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Search...

Cross species **NEW**

Promoter analysis

Homologous network

Network comparison

Rank by Unique TFBS

Rank genes

Rank by TF

Rank by GO

Regulatory Associations

Search for TFs

Search for Genes

Search for Associations

Pattern Matching

Search by DNA Motif

Find TF Binding Site(s)

Utilities

Search for Orthologs **NEW**

IUPAC Code Generation

ORF List ↔ Gene List

Retrieve

TF-Consensus List

Regulation Matrix

Upstream Sequence

About

Contact/Credits

How to cite

Acknowledgments

Search Transcription Factors

Yap5p-935

Yap7p-708

Rpn4p-673

Rpn4p-628

Yap6p-411

Yap7p-383

Yap6p-366

Rpn4p-282

☒ Yap1p

☒ Yap7p

☒ Yap6p

☒ Rpn4p

☒ Yap5p

☒ Rpn4p

Select: [All](#) [None](#) [Common](#) ☐ Consider only documented TFs

Redisplay

Transcription Factor	Consensus	Position	Strand
Target Sequence: CTA1pr1kb (size 1000)			
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Yap1p	TTACAAA	-308	F
Yap1p Yap7p	TTAGTAA	-756	F
Yap1p Yap7p	TTAGTAA	-383	F
Yap1p	VTTACWAAB	-309	F
Yap1p	VTTACWAAB	-375	R
Yap1p	VTTACWAAB	-748	R
Yap1p	VDTASTAA	-757	F
Yap1p	VDTASTAA	-384	F
Yap1p	VDTASTAA	-375	R
Yap1p	VDTASTAA	-748	R
Yap6p	AATKACV	-411	F
Yap6p	AATKACV	-310	F
Yap5p	TTACKAA	-376	R
Yap5p	TTACKAA	-749	R
Yap5p	ATTACTA	-748	R
Yap7p	TTACTAA	-376	R
Yap7p	TTACTAA	-749	R
Yap7p	ATTACTA	-748	R
Yap7p	MTKASTMA	-757	F
Yap7p	MTKASTMA	-384	F
Yap7p	MTKASTMA	-748	R
Yap7p	DATTACTAAD	-747	R
Yap7p	TTASTAAK	-756	F
Yap7p	TTASTAAK	-376	R
Yap7p	TTASTAAK	-749	R
Yap7p	KTACTAAD	-376	R
Yap7p	KTACTAAD	-749	R
Hap5p	CCAAT	-935	F
Rpn4p	TTGCAAA	-327	R

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WSXHTML 1.0

WSXCSS 2.0

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pathoyeasttract.org/cglabrata/findregulators.php

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