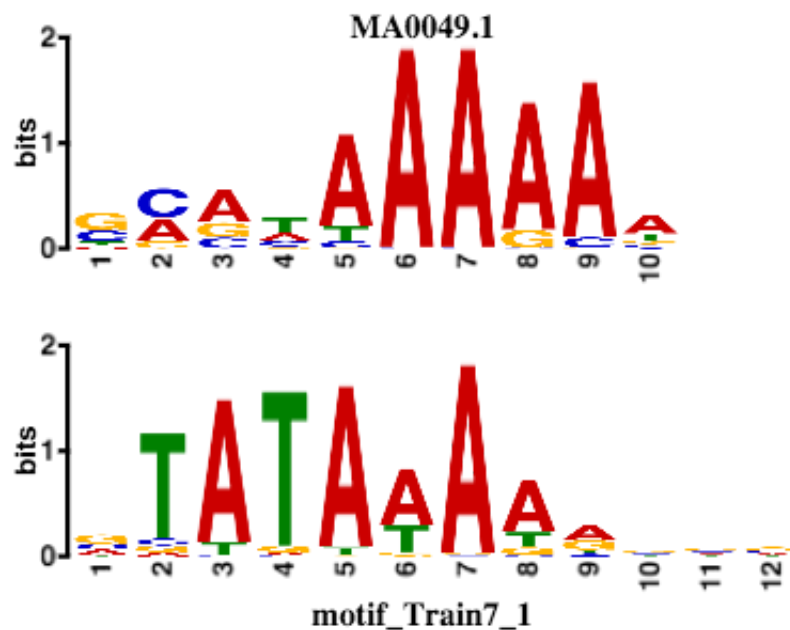


## Summary

<b>Name</b>	<a href="#">MA0049.1 (hb)</a>
<b>Database</b>	JASPAR CORE 2014
<b><i>p</i>-value</b>	8.74e-03
<b><i>E</i>-value</b>	8.56e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	0
<b>Orientation</b>	Normal

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## Alignment

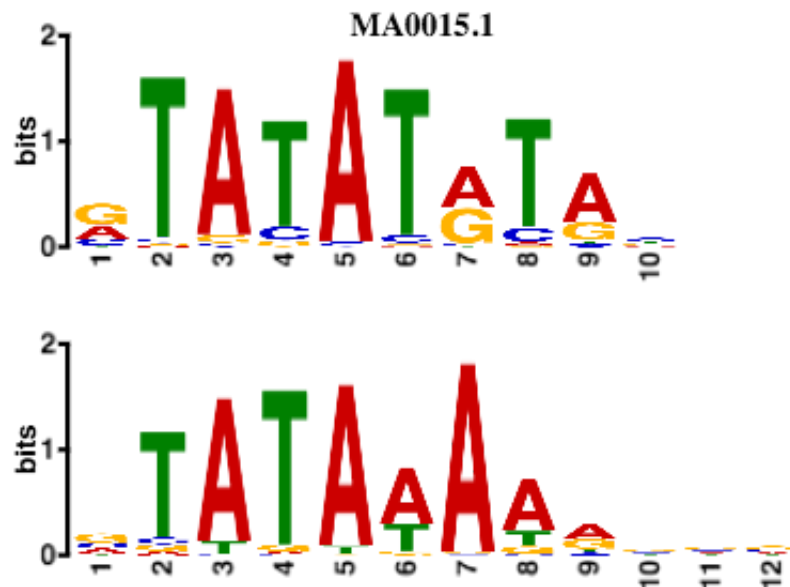


## Summary

<b>Name</b>	<a href="#">MA0015.1 (Cf2 II)</a>
<b>Database</b>	JASPAR CORE 2014
<b><i>p</i>-value</b>	9.98e-03
<b><i>E</i>-value</b>	9.77e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	0
<b>Orientation</b>	Normal

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## Alignment



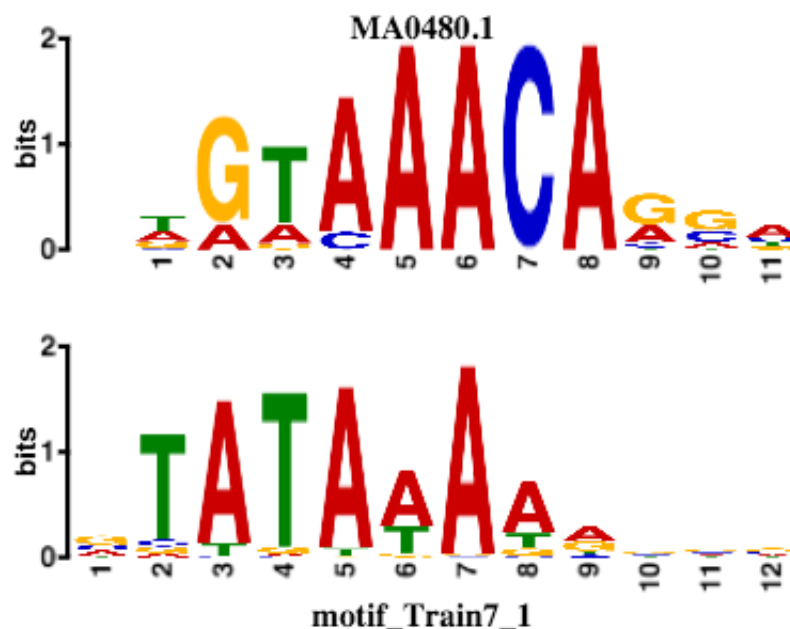
## motif\_Train7\_1

## Summary

<b>Name</b>	<a href="#">MA0480.1 (Foxo1)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	1.01e-02
<b>E-value</b>	9.86e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	11
<b>Offset</b>	-1
<b>Orientation</b>	Reverse Complement

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## Alignment



## MATCHES TO MOTIF\_Train3\_19 (22-TGAAACGAGA)

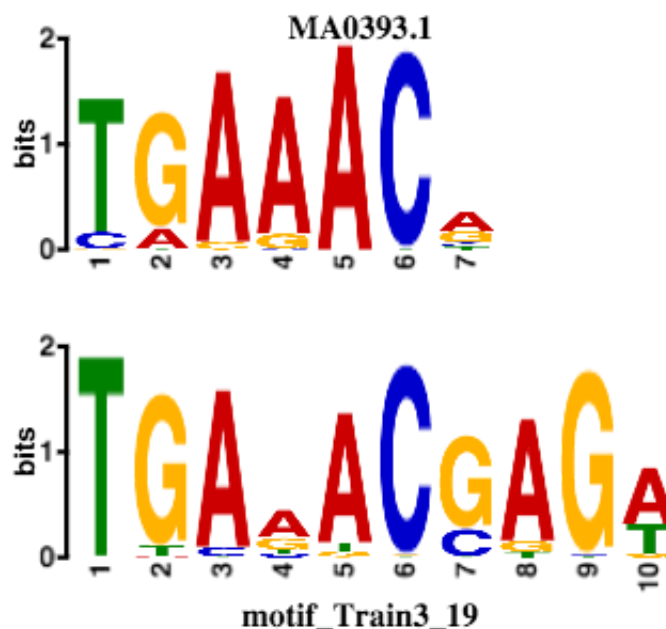
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## Summary

<b>Name</b>	<a href="#">MA0393.1 (STE12)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	5.56e-04
<b>E-value</b>	5.44e-01
<b>q-value</b>	1.00e+00
<b>Overlap</b>	7
<b>Offset</b>	0
<b>Orientation</b>	Normal

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## Alignment



## Summary

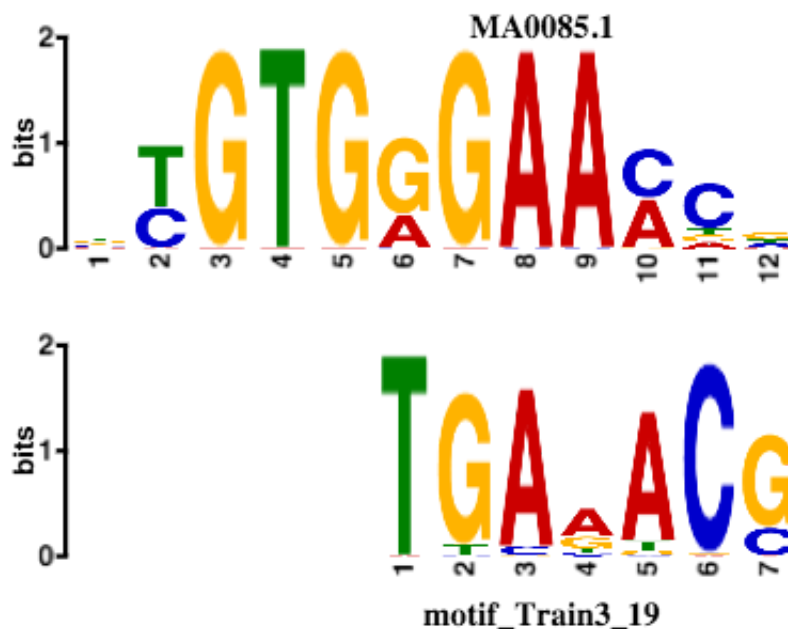
<b>Name</b>	<a href="#">MA0085.1 (Su(H))</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	3.33e-03
<b>E-value</b>	3.26e+00
<b>q-value</b>	1.00e+00

## Alignment



**Overlap** 10  
**Offset** 5  
**Orientation** Normal

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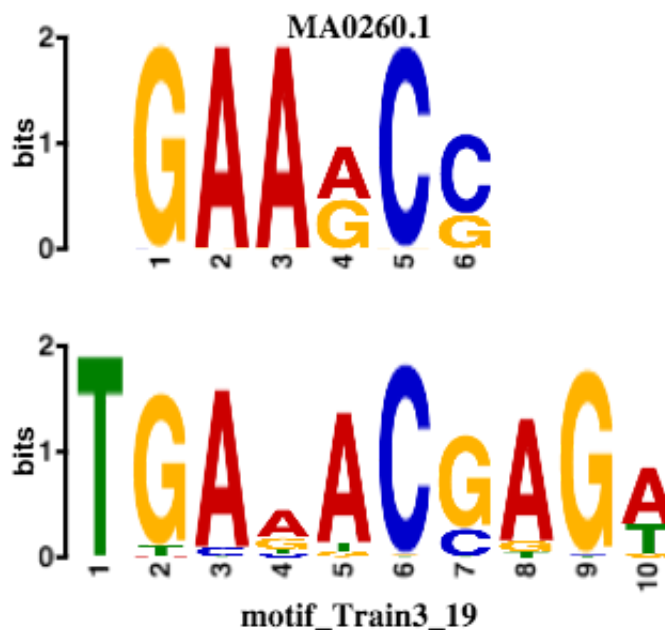
## Summary

**Name** [MA0260.1 \(che-1\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 3.54e-03  
**E-value** 3.47e+00  
**q-value** 1.00e+00

**Overlap** 6  
**Offset** -1  
**Orientation** Normal

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## Alignment



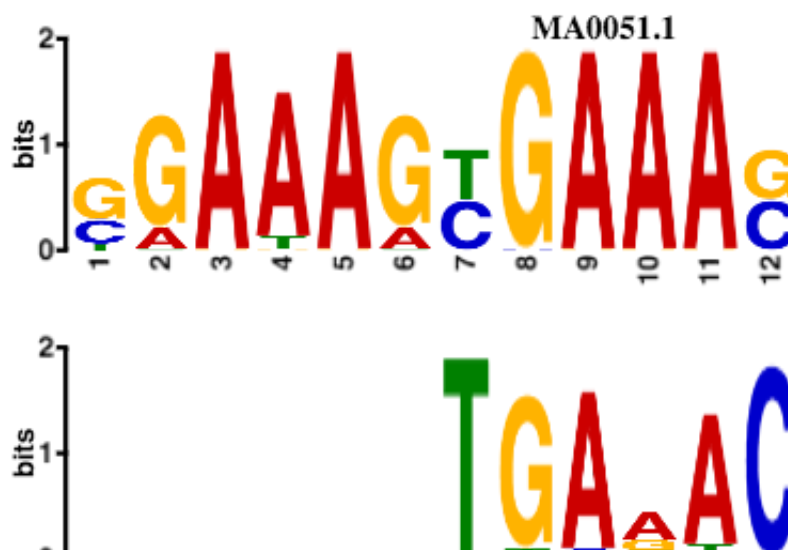
## Summary

**Name** [MA0051.1 \(IRF2\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 4.17e-03  
**E-value** 4.08e+00  
**q-value** 1.00e+00

**Overlap** 10  
**Offset** 6  
**Orientation** Normal

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## Alignment

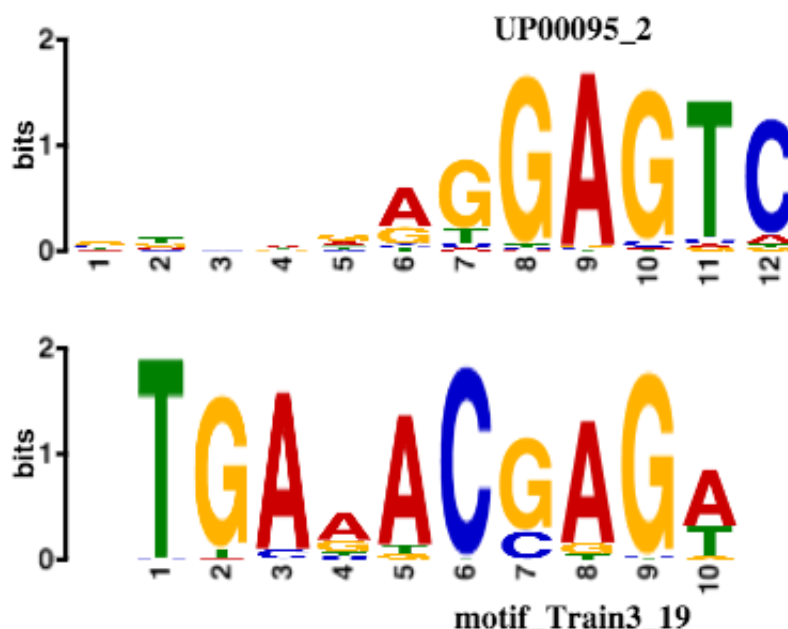


## Summary

## Alignment

<b>Name</b>	<a href="#">UP00095_2</a> (Zfp691_secondary)
<b>Database</b>	uniprobe mouse
<b>p-value</b>	8.24e-03
<b>E-value</b>	8.07e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	1
<b>Orientation</b>	Reverse Complement

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## MATCHES TO MOTIF\_Train8\_2 (1-GTATAAAARSVG)

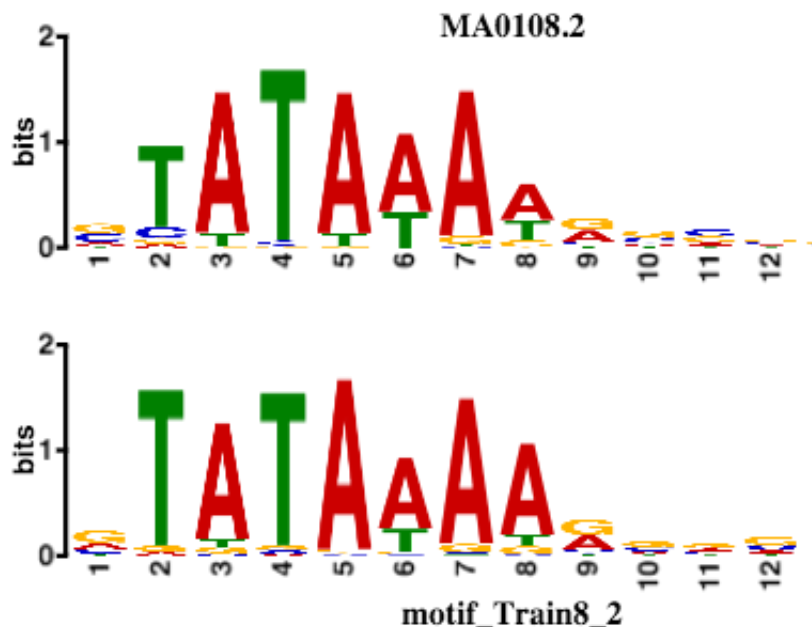
[Previous](#) [Next](#) [Top](#)

## Summary

## Alignment

<b>Name</b>	<a href="#">MA0108.2 (TBP)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	2.86e-08
<b>E-value</b>	2.80e-05
<b>q-value</b>	5.61e-05
<b>Overlap</b>	12
<b>Offset</b>	0
<b>Orientation</b>	Normal

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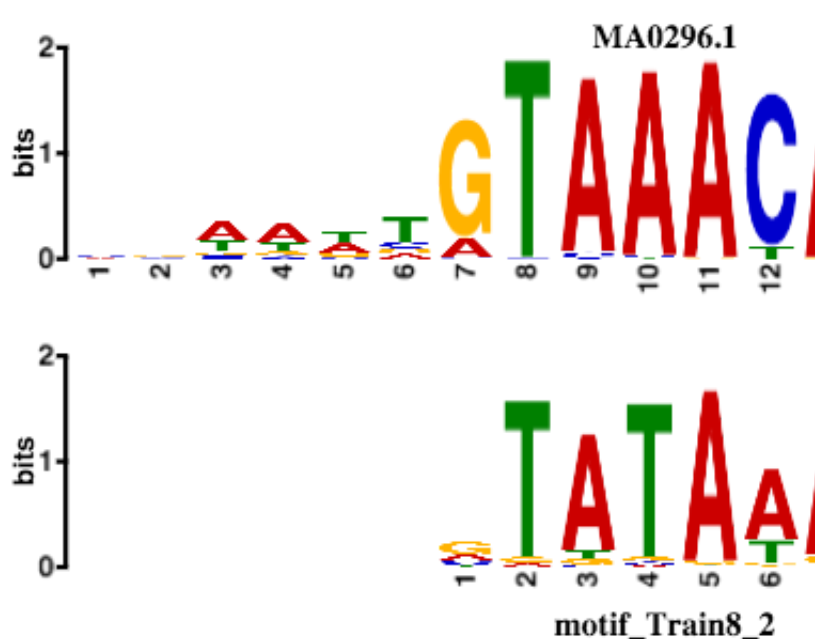
## Summary

## Alignment

**Name** [MA0296.1 \(FKH1\)](#)

<b>Database</b>	JASPAR CORE 2014
<b><i>p</i>-value</b>	2.99e-03
<b><i>E</i>-value</b>	2.93e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	6
<b>Orientation</b>	Normal

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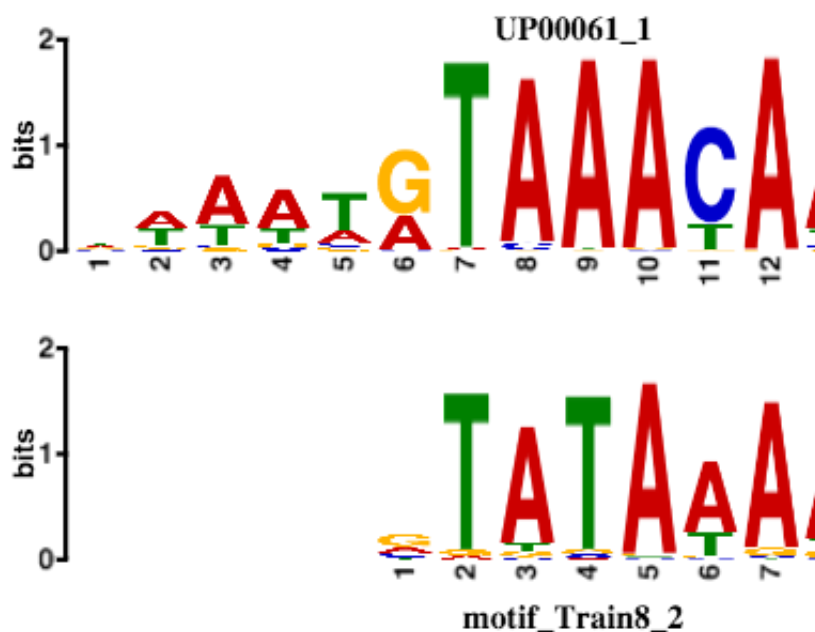


## Summary

<b>Name</b>	<a href="#">UP00061_1</a> (Foxl1_primary)
<b>Database</b>	uniprobe mouse
<b><i>p</i>-value</b>	3.84e-03
<b><i>E</i>-value</b>	3.76e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	5
<b>Orientation</b>	Normal

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## Alignment

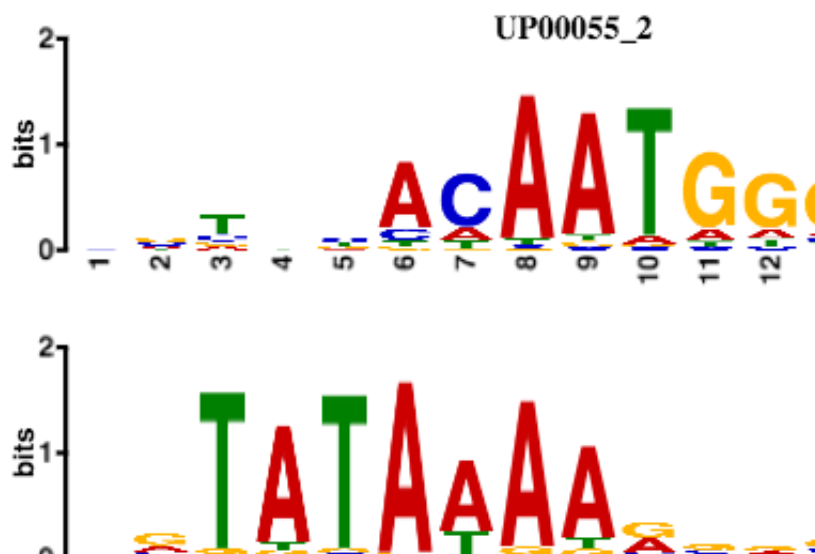


## Summary

<b>Name</b>	<a href="#">UP00055_2</a> (Hbp1_secondary)
<b>Database</b>	uniprobe mouse
<b><i>p</i>-value</b>	4.01e-03
<b><i>E</i>-value</b>	3.93e+00
<b><i>q</i>-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	1
<b>Orientation</b>	Reverse Complement

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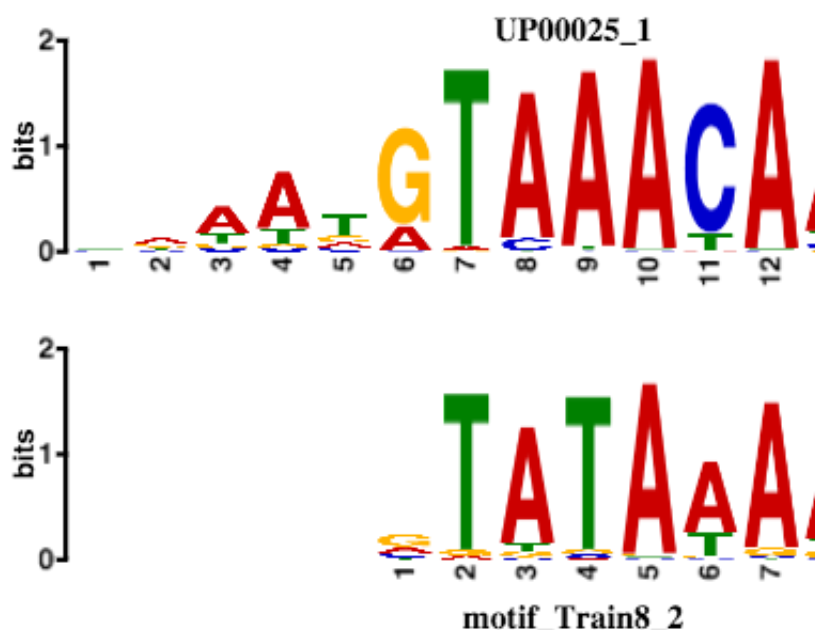
## Alignment



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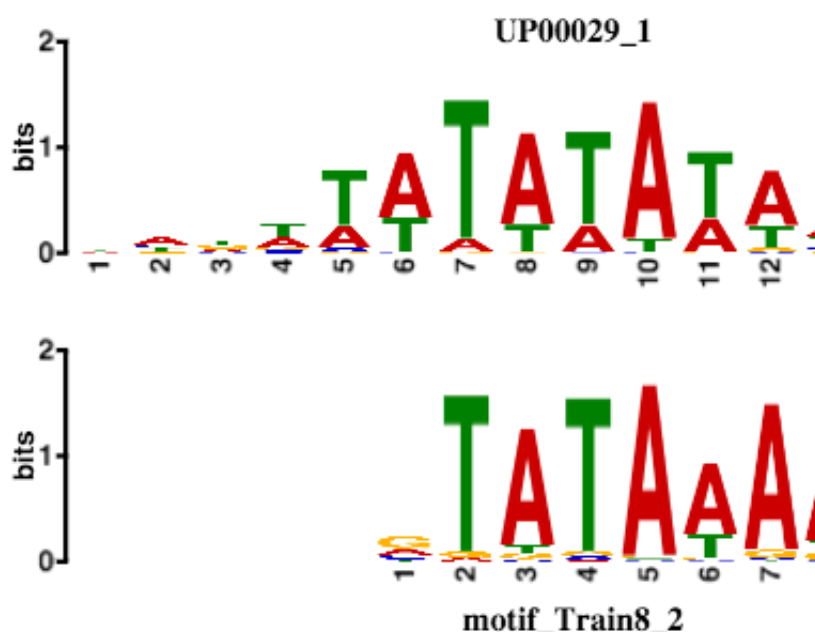
**Name** [UP00025\\_1 \(Foxk1\\_primary\)](#)  
**Database** uniprobe mouse  
**p-value** 5.01e-03  
**E-value** 4.91e+00  
**q-value** 1.00e+00

**Overlap** 12  
**Offset** 5  
**Orientation** Normal

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**Name** [UP00029\\_1 \(Tbp\\_primary\)](#)  
**Database** uniprobe mouse  
**p-value** 6.98e-03  
**E-value** 6.84e+00  
**q-value** 1.00e+00

**Overlap** 11  
**Offset** 5  
**Orientation** Reverse Complement

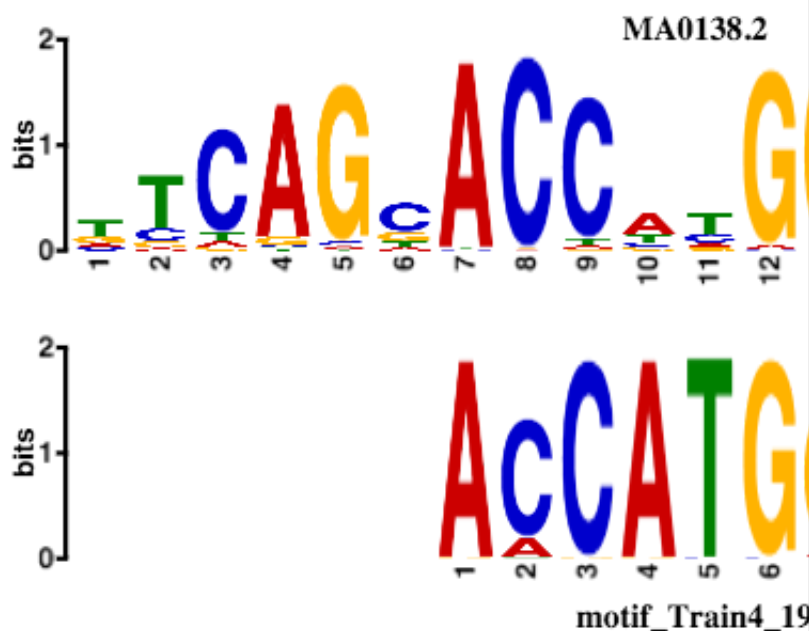
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**Name** [MA0138.2 \(REST\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 3.79e-06

**Alignment**

**E-value** 3.71e-03  
**q-value** 7.42e-03  
**Overlap** 8  
**Offset** 6  
**Orientation** Normal

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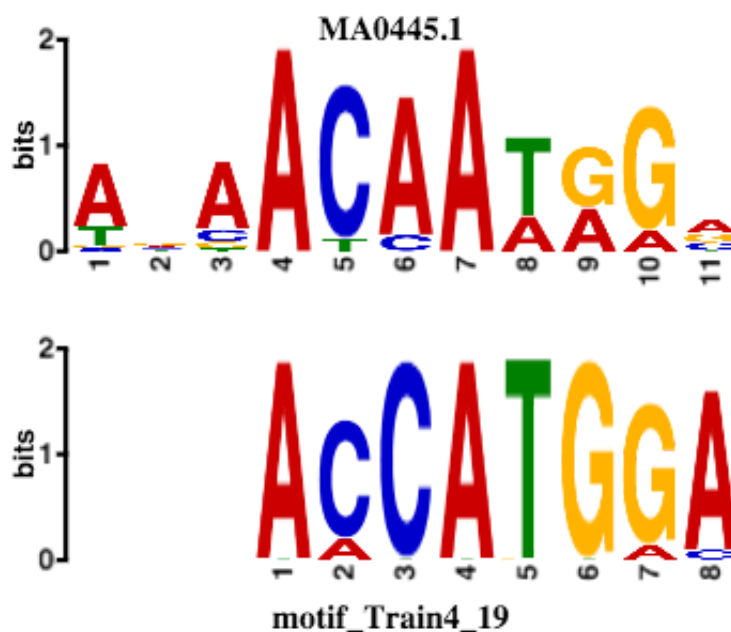
## Summary

**Name** [MA0445.1 \(D\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 1.21e-03  
**E-value** 1.18e+00  
**q-value** 9.31e-01

**Overlap** 8  
**Offset** 3  
**Orientation** Reverse Complement

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## Alignment



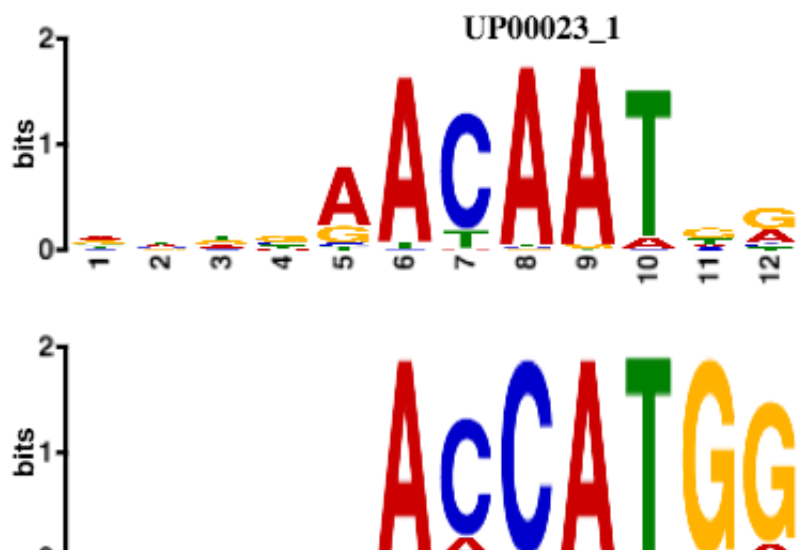
## Summary

**Name** [UP00023\\_1](#)  
**Database** uniprobe mouse  
**p-value** 1.64e-03  
**E-value** 1.60e+00  
**q-value** 9.31e-01

**Overlap** 8  
**Offset** 5  
**Orientation** Normal

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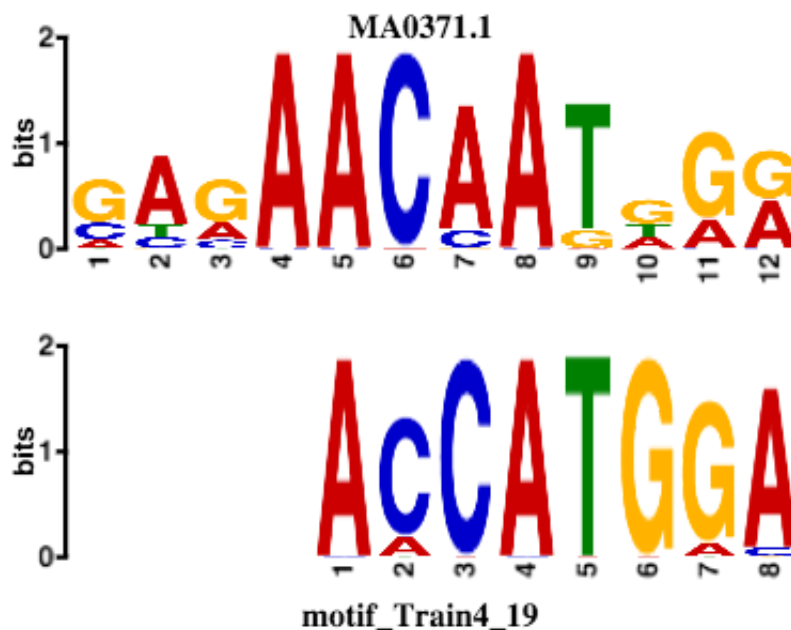
## Alignment



## Summary

<b>Name</b>	<a href="#">MA0371.1 (ROX1)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	1.90e-03
<b>E-value</b>	1.86e+00
<b>q-value</b>	9.31e-01
<b>Overlap</b>	8
<b>Offset</b>	4
<b>Orientation</b>	Reverse Complement
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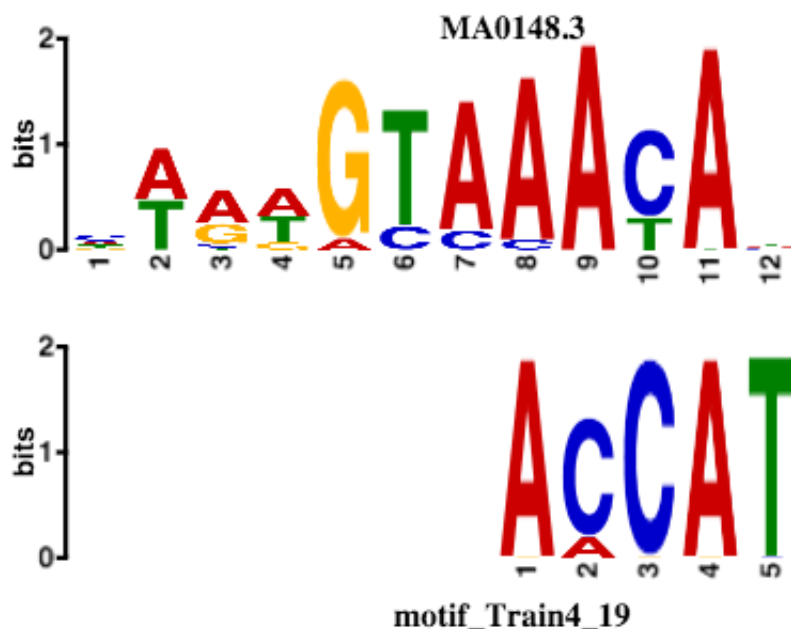
## Alignment



## Summary

<b>Name</b>	<a href="#">MA0148.3 (FOXA1)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	2.81e-03
<b>E-value</b>	2.76e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	8
<b>Offset</b>	7
<b>Orientation</b>	Reverse Complement
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## Alignment



## Summary

<b>Name</b>	<a href="#">UP00004_2</a> <a href="#">(Sox14 secondary)</a>
<b>Database</b>	uniprobe mouse
<b>p-value</b>	5.39e-03
<b>E-value</b>	5.28e+00

## Alignment



**E-value** 5.20e+00  
**q-value** 1.00e+00

**Overlap** 8  
**Offset** 5  
**Orientation** Normal

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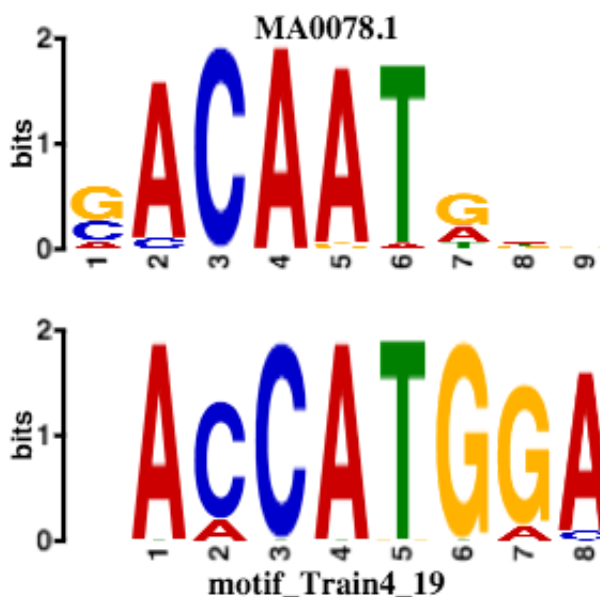
## Summary

**Name** [MA0078.1 \(Sox17\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 5.89e-03  
**E-value** 5.77e+00  
**q-value** 1.00e+00

**Overlap** 8  
**Offset** 1  
**Orientation** Reverse Complement

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## Alignment



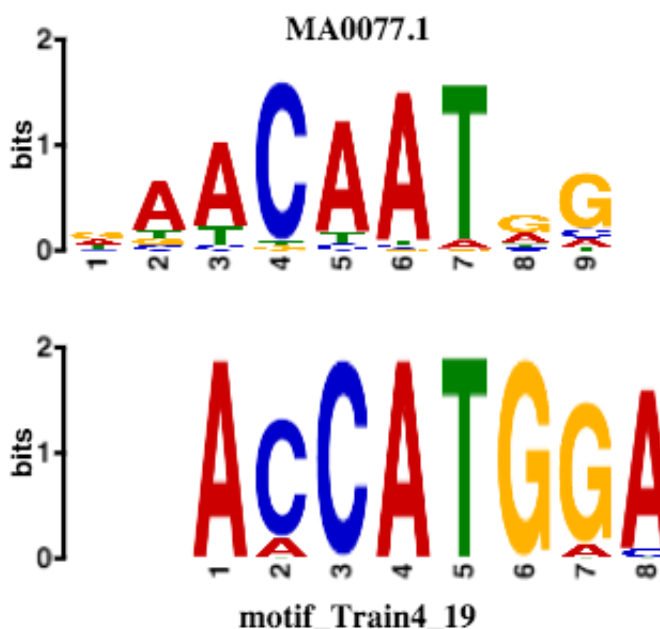
## Summary

**Name** [MA0077.1 \(SOX9\)](#)  
**Database** JASPAR CORE 2014  
**p-value** 5.89e-03  
**E-value** 5.77e+00  
**q-value** 1.00e+00

**Overlap** 7  
**Offset** 2  
**Orientation** Reverse Complement

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## Alignment



## Summary

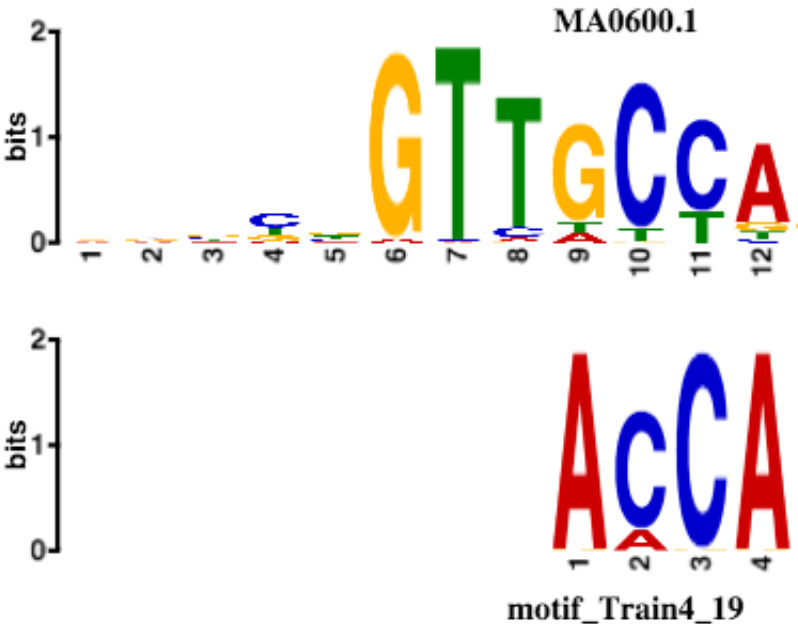
**Name** [MA0600.1 \(RFX2\)](#)  
**Database** JASPAR CORE

## Alignment



Database	2014
p-value	7.22e-03
E-value	7.07e+00
q-value	1.00e+00
Overlap	8
Offset	8
Orientation	Reverse Complement

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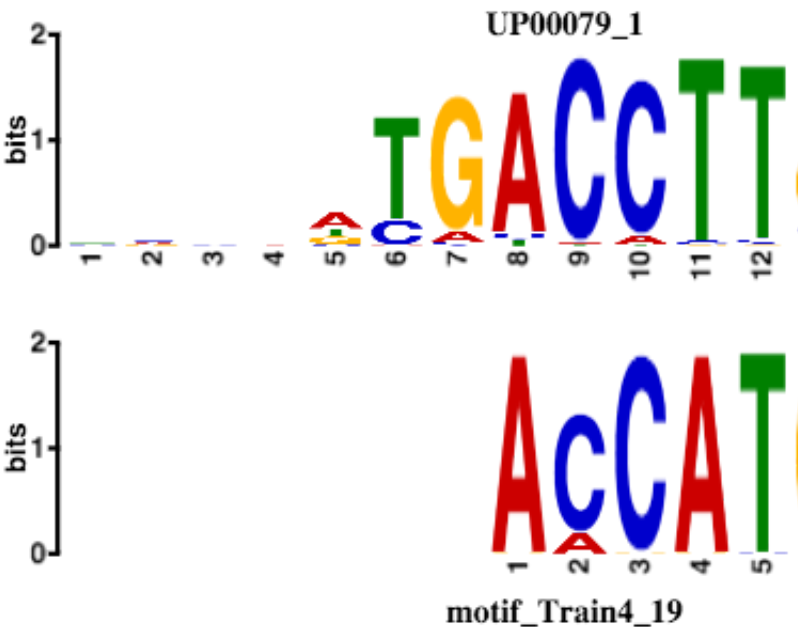


Summary

Name	<a href="#">UP00079_1 (Esrra_primary)</a>
Database	uniprobe mouse
p-value	7.75e-03
E-value	7.59e+00
q-value	1.00e+00
Overlap	8
Offset	7
Orientation	Reverse Complement

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Alignment



Summary

Name	<a href="#">MA0143.3 (Sox2)</a>
Database	JASPAR CORE 2014

Alignment

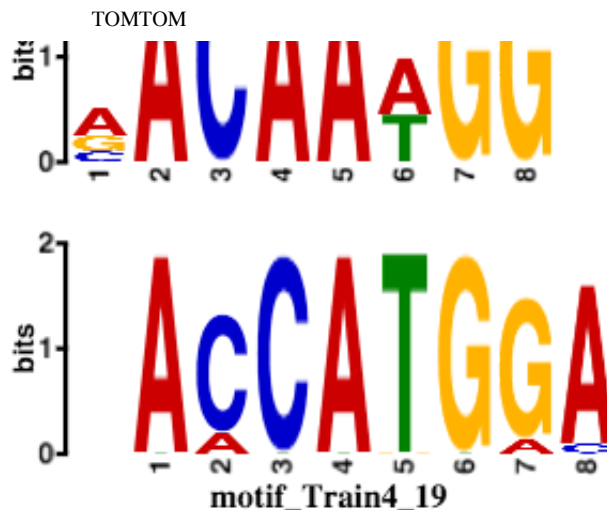


2014

**p-value** 8.09e-03  
**E-value** 7.92e+00  
**q-value** 1.00e+00

**Overlap** 7  
**Offset** 1  
**Orientation** Reverse  
 Complement

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## Summary

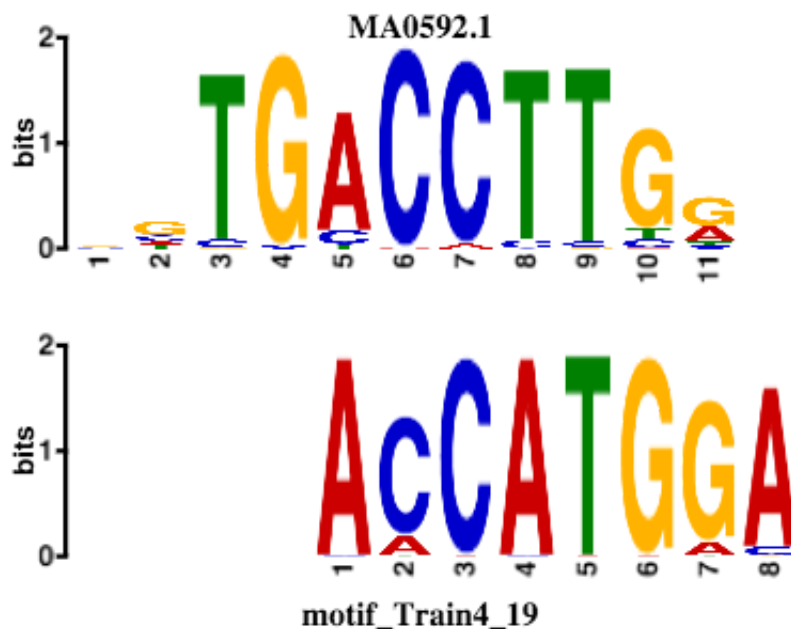
## Alignment

**Name** [MA0592.1 \(ESRRA\)](#)  
**Database** JASPAR CORE  
 2014

**p-value** 8.99e-03  
**E-value** 8.80e+00  
**q-value** 1.00e+00

**Overlap** 7  
**Offset** 4  
**Orientation** Reverse  
 Complement

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## Summary

## Alignment

**Name** [MA0515.1 \(Sox6\)](#)  
**Database** JASPAR CORE  
 2014

**p-value** 9.20e-03  
**E-value** 9.00e+00



**q-value** 1.00e+00

**Overlap** 7

**Offset** 3

**Orientation** Reverse Complement

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## Summary

**Name** [MA0509.1 \(Rfx1\)](#)

**Database** JASPAR CORE 2014

**p-value** 9.64e-03

**E-value** 9.44e+00

**q-value** 1.00e+00

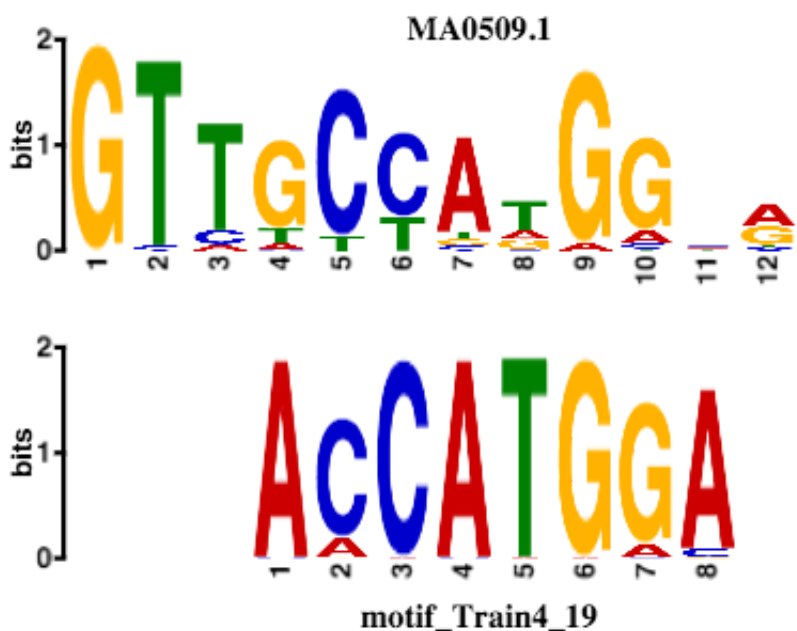
**Overlap** 8

**Offset** 3

**Orientation** Reverse Complement

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## Alignment



## MATCHES TO MOTIF\_10\_1 (1-RTATAAAAASVS)

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

## Summary

**Name** [MA0108.2 \(TBP\)](#)

**Database** JASPAR CORE 2014

**p-value** 1.60e-09

**E-value** 1.56e-06

**q-value** 3.13e-06

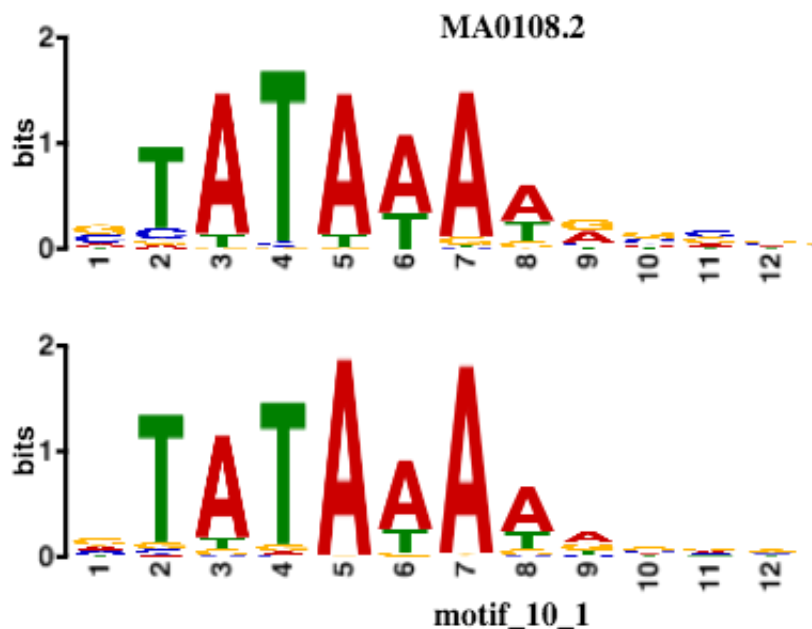
**Overlap** 12

**Offset** 0

**Orientation** Normal

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## Alignment



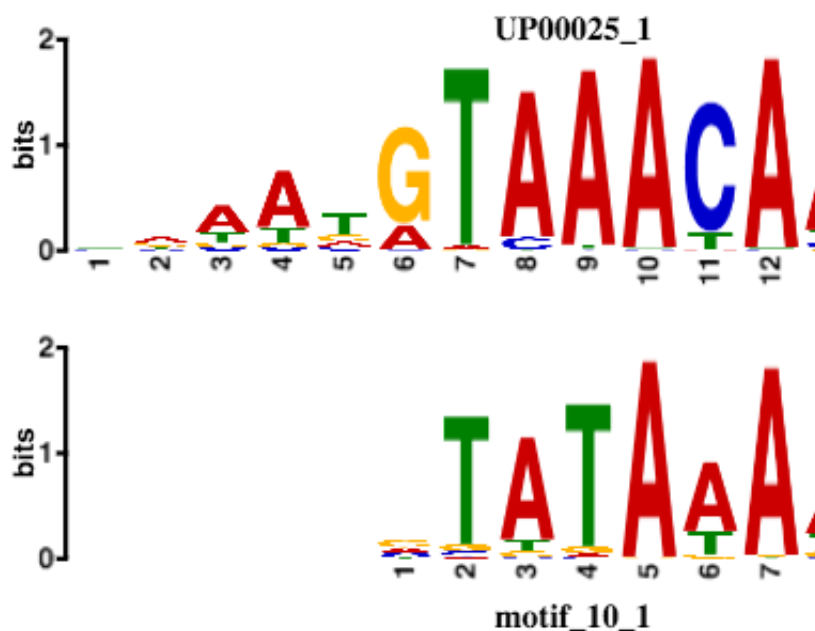
## Summary

## Alignment

**Name** [UP00025\\_1](#)  
[\(Foxk1\\_primary\)](#)  
**Database** uniprobe mouse  
**p-value** 1.60e-03  
**E-value** 1.57e+00  
**q-value** 8.34e-01

**Overlap** 12  
**Offset** 5  
**Orientation** Normal

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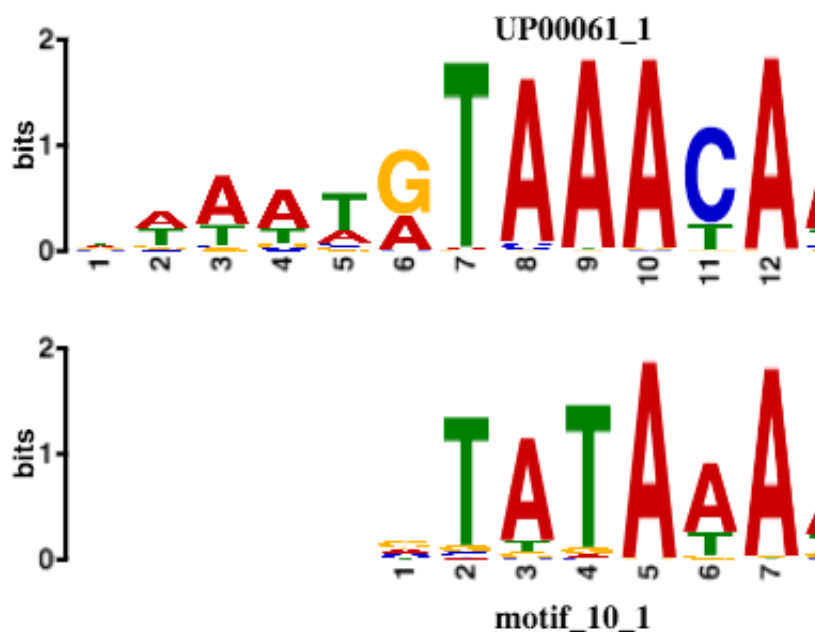
## Summary

**Name** [UP00061\\_1](#)  
[\(Foxl1\\_primary\)](#)  
**Database** uniprobe mouse  
**p-value** 1.60e-03  
**E-value** 1.57e+00  
**q-value** 8.34e-01

**Overlap** 12  
**Offset** 5  
**Orientation** Normal

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## Alignment



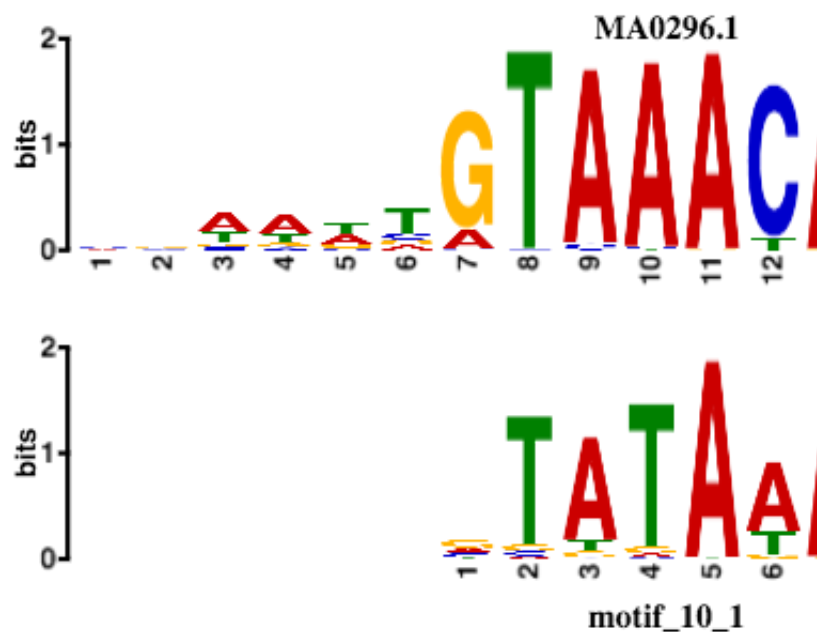
## Summary

**Name** [MA0296.1 \(FKH1\)](#)  
**Database** JASPAR CORE  
 2014  
**p-value** 1.70e-03  
**E-value** 1.67e+00  
**q-value** 8.34e-01

**Overlap** 12  
**Offset** 6  
**Orientation** Normal

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## Alignment

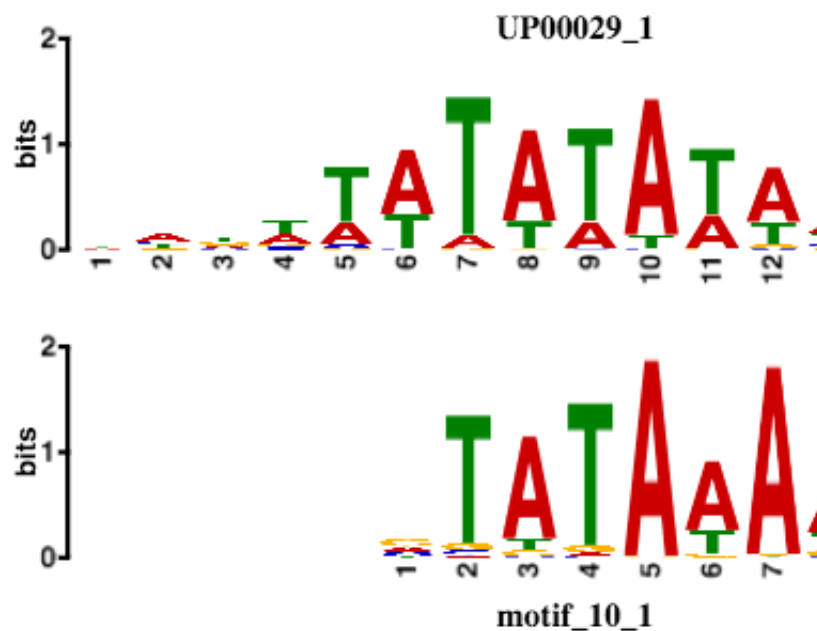


### Summary

<b>Name</b>	<a href="#">UP00029_1</a> (Tbp_primary)
<b>Database</b>	uniprobe mouse
<b>p-value</b>	3.01e-03
<b>E-value</b>	2.94e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	11
<b>Offset</b>	5
<b>Orientation</b>	Reverse Complement

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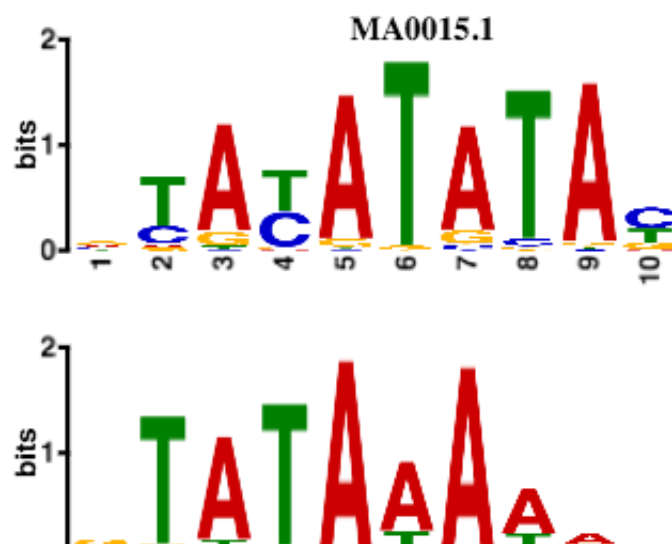
### Alignment



### Summary

<b>Name</b>	<a href="#">MA0015.1</a> (Cf2_II)
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	4.76e-03
<b>E-value</b>	4.66e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	0
<b>Orientation</b>	Reverse Complement

### Alignment



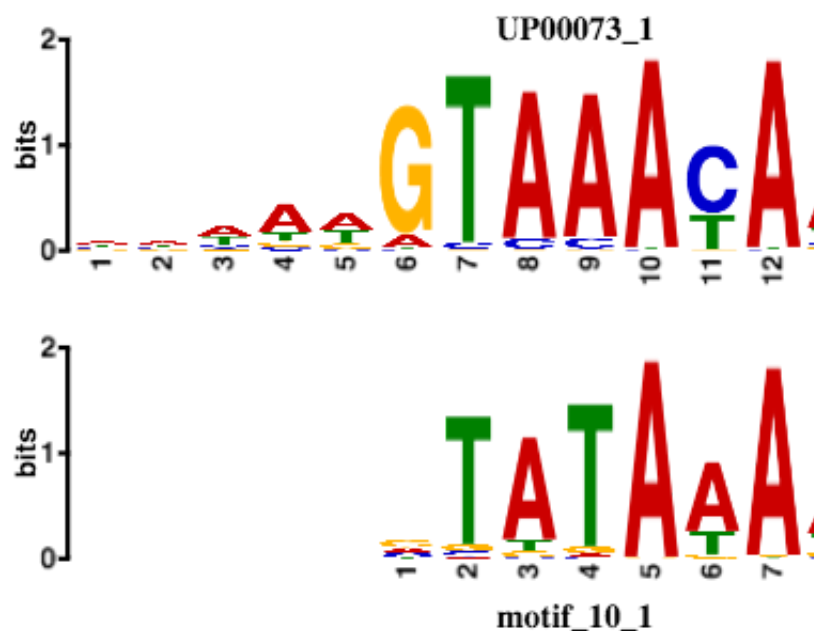
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## Summary

<b>Name</b>	<a href="#">UP00073_1</a> (Foxa2_primary)
<b>Database</b>	uniprobe mouse
<b>p-value</b>	5.94e-03
<b>E-value</b>	5.82e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	5
<b>Orientation</b>	Normal

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## Alignment

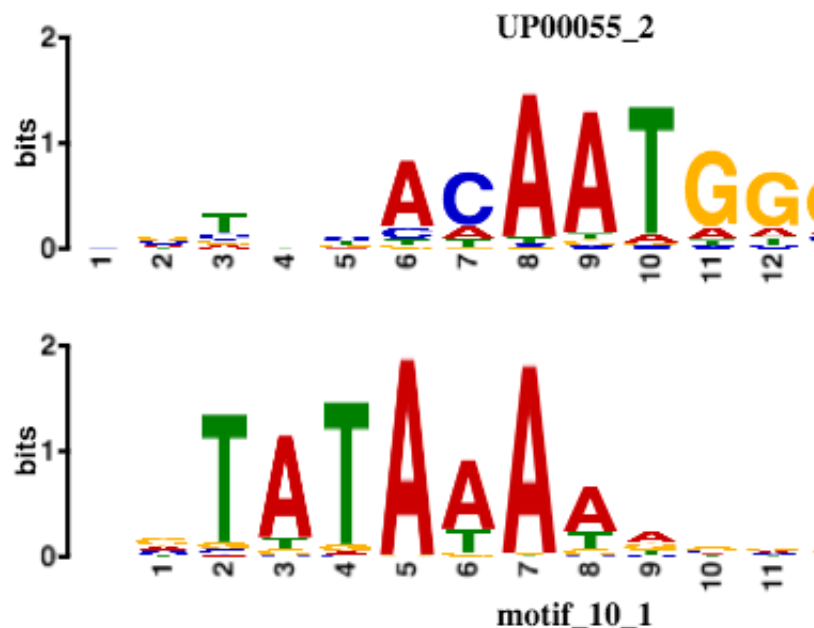


## Summary

<b>Name</b>	<a href="#">UP00055_2</a> (Hbp1_secondary)
<b>Database</b>	uniprobe mouse
<b>p-value</b>	7.69e-03
<b>E-value</b>	7.52e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	12
<b>Offset</b>	1
<b>Orientation</b>	Reverse Complement

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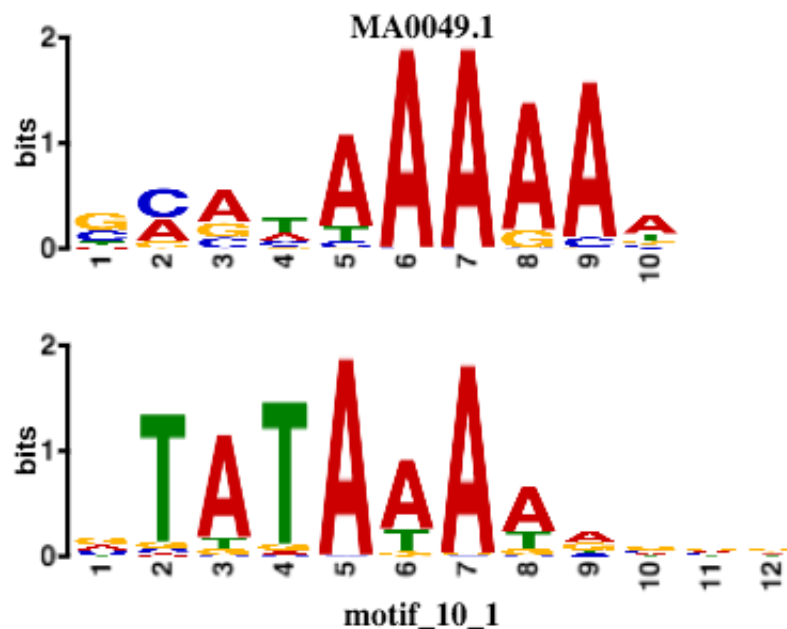
## Alignment



## Summary

## Alignment

<b>Name</b>	<a href="#">MA0049.1 (hb)</a>
<b>Database</b>	JASPAR CORE 2014
<b>p-value</b>	1.01e-02
<b>E-value</b>	9.89e+00
<b>q-value</b>	1.00e+00
<b>Overlap</b>	10
<b>Offset</b>	0
<b>Orientation</b>	Normal
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## 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

<b>Strand Handling</b>	Motifs may be reverse complemented before comparison to find a better match.
<b>Distance Measure</b>	Pearson correlation coefficient
<b>Match Threshold</b>	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 -eval -thresh 10.0 motif_group7.txt
```

Result calculation took 9.832 seconds





