




















































































Motifs Found

+		
+		A-box
+		AAGAA-motif
+		ABRE
+		ABRE3a
+		ABRE4
+		ARE
+		ATCT-motif
+		AT~TATA-box
+		Box 4
+		CAAT-box
+		CCGTCC motif
+		CCGTCC-box
+		CGTCA-motif
+		CTAG-motif
+		DRE1
+		G-box
+		GATA-motif
+		GCN4_motif
+		GT1-motif
+		I-box
+		LAMP-element
+		MBS
+		MRE
+		MYB
+		MYB-like sequence
+		MYC
+		Myb
+		Myb-binding site
+		O2-site
+		P-box
+		TATA
+		TATA-box
+		TCA
+		TCA-element
+		TCCC-motif
+		TGACG-motif
+		Unnamed__1
+		Unnamed__16
+		Unnamed__4
+		Unnamed__6
+		W box








































- +  WRE3
- +  WUN-motif
- +  Y-box
- +  as-1

Motifs Found









































- + 
- +  4cl-CMA2b
- +  AAGAA-motif
- +  ABRE
- +  ABRE3a
- +  ABRE4
- +  ARE
- +  AT-rich sequence
- +  ATC-motif
- +  AT~TATA-box
- +  Box 4
- +  CAAT-box
- +  CARE
- +  CAT-box
- +  CGTCA-motif
- +  G-box
- +  GC-motif
- +  GCN4_motif
- +  GT1-motif
- +  LAMP-element
- +  LTR
- +  MBS
- +  MBSI
- +  MRE
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  Myb-binding site
- +  Myc
- +  STRE
- +  TATA-box
- +  TC-rich repeats
- +  TCA-element
- +  TCT-motif
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__2
- +  Unnamed__4
- +  W box
- +  WUN-motif
- +  as-1


+  chs-Unit 1 ml

Motifs Found






































- + 
- +  AAGAA-motif
- +  ABRE
- +  ARE
- +  ATC-motif
- +  AT~TATA-box
- +  Box 4
- +  Box III
- +  CAAT-box
- +  CGTCA-motif
- +  DRE core
- +  ERE
- +  G-Box
- +  G-box
- +  GARE-motif
- +  GT1-motif
- +  I-box
- +  MRE
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  Myb-binding site
- +  O2-site
- +  STRE
- +  TATA-box
- +  TATC-box
- +  TC-rich repeats
- +  TCA-element
- +  TCCC-motif
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  Unnamed__6
- +  W box
- +  WRE3
- +  as-1
- +  box S
- +  chs-CMA1a

Motifs Found









































+		
+		A-box
+		AAGAA-motif
+		ABRE
+		AC-I
+		ARE
+		ATCT-motif
+		AT~TATA-box
+		Box 4
+		CAAT-box
+		CAT-box
+		CCAAT-box
+		CCGTCC motif
+		CCGTCC-box
+		CGTCA-motif
+		ERE
+		G-Box
+		GA-motif
+		GATA-motif
+		GT1-motif
+		LTR
+		MRE
+		MYB
+		MYB recognition site
+		MYB-like sequence
+		Myc
+		O2-site
+		P-box
+		STRE
+		TATA
+		TATA-box
+		TC-rich repeats
+		TCT-motif
+		TGA-element
+		TGACG-motif
+		Unnamed__2
+		Unnamed__4
+		W box
+		WRE3
+		WUN-motif
+		as-1
+		box S

+  chs-Unit 1 ml









































Motifs Found



- + 
- +  AAGAA-motif
- +  AE-box
- +  AP-1
- +  ARE
- +  AT~TATA-box
- +  Box 4
- +  Box II -like sequence
- +  CAAT-box
- +  CAT-box
- +  CGTCA-motif
- +  CTAG-motif
- +  DRE core
- +  GT1-motif
- +  MBS
- +  MRE
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  P-box
- +  STRE
- +  TATA-box
- +  TATC-box
- +  TCCC-motif
- +  TCT-motif
- +  TGA-element
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  W box
- +  WRE3
- +  WUN-motif
- +  as-1
- +  box S
- +  chs-CMA1a
- +  circadian

Motifs Found









































- + 
- +  3-AF3 binding site
- +  4cl-CMA1b
- +  AAGAA-motif
- +  ABRE
- +  ABRE3a
- +  ABRE4
- +  ACE
- +  ARE
- +  AT-rich sequence
- +  AT~TATA-box
- +  Box 4
- +  CAAT-box
- +  CAT-box
- +  CGTCA-motif
- +  DRE core
- +  ERE
- +  G-Box
- +  G-box
- +  GATA-motif
- +  GCN4_motif
- +  GT1-motif
- +  LTR
- +  MRE
- +  MSA-like
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  STRE
- +  TATA
- +  TATA-box
- +  TC-rich repeats
- +  TCA
- +  TCA-element
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  Unnamed__6
- +  as-1
- +  dOCT


Motifs Found

+		
+		AAGAA-motif
+		ABRE
+		ABRE3a
+		ABRE4
+		AE-box
+		ARE
+		ATC-motif
+		AT~ABRE
+		AT~TATA-box
+		CAAT-box
+		CAT-box
+		CGTCA-motif
+		DRE core
+		DRE1
+		G-box
+		GA-motif
+		GT1-motif
+		Gap-box
+		I-box
+		LTR
+		MBS
+		MRE
+		MYB
+		MYB-like sequence
+		MYC
+		Myb
+		Myb-binding site
+		O2-site
+		P-box
+		STRE
+		TATA-box
+		TC-rich repeats
+		TCA
+		TCT-motif
+		TGA-element
+		TGACG-motif
+		Unnamed__1
+		Unnamed__4
+		W box
+		WRE3
+		WUN-motif




































- +  as-1
- +  circadian

Motifs Found



































+		
+		A-box
+		AAGAA-motif
+		ABRE
+		ABRE3a
+		ABRE4
+		AC-II
+		ARE
+		AT-rich element
+		AT~TATA-box
+		Box 4
+		CAAT-box
+		CAT-box
+		CCGTCC motif
+		CCGTCC-box
+		G-Box
+		G-box
+		GA-motif
+		GC-motif
+		GT1-motif
+		LAMP-element
+		LTR
+		MRE
+		MYB
+		MYB-like sequence
+		MYC
+		Myb
+		P-box
+		STRE
+		TATA
+		TATA-box
+		TATC-box
+		TCA-element
+		TCT-motif
+		TGA-element
+		Unnamed__1
+		Unnamed__2
+		Unnamed__4
+		W box
+		WRE3
+		WUN-motif
+		box S

+  chs-CMA1a





































Motifs Found

- + 
- +  AAGAA-motif
- +  ABRE
- +  ARE
- +  AT~TATA-box
- +  Box 4
- +  CAAT-box
- +  CAT-box
- +  CGTCA-motif
- +  ERE
- +  G-Box
- +  G-box
- +  GA-motif
- +  GATA-motif
- +  GT1-motif
- +  LTR
- +  MRE
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  P-box
- +  TATA
- +  TATA-box
- +  TC-rich repeats
- +  TCA
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__2
- +  Unnamed__4
- +  Unnamed__6
- +  W box
- +  WUN-motif
- +  as-1
- +  circadian









































Motifs Found






- + 
- +  AAGAA-motif
- +  ABRE
- +  AP-1
- +  ARE
- +  AT-rich element
- +  Box 4
- +  CAAT-box
- +  CAT-box
- +  CGTCA-motif
- +  DRE1
- +  ERE
- +  G-Box
- +  G-box
- +  GATA-motif
- +  GT1-motif
- +  LTR
- +  MBS
- +  MBSI
- +  MYB
- + MYB-like sequence
- +  MYC
- +  Myb
- +  Myb-binding site
- +  O2-site
- +  P-box
- +  STRE
- +  TATA-box
- +  TCA
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  Unnamed__6
- +  WUN-motif
- +  as-1

Motifs Found











































+		
+		A-box
+		ABRE
+		ACE
+		ARE
+		AT-rich element
+		ATCT-motif
+		AT~TATA-box
+		Box 4
+		CAAT-box
+		CCAAT-box
+		CCGTCC motif
+		CCGTCC-box
+		DRE1
+		ERE
+		G-Box
+		G-box
+		GATA-motif
+		GT1-motif
+		I-box
+		LTR
+		MRE
+		MYB
+		MYB recognition site
+		MYC
+		Myb
+		Myb-binding site
+		STRE
+		TATA-box
+		TATC-box
+		TCT-motif
+		TGA-element
+		Unnamed__4
+		Unnamed__6
+		WRE3
+		WUN-motif
+		chs-CMA1a














Motifs Found

+		
+		AAGAA-motif
+		ABRE
+		ABRE3a
+		ABRE4
+		ACE
+		AE-box
+		ARE
+		AT~TATA-box
+		AuxRR-core
+		Box 4
+		Box II
+		CAAT-box
+		CGTCA-motif
+		ERE
+		G-box
+		GARE-motif
+		GATA-motif
+		GT1-motif
+		LTR
+		MBS
+		MRE
+		MYB
+		MYB-like sequence
+		MYC
+		Myb
+		Myb-binding site
+		O2-site
+		RY-element
+		STRE
+		TATA-box
+		TC-rich repeats
+		TCA
+		TCT-motif
+		TGA-element
+		TGACG-motif
+		Unnamed__1
+		Unnamed__10
+		Unnamed__12
+		Unnamed__14
+		Unnamed__4
+		Unnamed__8






































- +  W box
- +  WRE3
- +  as-1
- +  box S
- +  chs-CMA2a

Motifs Found








































+		
+		3-AF1 binding site
+		A-box
+		AAGAA-motif
+		ABRE
+		ABRE3a
+		ABRE4
+		AE-box
+		ARE
+		AT-rich sequence
+		ATC-motif
+		AT~TATA-box
+		AuxRR-core
+		Box 4
+		Box III
+		CAAT-box
+		CCGTCC motif
+		CCGTCC-box
+		CGTCA-motif
+		ERE
+		F-box
+		G-Box
+		G-box
+		GA-motif
+		GATA-motif
+		GCN4_motif
+		GT1-motif
+		HD-Zip 3
+		I-box
+		LTR
+		MBS
+		MRE
+		MYB
+		MYB-like sequence
+		MYC
+		Myb
+		Myb-binding site
+		O2-site
+		P-box
+		STRE
+		TATA
+		TATA-box

- +  TCA-element
- +  TCT-motif
- +  TGA-box
- +  TGA-element
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__2
- +  Unnamed__4
- +  W box
- +  WUN-motif
- +  as-1
- +  chs-CMA1a
- +  circadian










































Motifs Found





- + 
- +  AAGAA-motif
- +  ABRE
- +  ABRE3a
- +  ABRE4
- +  ARE
- +  AT~TATA-box
- +  Box 4
- +  CAAT-box
- +  CGTCA-motif
- +  DRE core
- +  G-box
- +  GA-motif
- +  GATA-motif
- +  GCN4_motif
- +  GT1-motif
- +  LAMP-element
- +  MBS
- +  MBSI
- +  MRE
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  P-box
- +  TATA
- +  TATA-box
- +  TCA
- +  TCA-element
- +  TGA-element
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  W box
- +  as-1
- +  box S
- +  chs-CMA1a

Motifs Found





































- + 
- +  AAGAA-motif
- +  ABRE
- +  ABRE3a
- +  ABRE4
- +  AT-rich sequence
- +  AT~TATA-box
- +  AuxRR-core
- +  Box 4
- +  CAAT-box
- +  CAT-box
- +  CCAAT-box
- +  CGTCA-motif
- +  G-Box
- +  G-box
- +  GATA-motif
- +  GT1-motif
- +  MBS
- +  MYB
- +  MYB recognition site
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  Myb-binding site
- +  P-box
- +  STRE
- +  TATA-box
- +  TCA
- +  TCA-element
- +  TCT-motif
- +  TGA-element
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  Unnamed__6
- +  W box
- +  WUN-motif
- +  as-1
- +  chs-CMA1a

Motifs Found






































+		
+		ABRE
+		ABRE3a
+		ABRE4
+		AC-I
+		AC-II
+		ARE
+		AT-rich sequence
+		ATCT-motif
+		AT~TATA-box
+		Box 4
+		Box II -like sequence
+		CAAT-box
+		CARE
+		CGTCA-motif
+		CTAG-motif
+		ERE
+		G-Box
+		G-box
+		GARE-motif
+		GT1-motif
+		Gap-box
+		MBS
+		MRE
+		MYB
+		MYC
+		Myb
+		Myb-binding site
+		O2-site
+		P-box
+		SARE
+		STRE
+		TATA
+		TATA-box
+		TC-rich repeats
+		TCA
+		TCA-element
+		TCCC-motif
+		TCT-motif
+		TGACG-motif
+		Unnamed__1
+		Unnamed__4

- +  WRE3
- +  WUN-motif
- +  as-1
- +  chs-Unit 1 m1

Motifs Found

- + 
- +  AAGAA-motif
- +  ABRE
- +  ABRE3a
- +  ABRE4
- +  ARE
- +  AT~TATA-box
- +  Box 4
- +  CAAT-box
- +  CAT-box
- +  CGTCA-motif
- +  DRE core
- +  G-Box
- +  G-box
- +  GA-motif
- +  GARE-motif
- +  GATA-motif
- +  I-box
- +  LAMP-element
- +  MBS
- +  MYB
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  Myb-binding site
- +  Myc
- +  P-box
- +  RY-element
- +  STRE
- +  TATA-box
- +  TCT-motif
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  as-1
- +  circadian

Motifs Found

- + 
- +  ABRE
- +  AE-box
- +  ARE
- +  AT~TATA-box
- +  AuxRR-core
- +  Box 4
- +  CAAT-box
- +  CCAAT-box
- +  CGTCA-motif
- +  ERE
- +  F-box
- +  G-box
- +  GA-motif
- +  GATA-motif
- +  GT1-motif
- +  LTR
- +  MBS
- +  MRE
- +  MYB
- + MYB recognition site
- +  MYB-like sequence
- +  MYC
- +  Myb
- +  STRE
- +  TATA-box
- +  TC-rich repeats
- +  TCA
- +  TCT-motif
- +  TGA-box
- +  TGACG-motif
- +  Unnamed__1
- +  Unnamed__4
- +  W box
- +  WRE3
- +  WUN-motif
- +  as-1
- +  re2f-1