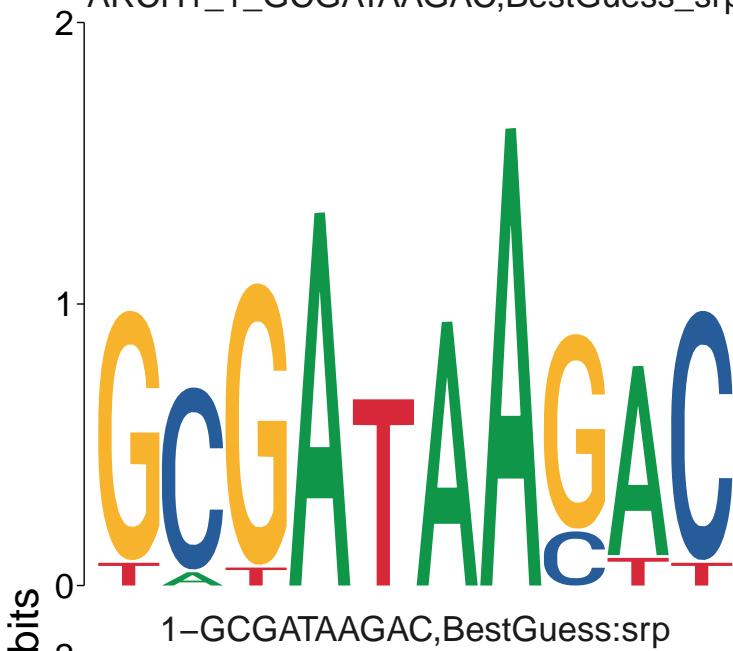
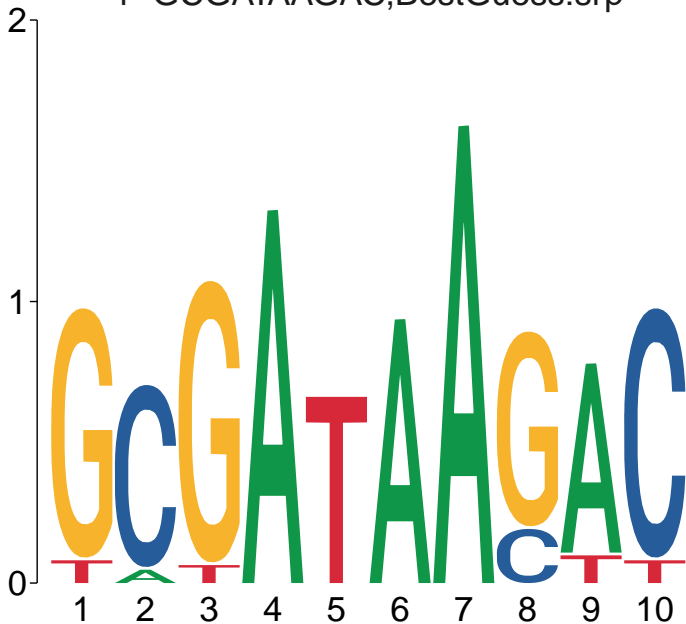


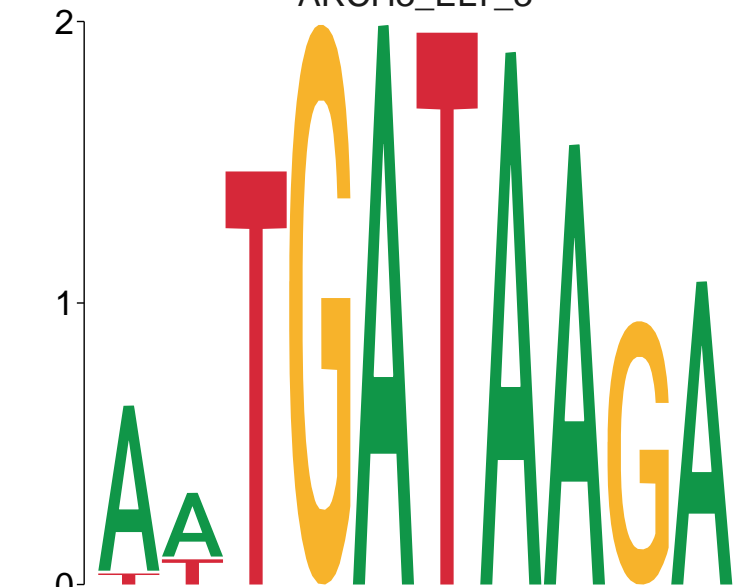
ARCH1_1_GCGATAAGAC,BestGuess_srp



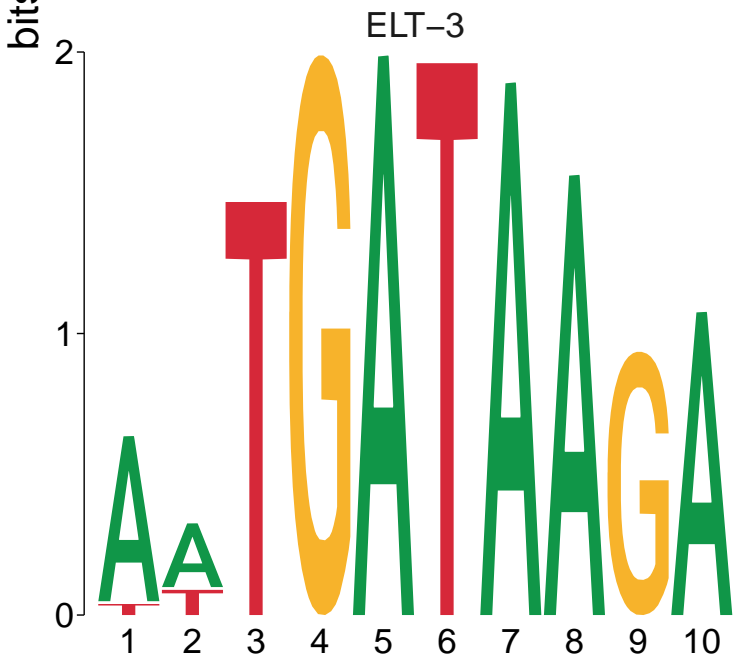
1-GCGATAAGAC,BestGuess:srp



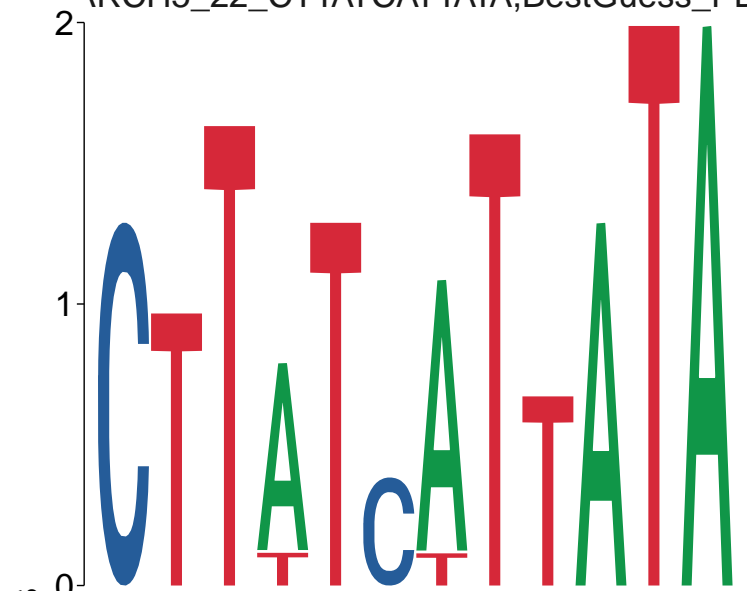
ARCH3_ELT_3



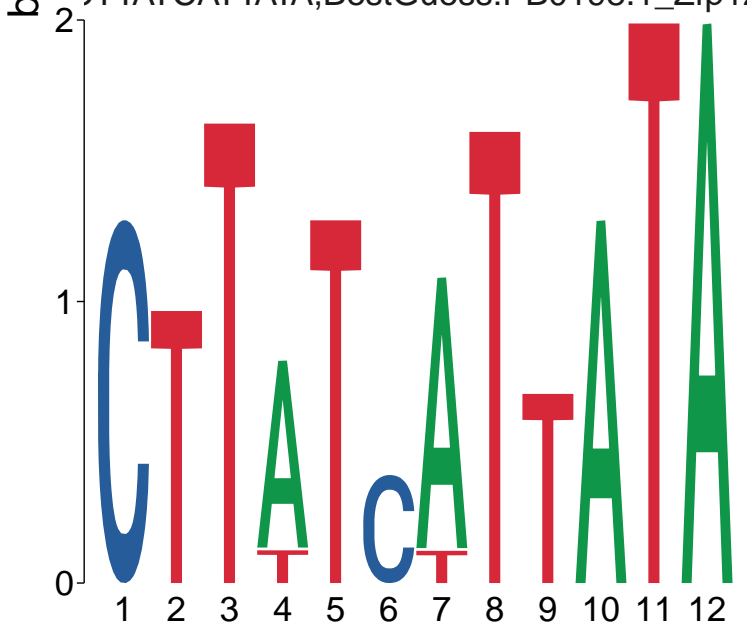
ELT-3



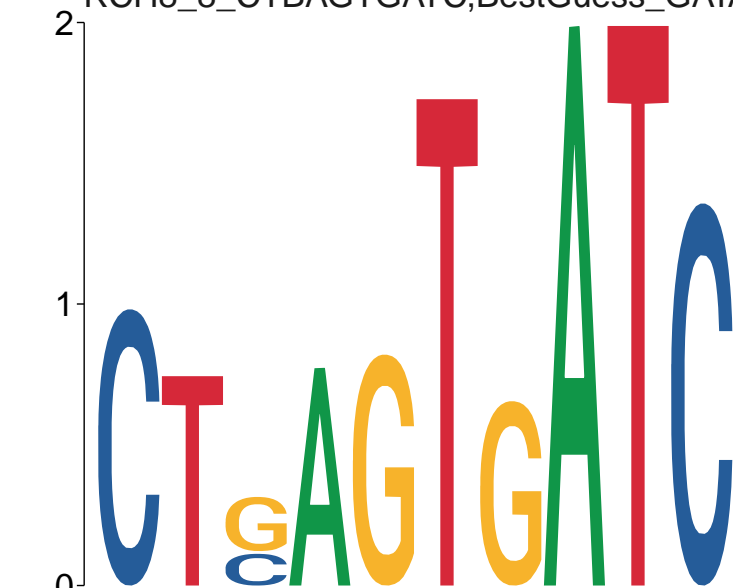
ARCH5_22_CTTATCATTATA,BestGuess_PE



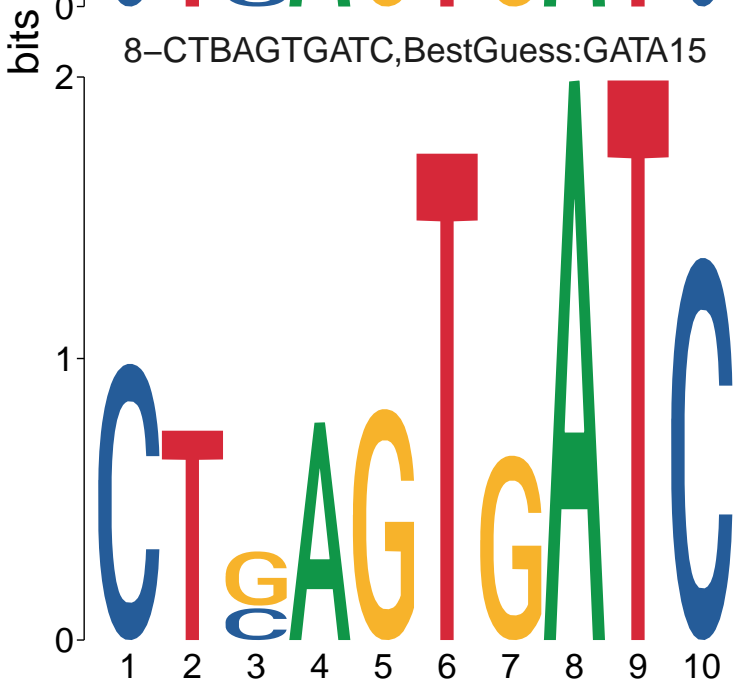
CTTATCATTATA,BestGuess:PB0198.1_Zfp1



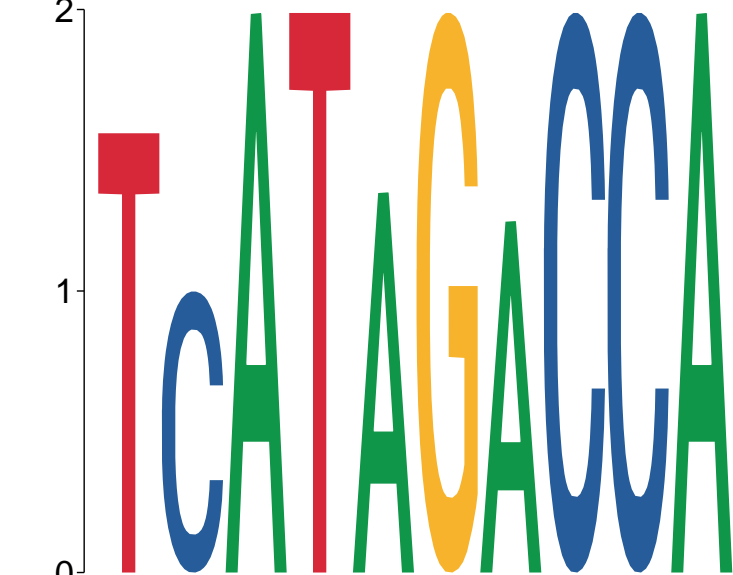
RCH8_8_CT BAGTGATC, BestGuess_GATA



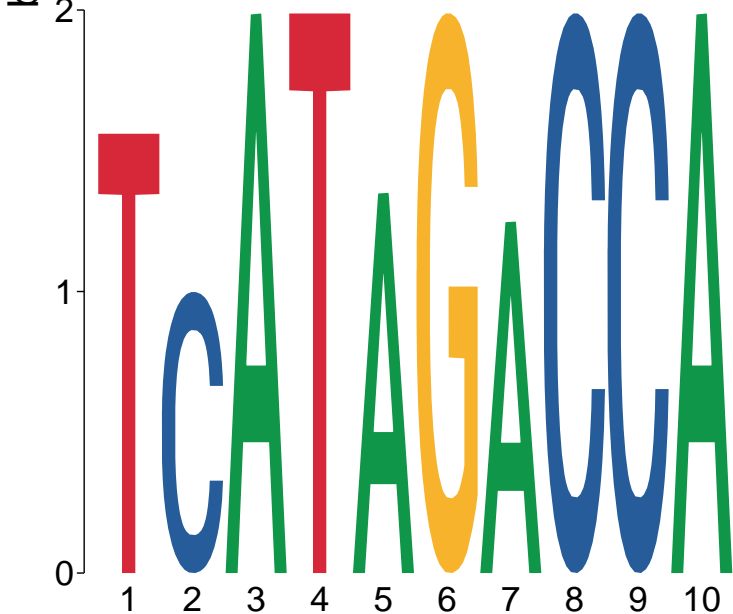
8-CT BAGTGATC, BestGuess:GATA15



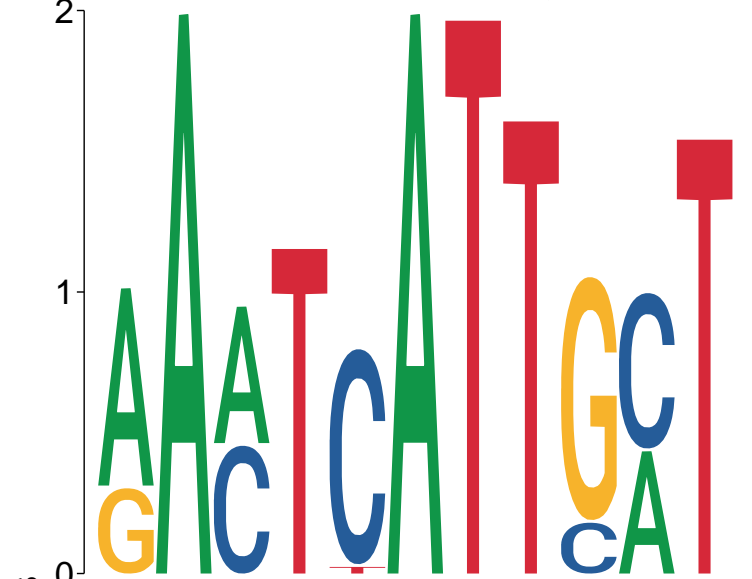
ARCH9_7_TCATAGACCA,BestGuess_NFYF



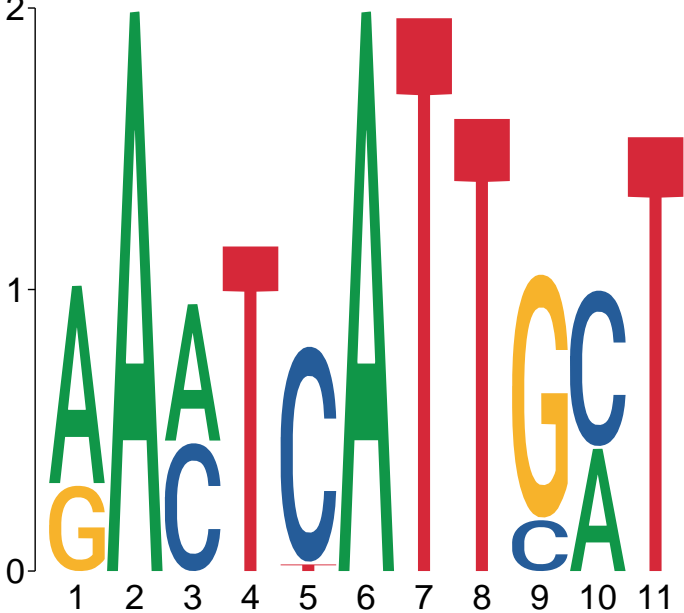
7-TCATAGACCA,BestGuess:NFYB



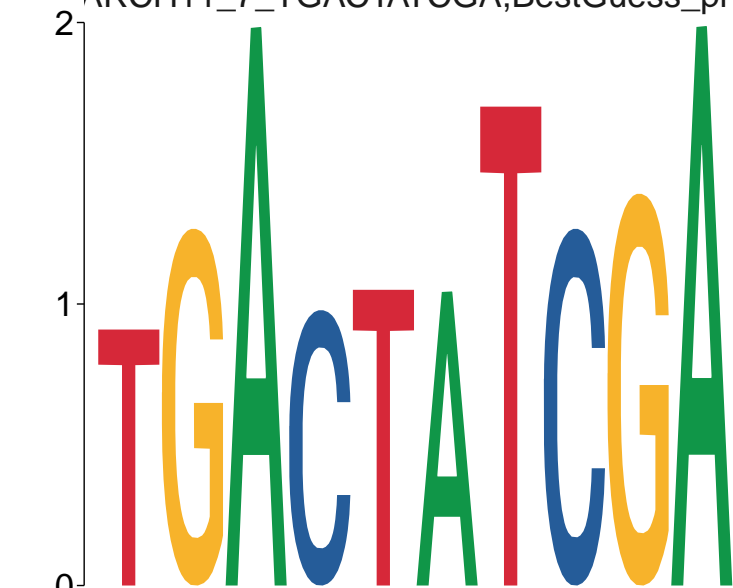
RCH10_22_AAMTCATTGMTCT, BestGuess_



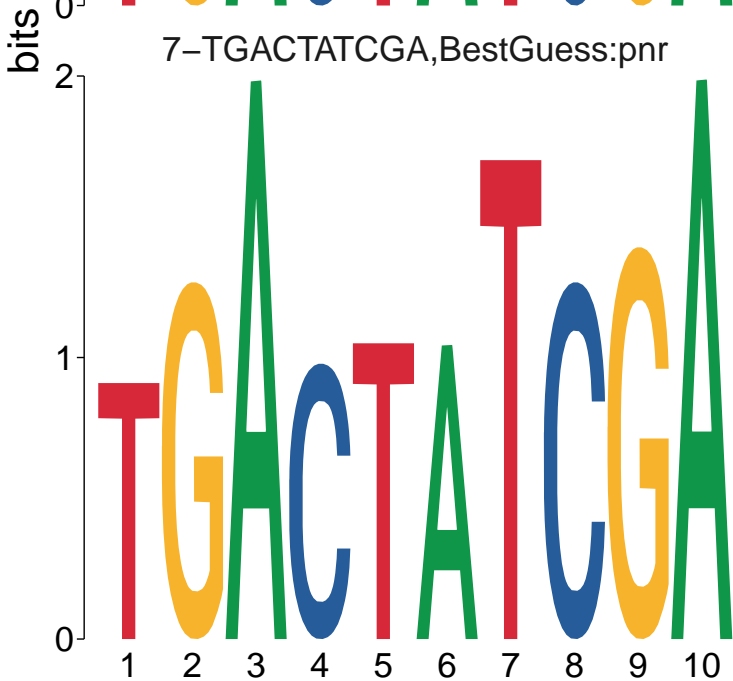
22-AAMTCATTGMTCT, BestGuess:ATHB20



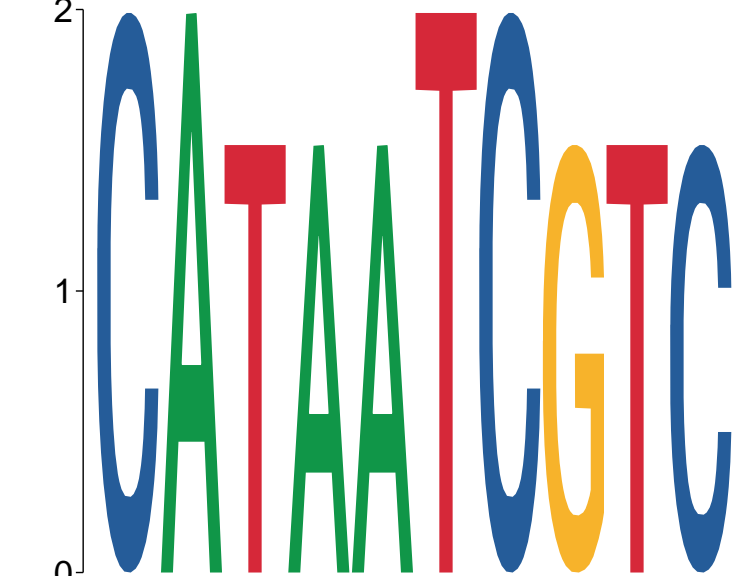
ARCH11_7_TGACTATCGA,BestGuess_pnr



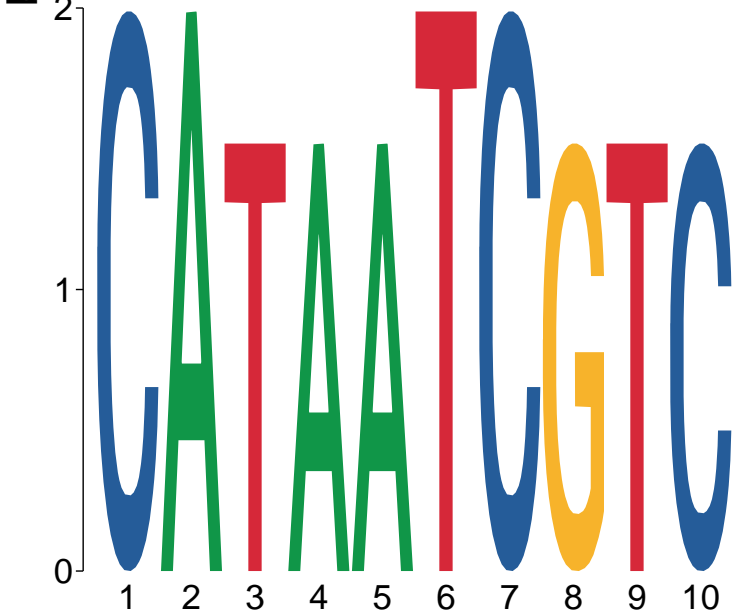
7-TGACTATCGA,BestGuess:pnr



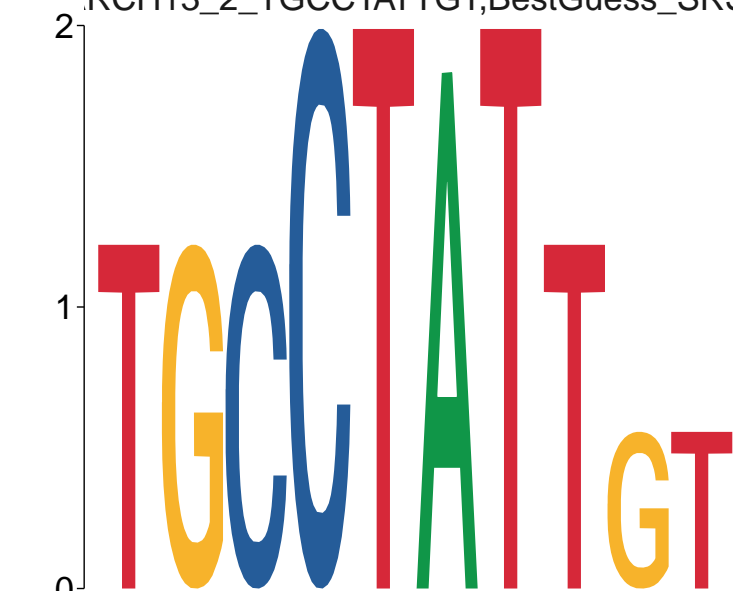
ARCH12_5_CATAATCGTC,BestGuess_ems



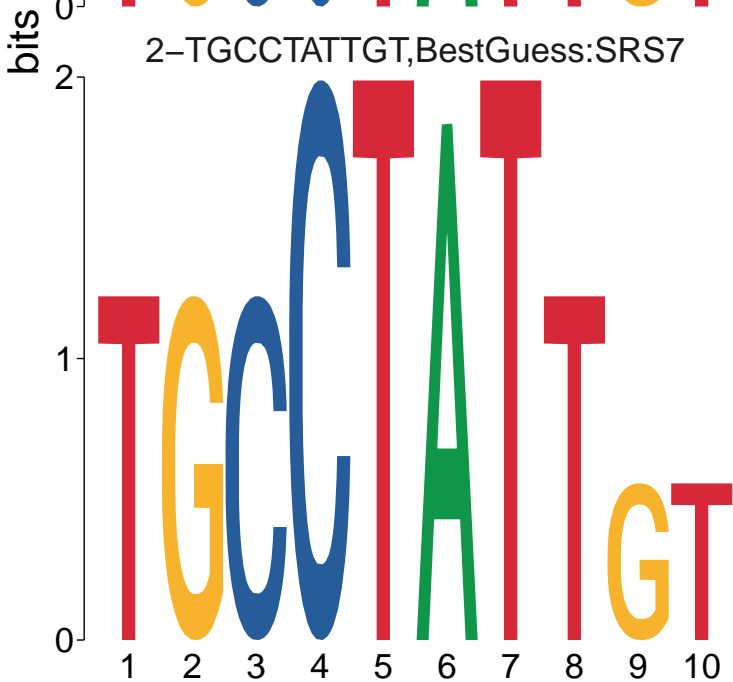
5-CATAATCGTC,BestGuess:ems



RCH13_2_TGCCTATTGT,BestGuess_SRS

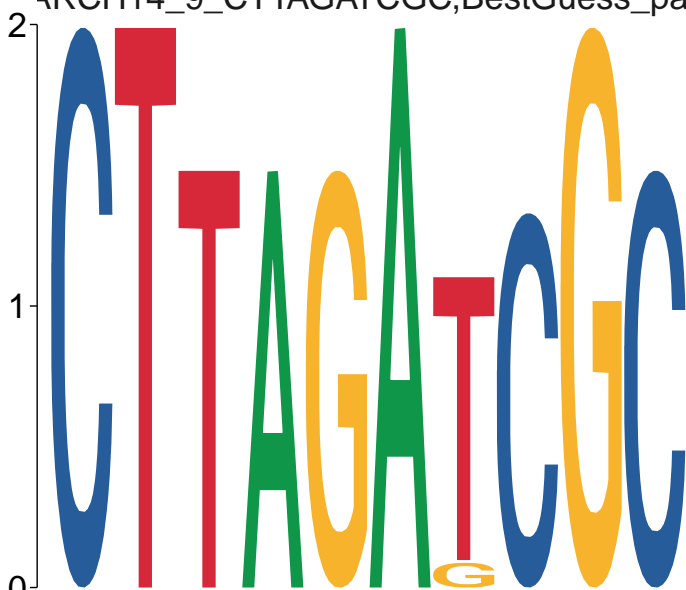


2-TGCCTATTGT,BestGuess:SRS7

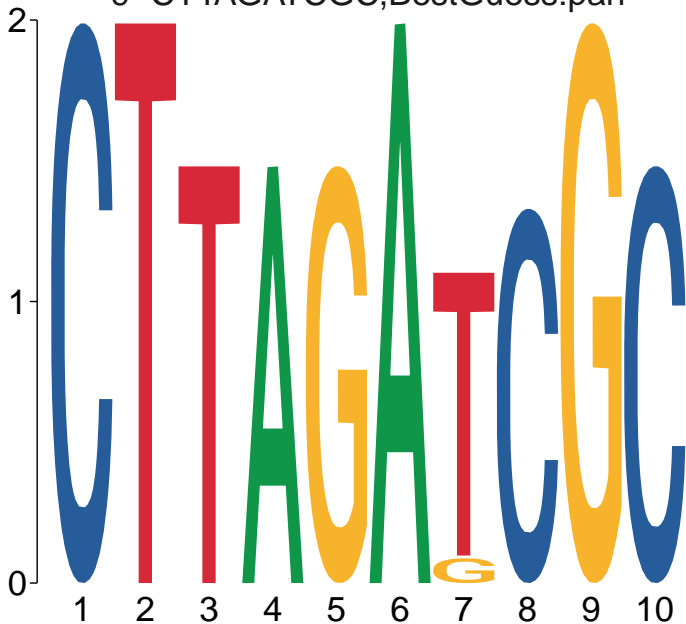


ARCH14_9_CTTAGATCGC,BestGuess_par

bits

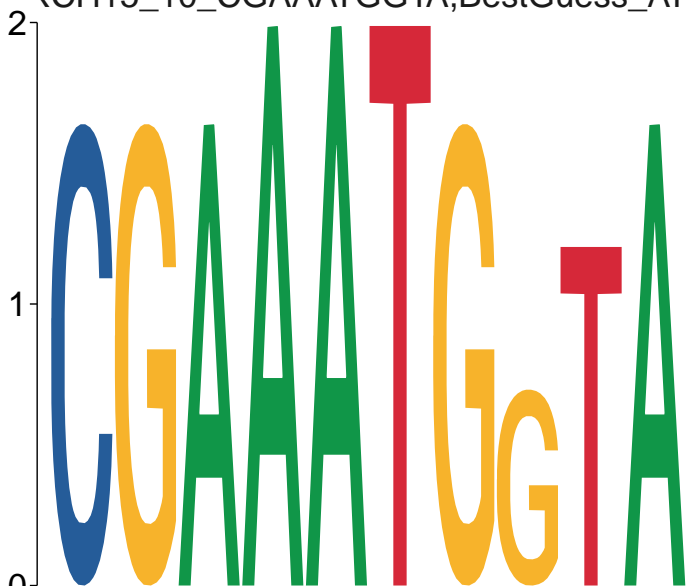


9-CTTAGATCGC,BestGuess:pan

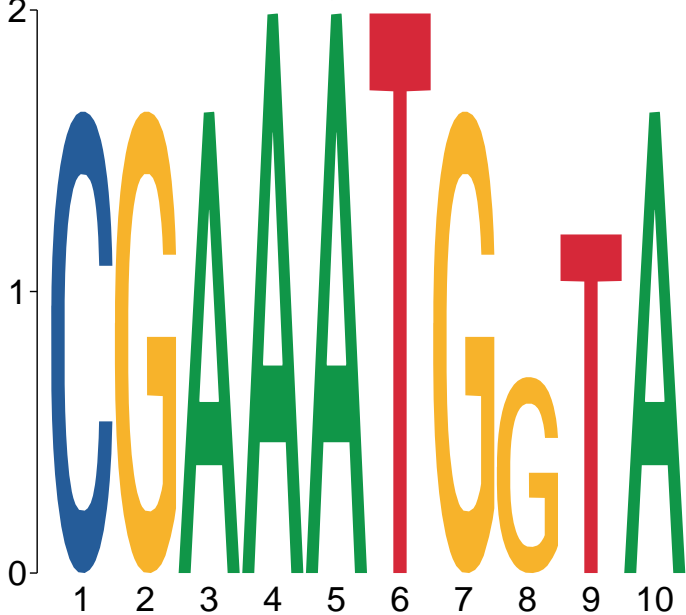


RCH15_10_CGAAATGGTA,BestGuess_AT2

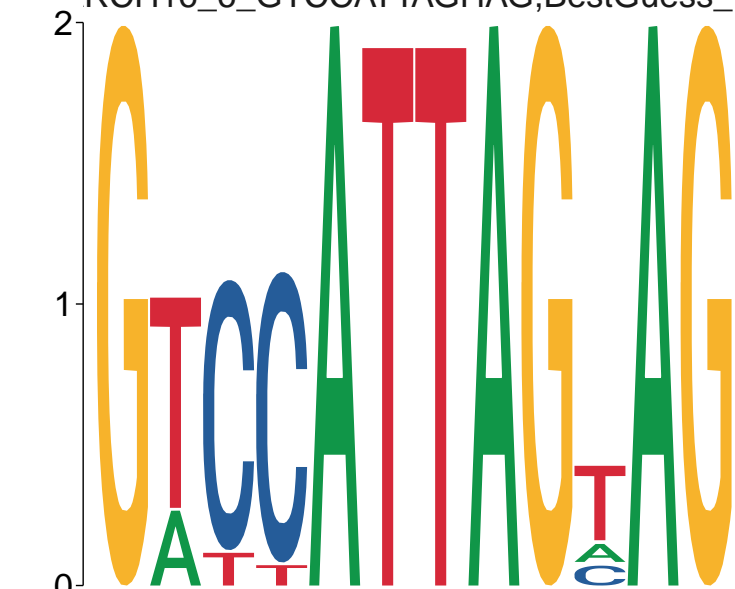
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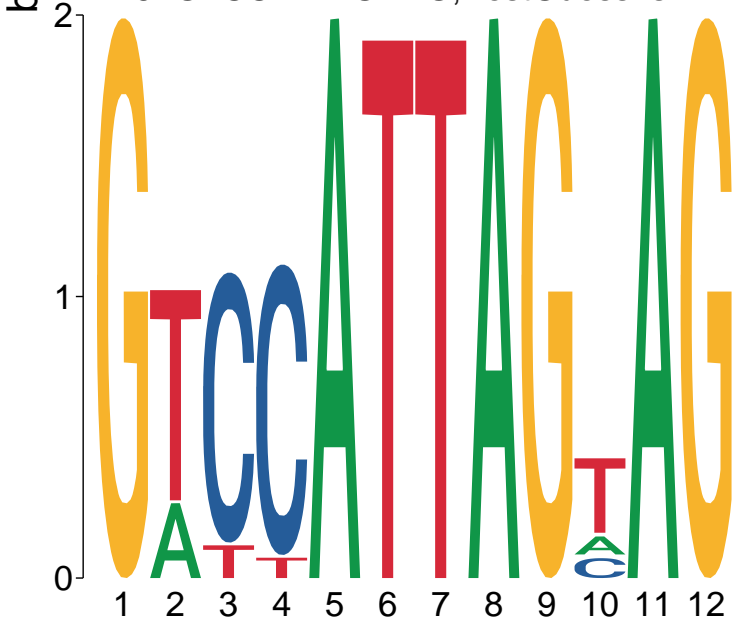
10-CGAAATGGTA,BestGuess:AT2G31460



RCH16_6_GTCCATTAGHAG,BestGuess_I:



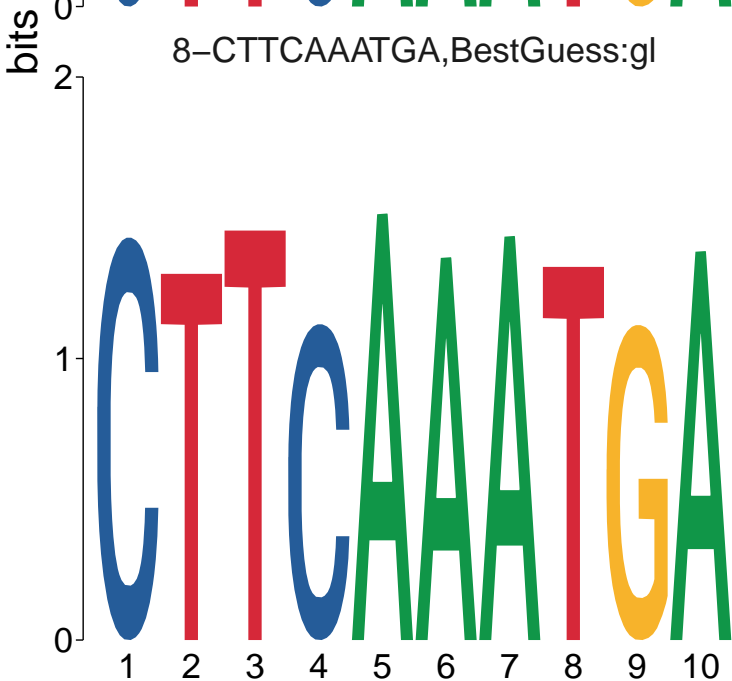
6-GTCCATTAGHAG,BestGuess:Isl1



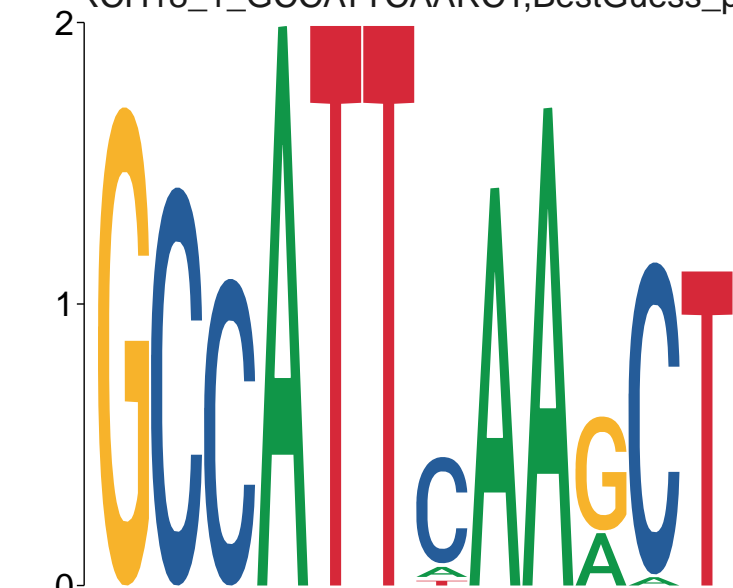
ARCH17_8_CTTCAAATGA,BestGuess_gl



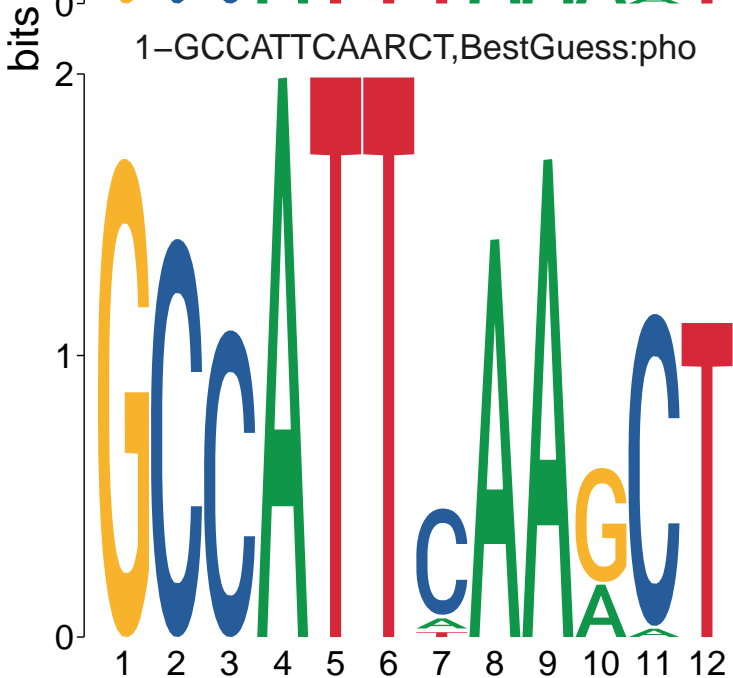
8-CTTCAAATGA,BestGuess:gl



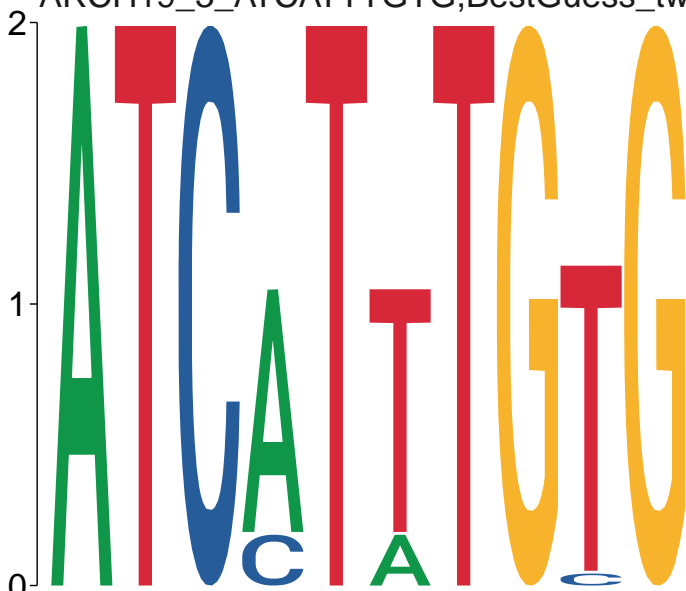
RCH18_1_GCCATTCAARCT,BestGuess_pl



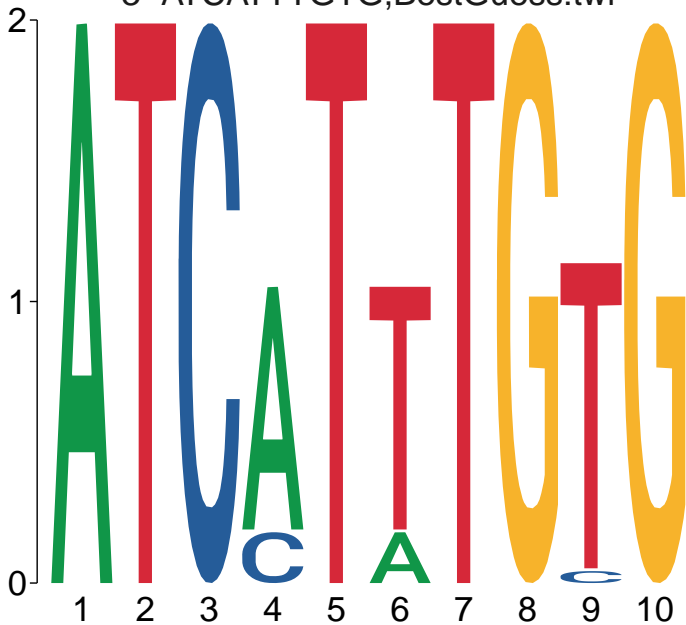
1-GCCATTCAARCT,BestGuess:pho



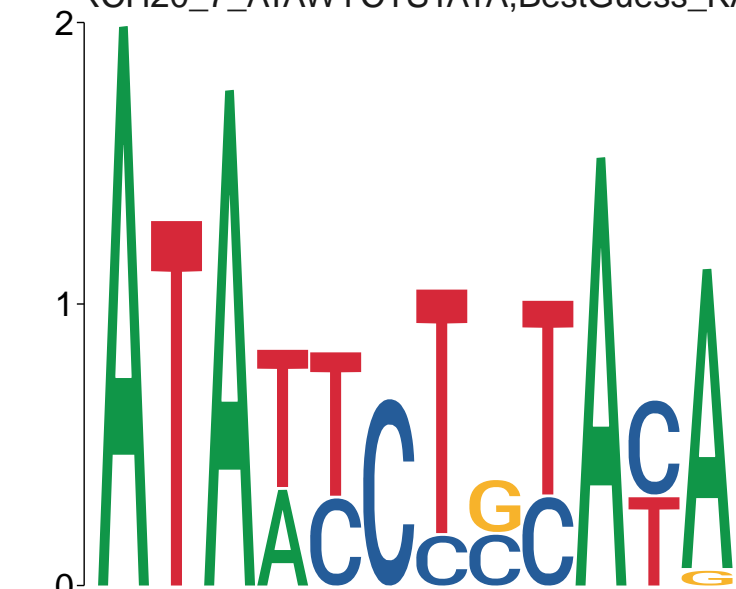
ARCH19_3_ATCATTGTG,BestGuess_tw



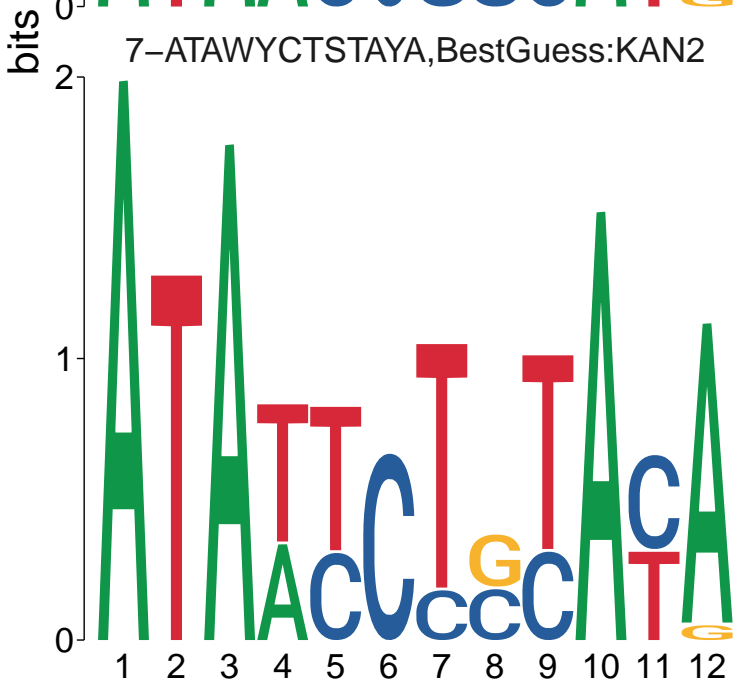
3-ATCATTGTG,BestGuess:tw



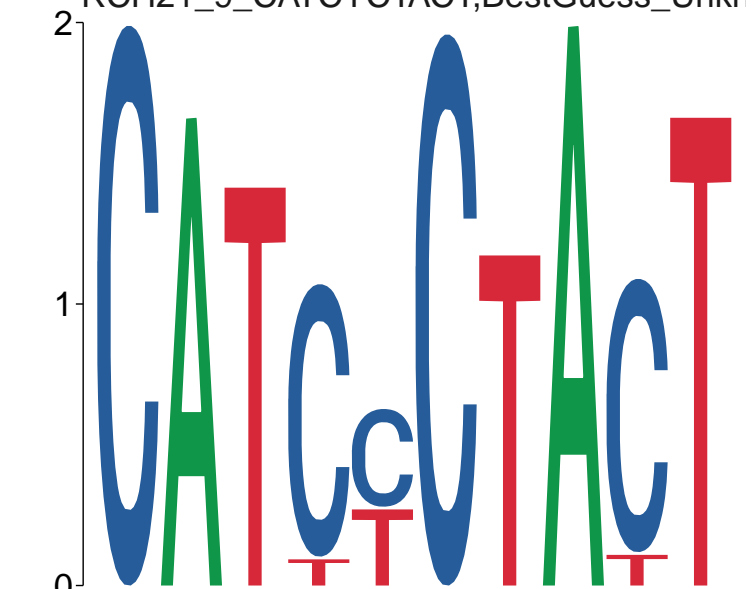
RCH20_7_ATAWYCTSTAYA,BestGuess_KA



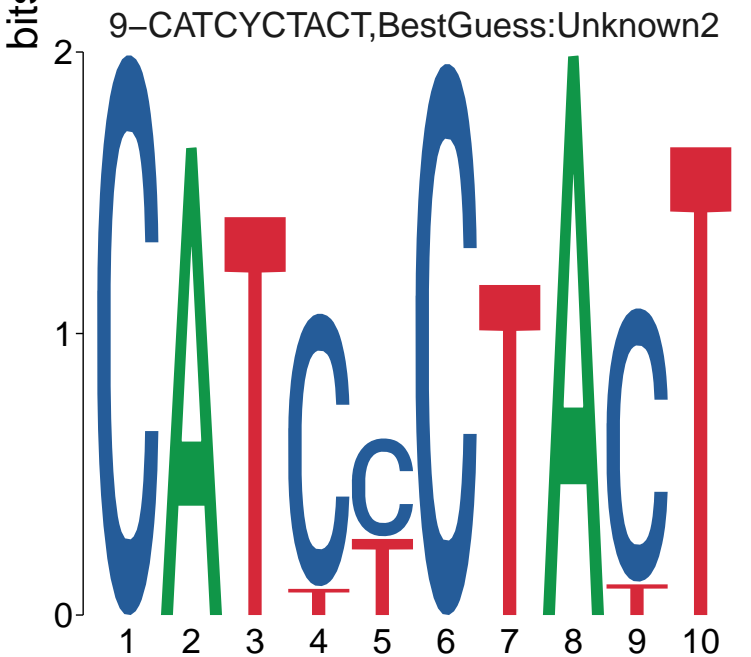
7-ATAWYCTSTAYA,BestGuess:KAN2



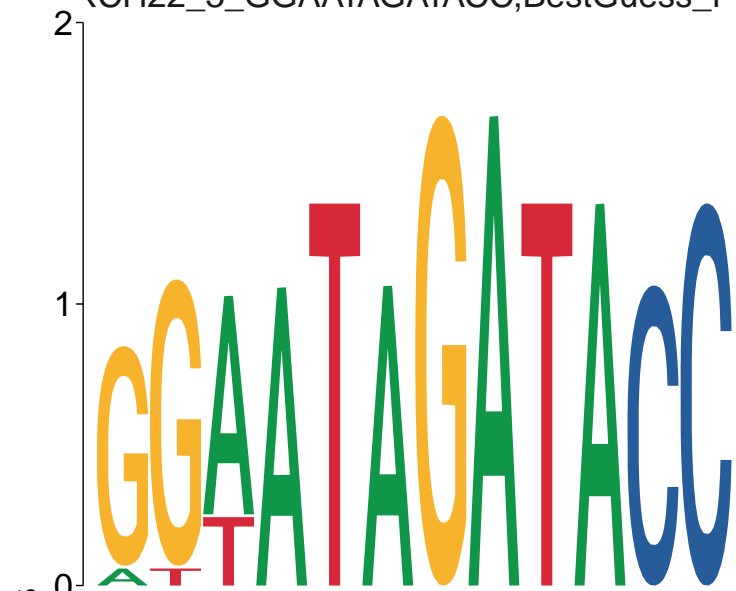
RCH21_9_CATCYCTACT,BestGuess_Unkn



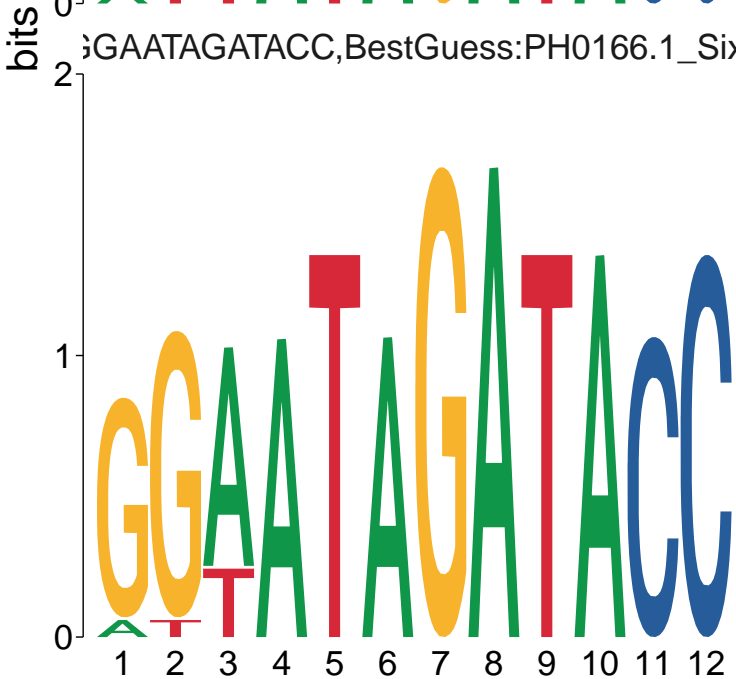
9-CATCYCTACT,BestGuess:Unknown2



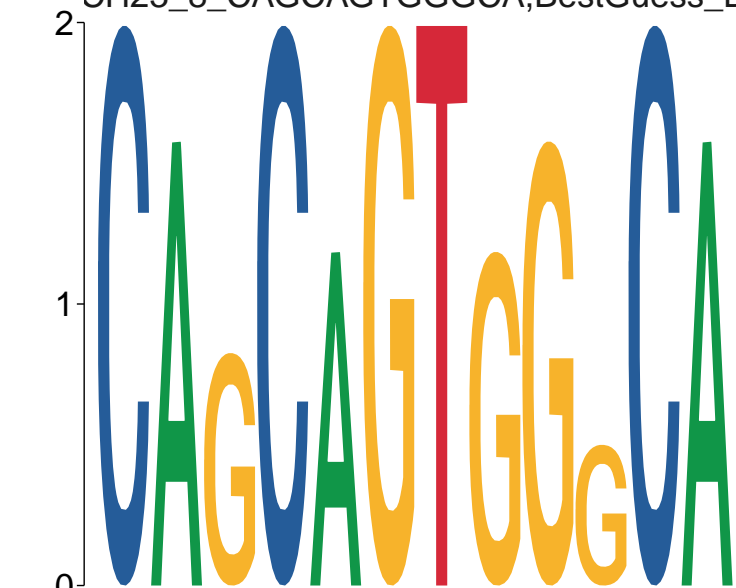
RCH22_3_GGAATAGATACC,BestGuess_Pf



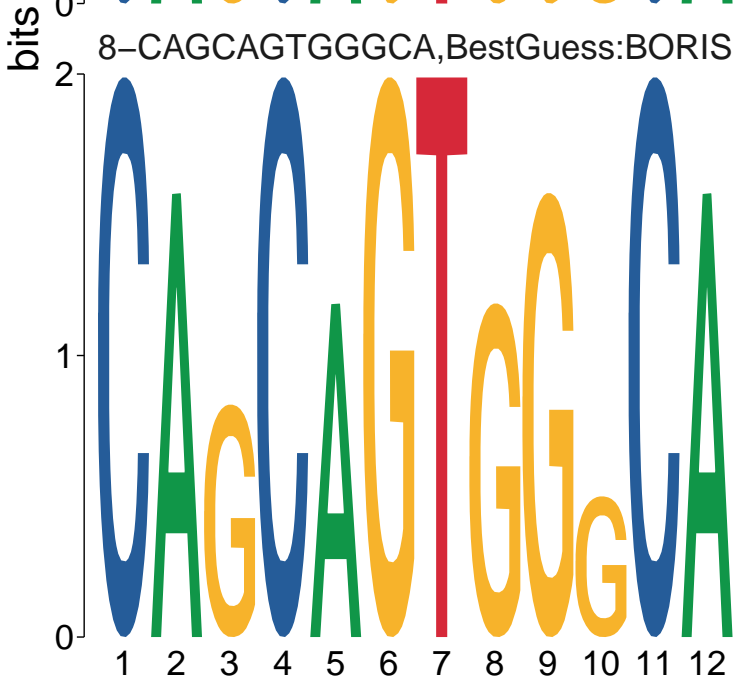
GGAATAGATACC,BestGuess:PH0166.1_Six



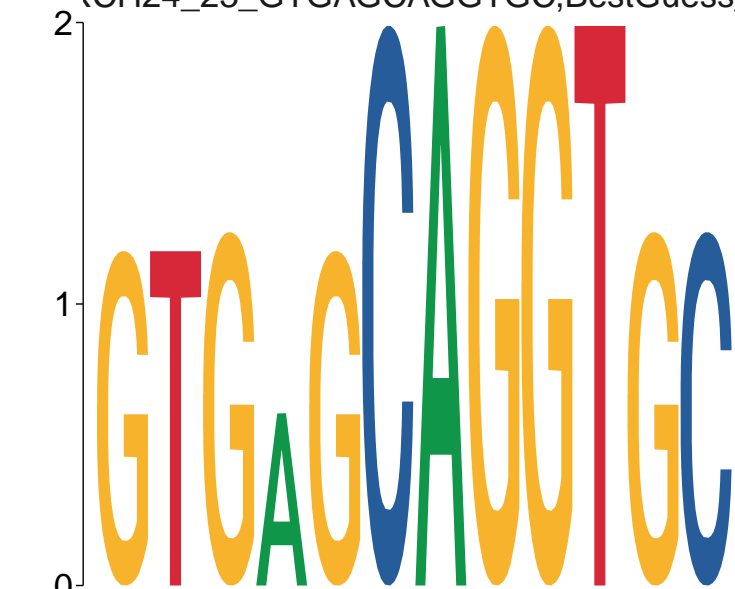
CH23_8_CAGCAGTGGGCA,BestGuess_B



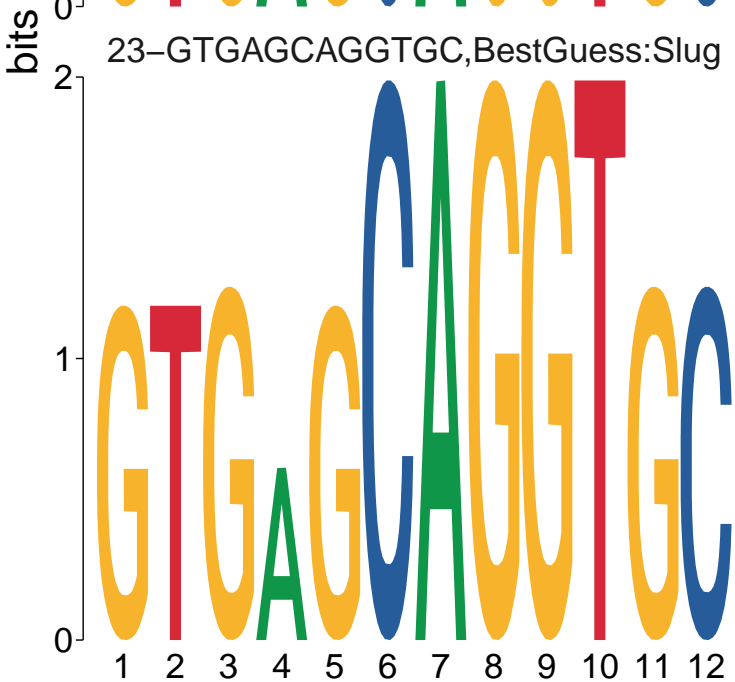
8-CAGCAGTGGGCA,BestGuess:BORIS



CH24_23_GTGAGCAGGTGC,BestGuess_



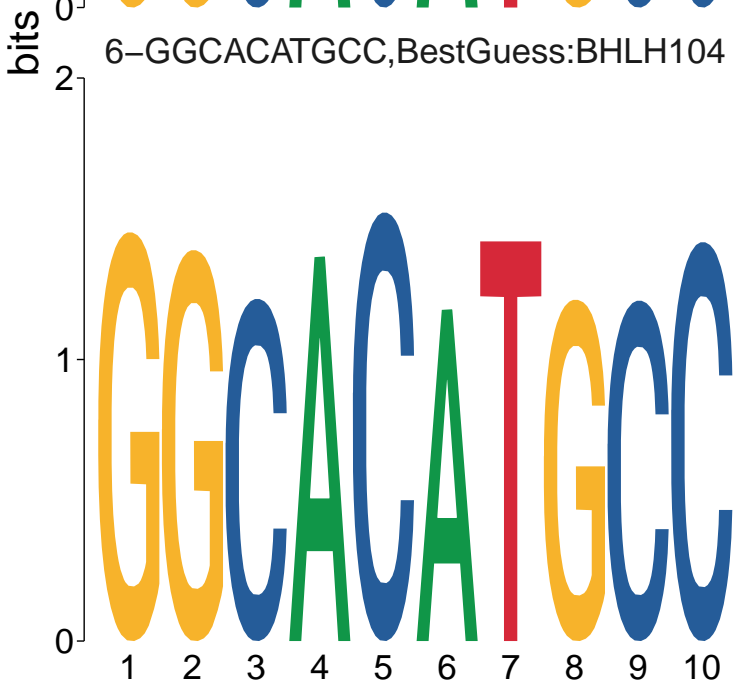
23-GTGAGCAGGTGC,BestGuess:Slug



CH25_6_GGCACATGCC,BestGuess_BHL

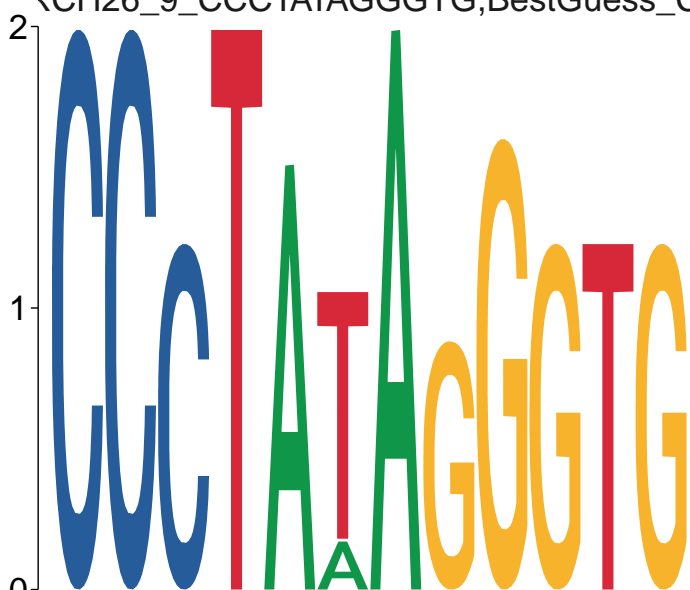


6-GGCACATGCC,BestGuess:BHLH104

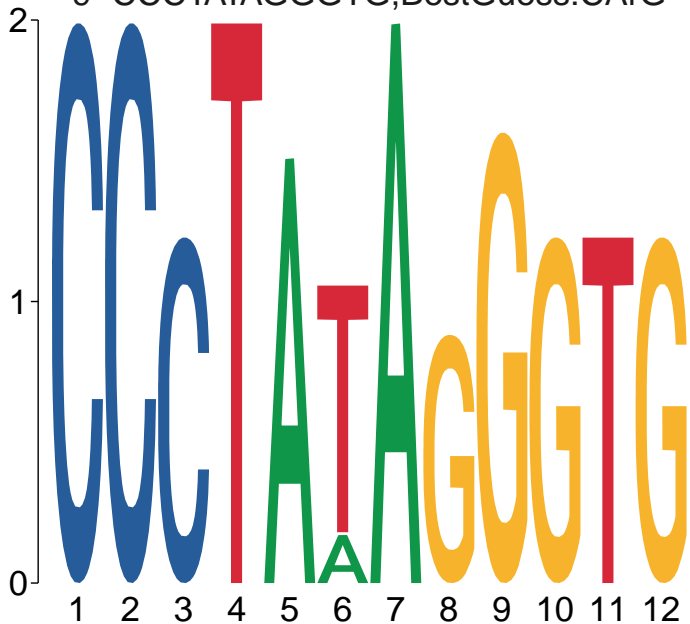


RCH26_9_CCCTATAGGGTG,BestGuess_C

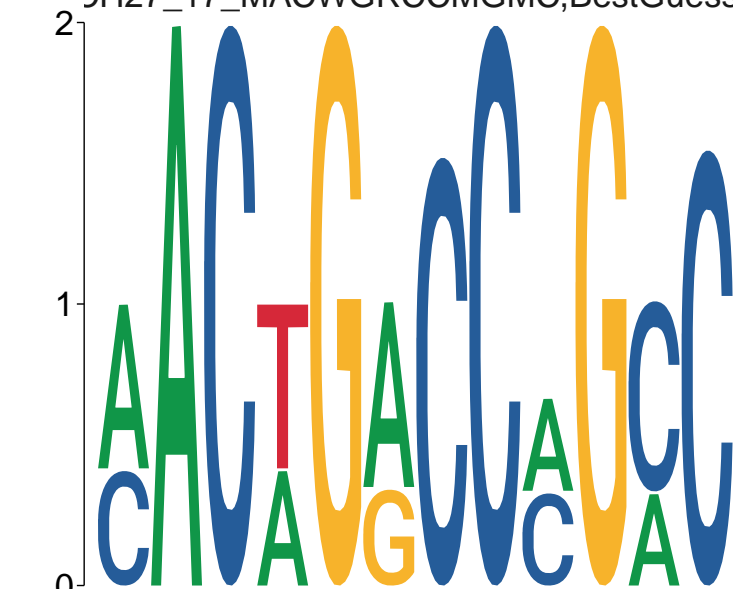
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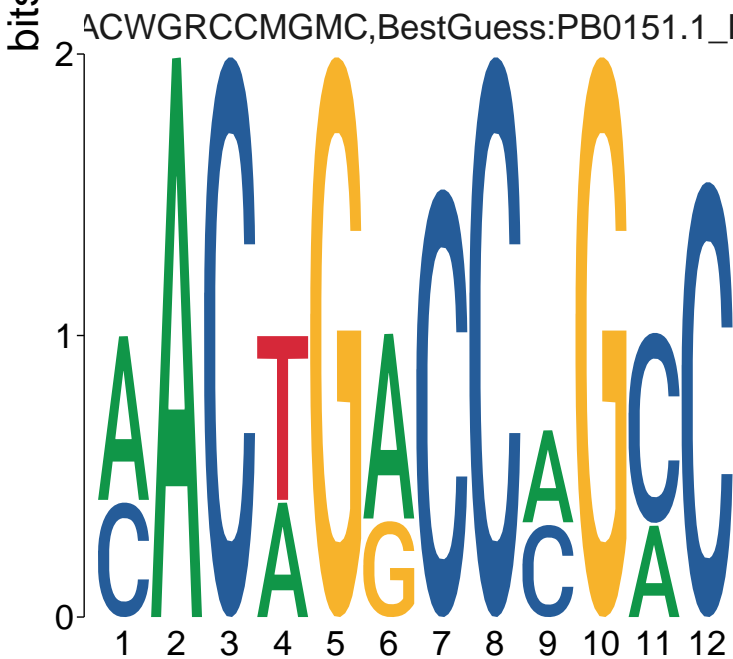
9-CCCTATAGGGTG,BestGuess:CArG



CH27_17_MACWGRCCMGMCMC,BestGuess_



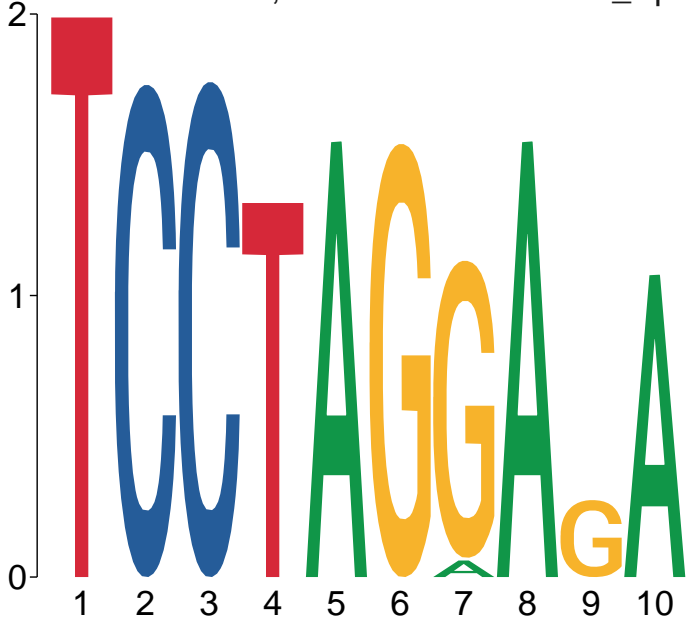
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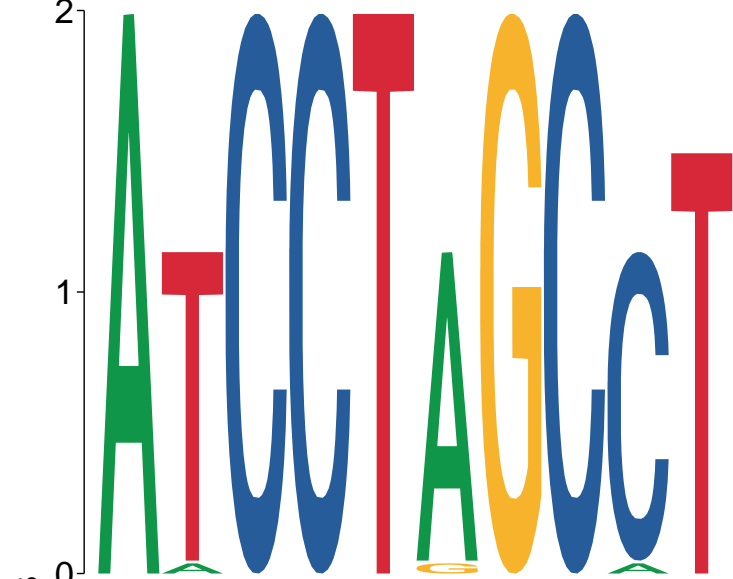
RCH28_10_TCCTAGGAGA,BestGuess_PBI



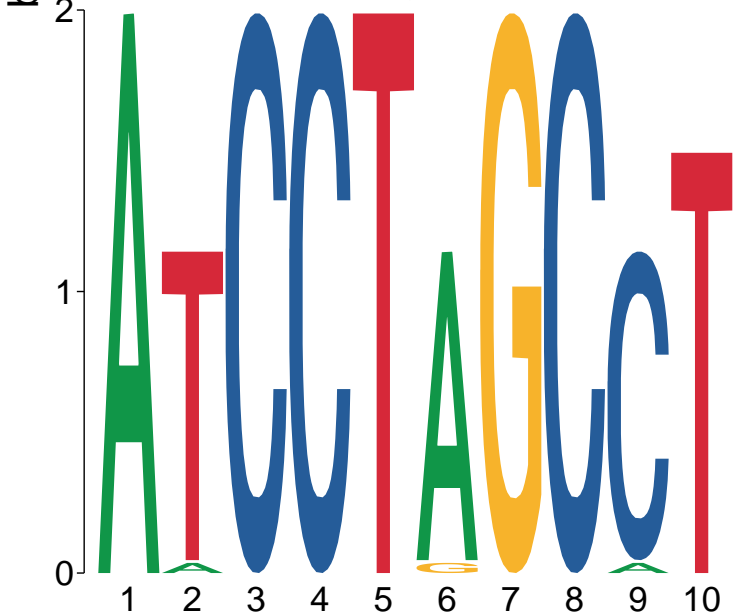
TCCTAGGAGA,BestGuess:PB0181.1_Spde



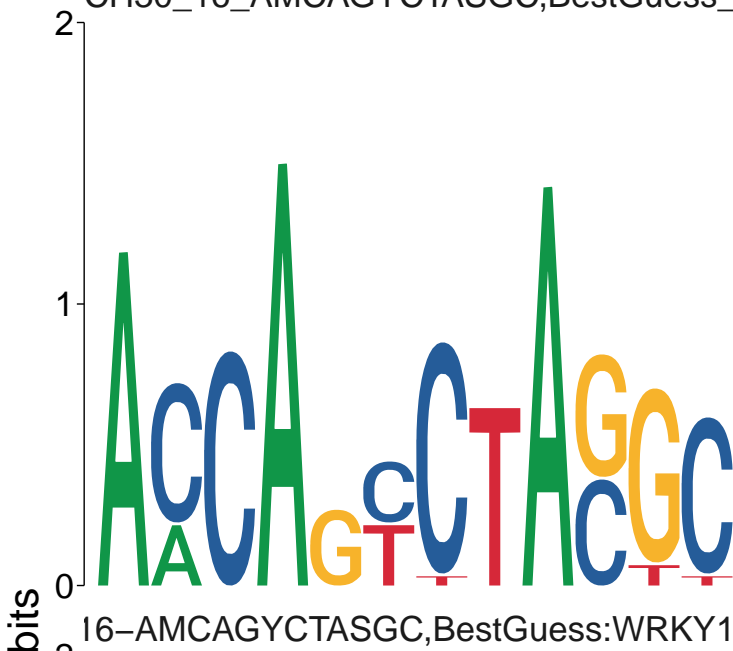
RCH29_23_ATCCTAGCCT,BestGuess_G3E



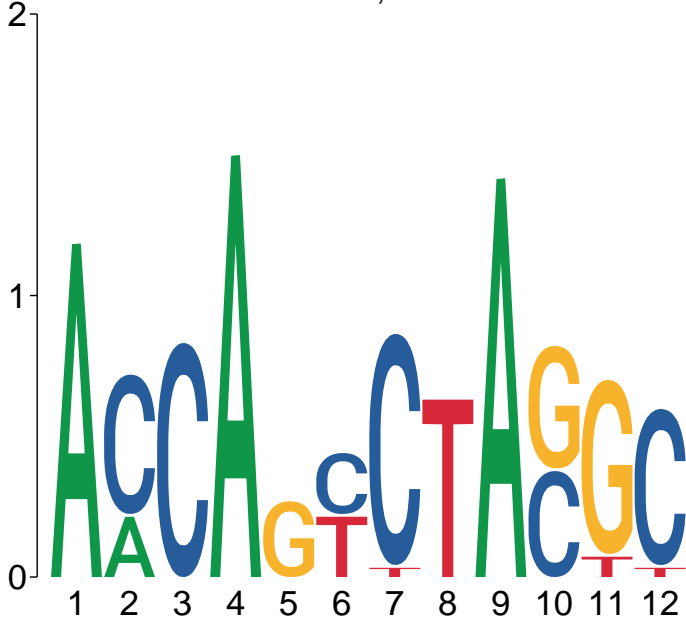
23-ATCCTAGCCT,BestGuess:G3BP2



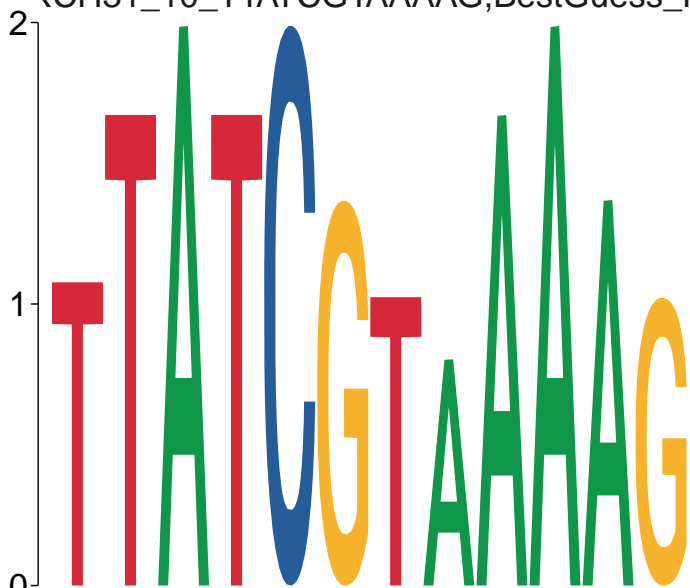
CH30_16_AMCAGYCTASGC,BestGuess_\



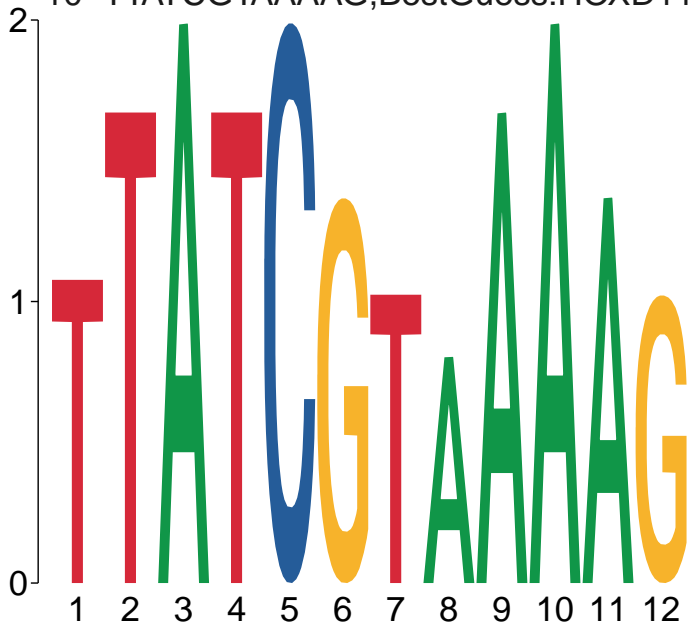
16-AMCAGYCTASGC,BestGuess:WRKY17



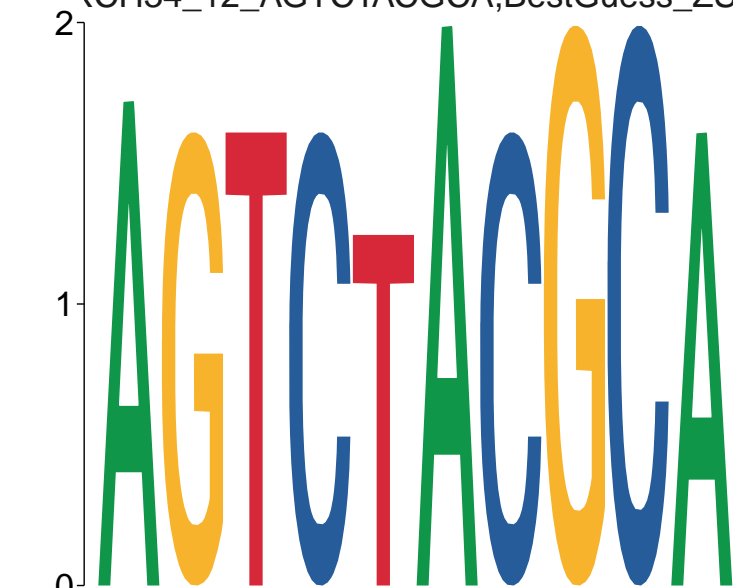
RCH31_10_TTATCGTAAAAG,BestGuess_1



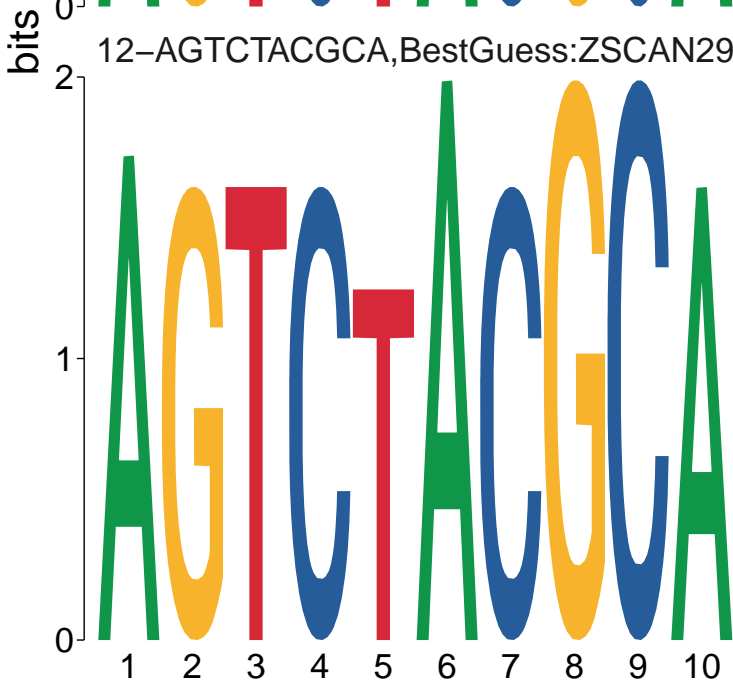
10-TTATCGTAAAAG,BestGuess:HOXD11



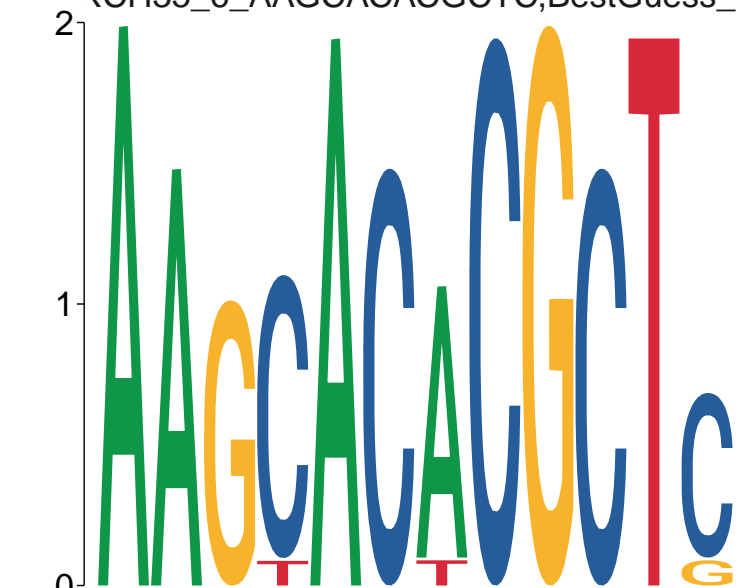
12-AGTCTACGCA,BestGuess_ZSCAN29



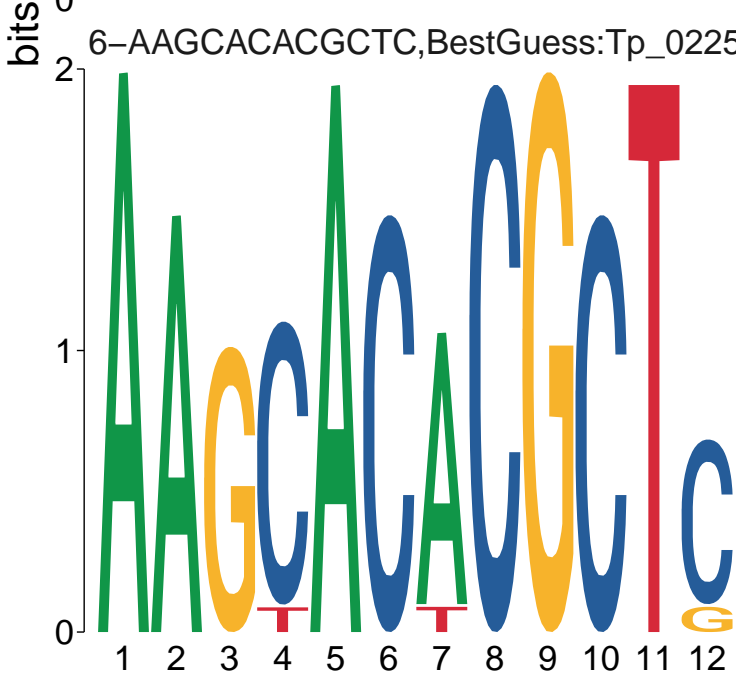
12-AGTCTACGCA,BestGuess:ZSCAN29



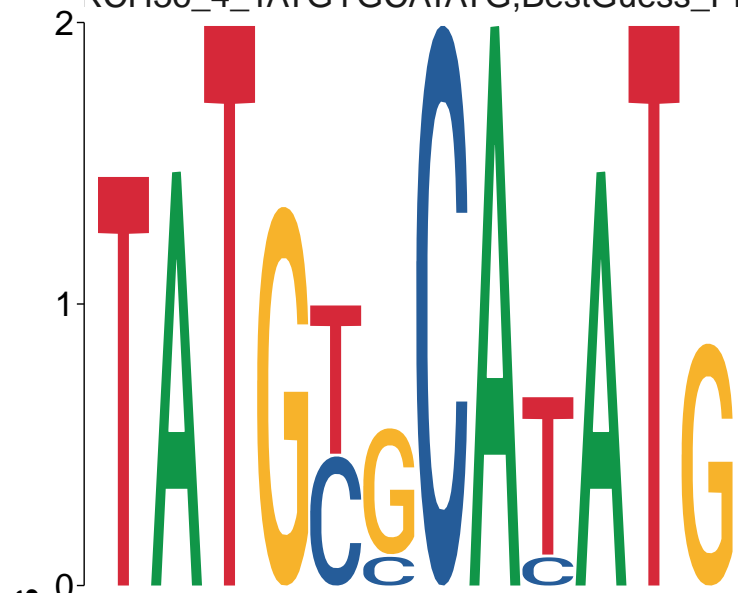
RCH35_6_AAGCACACGCTC,BestGuess_1



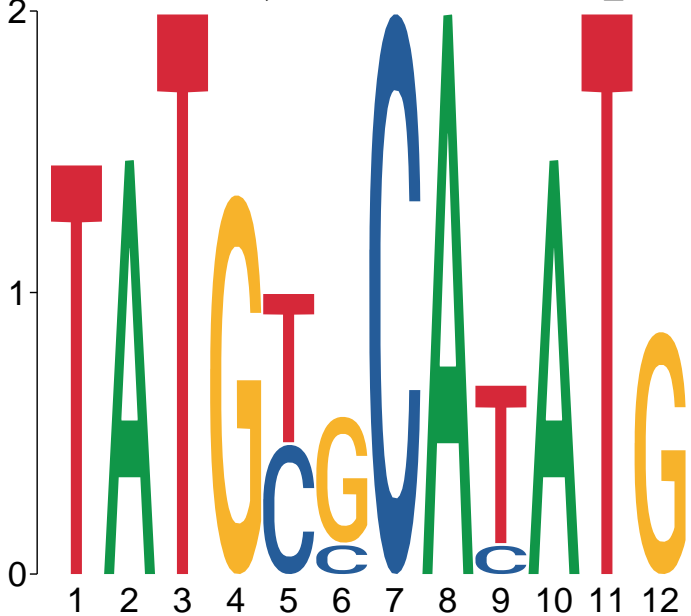
6-AAGCACACGCTC,BestGuess:Tp_0225



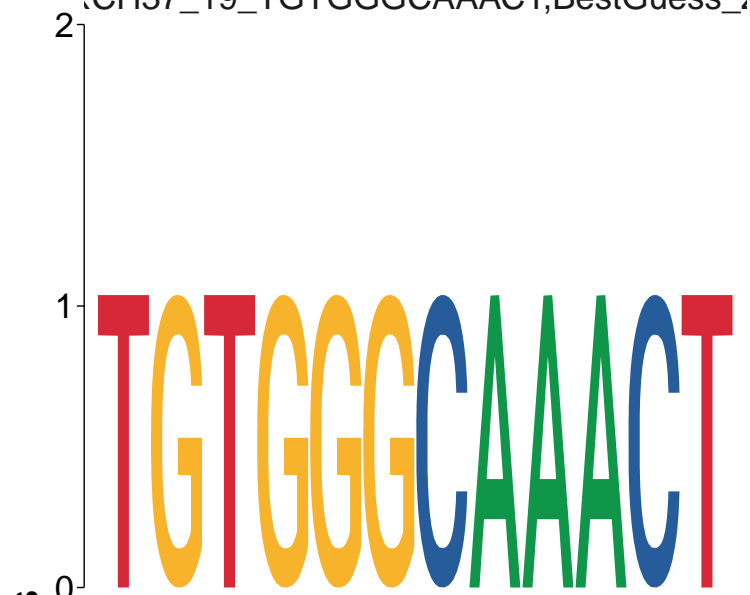
RCH36_4_TATGYGCATATG,BestGuess_PE



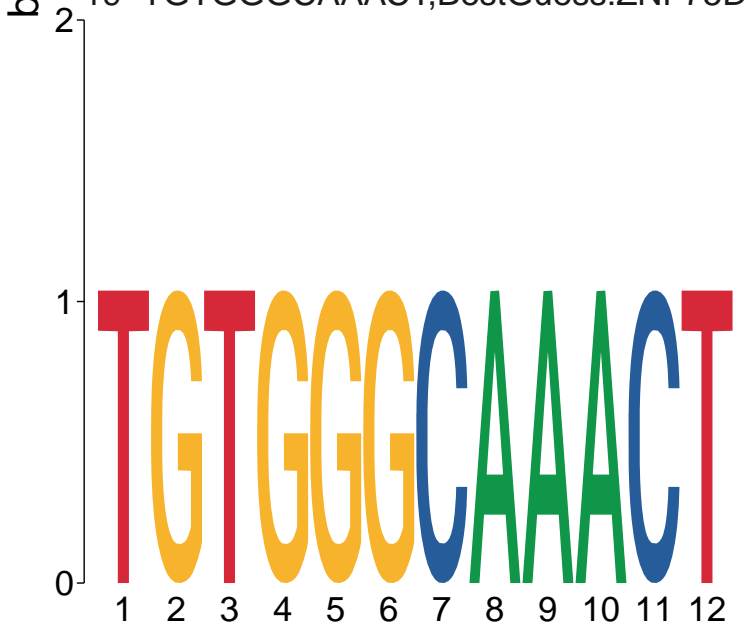
TGYGCATATG,BestGuess:PB0104.1_Zsca



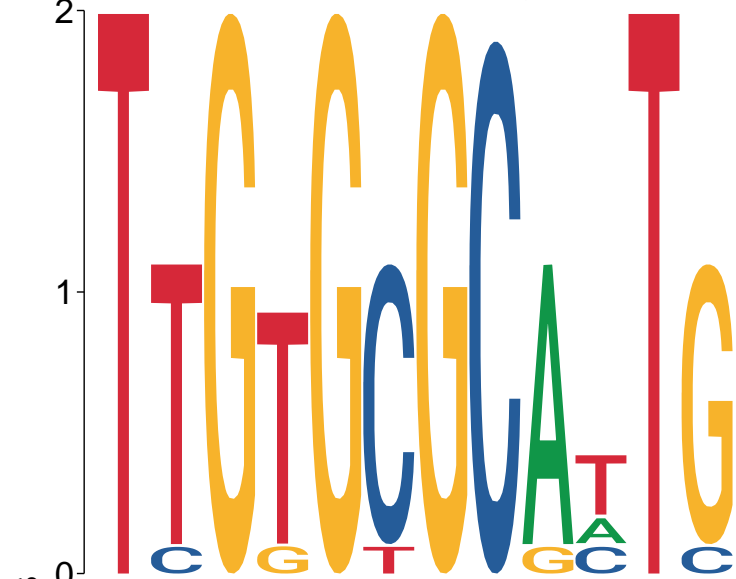
CH37_19_TGTGGGCAAAC, BestGuess_4



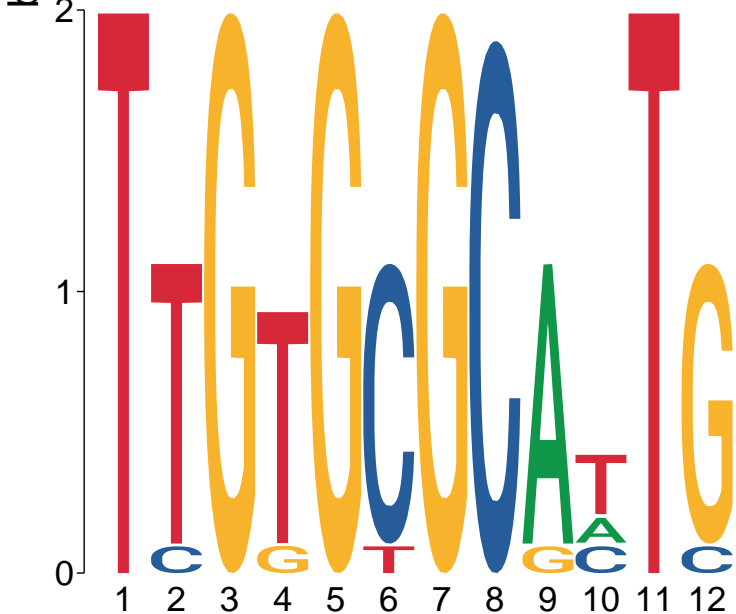
19-TGTGGGCAAAC, BestGuess:ZNF75D



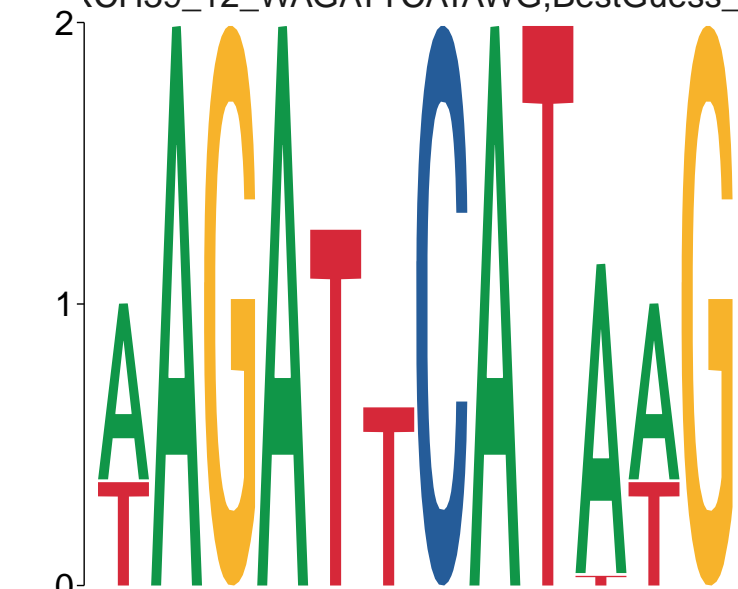
CH38_2_TTGTGCGCAHTG,BestGuess_R



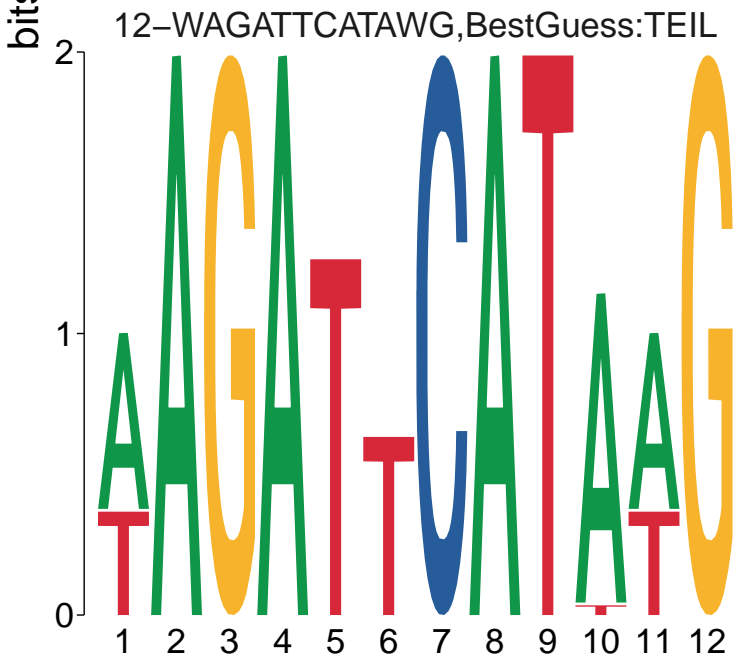
2-TTGTGCGCAHTG,BestGuess:RCS1



CH39_12_WAGATTCATAWG, BestGuess_



12-WAGATTCATAWG, BestGuess:TEIL



RCH40_16_TATAATAGCATG,BestGuess_G

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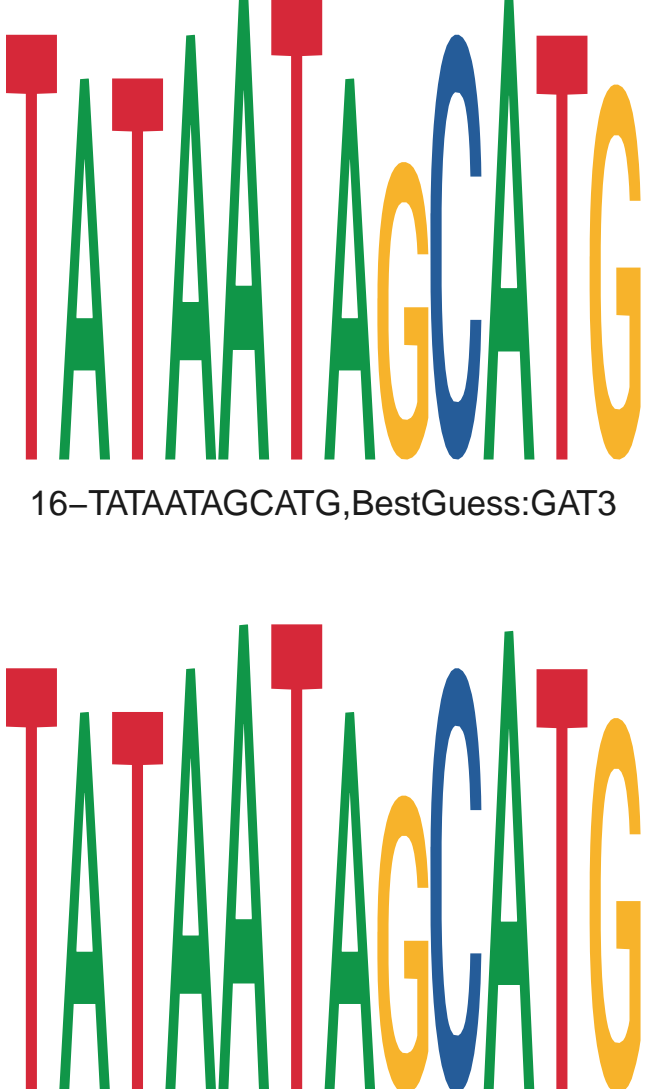
16-TATAATAGCATG,BestGuess:GAT3

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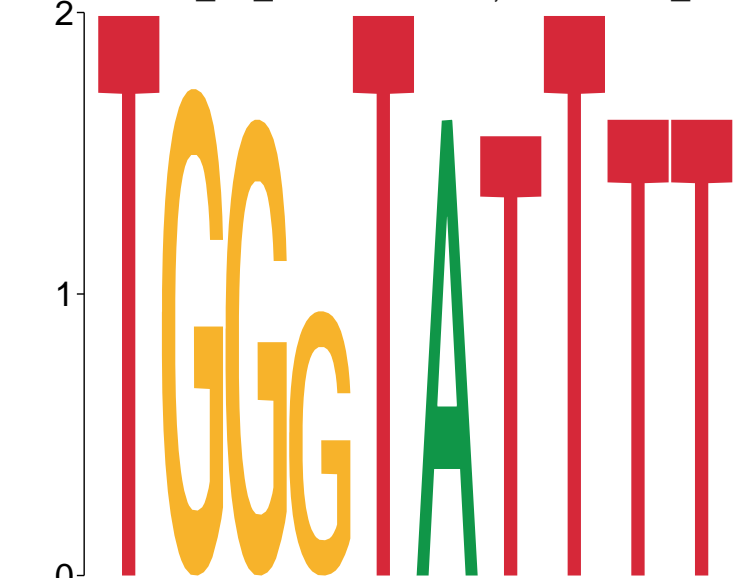
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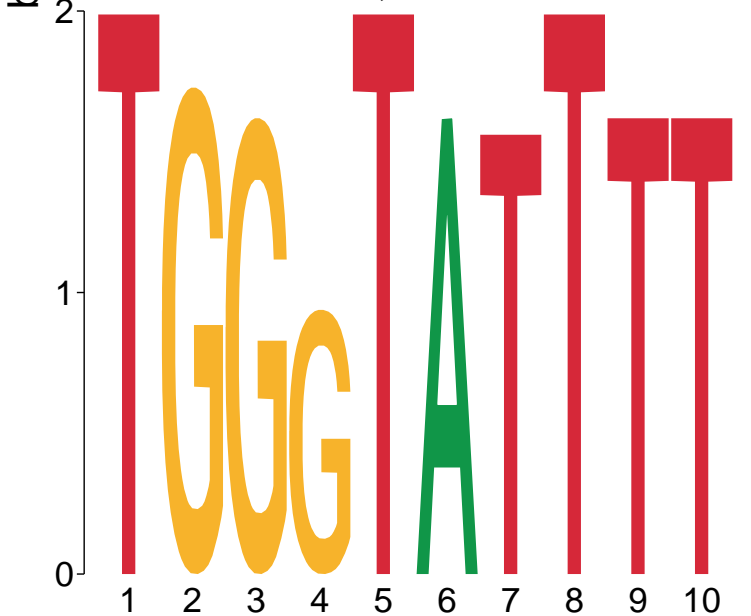
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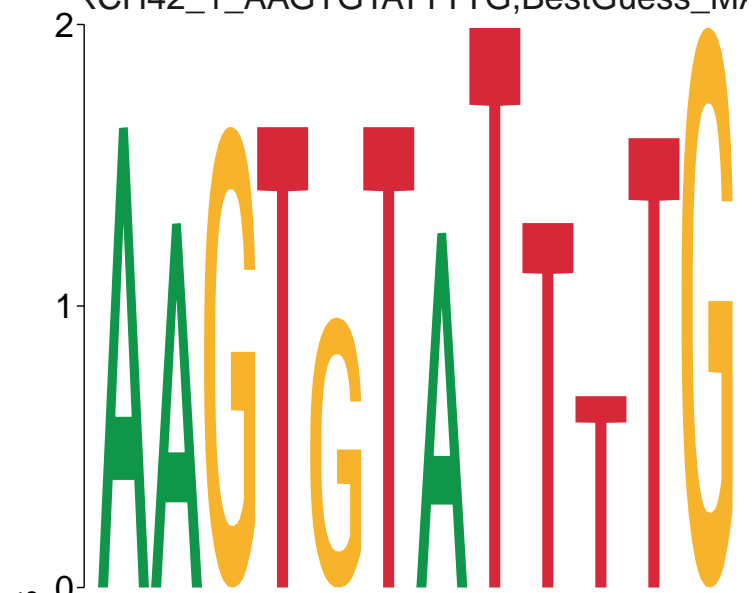
RCH41_20_TGGGTATTTT,BestGuess_Unk



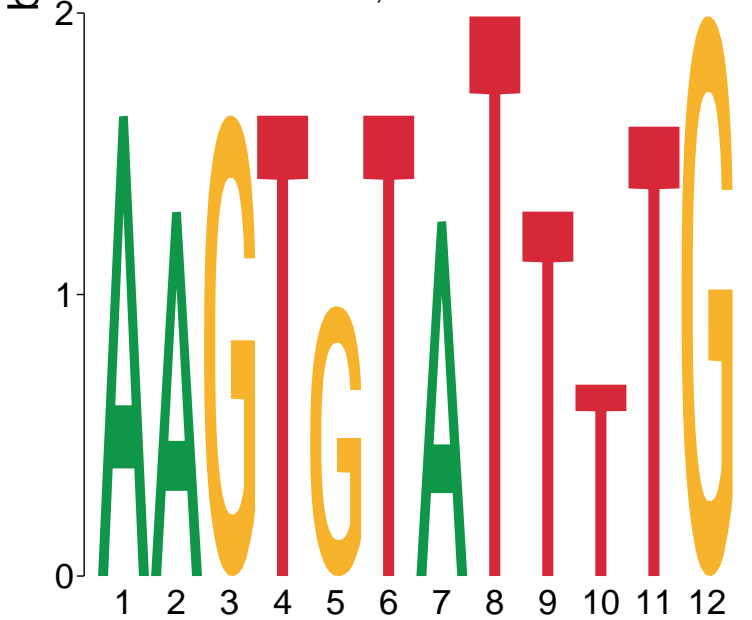
20-TGGGTATTTT,BestGuess:Unknown4



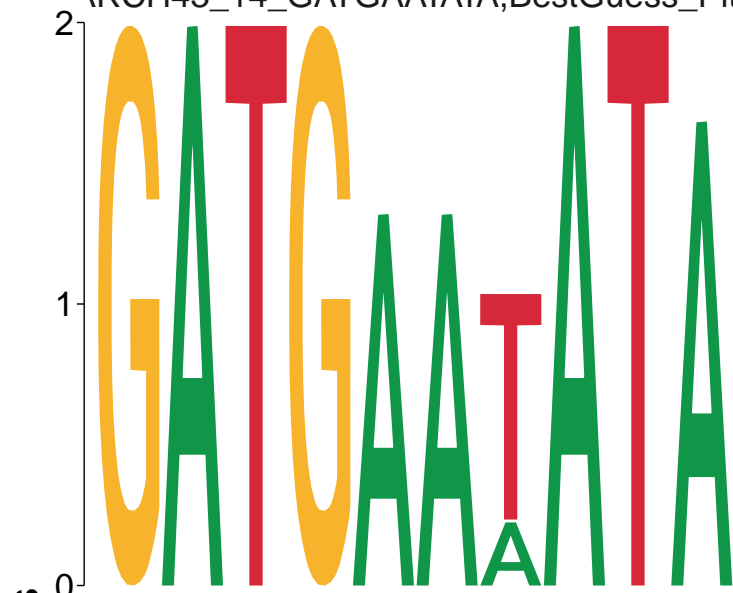
RCH42_1_AAGTGTATTTTG,BestGuess_M/



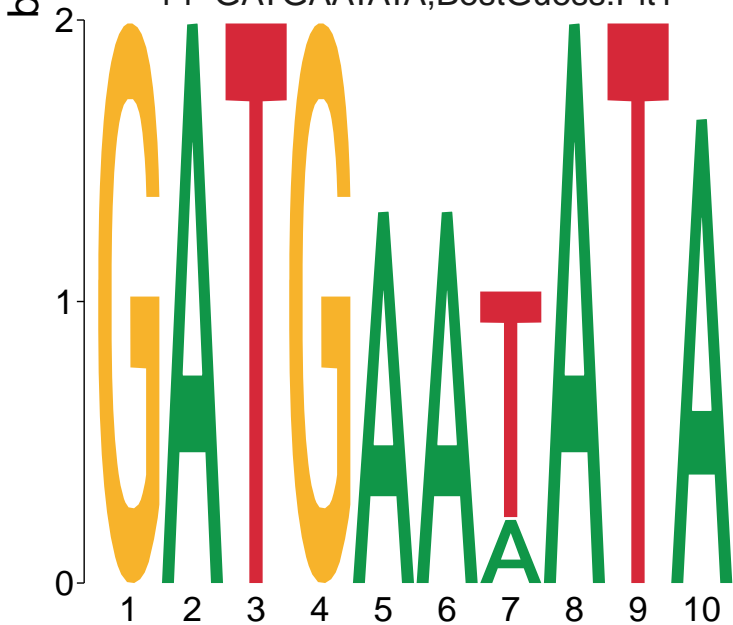
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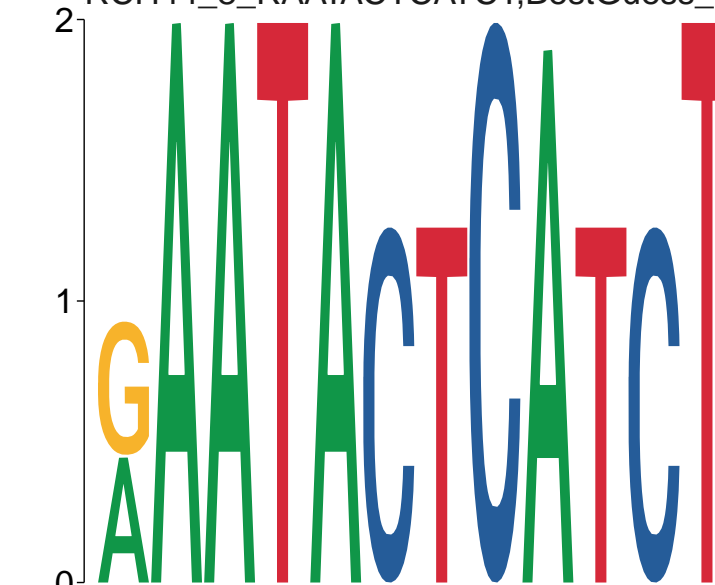
ARCH43_14_GATGAATATA,BestGuess_Pit'



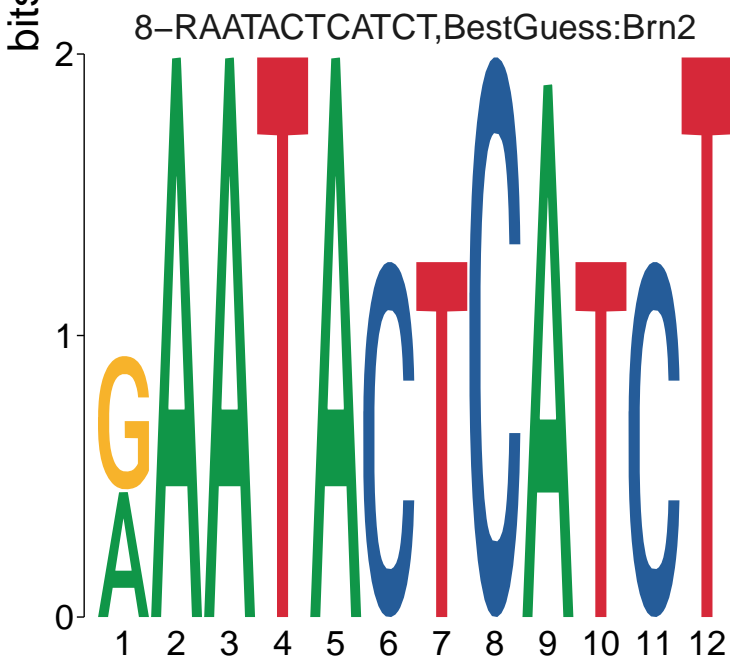
14-GATGAATATA,BestGuess:Pit1



RCH44_8_RAATACTCATCT,BestGuess_Br



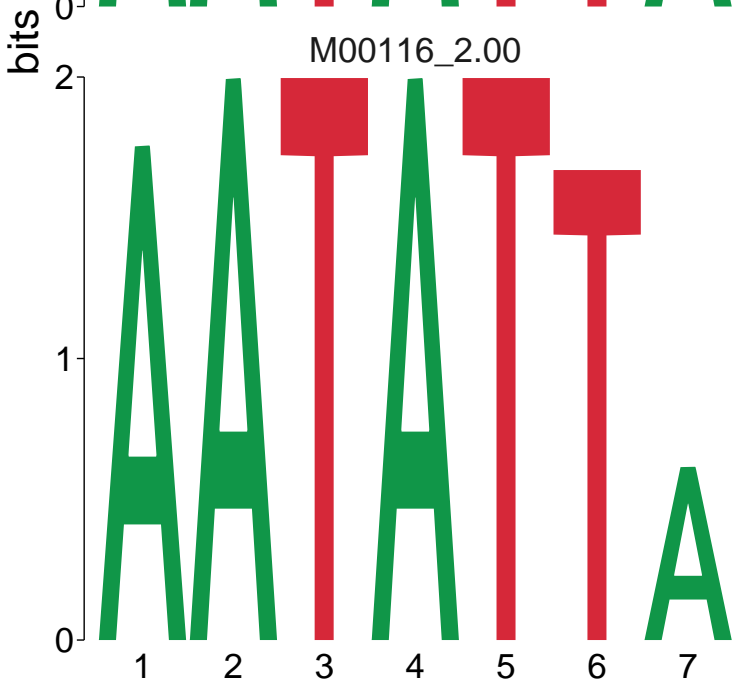
8-RAATACTCATCT,BestGuess:Brn2



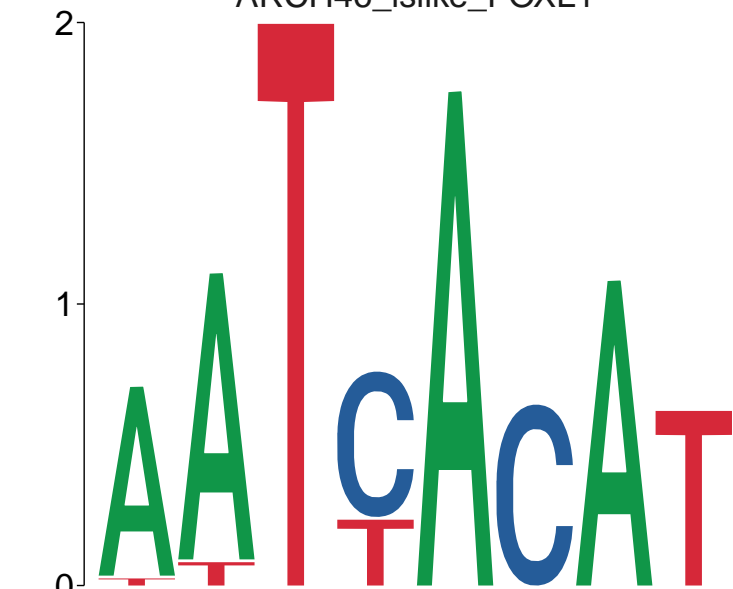
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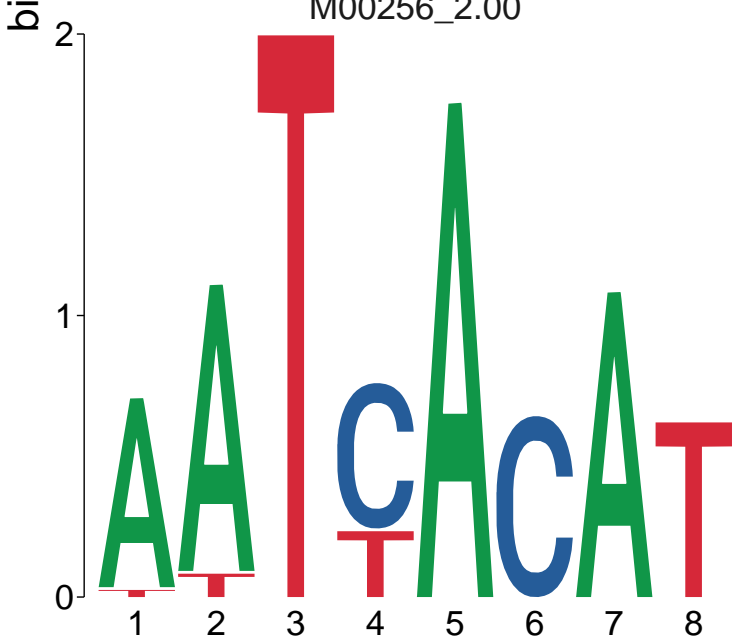
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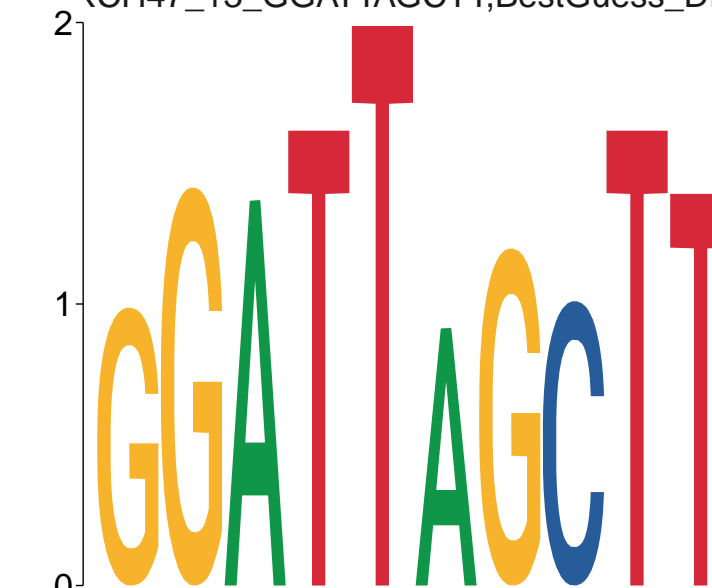
ARCH46_islike_FOXL1



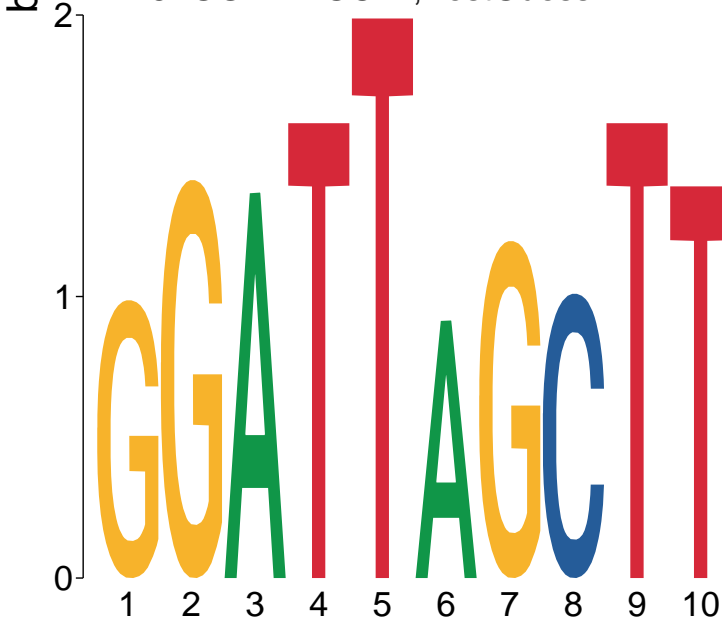
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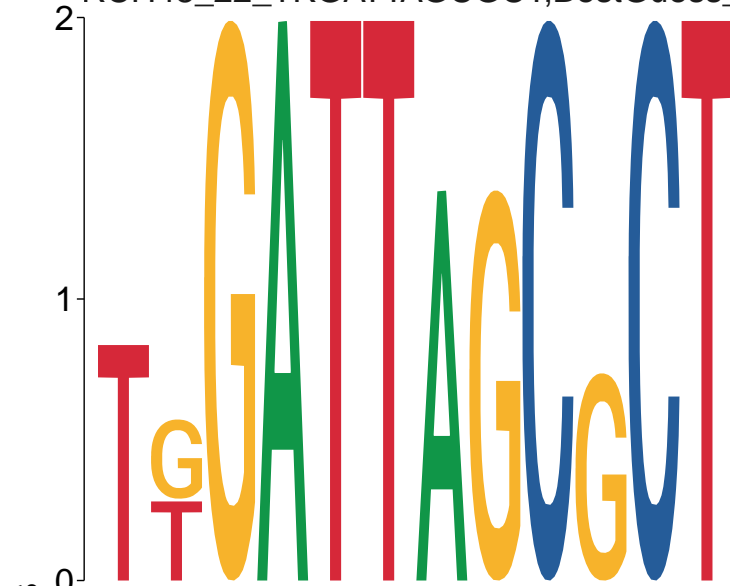
RCH47_13_GGATTAGCTT,BestGuess_DPF



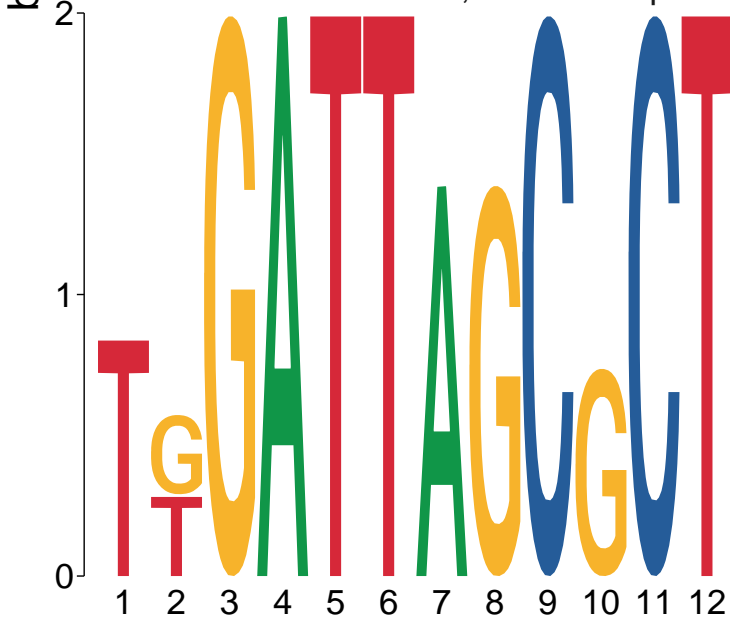
13-GGATTAGCTT,BestGuess:DPRX



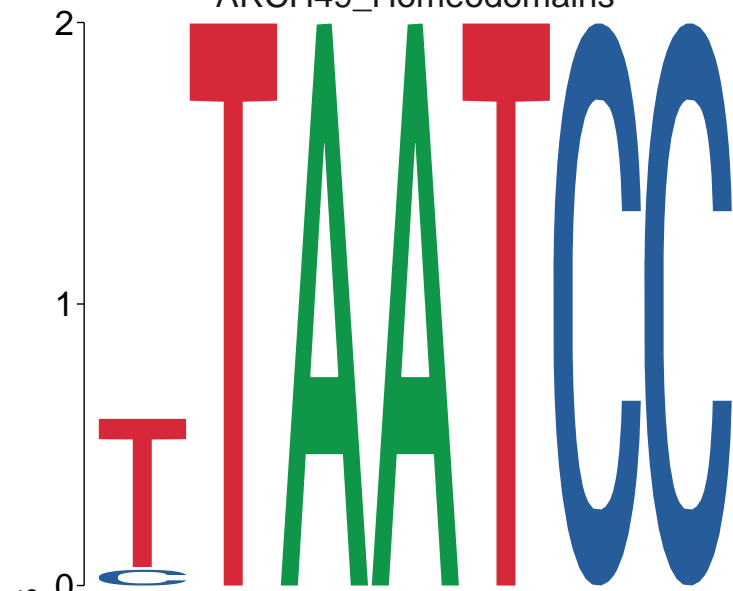
RCH48_22_TKGATTAGCGCT,BestGuess_



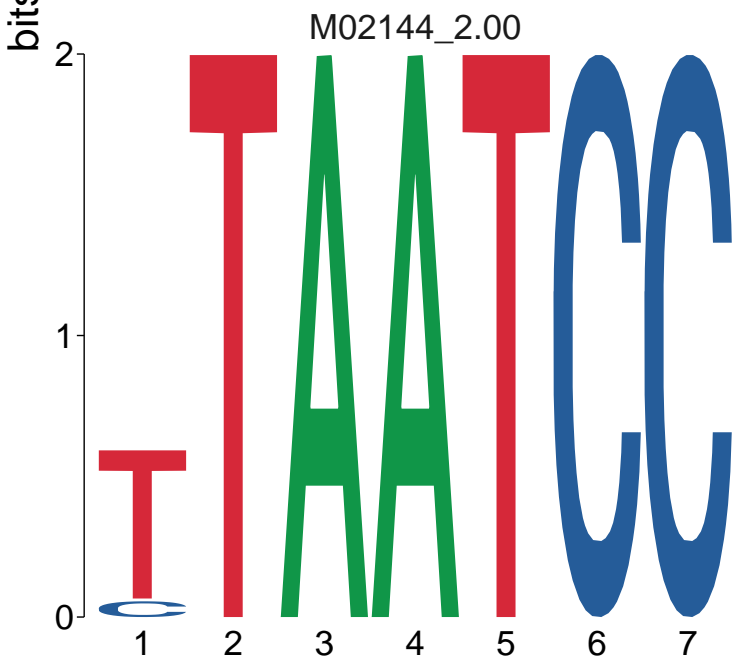
22-TKGATTAGCGCT,BestGuess:prd

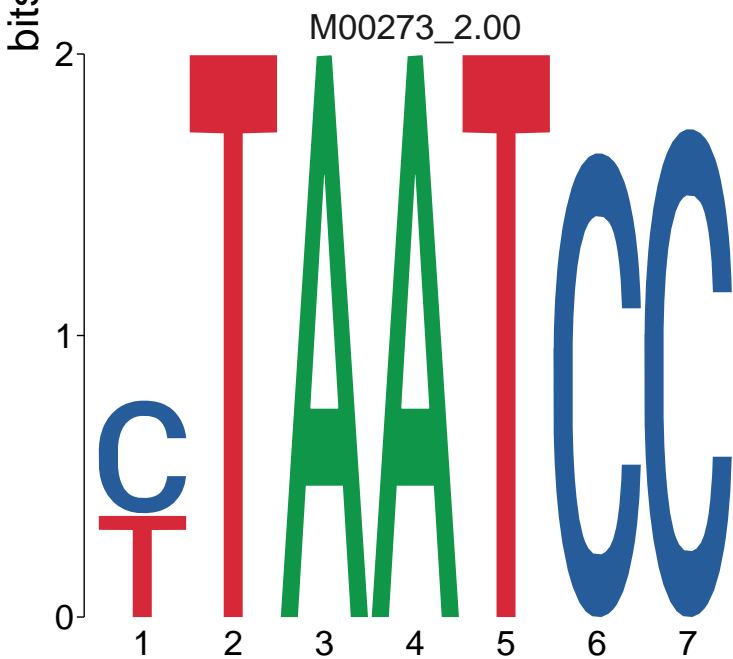
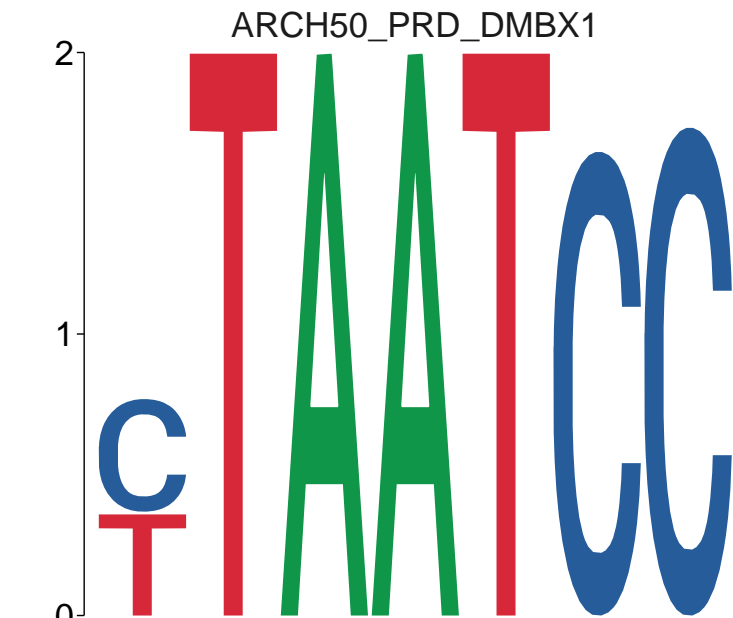


ARCH49_Homeodomains

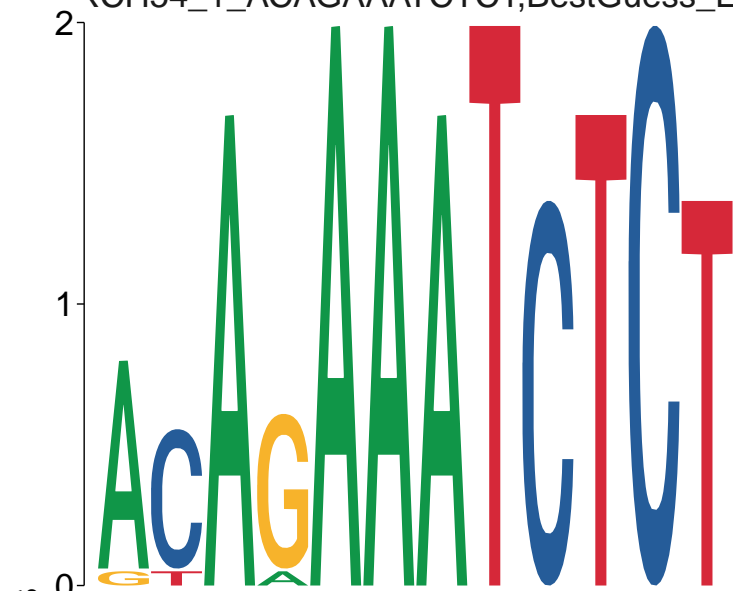


M02144_2.00

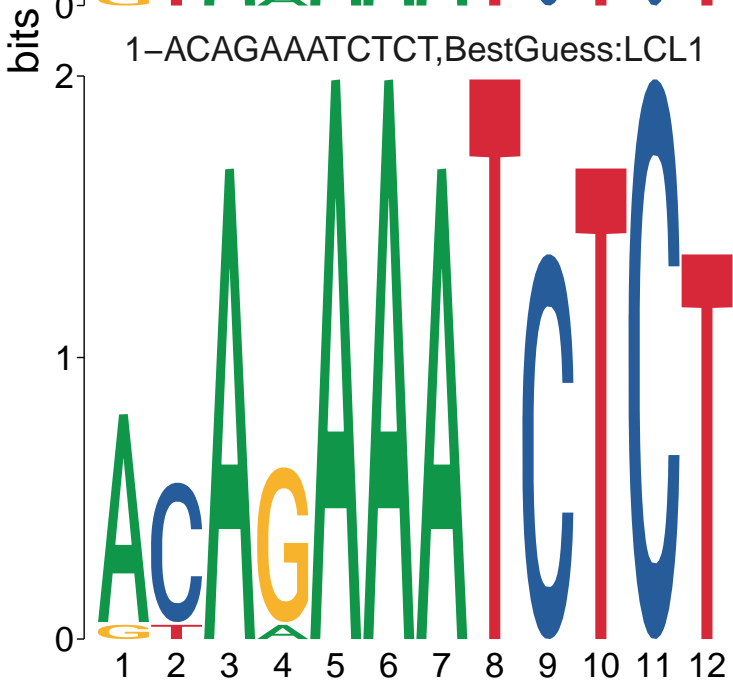




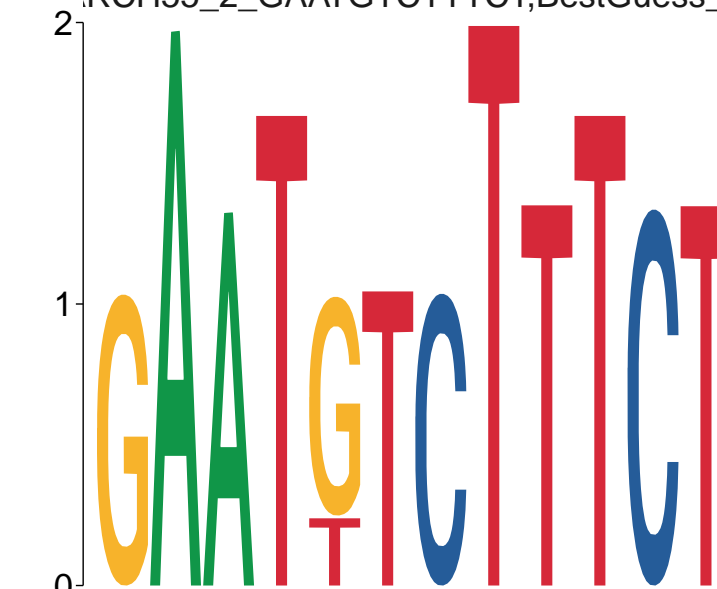
RCH54_1_ACAGAAATCTCT,BestGuess_LC



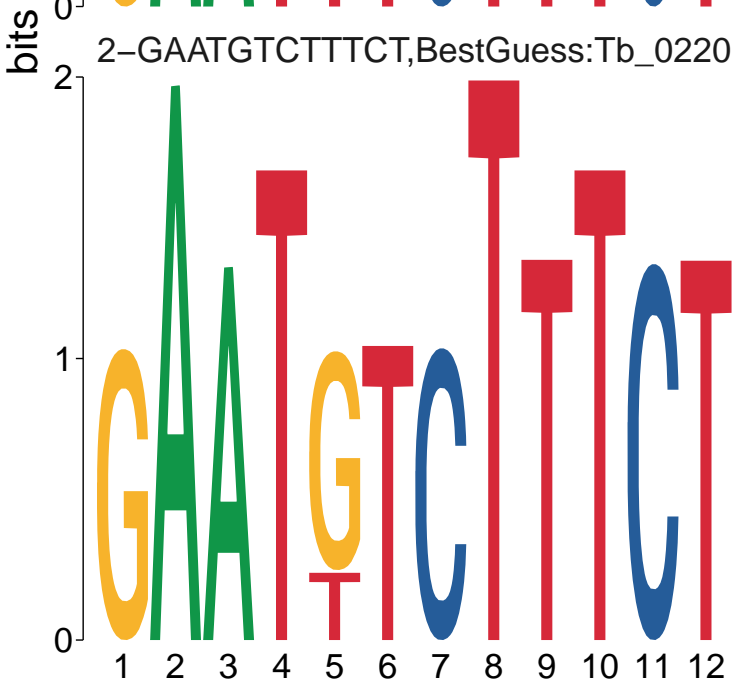
1-ACAGAAATCTCT,BestGuess:LCL1



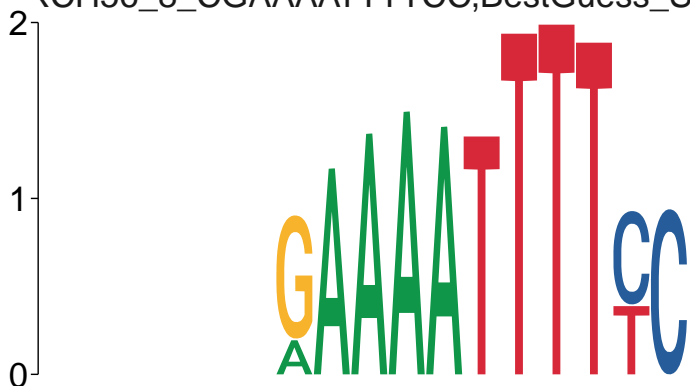
RCH55_2_GAATGTCTTTCT, BestGuess_T



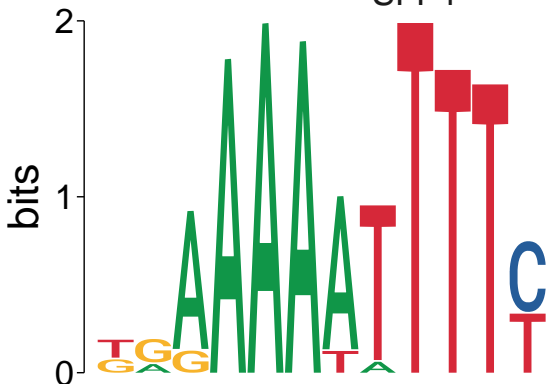
2-GAATGTCTTTCT, BestGuess:Tb_0220



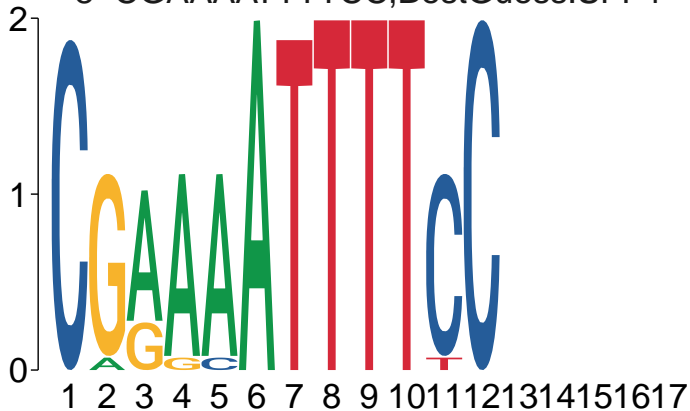
RCH56_8_CGAAAATTTTCC,BestGuess_Sf



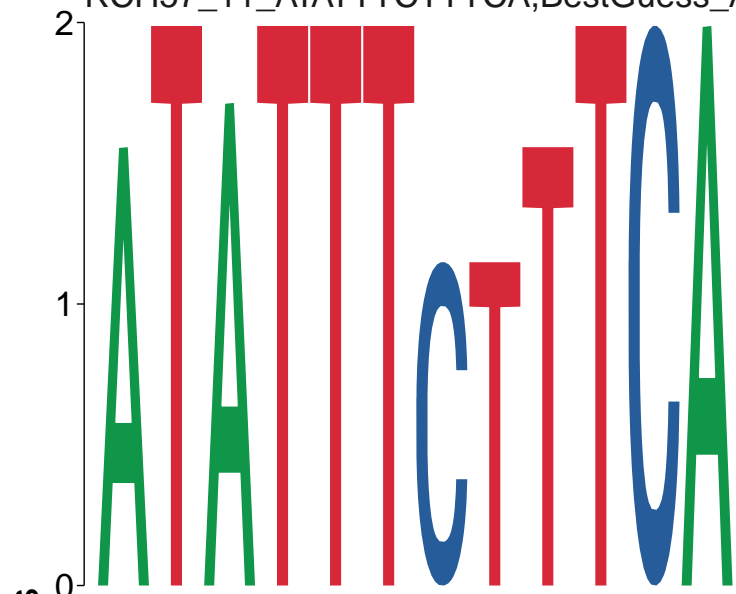
SFP1



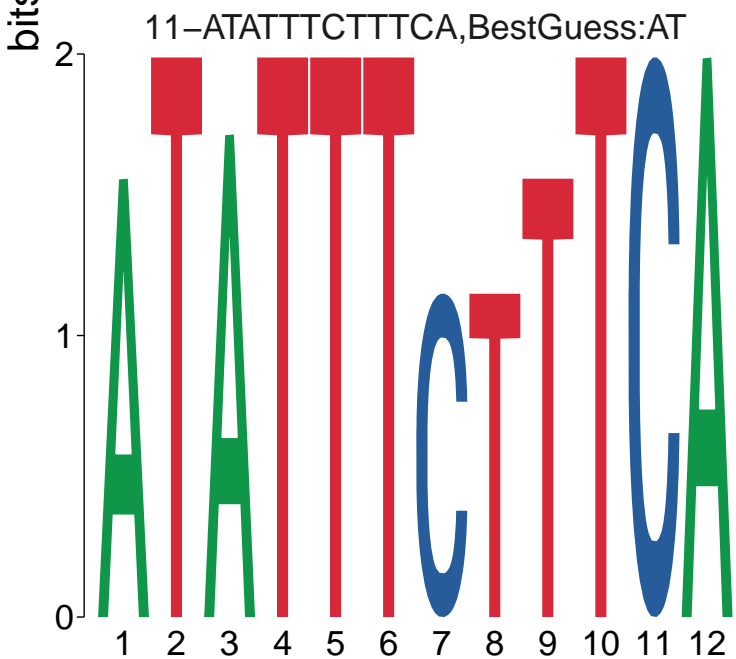
8-CGAAAATTTTCC,BestGuess:SFP1



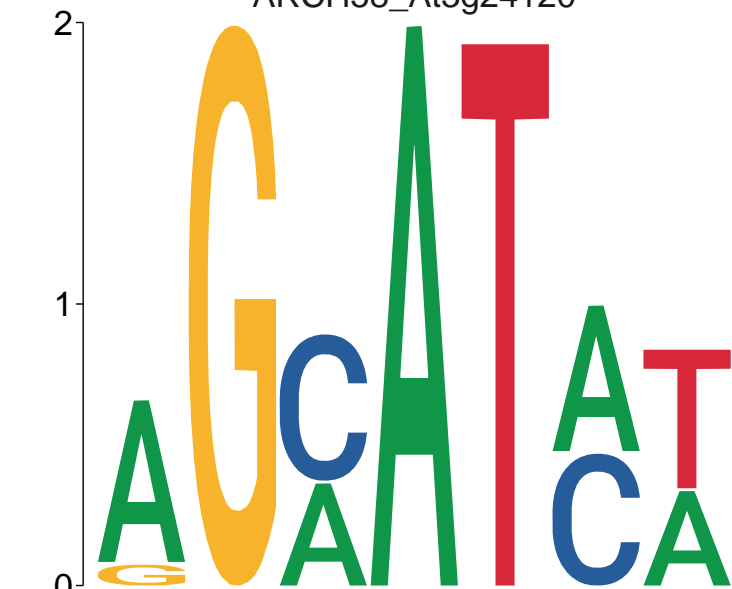
RCH57_11_ATATTTCTTTCA,BestGuess_A



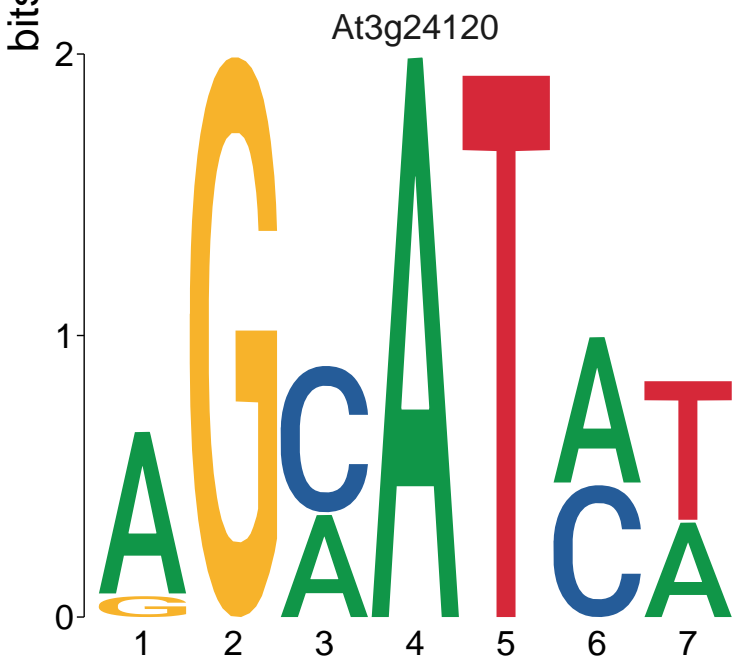
11-ATATTTCTTTCA,BestGuess:AT



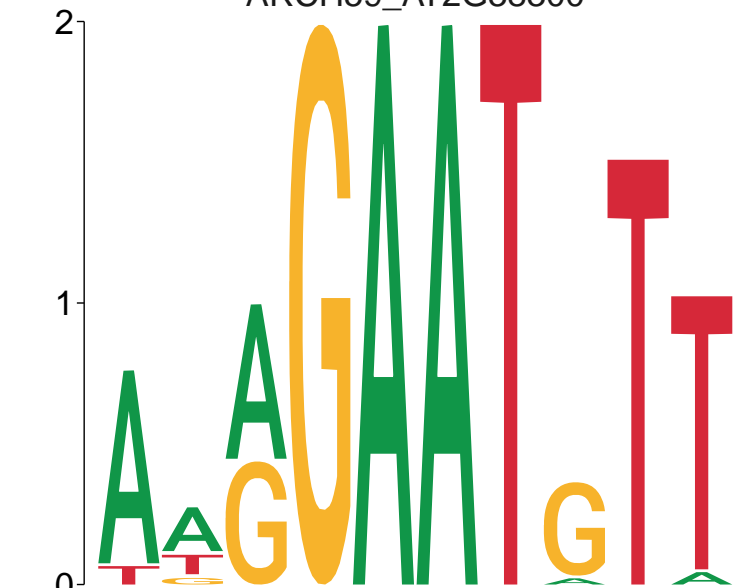
ARCH58_At3g24120



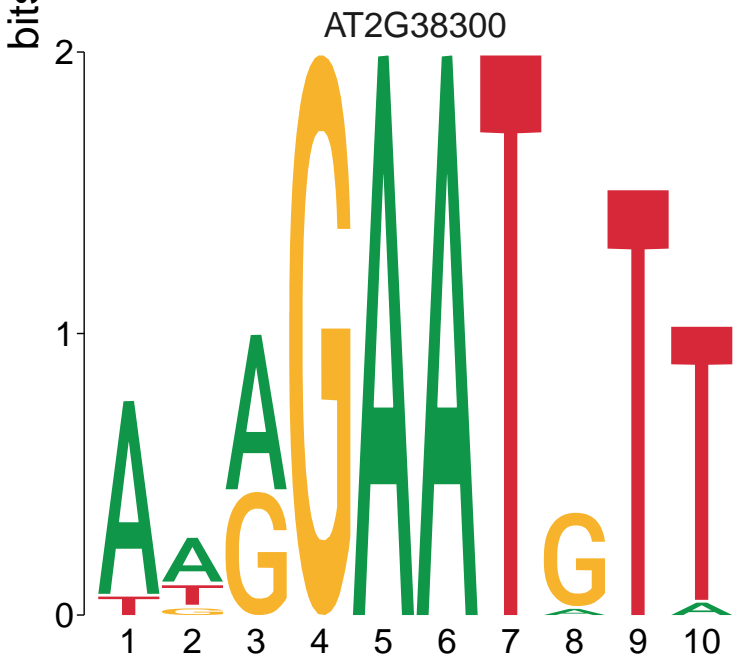
At3g24120



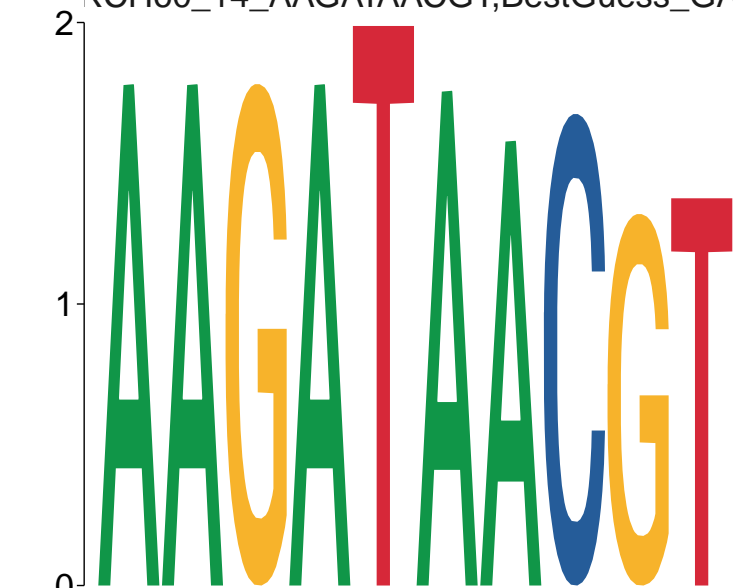
ARCH59_AT2G38300



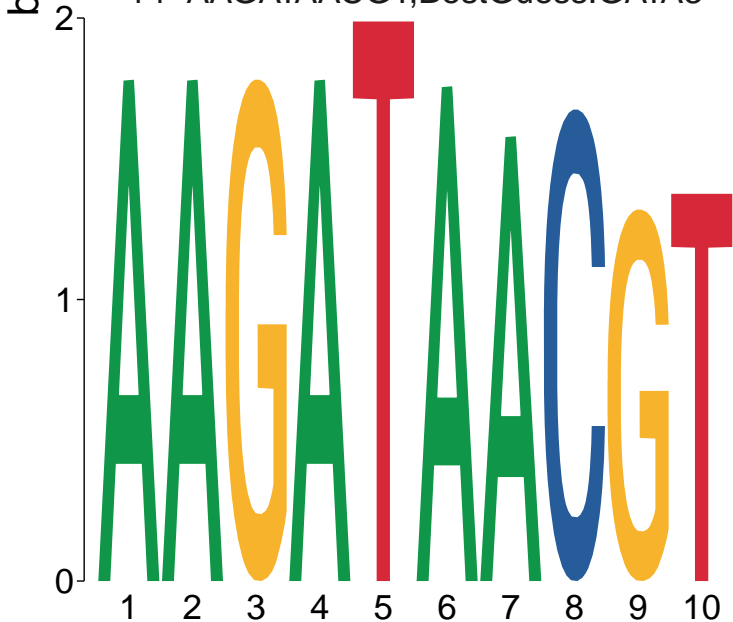
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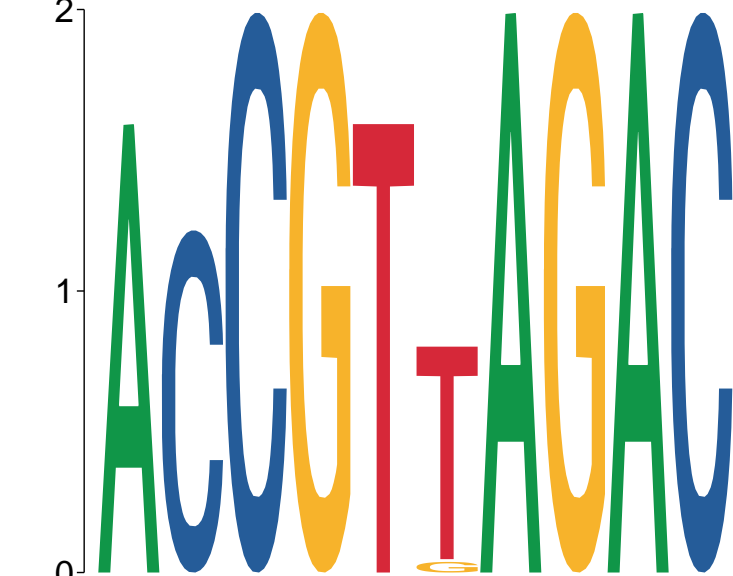
RCH60_14_AAGATAACGT,BestGuess_GA1



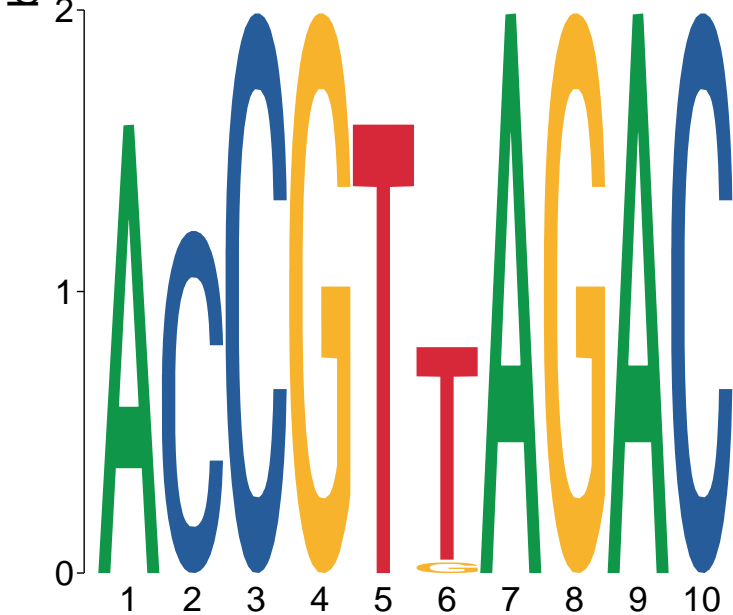
14-AAGATAACGT,BestGuess:GATA3



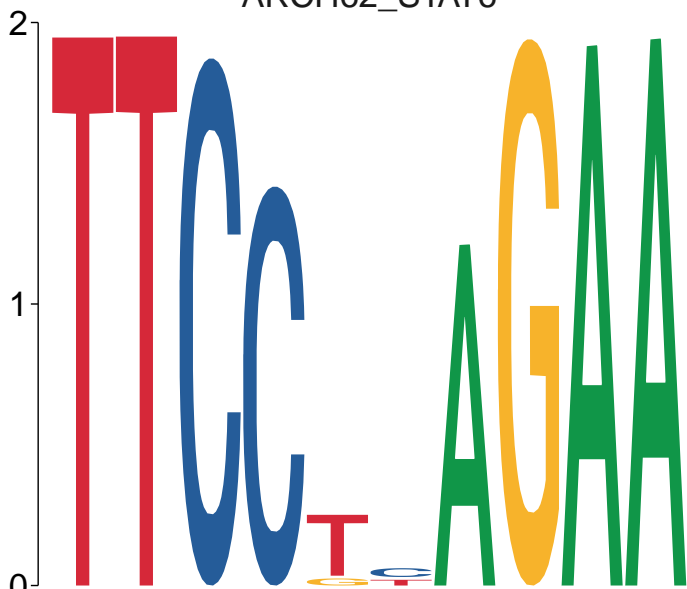
2RCH61_15_ACCGTTAGAC,BestGuess_OV0



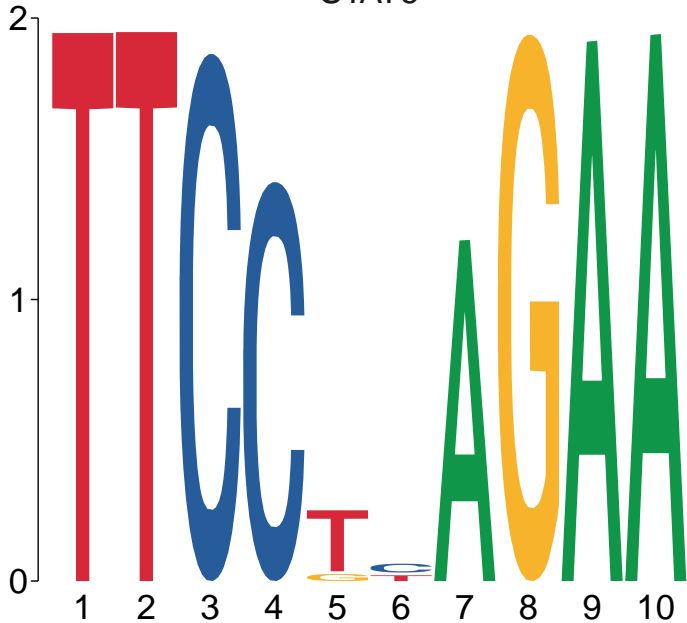
15-ACCGTTAGAC,BestGuess:OVOL2



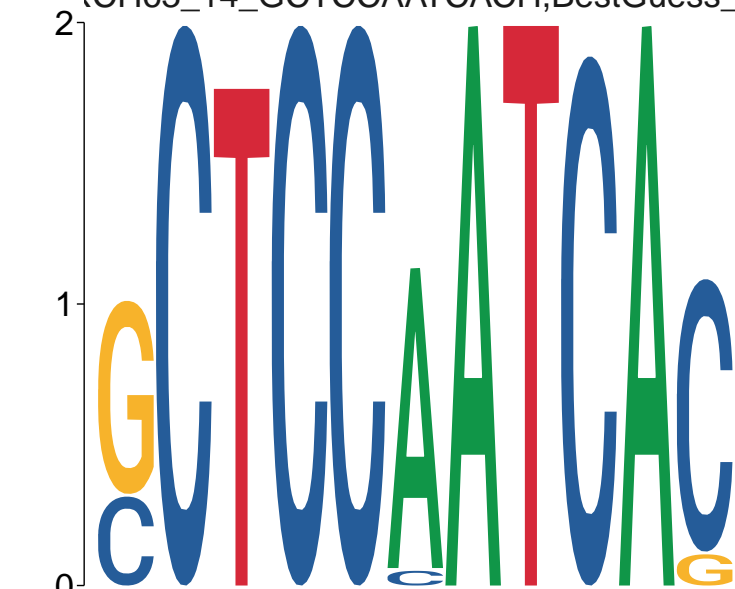
ARCH62_STAT6



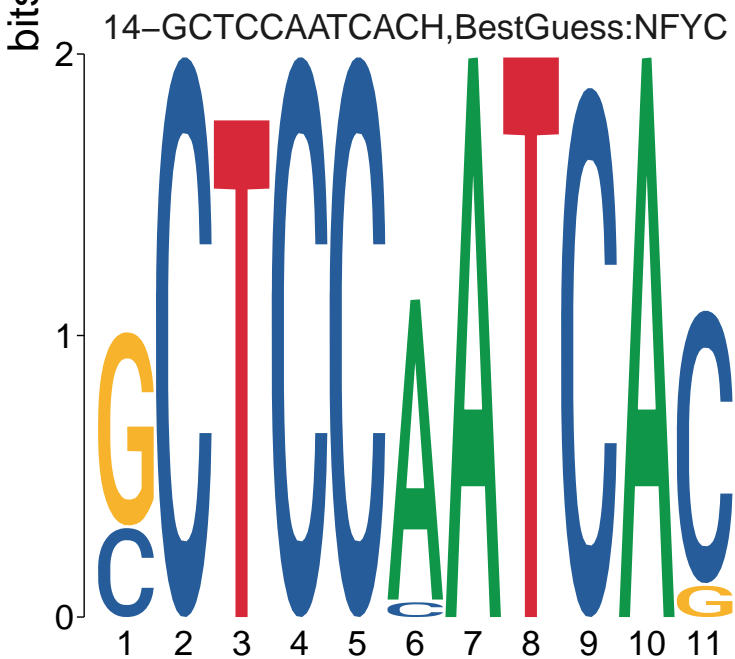
STAT6



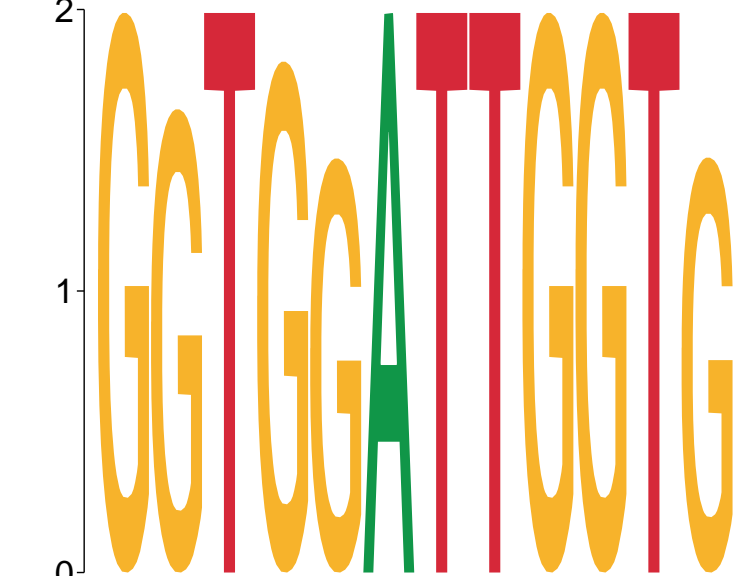
CH63_14_GCTCCAATCACH,BestGuess_I



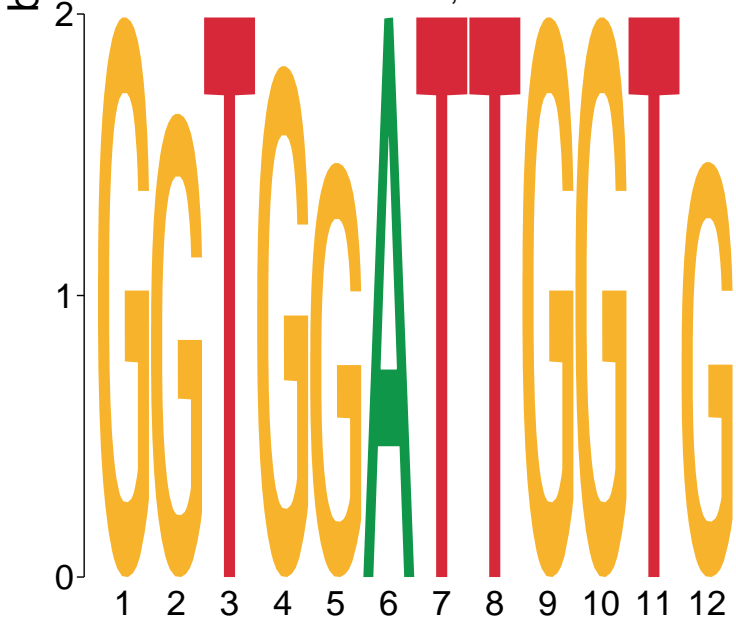
14-GCTCCAATCACH,BestGuess:NFYC



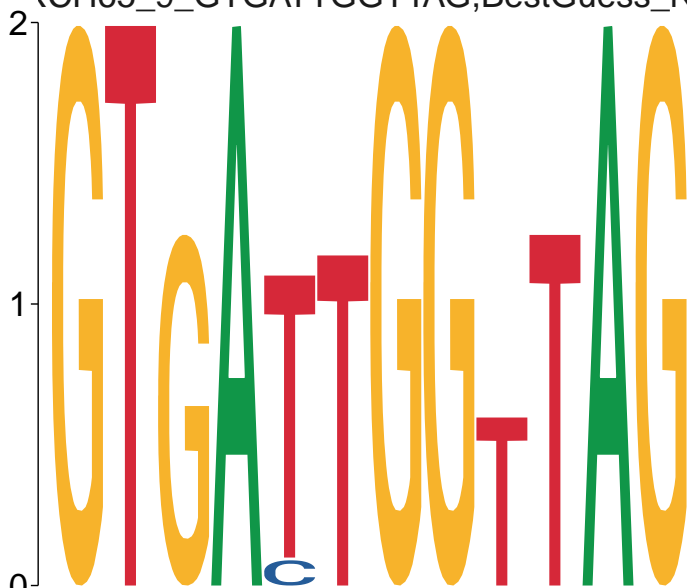
CH64_23_GGTGGATTGGTG,BestGuess_



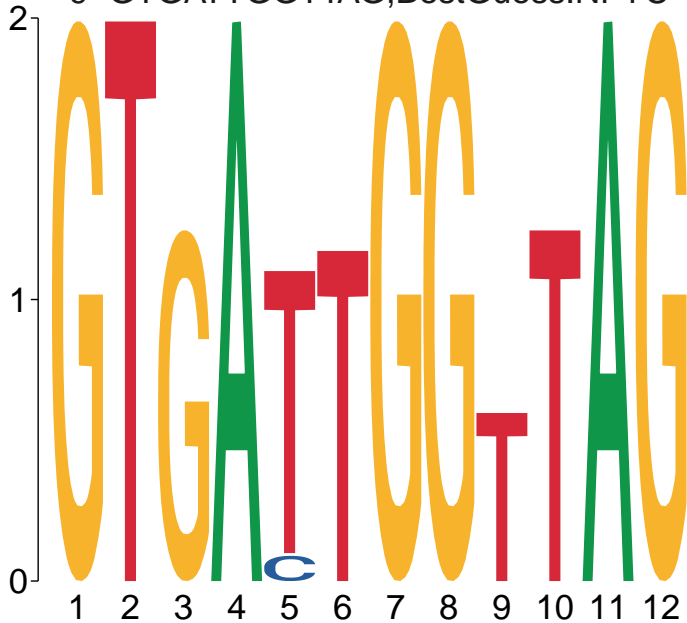
23-GGTGGATTGGTG,BestGuess:Foxh1



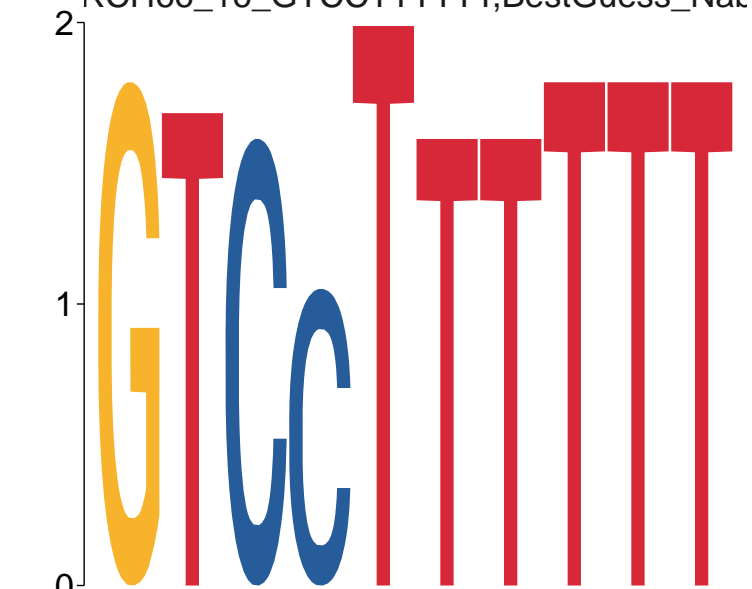
2CH65_9_GTGATTGGTTAG,BestGuess_NI



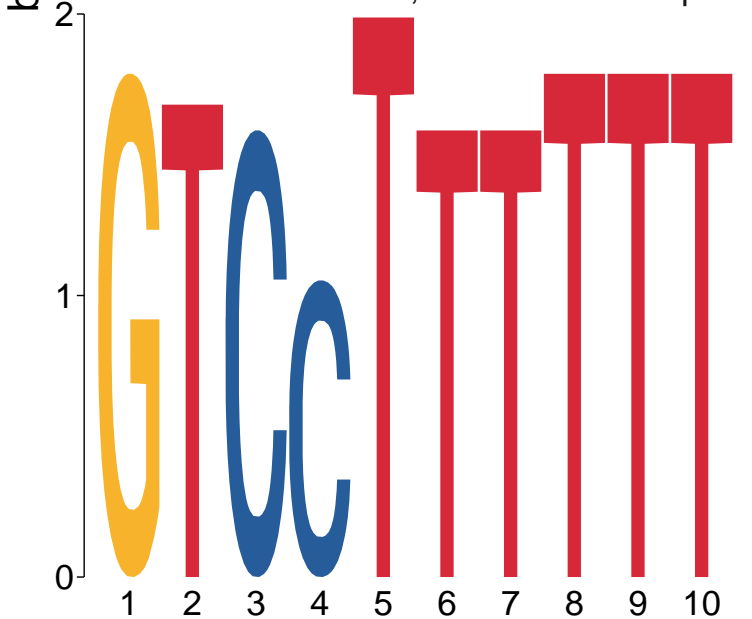
bits 9-GTGATTGGTTAG,BestGuess:NFYC



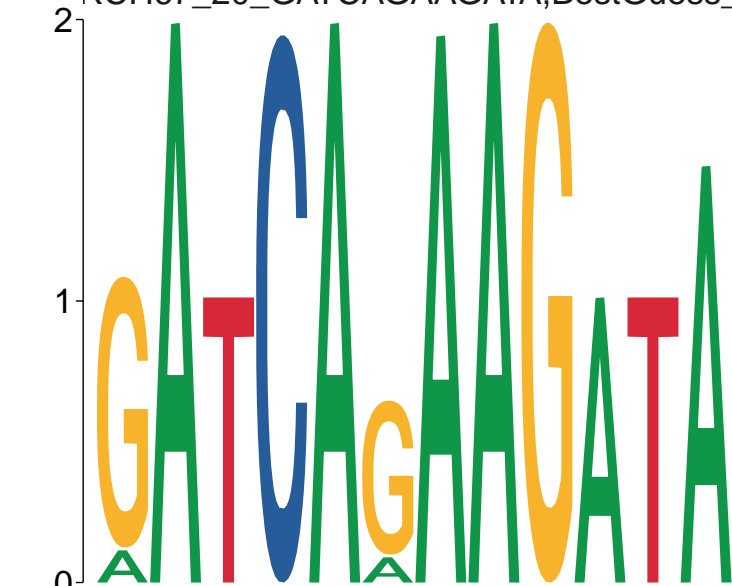
RCH66_16_GTCCTTTTTT,BestGuess_Nak



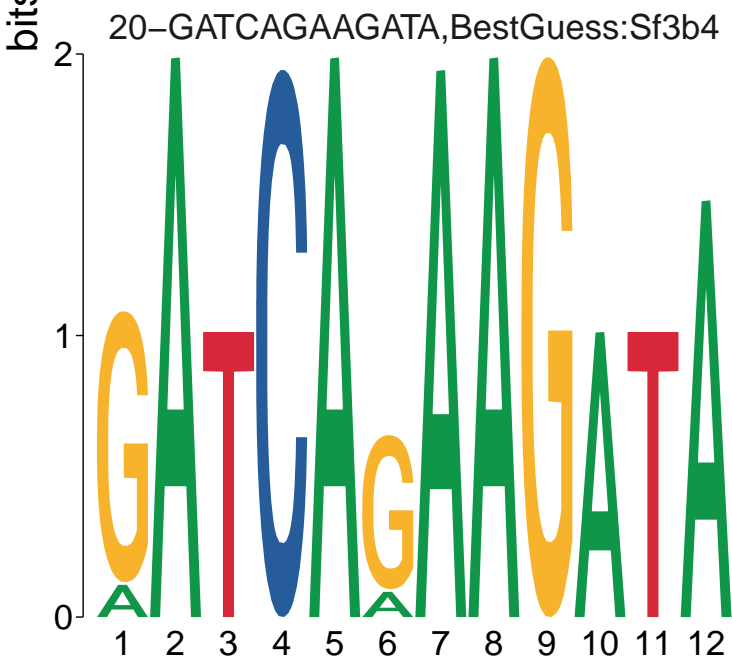
16-GTCCTTTTTT,BestGuess:Nab2p



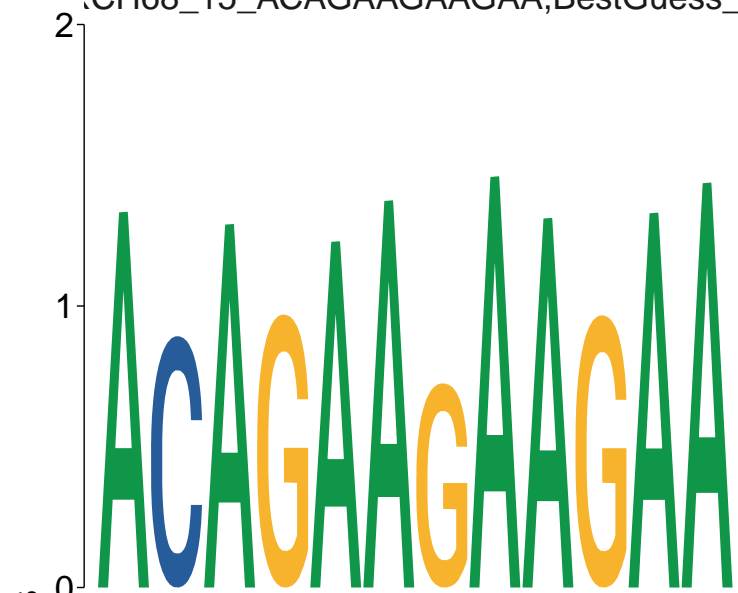
RCH67_20_GATCAGAAGATA,BestGuess_!



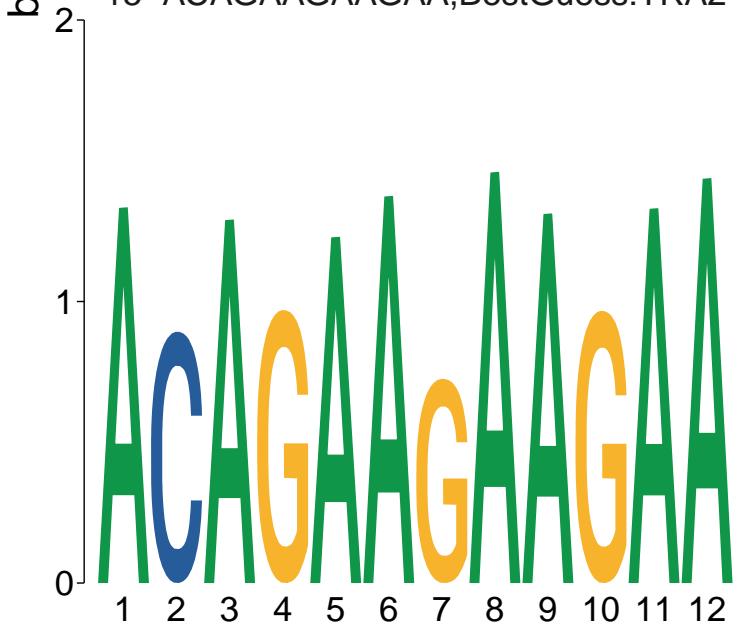
20-GATCAGAAGATA,BestGuess:Sf3b4



CH68_15_ACAGAAGAAGAA,BestGuess_



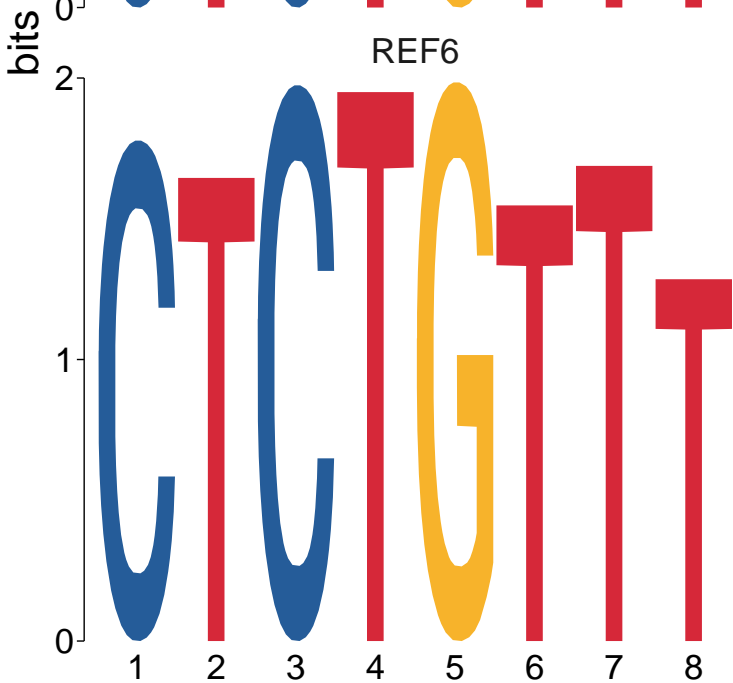
15-ACAGAAGAAGAA,BestGuess:TRA2



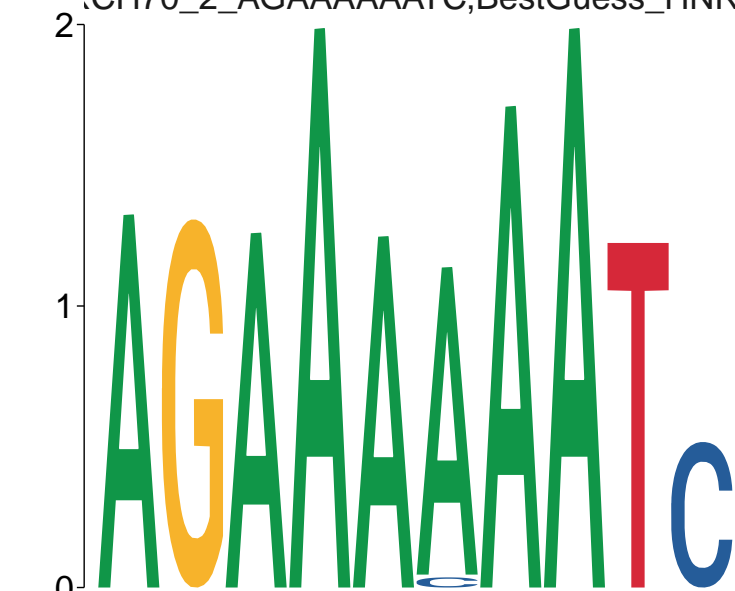
ARCH69_REF6



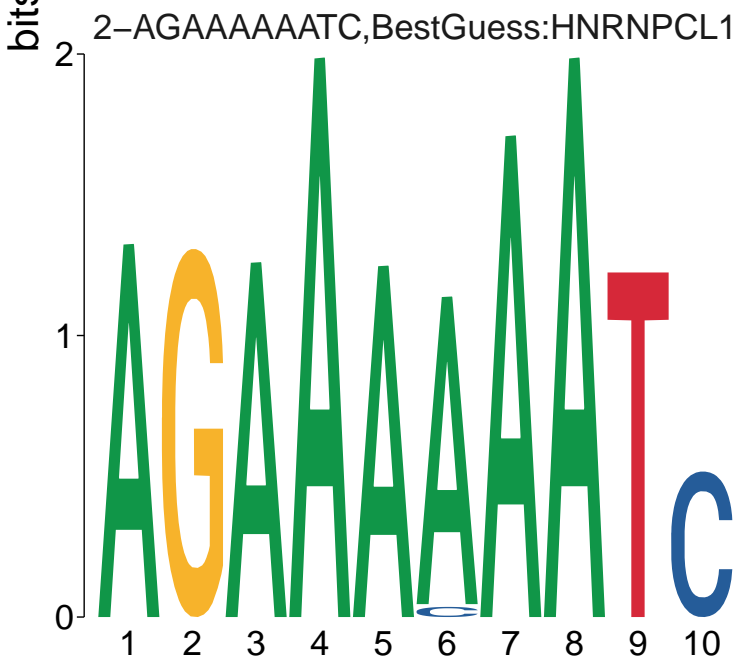
REF6



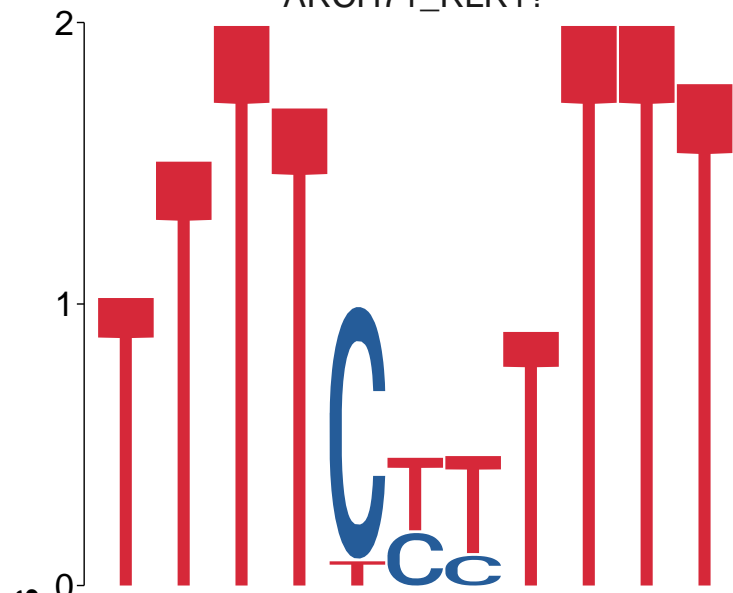
CH70_2_AGAAAAAATC,BestGuess_HNRI



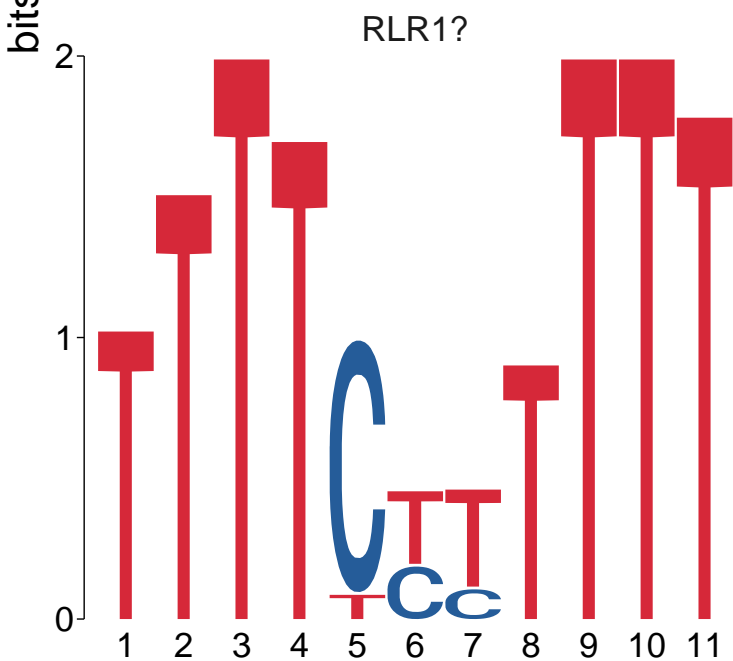
2-AGAAAAAATC,BestGuess:HNRPCL1



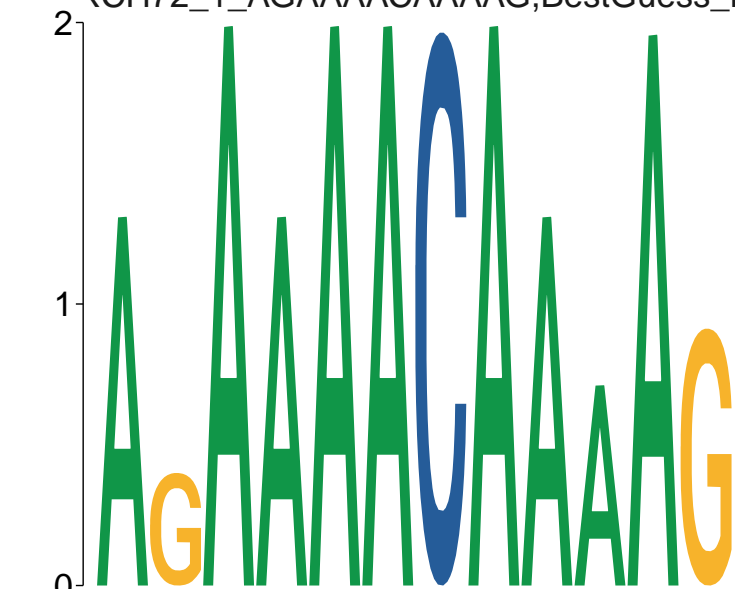
ARCH71_RLR1?



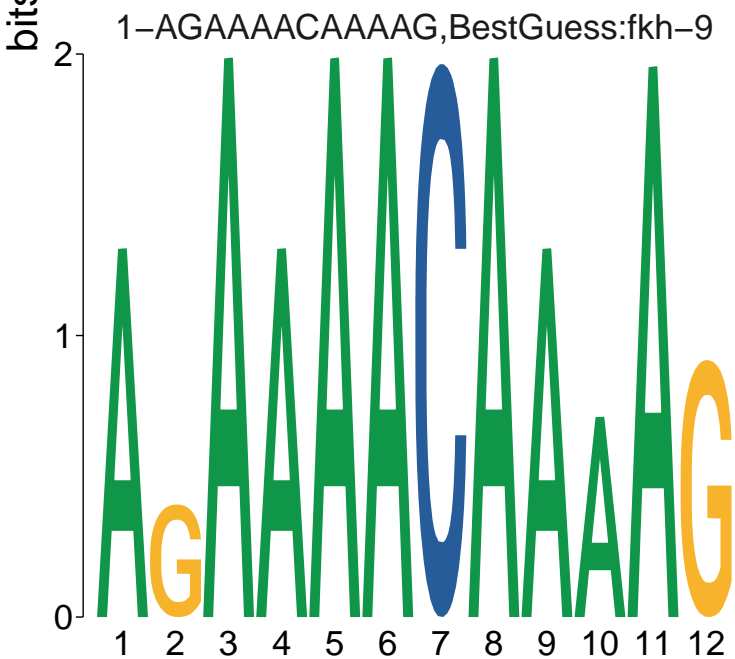
RLR1?



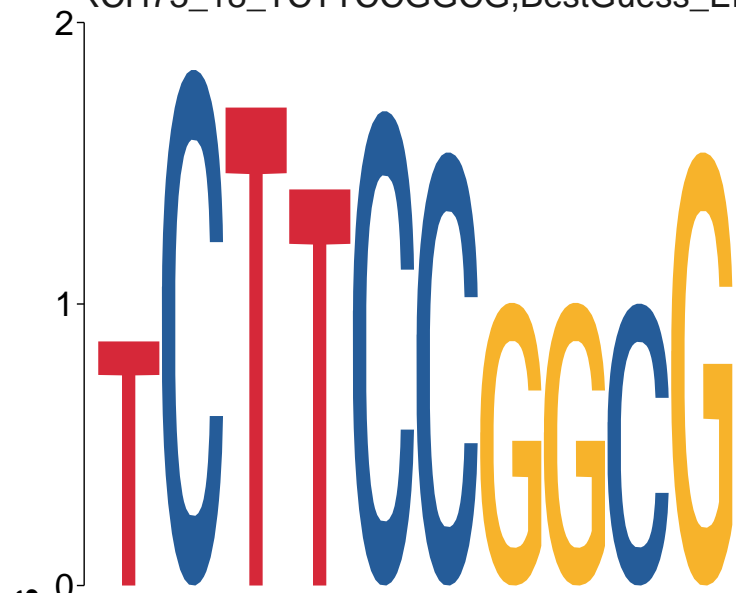
RCH72_1_AGAAAACAAAAG,BestGuess_fl



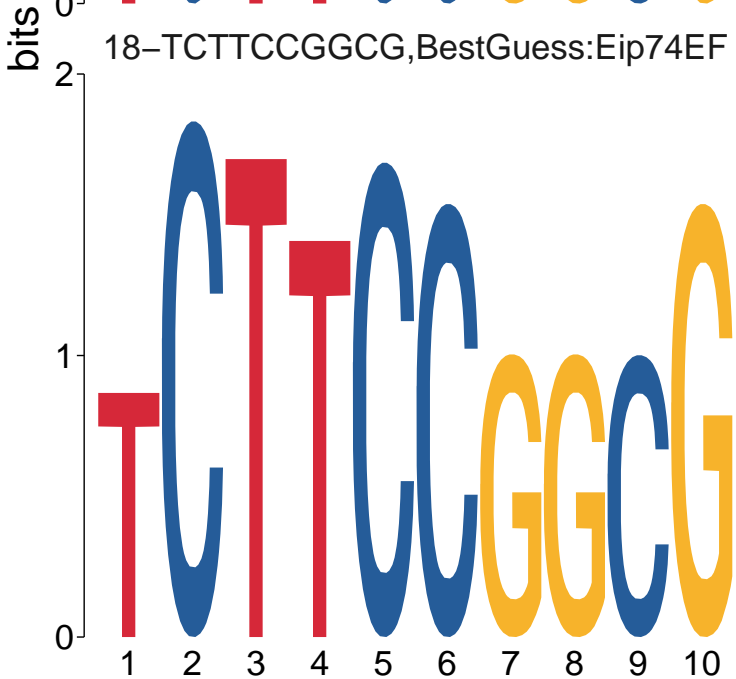
1-AGAAAACAAAAG,BestGuess:fkh-9



RCH73_18_TCTTCCGGCG,BestGuess_Eip



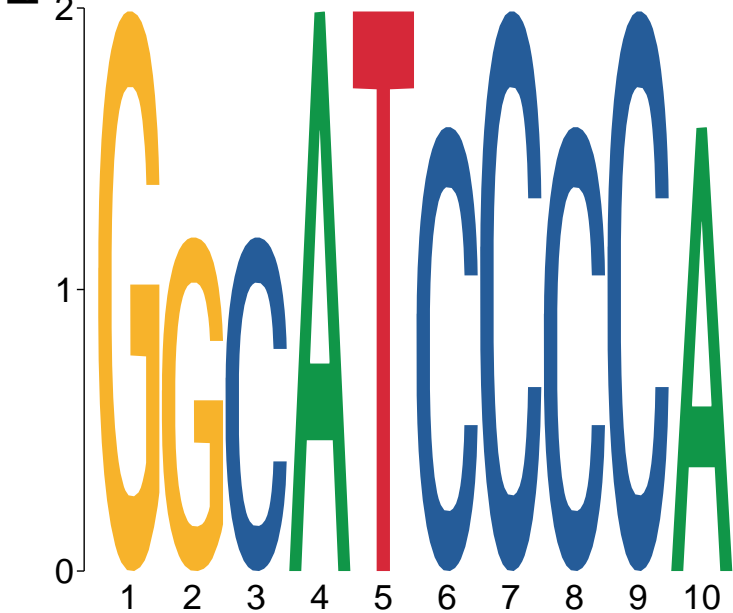
18-TCTTCCGGCG,BestGuess:Eip74EF



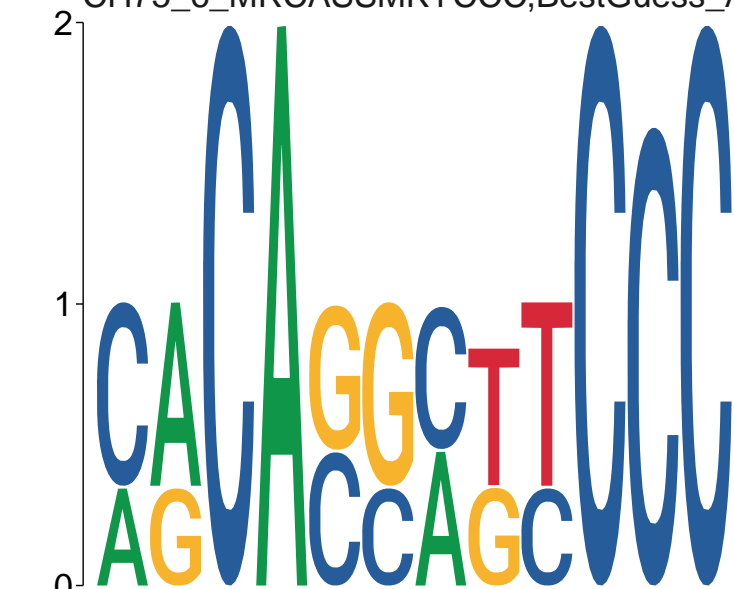
CH74_21_GGCATCCCCA,BestGuess_ESI



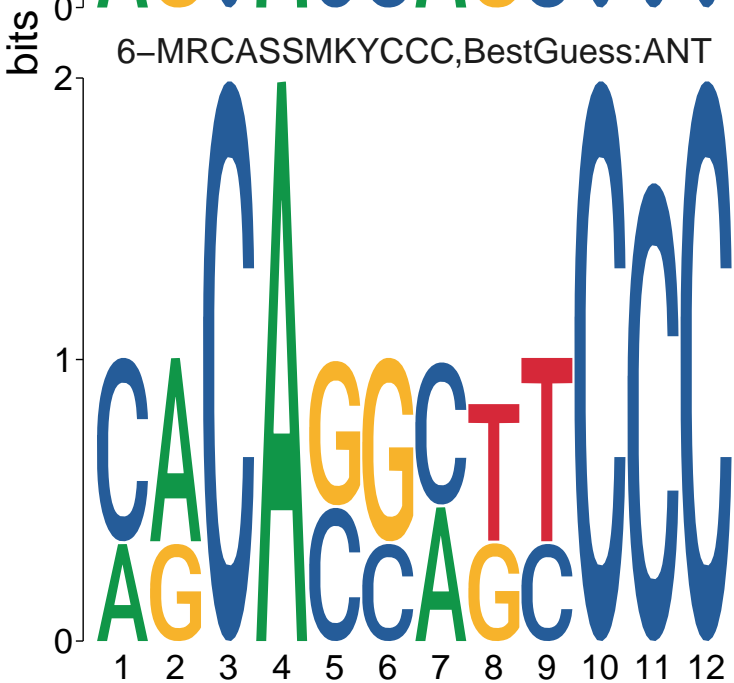
21-GGCATCCCCA,BestGuess:ESRP2



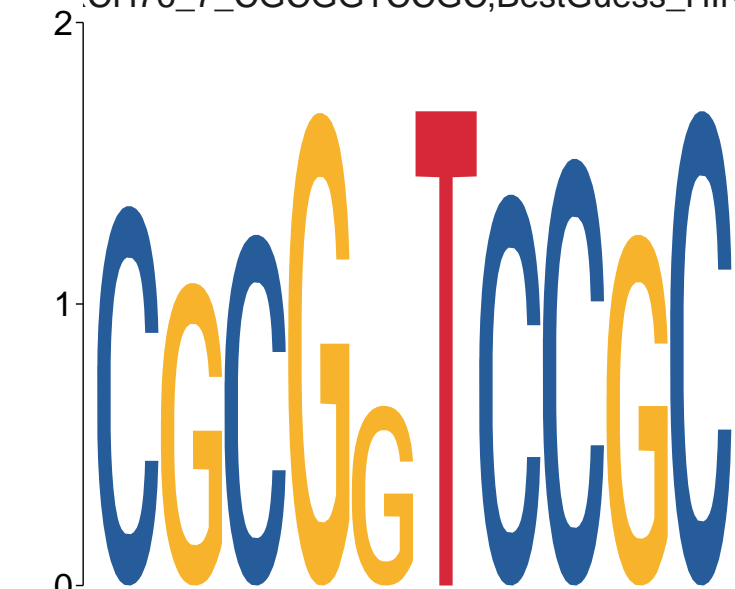
CH75_6_MRCASSMKYCCC,BestGuess_A



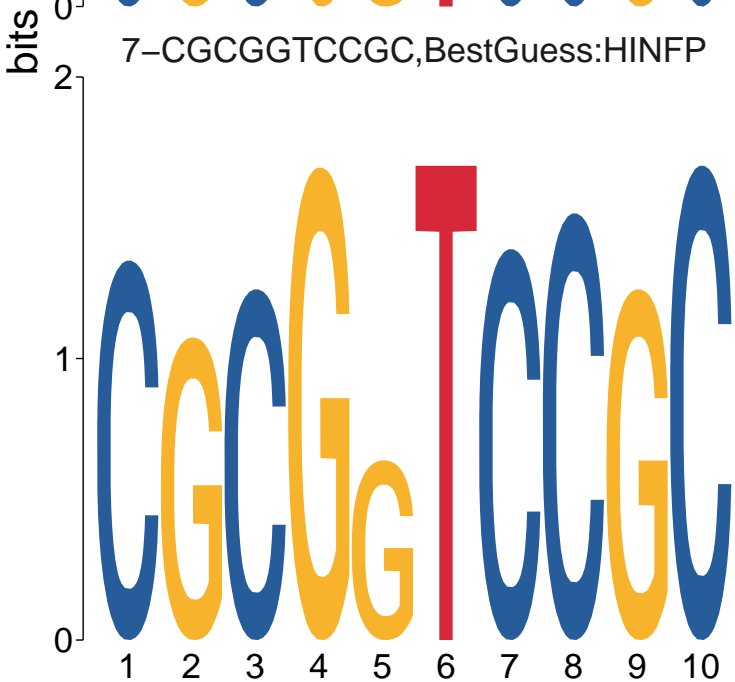
6-MRCASSMKYCCC,BestGuess:ANT



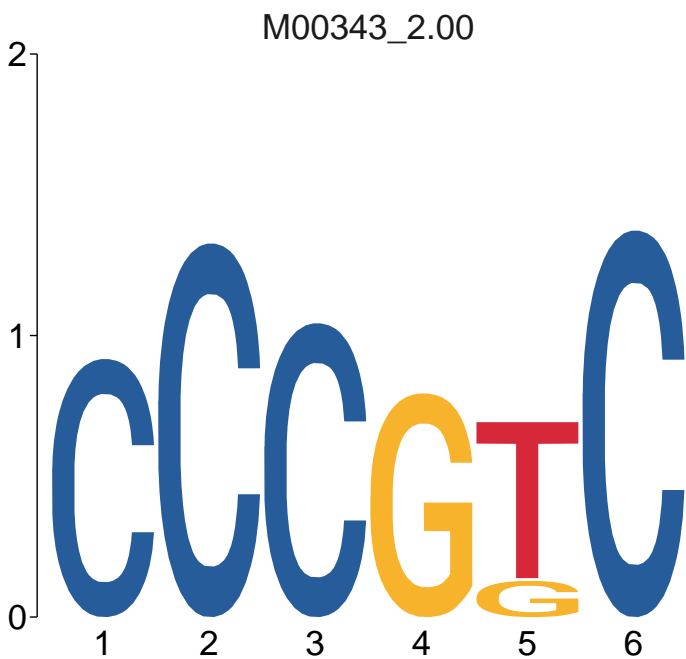
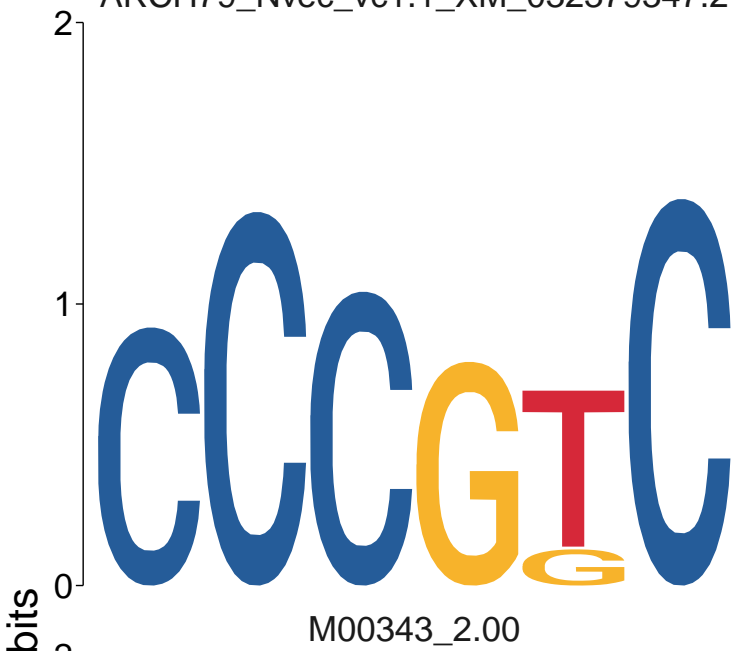
CH76_7_CGCGGTCCGC,BestGuess_HIN



7-CGCGGTCCGC,BestGuess:HINFP



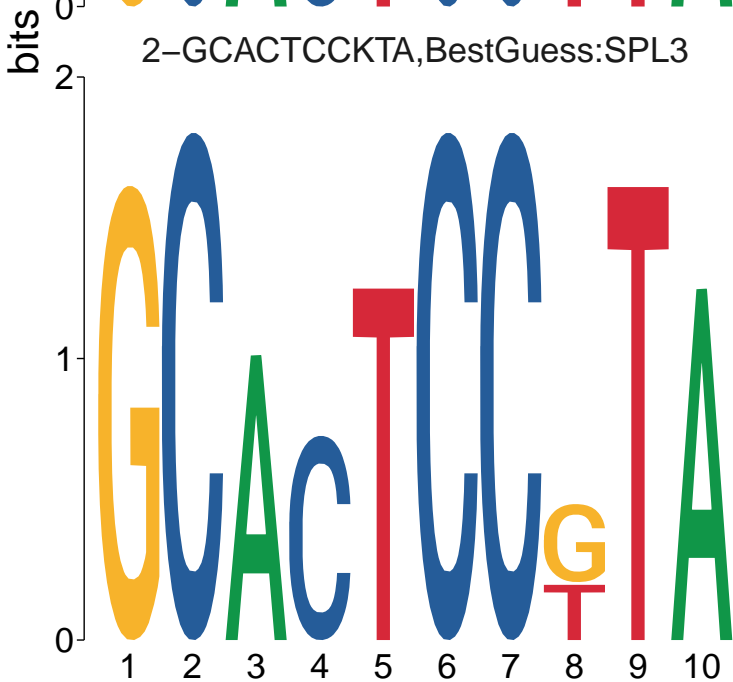
ARCH79_Nvec_vc1.1_XM_032379347.2



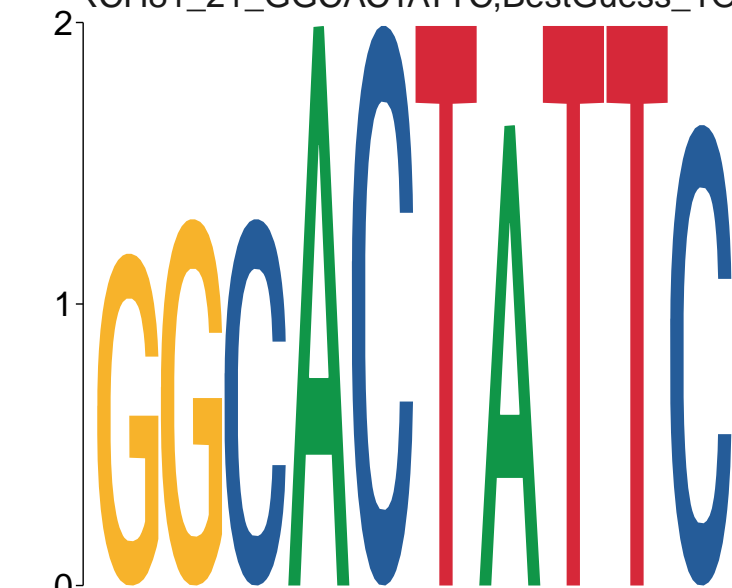
RCH80_2_GCACTCCKTA,BestGuess_SPL



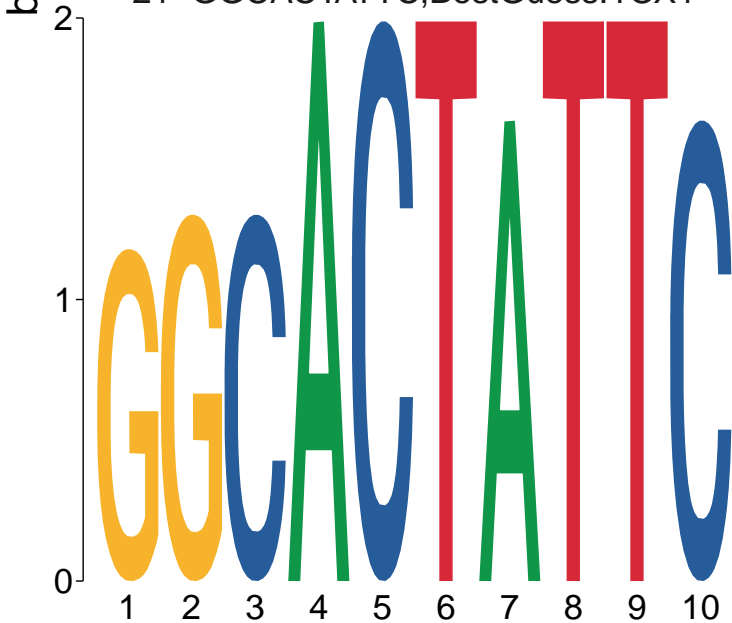
2-GCACTCCKTA,BestGuess:SPL3



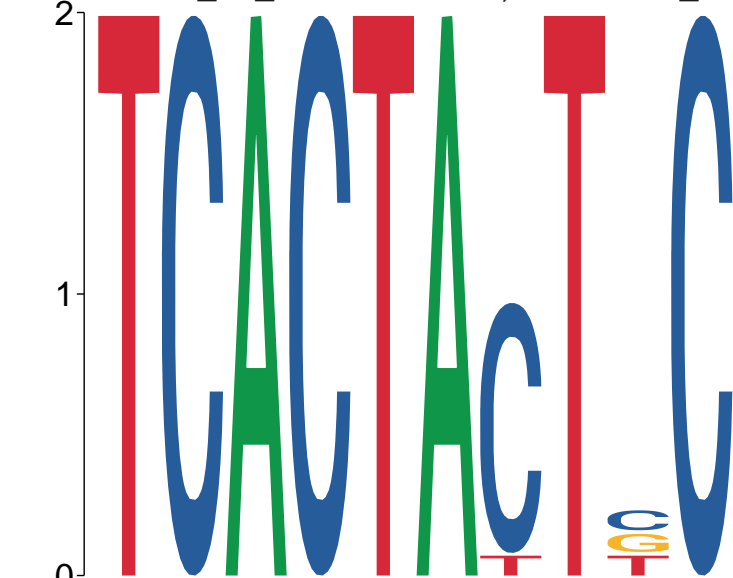
RCH81_21_GGCAC TATTC, BestGuess_YO)



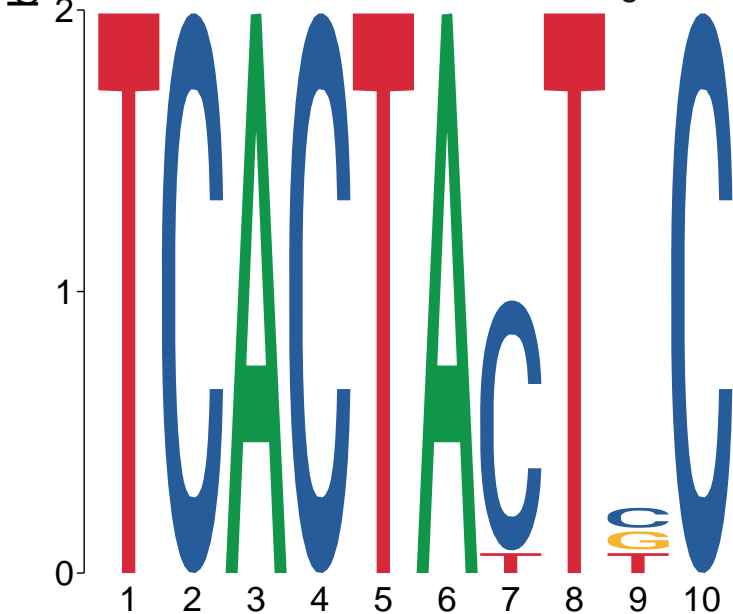
21-GGCAC TATTC, BestGuess:YOX1



RCH82_10_TCACTACTBC,BestGuess_At5

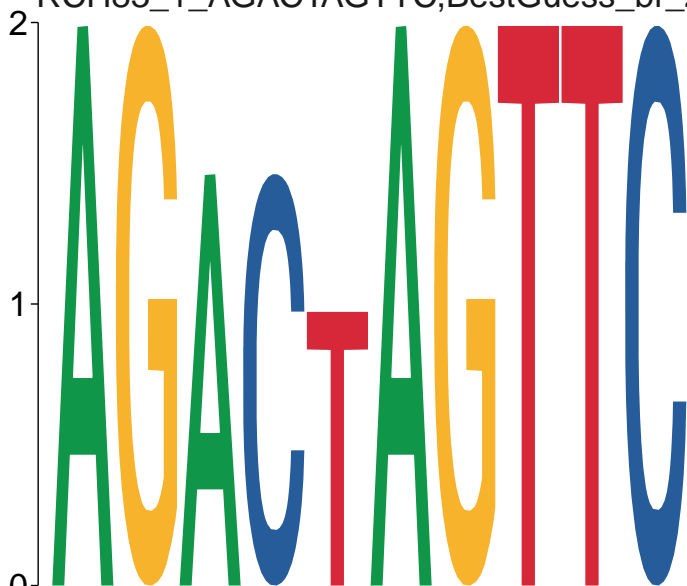


10-TCACTACTBC,BestGuess:At5g04390

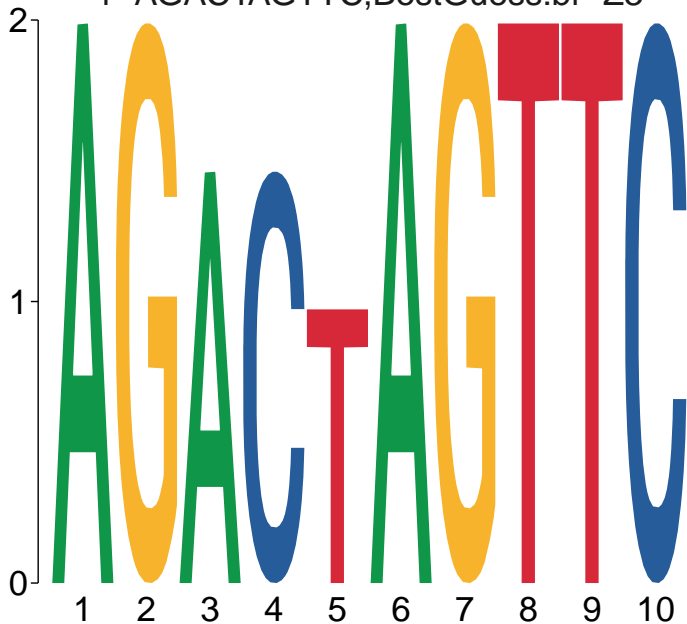


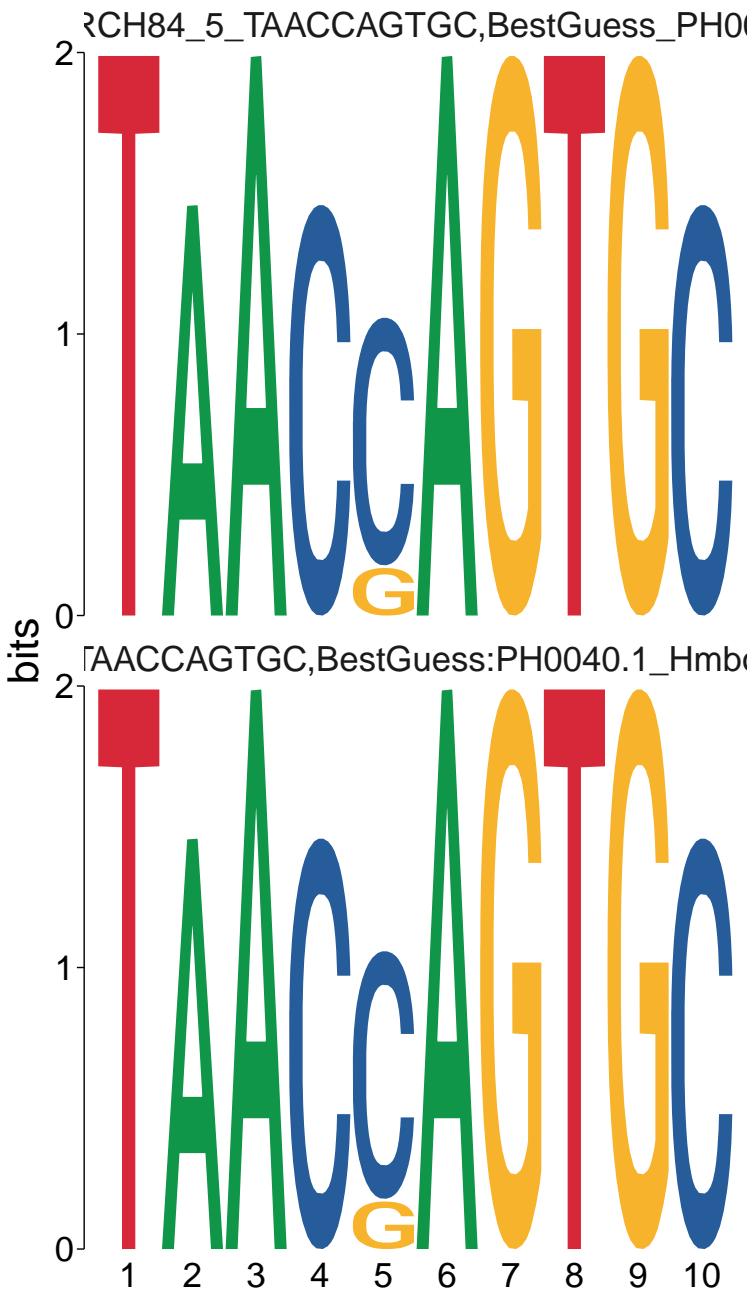
RCH83_1_AGACTAGTTC,BestGuess_br_Z

bits

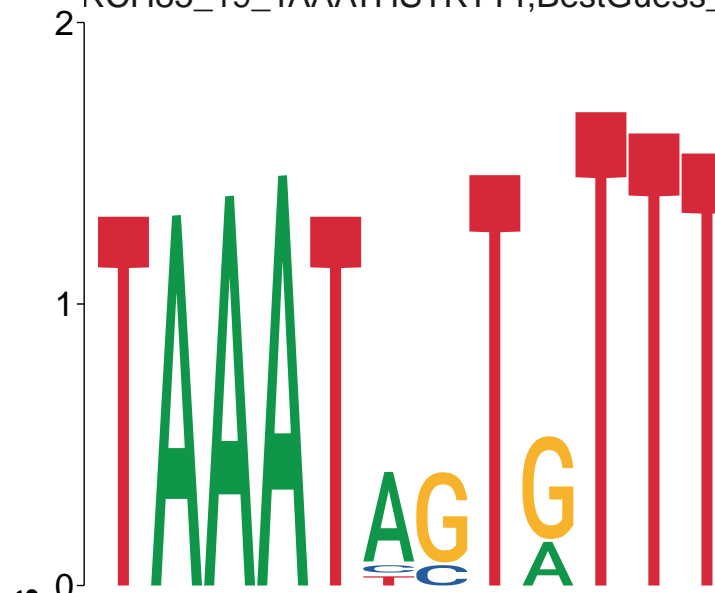


1-AGACTAGTTC,BestGuess:br-Z3

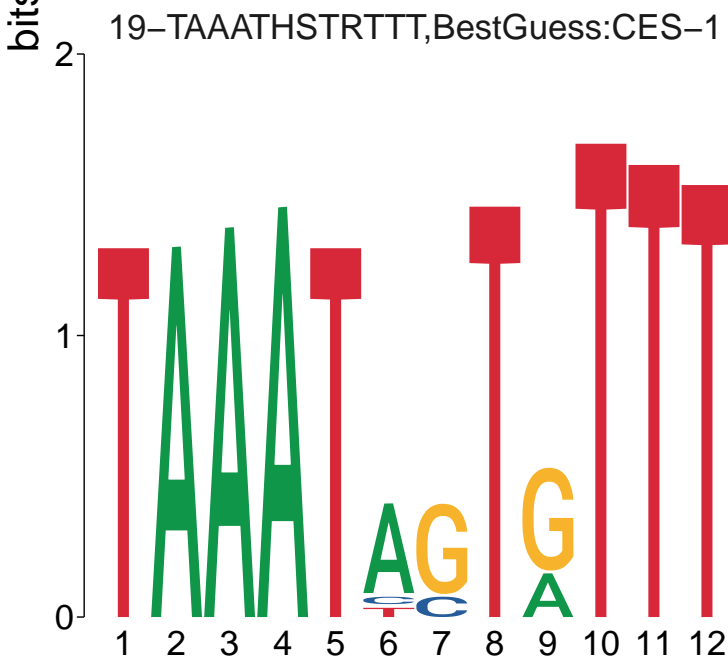




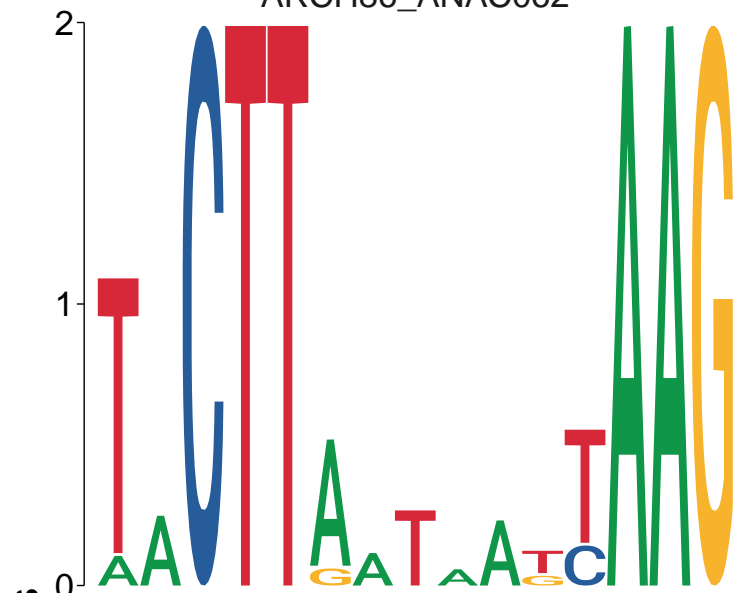
RCH85_19_TAAATHSTRTTT,BestGuess_C



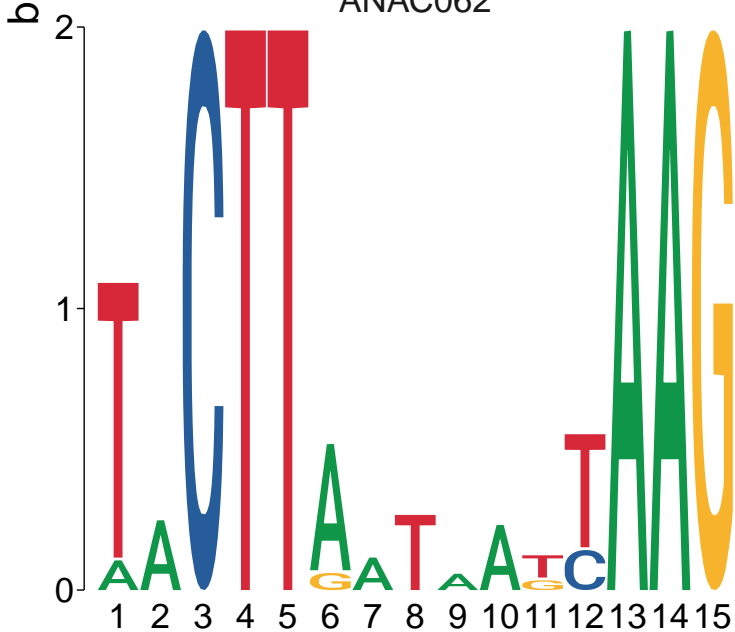
19-TAAATHSTRTTT,BestGuess:CES-1



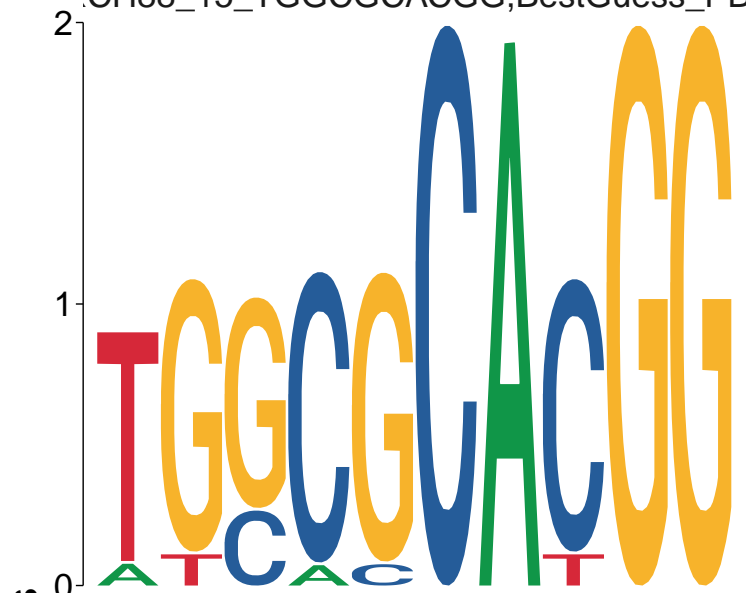
ARCH86_ANAC062



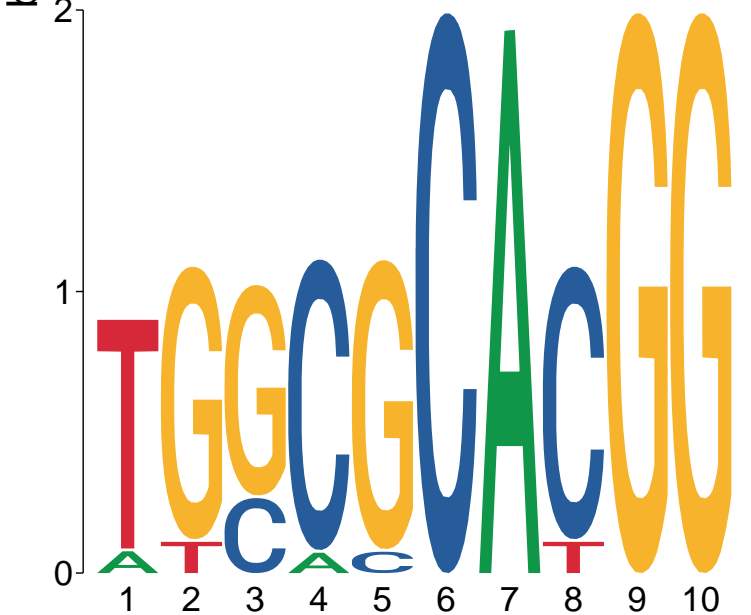
ANAC062



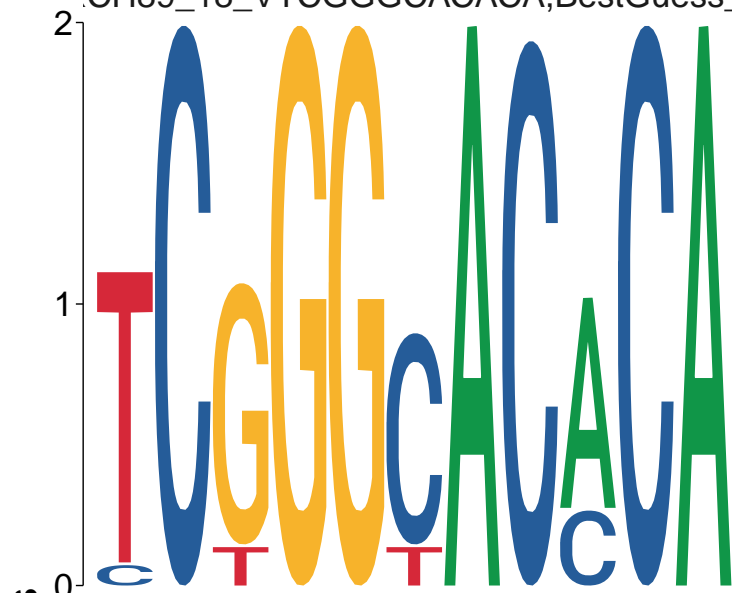
CH88_15_TGGCGCACGG,BestGuess_PB



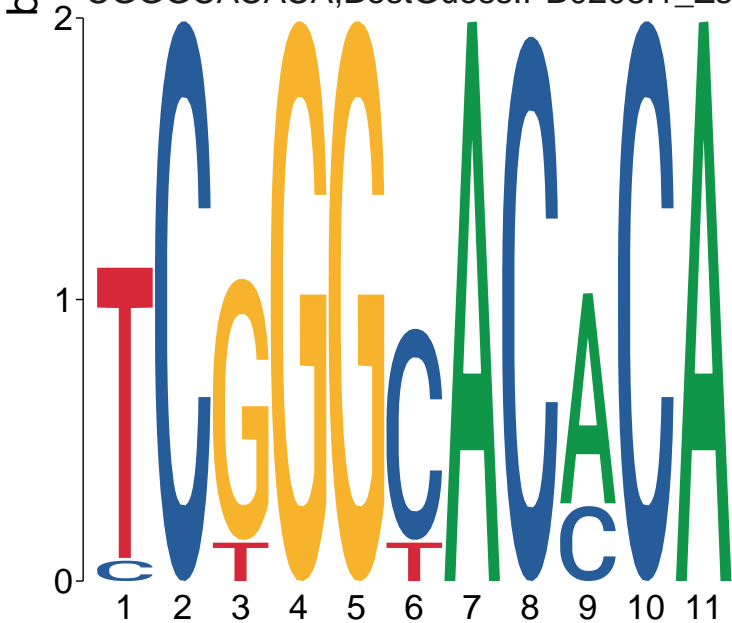
TGGCGCACGG,BestGuess:PB0044.1_Mtf



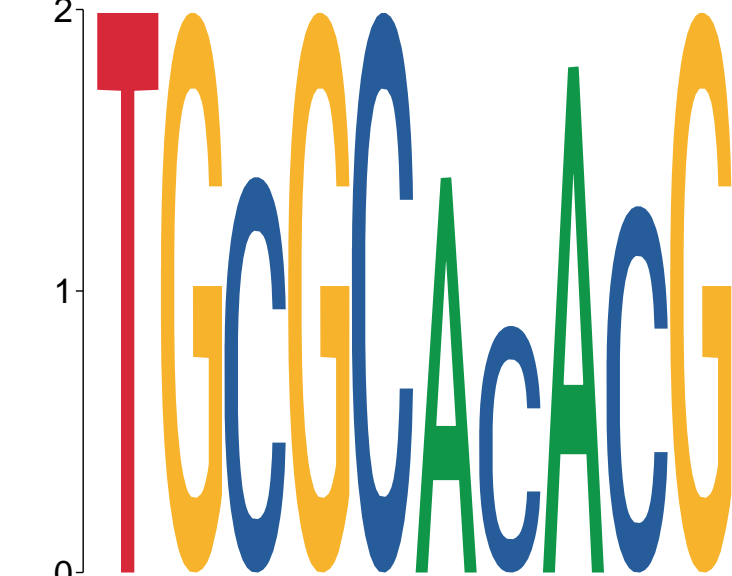
CH89_18_VTCGGGGCACACA,BestGuess_



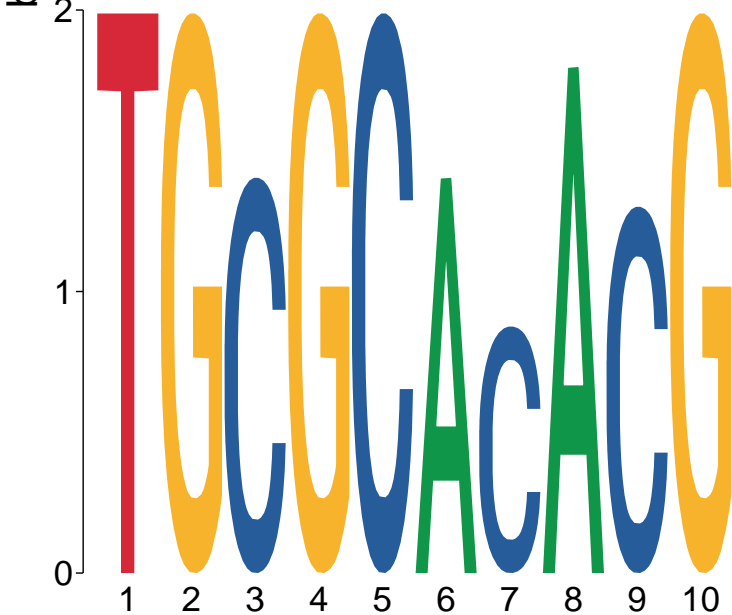
CGGGGCACACA,BestGuess:PB0208.1_Zsc



CH90_17_TGCGCACACG,BestGuess_NR



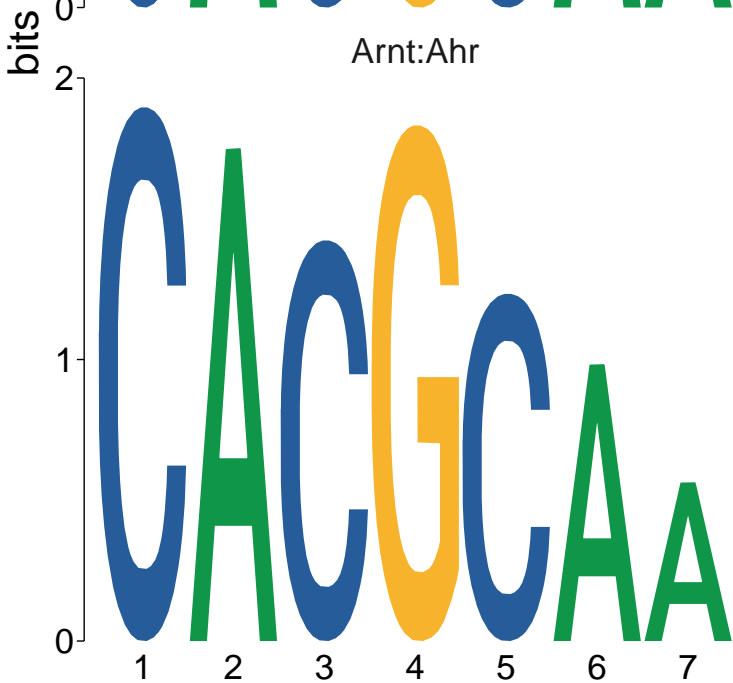
17-TGCGCACACG,BestGuess:NRF1



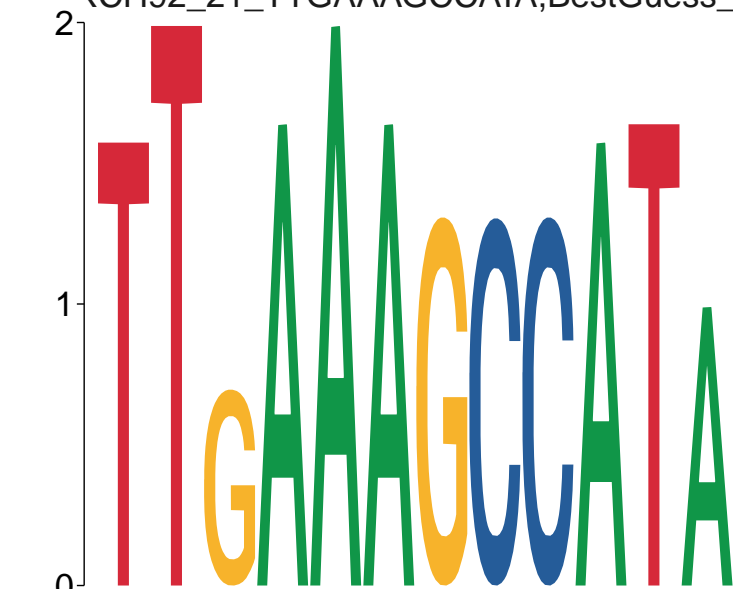
ARCH91_Arnt_Ahr



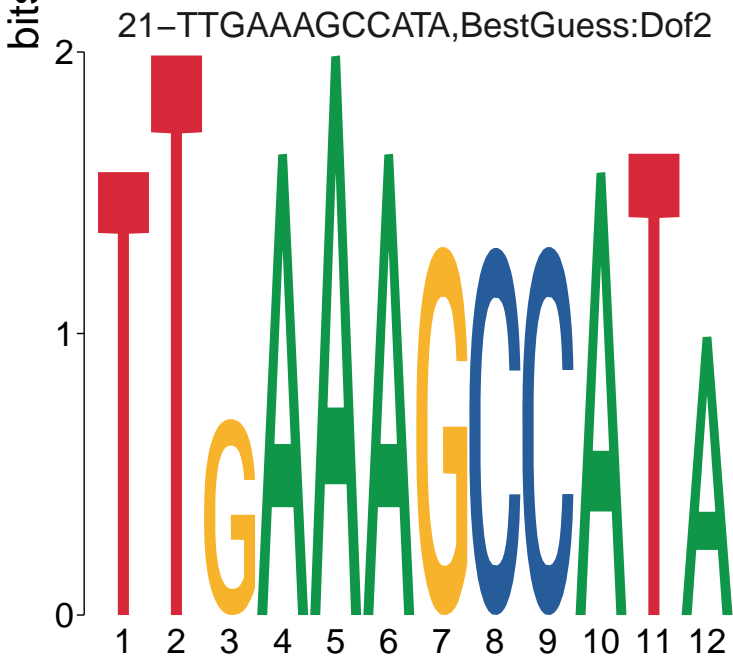
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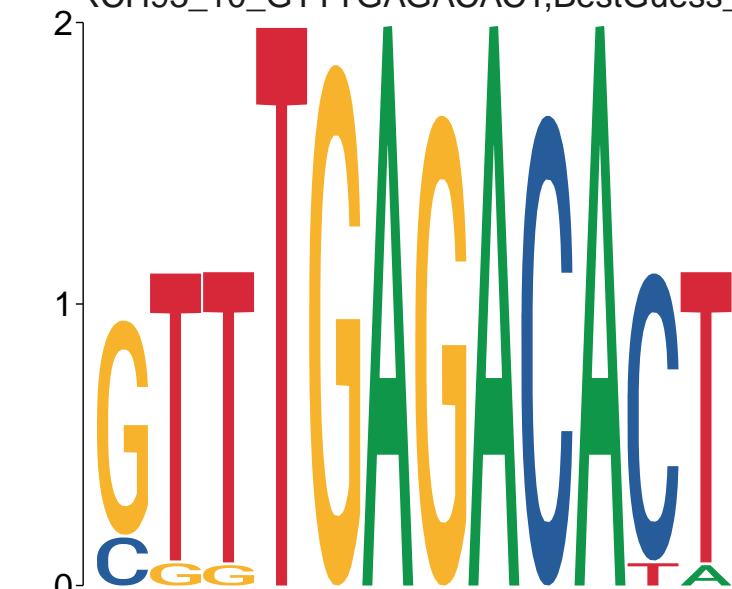
RCH92_21_TTGAAAGCCATA,BestGuess_L



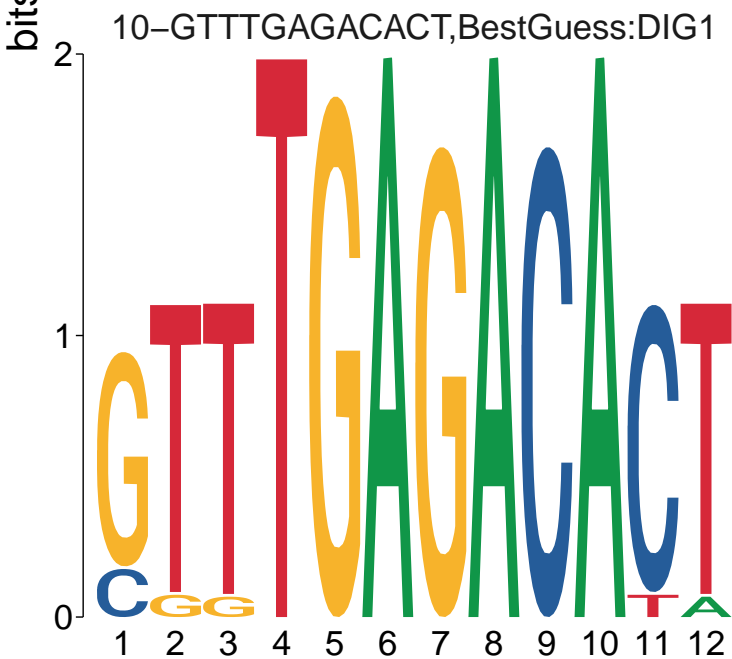
21-TTGAAAGCCATA,BestGuess:Dof2



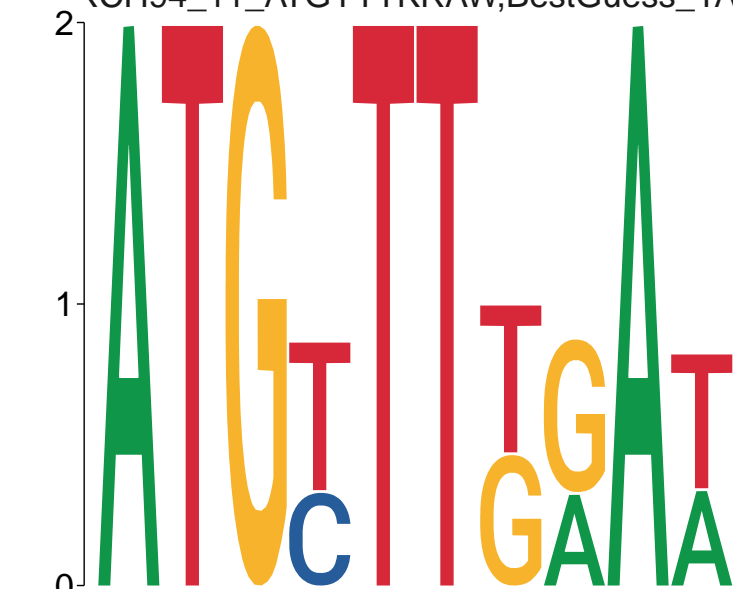
RCH93_10_GTTTGAGACACT, BestGuess_1



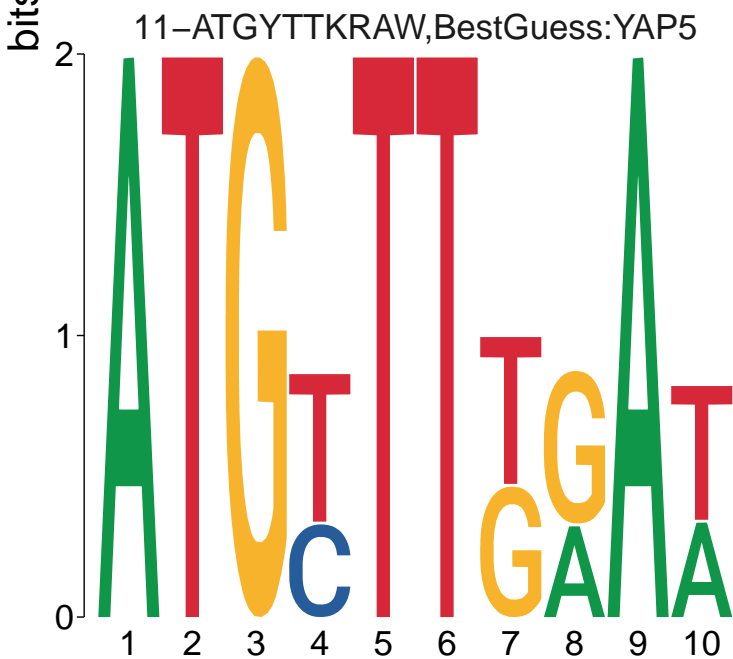
10-GTTTGAGACACT, BestGuess: DIG1



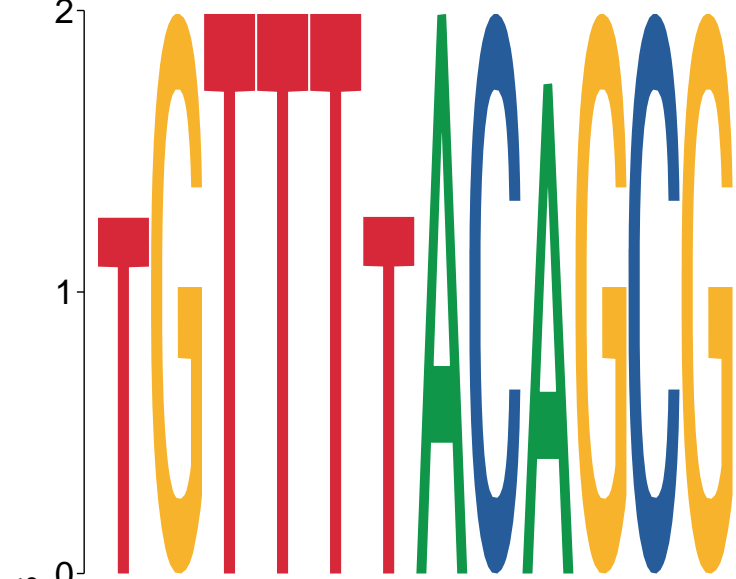
RCH94_11_ATGYTTKRAW,BestGuess_YAF



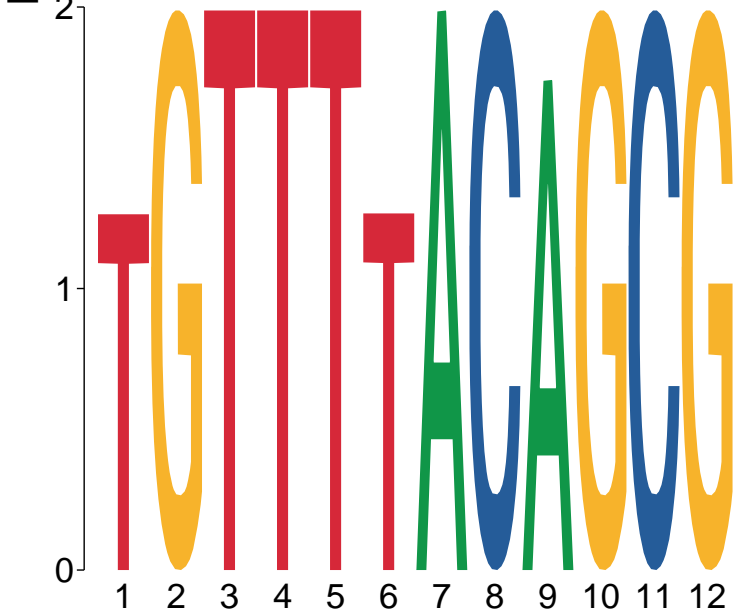
11-ATGYTTKRAW,BestGuess:YAP5



RCH95_4_TGTTTTACAGCG,BestGuess_RI



4-TGTTTTACAGCG,BestGuess:Rhox11



RCH96_17_ATATCTTACA,BestGuess_CC/

2

1

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bits

17-ATATCTTACA,BestGuess:CCA

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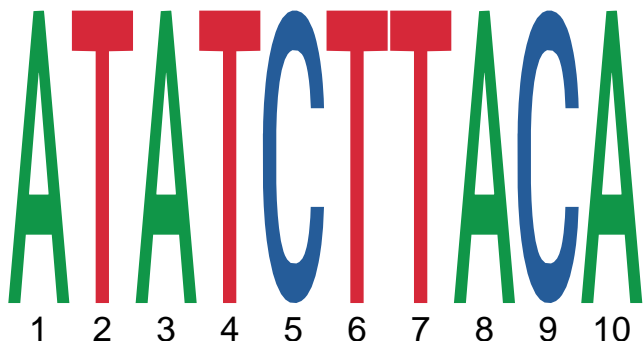
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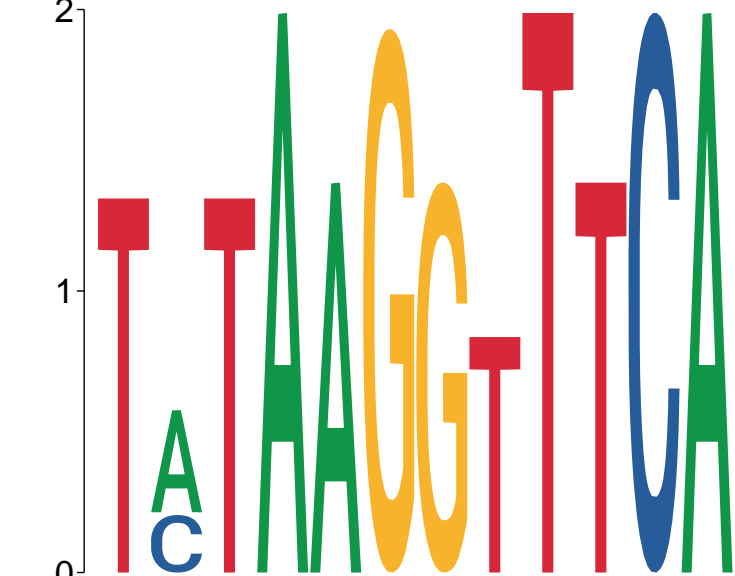
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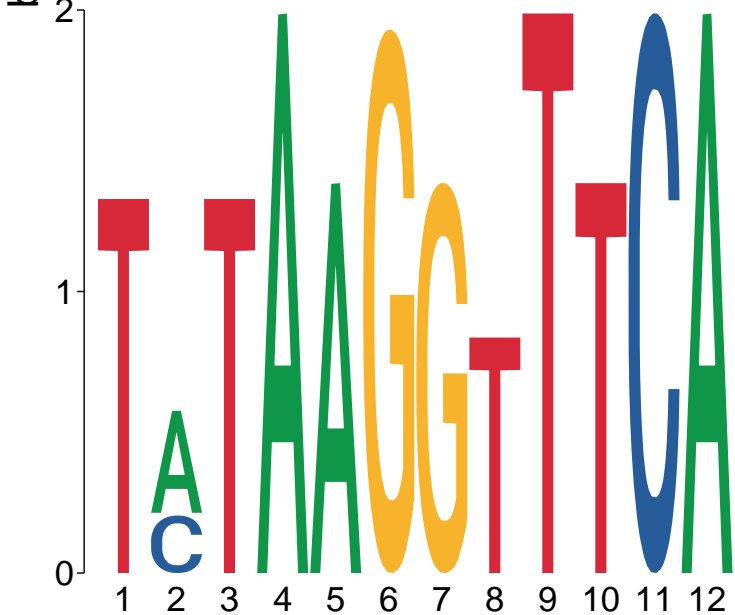
10



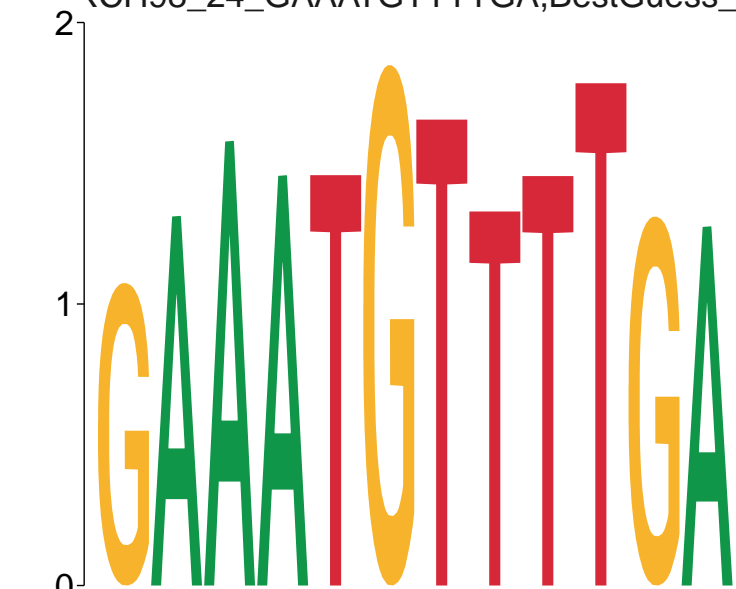
RCH97_5_TMTAAGGTTTCA,BestGuess_S



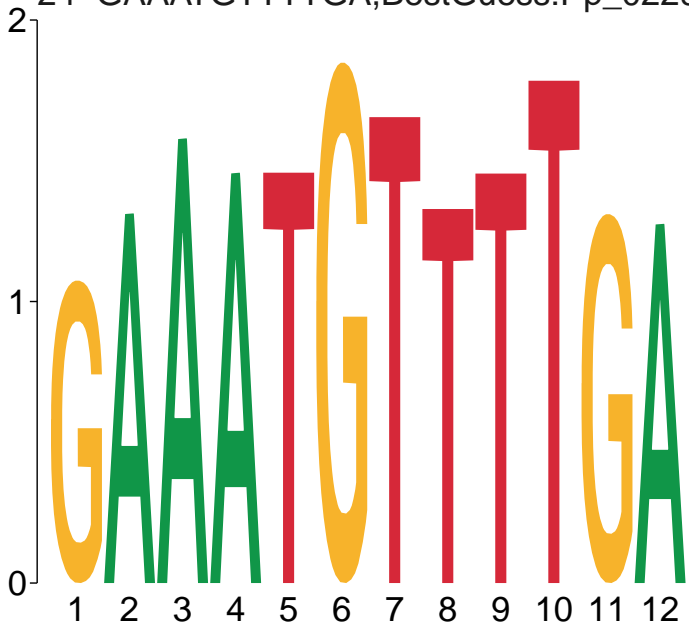
5-TMTAAGGTTTCA,BestGuess:SIX2



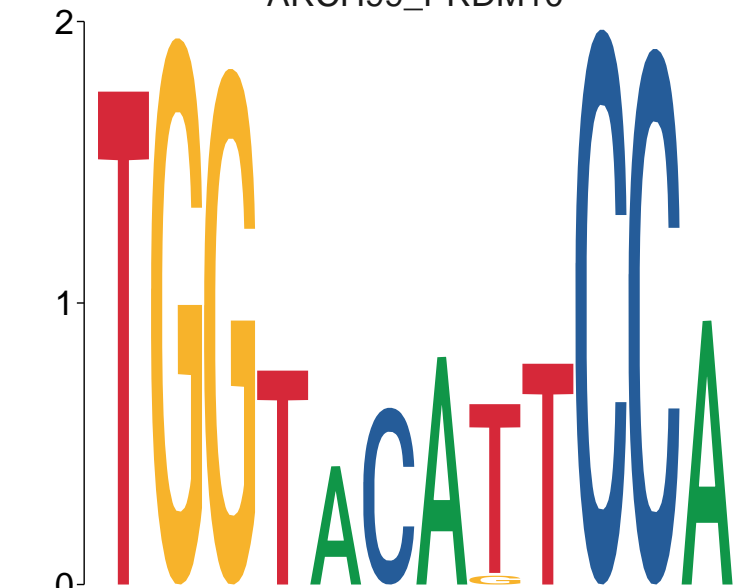
RCH98_24_GAAATGTTTTGA,BestGuess_f



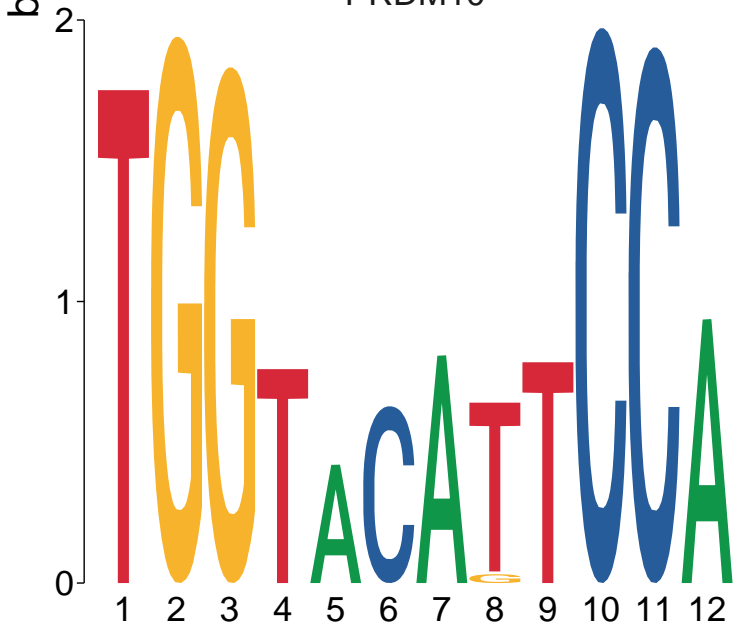
24-GAAATGTTTTGA,BestGuess:Pp_0229



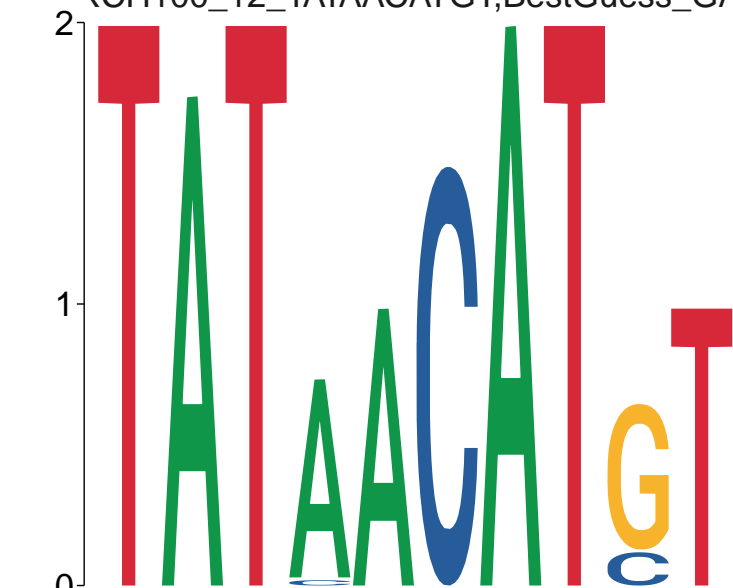
ARCH99_PRDM10



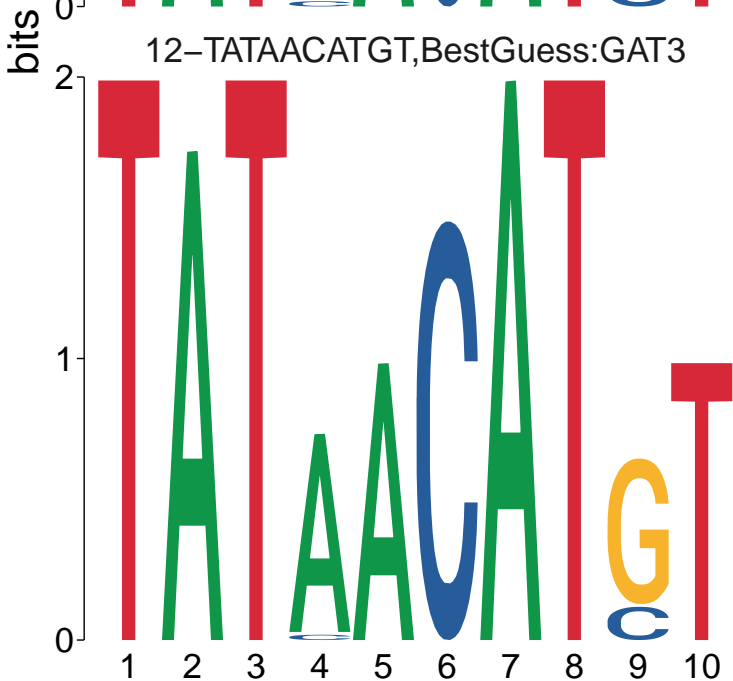
PRDM10



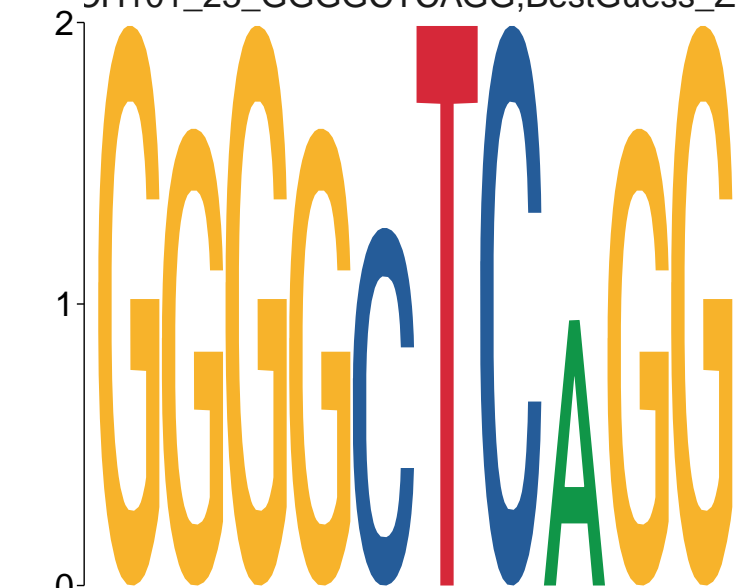
RCH100_12_TATAACATGT,BestGuess_GAT



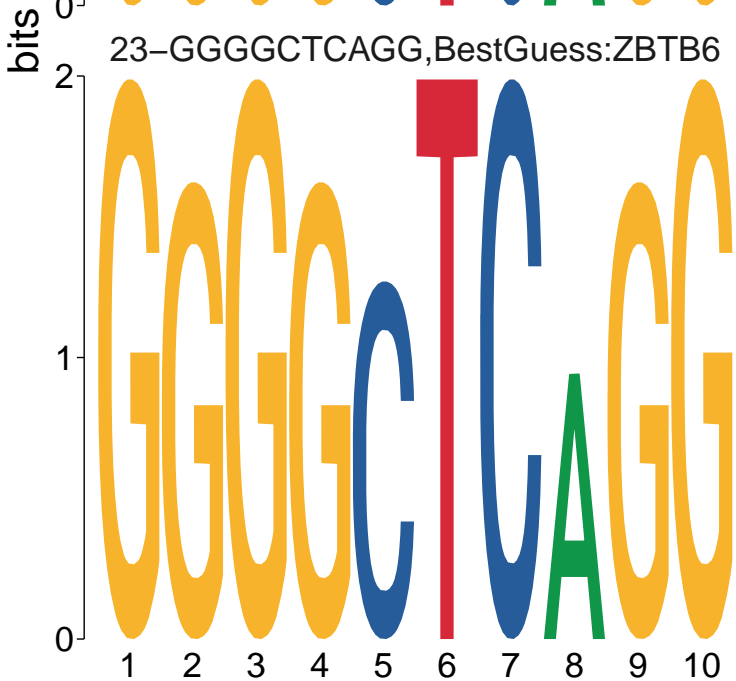
12-TATAACATGT,BestGuess:GAT3



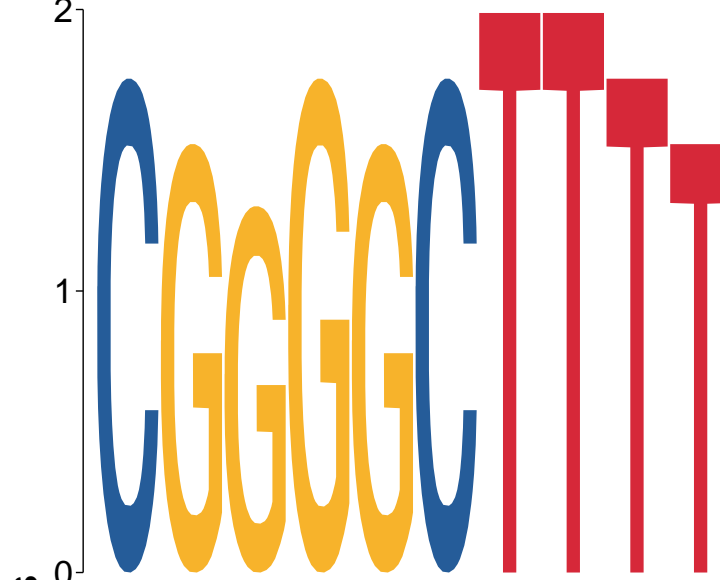
CH101_23_GGGGCTCAGG,BestGuess_ZE



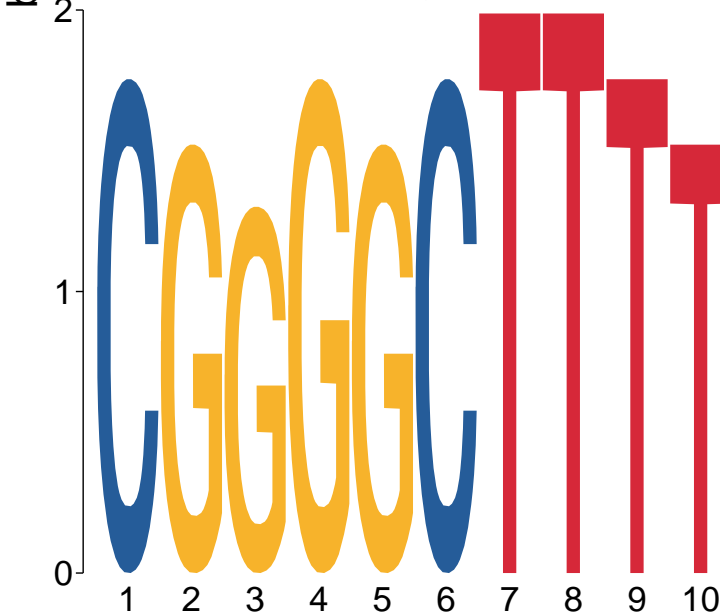
23-GGGGCTCAGG,BestGuess:ZBTB6



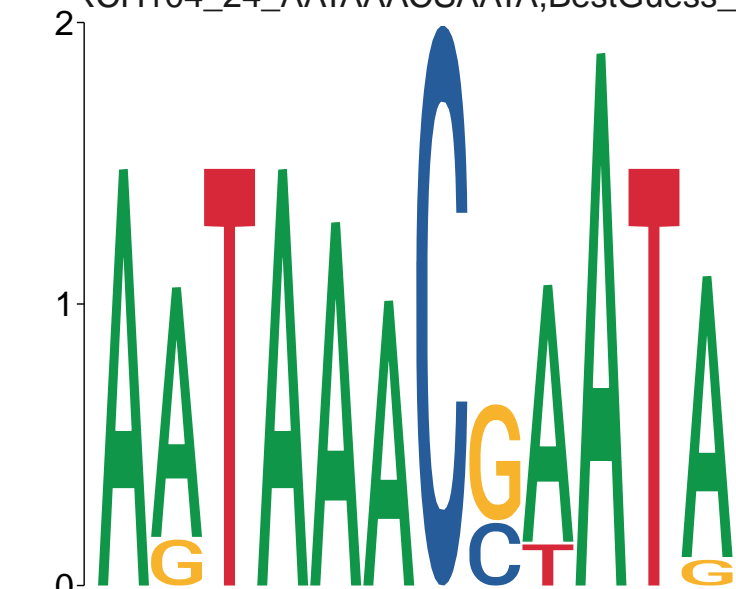
RCH102_3_CGGGGGCTTTT,BestGuess_Do



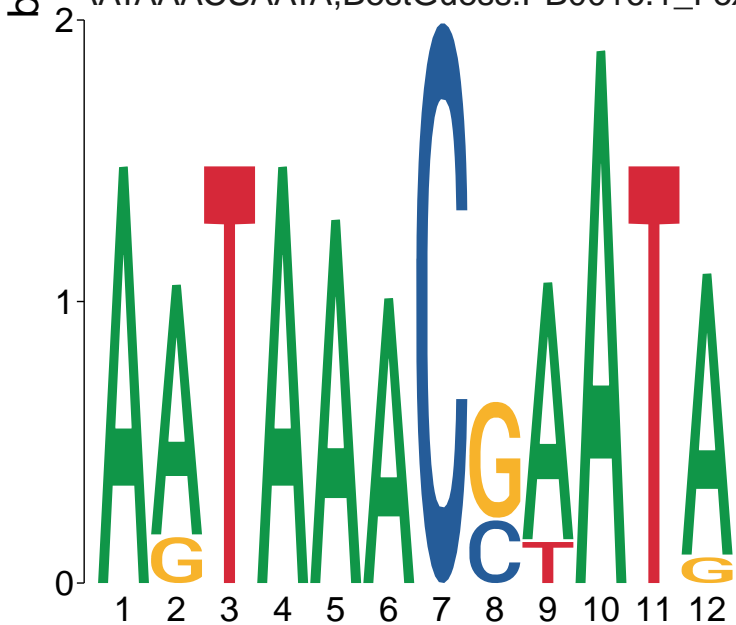
3-CGGGGGCTTTT,BestGuess:Dof2



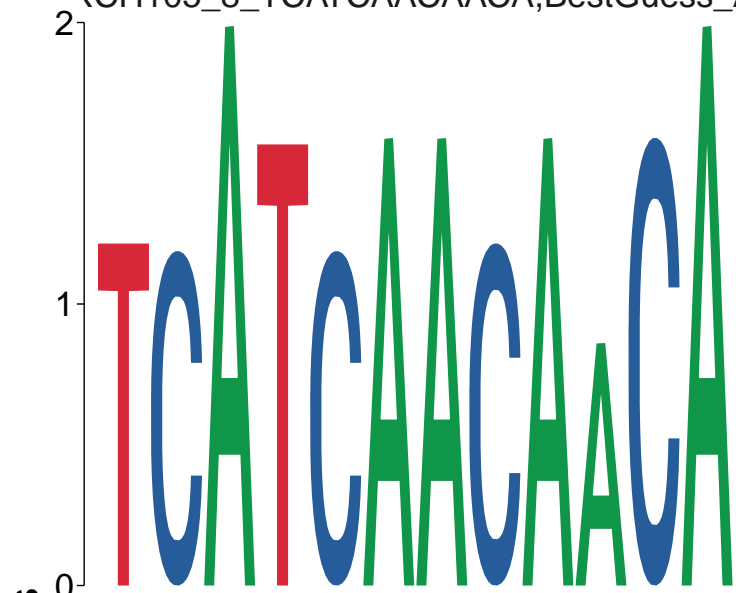
RCH104_24_AATAAACSAATA,BestGuess_F



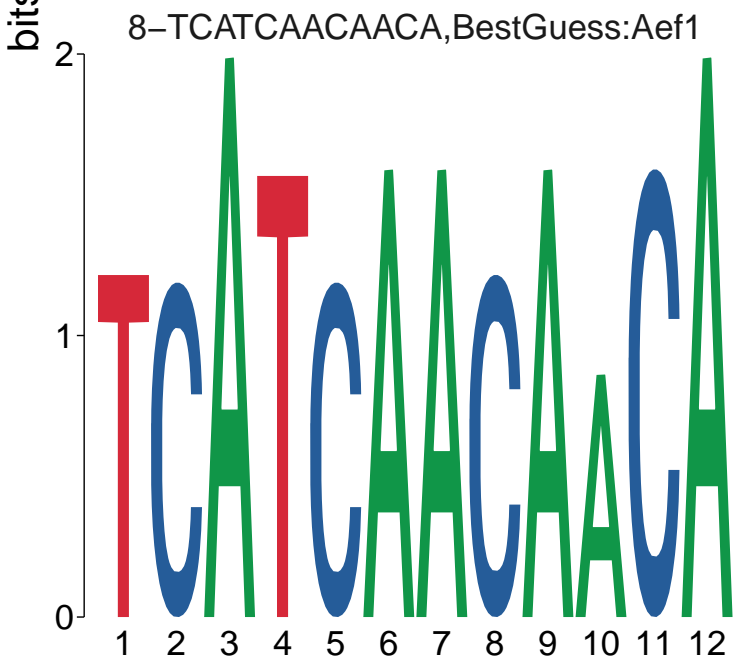
AATAAACSAATA,BestGuess:PB0016.1_Fox



2CH105_8_TCATCAACAACA, BestGuess_A



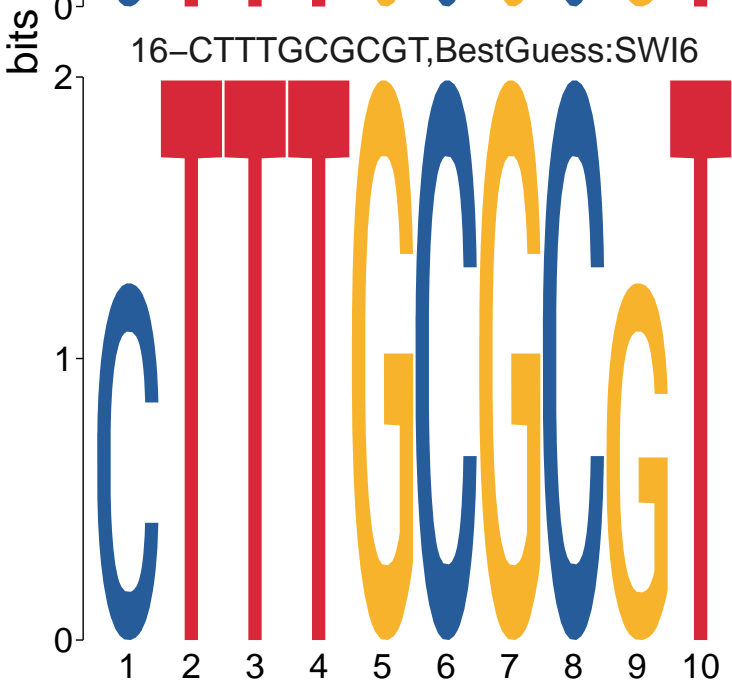
8-TCATCAACAACA, BestGuess:Aef1



CH108_16_CTTTGCGCGT,BestGuess_SV



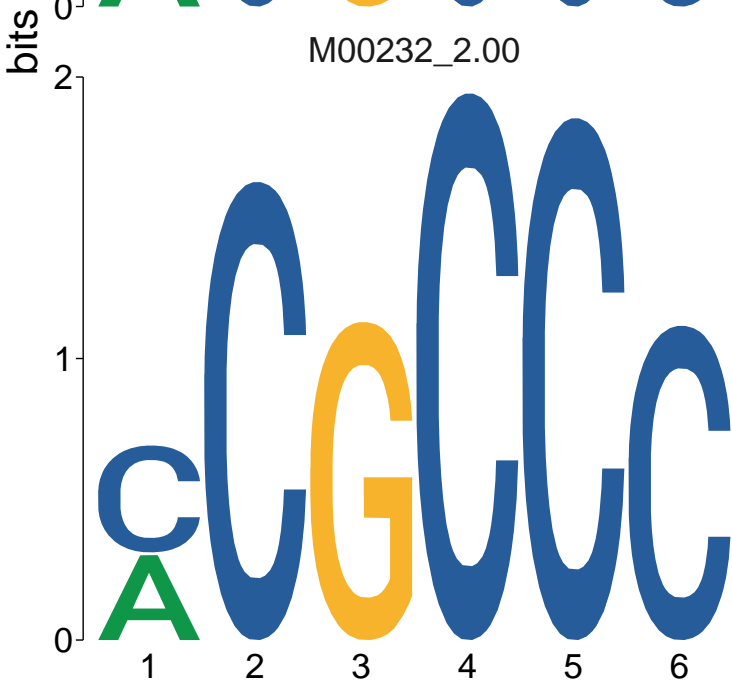
16-CTTTGCGCGT,BestGuess:SW16



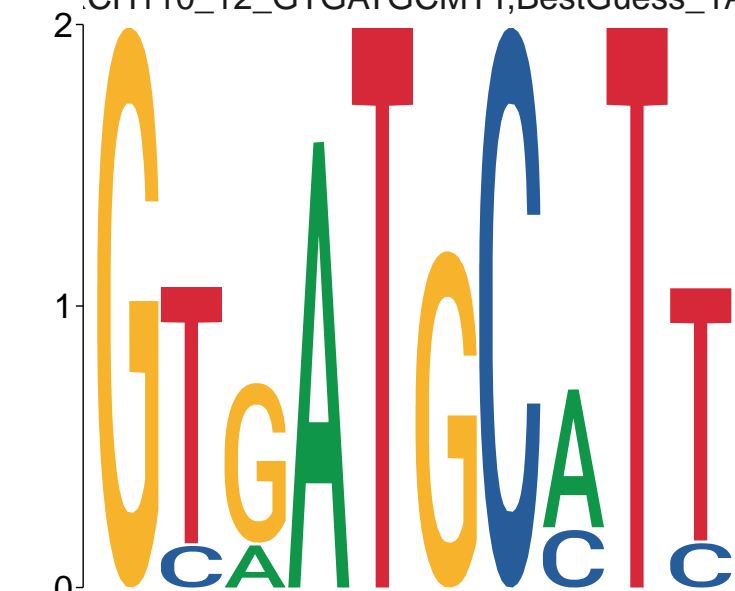
ARCH109_islike_EGR1_EGR2



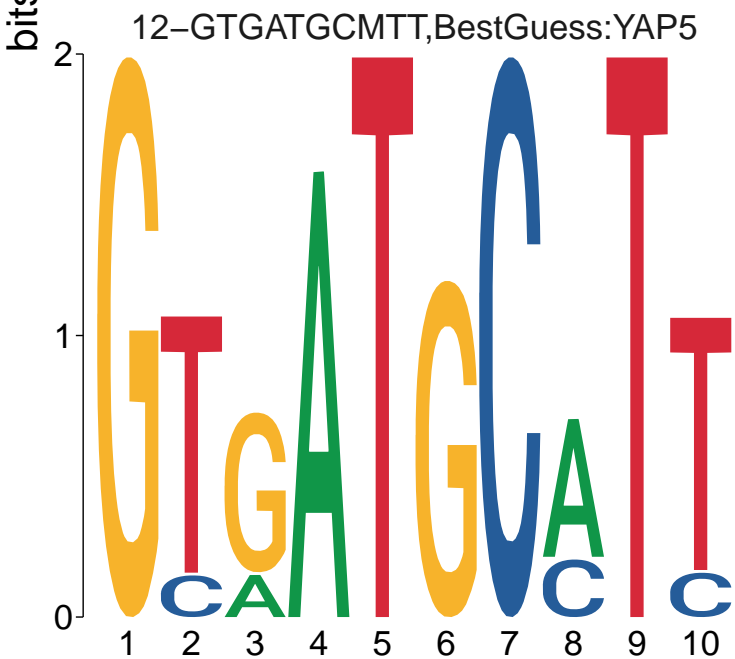
M00232_2.00



CH110_12_GTGATGCMTT,BestGuess_YA



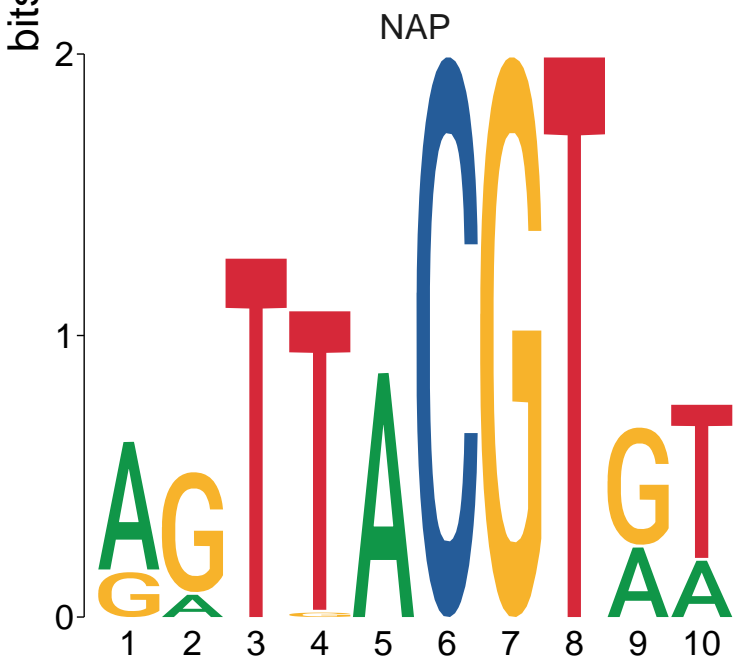
12-GTGATGCMTT,BestGuess:YAP5



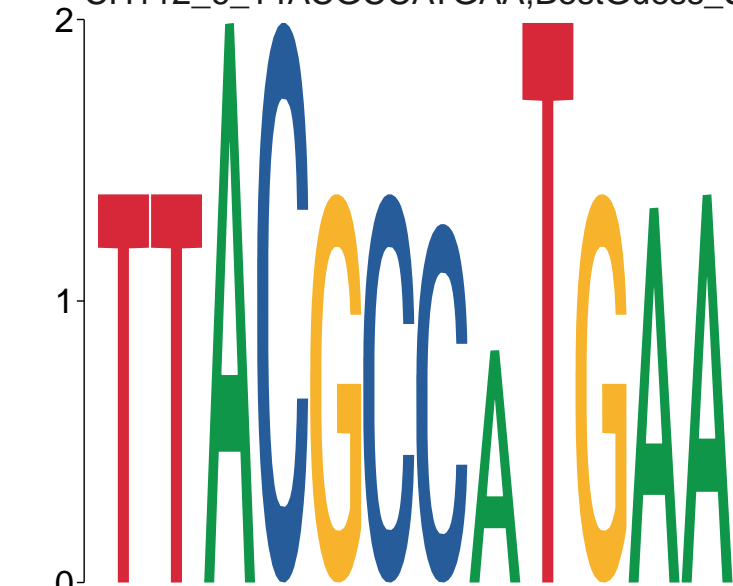
ARCH111_NAP



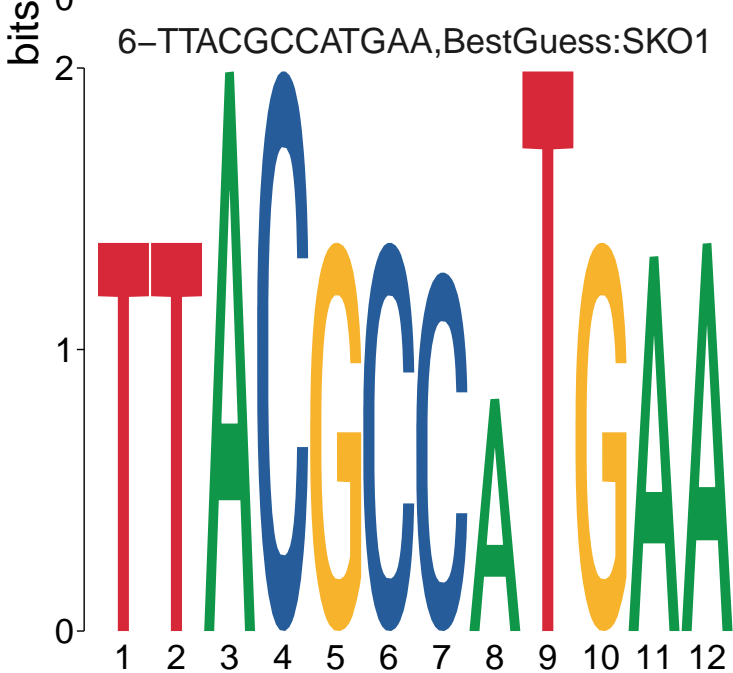
NAP



CH112_6_TTACGCCATGAA,BestGuess_S



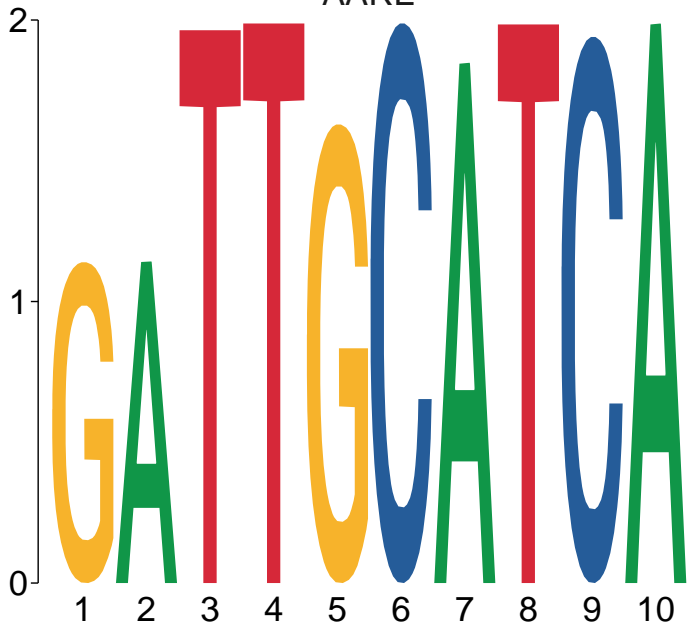
6-TTACGCCATGAA,BestGuess:SKO1



ARCH113_AARE



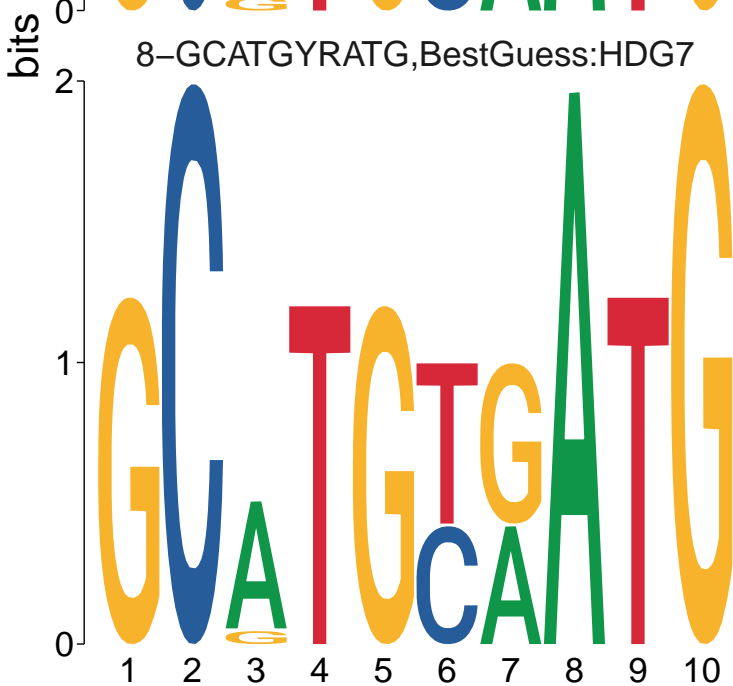
AARE



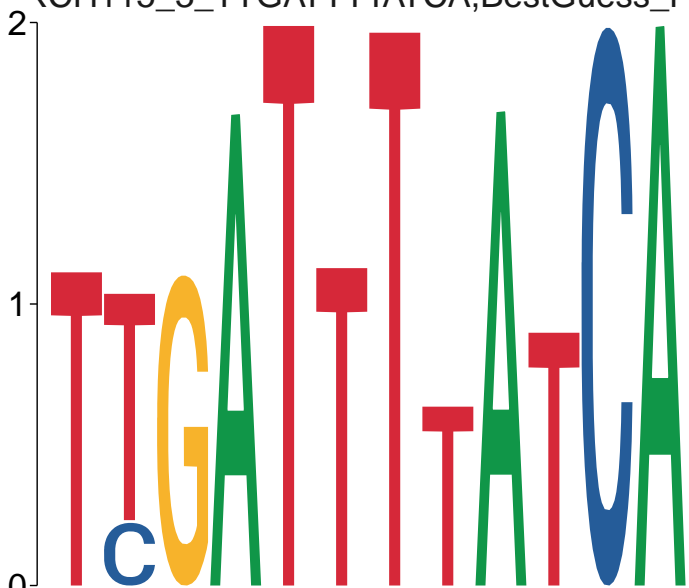
CH114_8_GCATGYRATG,BestGuess_HD(



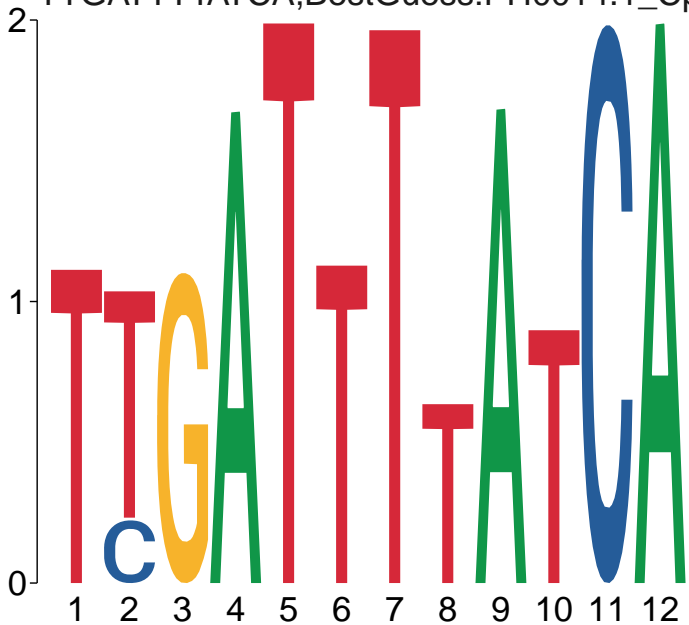
8-GCATGYRATG,BestGuess:HDG7



2RCH115_3_TTGATTTTATCA,BestGuess_Pf



TTGATTTTATCA,BestGuess:PH0014.1_Cp



CH116_25_ATCATCAAACCA,BestGuess_

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25-ATCATCAAACCA,BestGuess:pan

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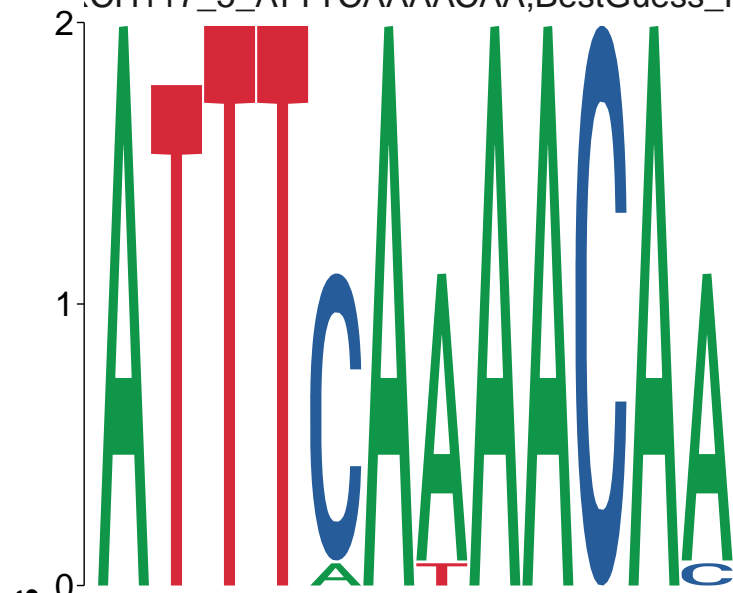
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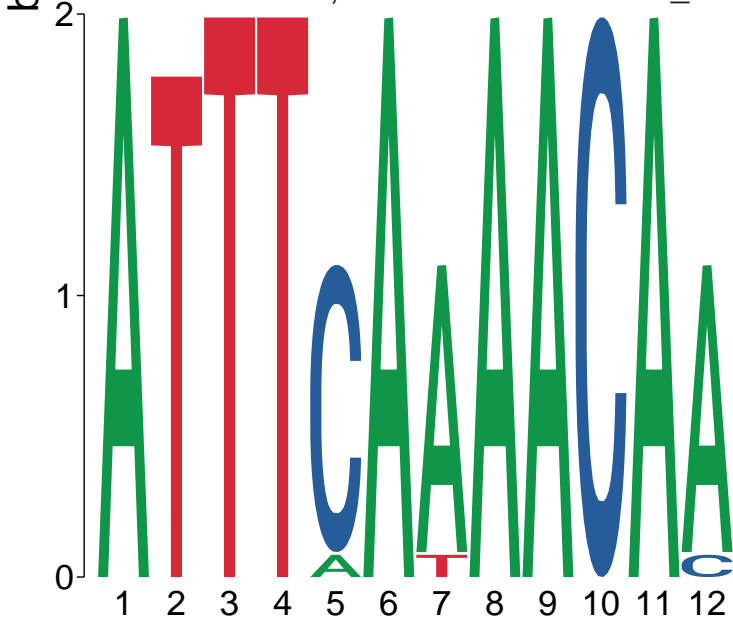
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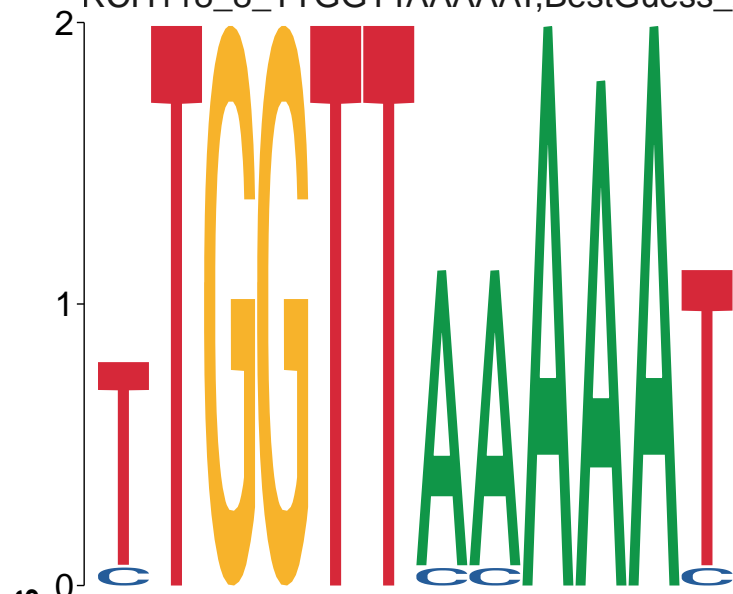
CH117_5_ATTTCAAACAA, BestGuess_P



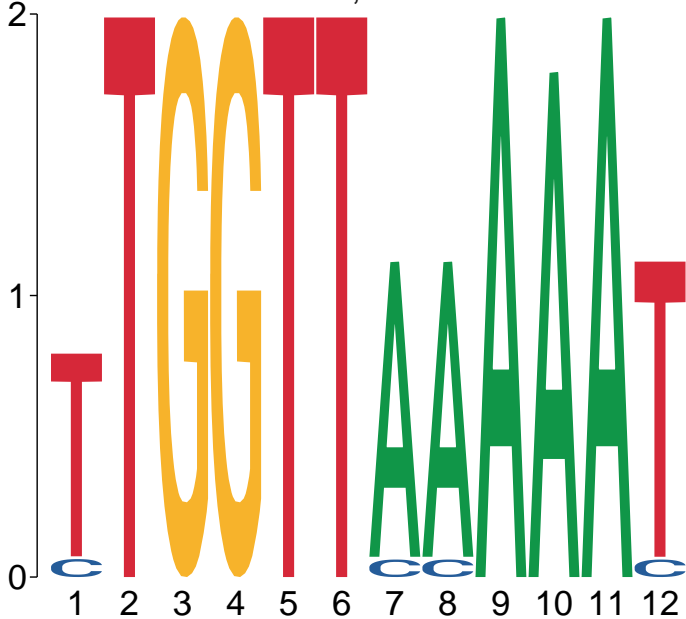
TTTCAAACAA, BestGuess:PB0123.1_Fox



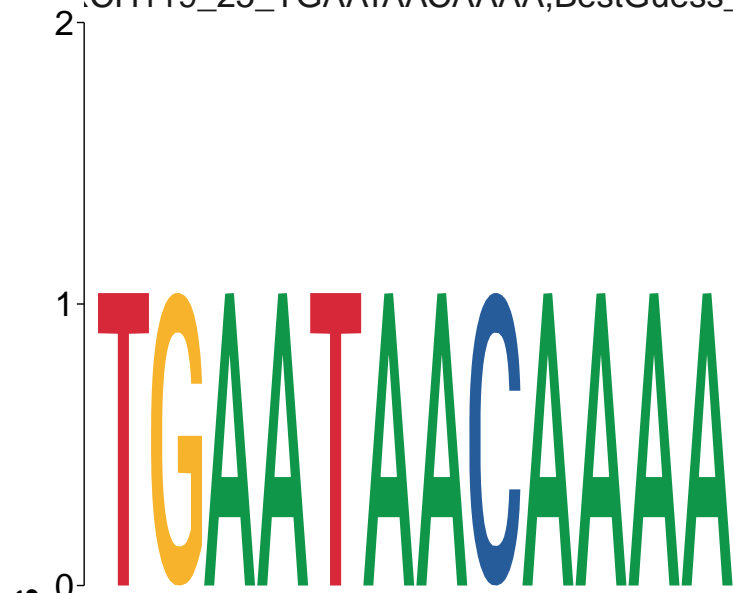
RCH118_8_TTGGTTAAAAAT, BestGuess_A



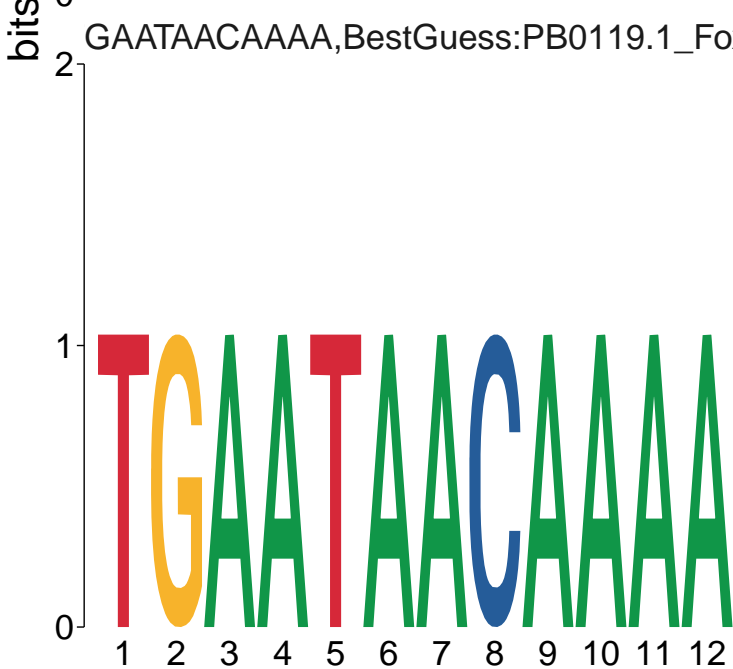
8-TTGGTTAAAAAT, BestGuess:AT-GTL1



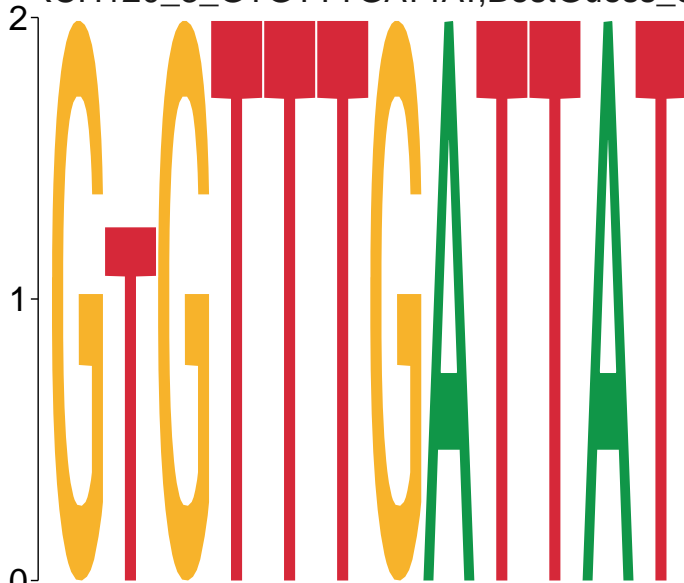
CH119_25_TGAATAACAAAA, BestGuess_



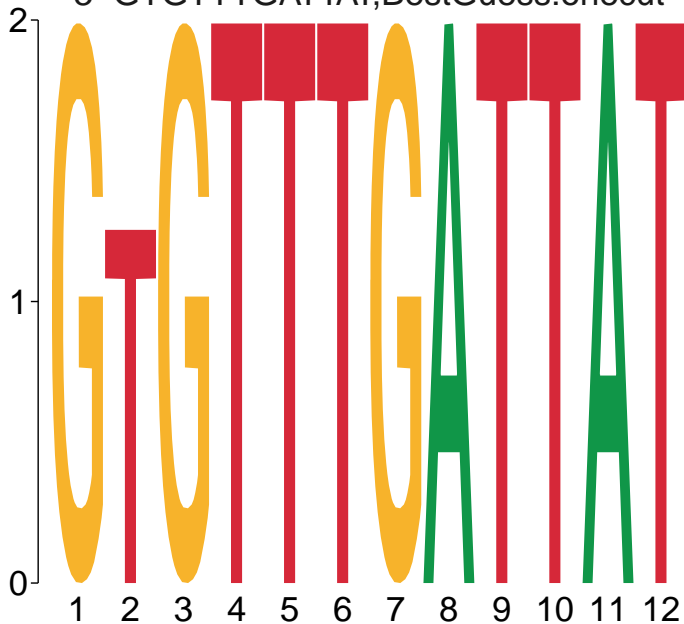
GAATAACAAAA, BestGuess:PB0119.1_Fox



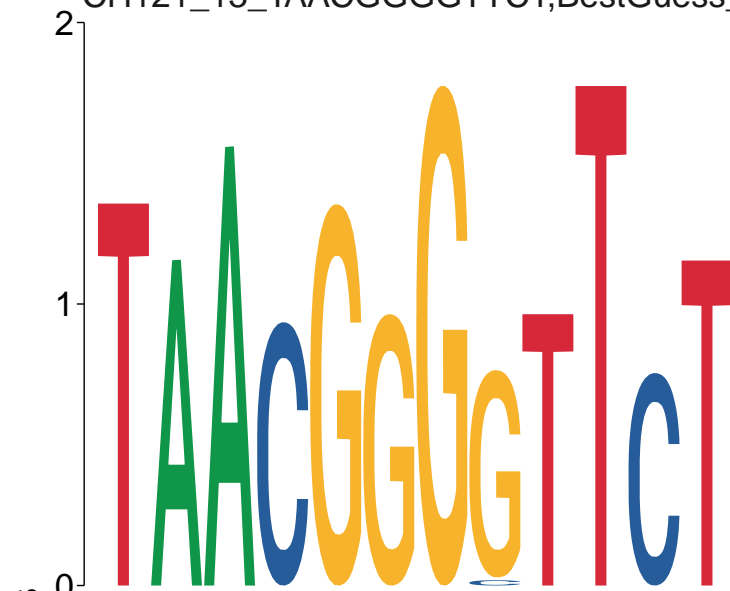
RCH120_3_GTGTTTGATTAT,BestGuess_oi



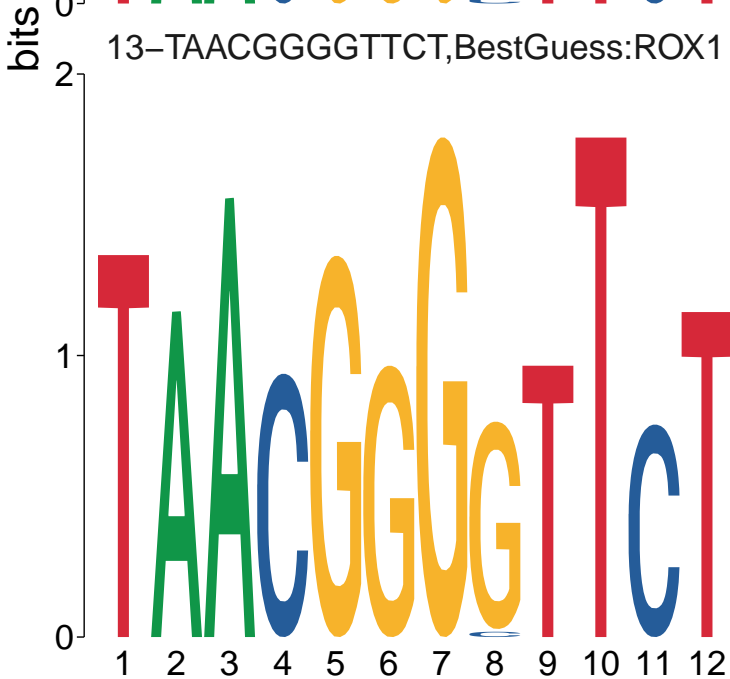
3-GTGTTTGATTAT,BestGuess:onecut



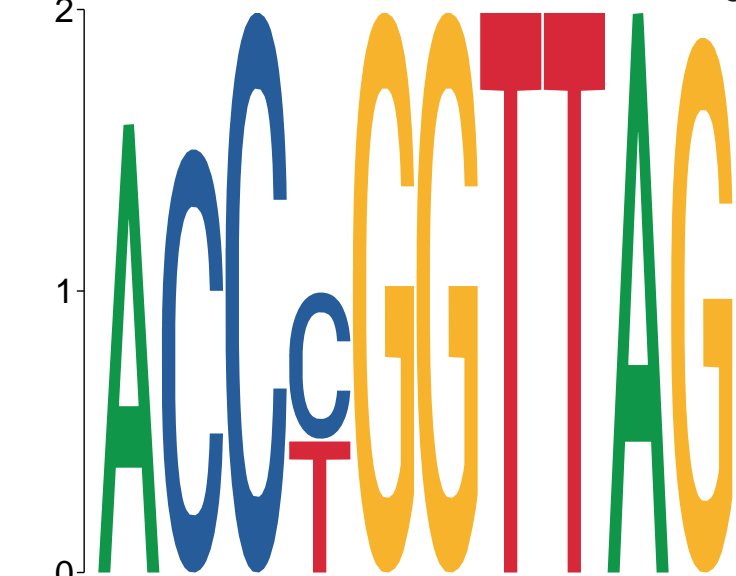
CH121_13_TAACGGGGTTCT,BestGuess_



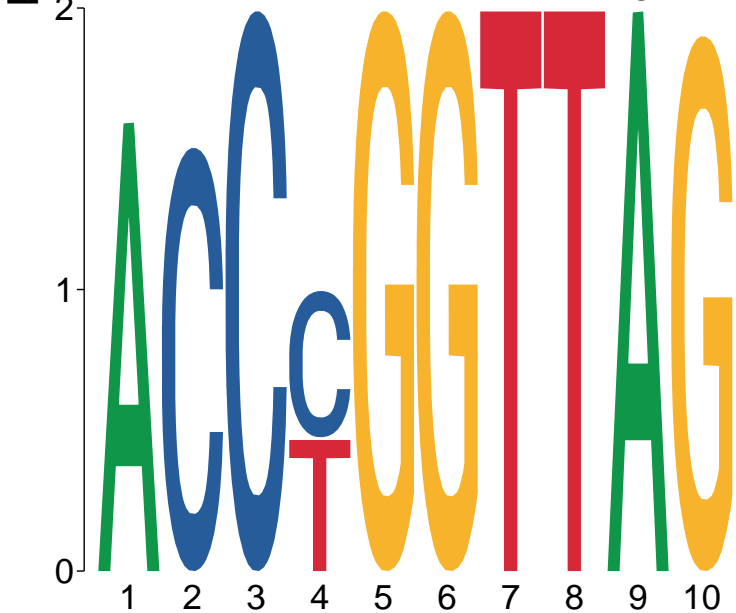
13-TAACGGGGTTCT,BestGuess:ROX1



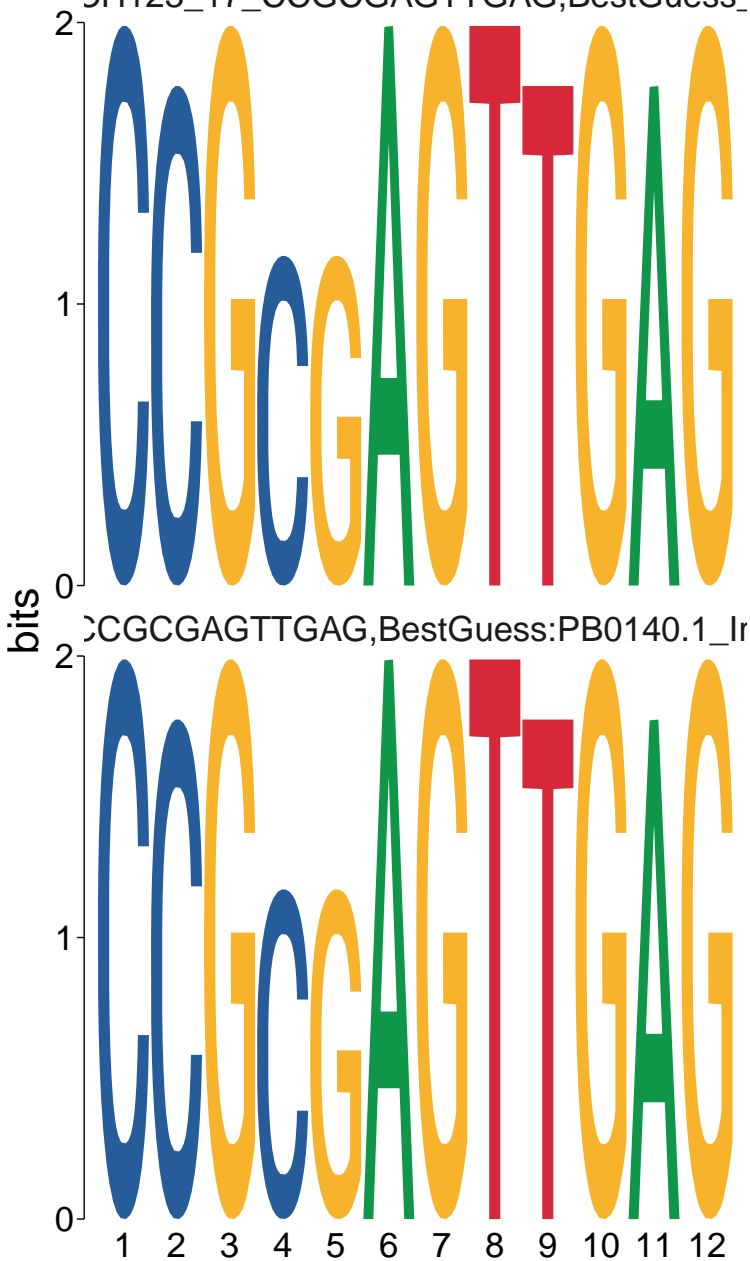
RCH122_12_ACCYGGTTAG,BestGuess_gi



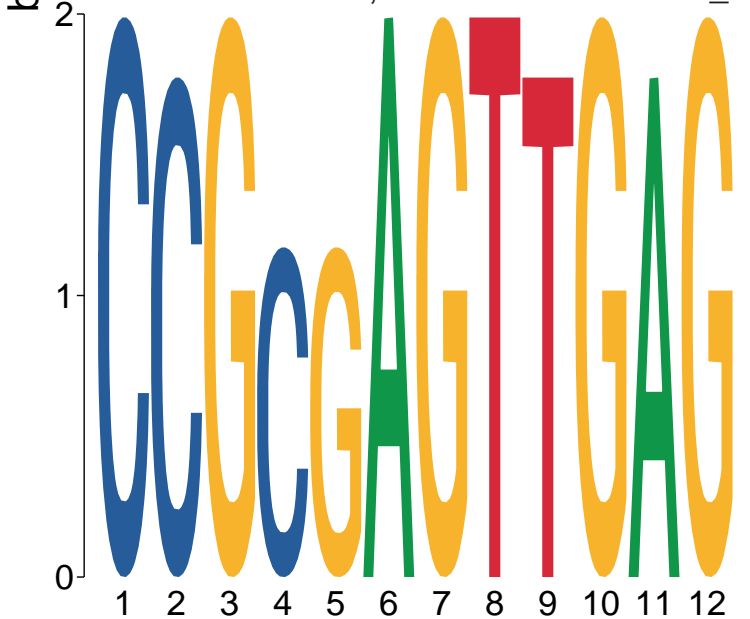
12-ACCYGGTTAG,BestGuess:grh



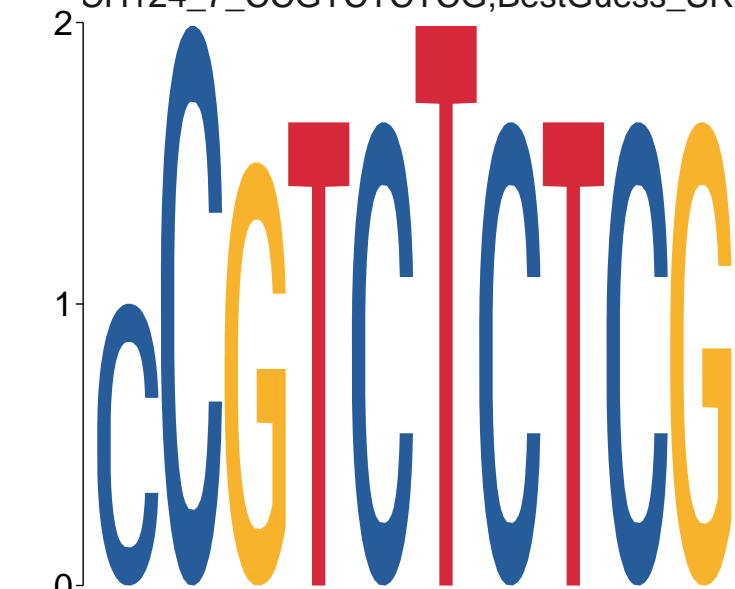
CH123_17_CCGCGAGTTGAG,BestGuess_



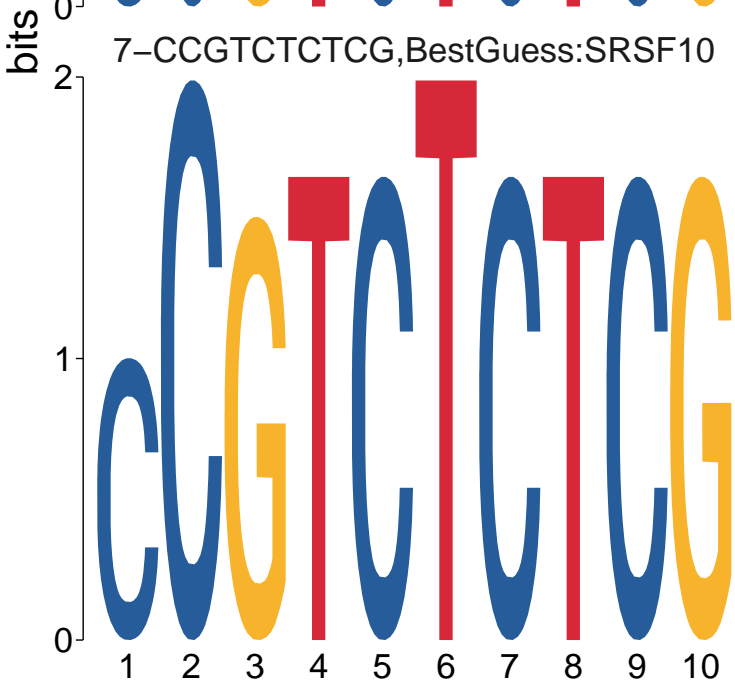
CCGCGAGTTGAG,BestGuess:PB0140.1_Ir



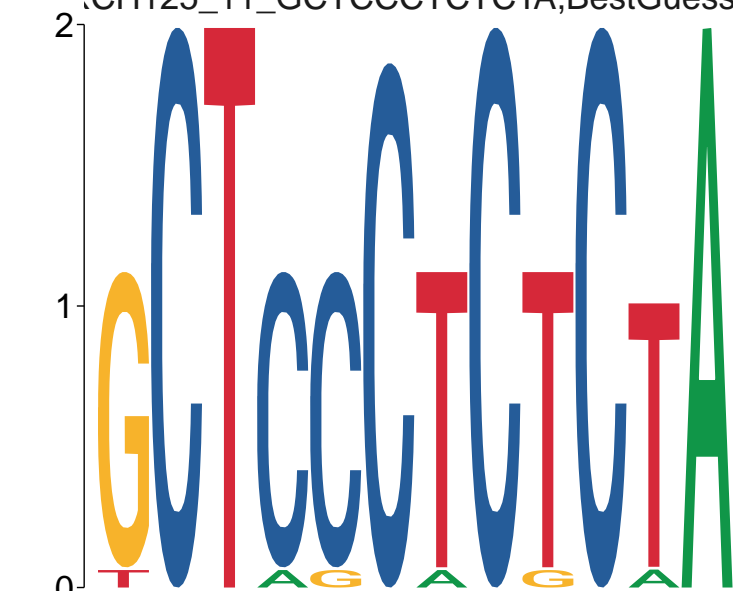
CH124_7_CCGTCTCTCTCG,BestGuess_SRS



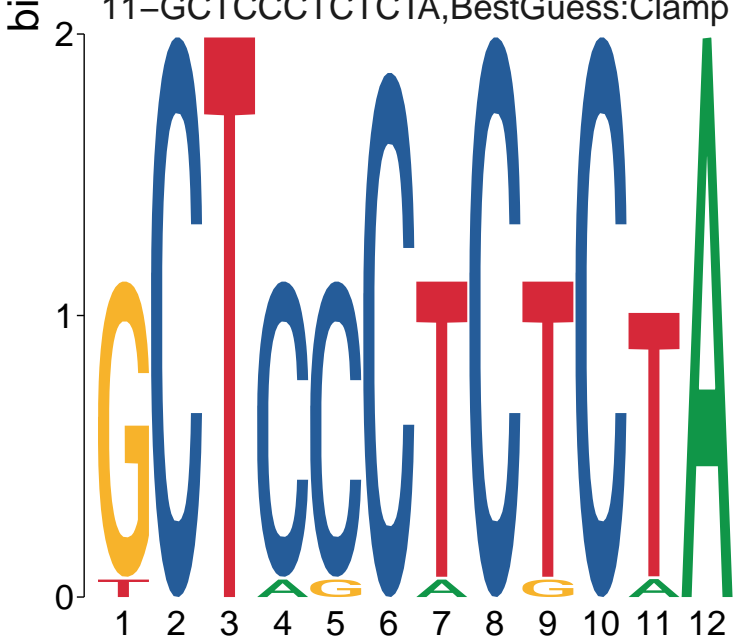
7-CCGTCTCTCTCG,BestGuess:SRSF10



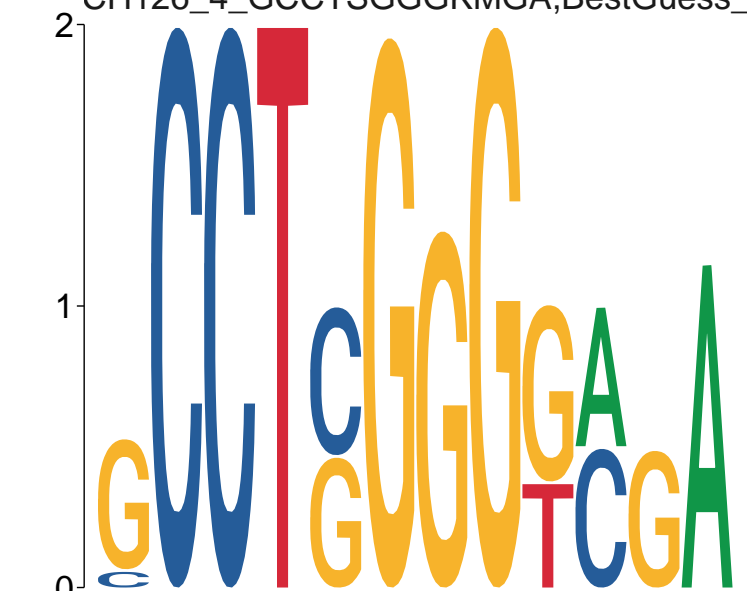
CH125_11_GCTCCCTCTCTA,BestGuess_



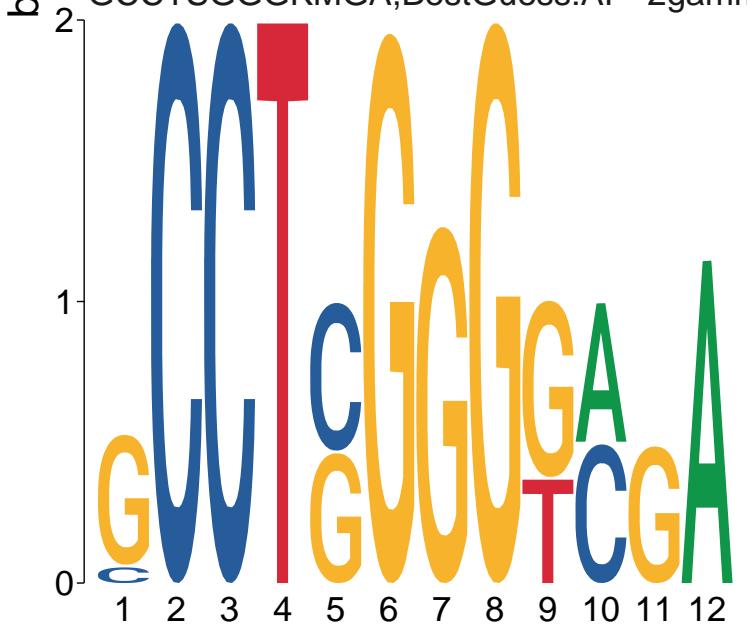
11-GCTCCCTCTCTA,BestGuess:Clamp



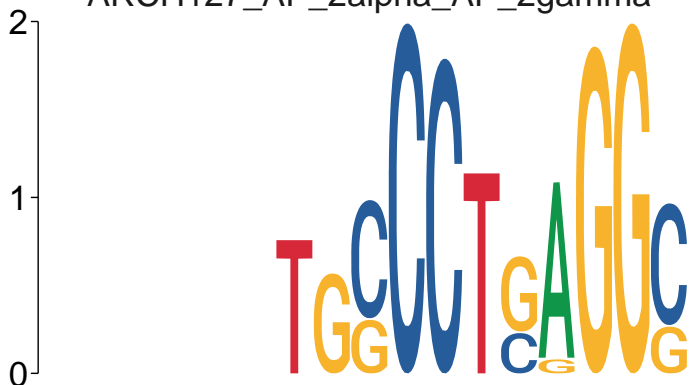
CH126_4_GCCTSGGGKMGGA,BestGuess_



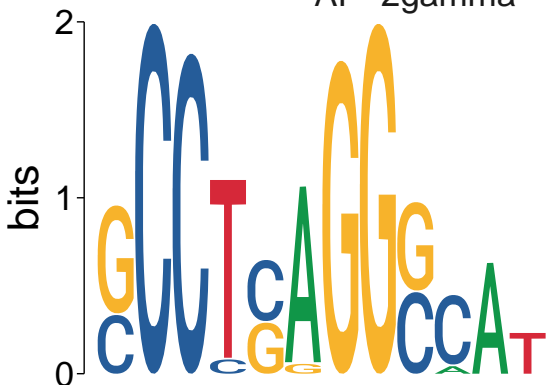
GCCTSGGGKMGGA,BestGuess:AP-2gamr



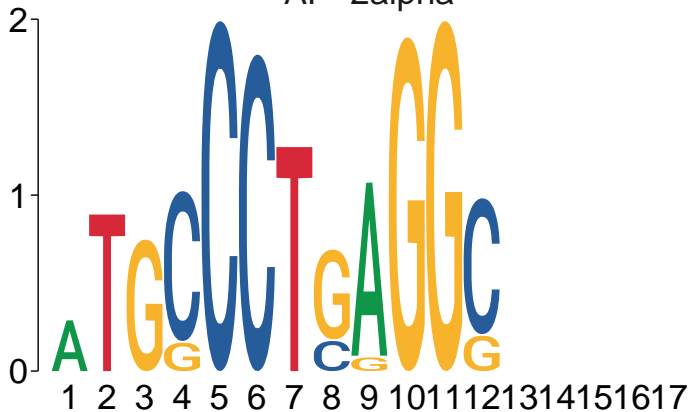
ARCH127_AP_2alpha_AP_2gamma



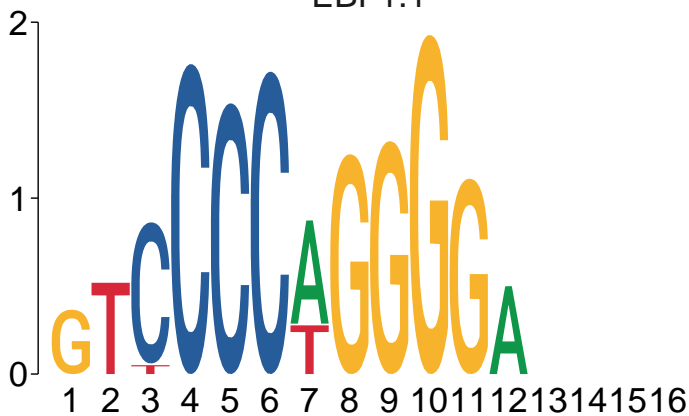
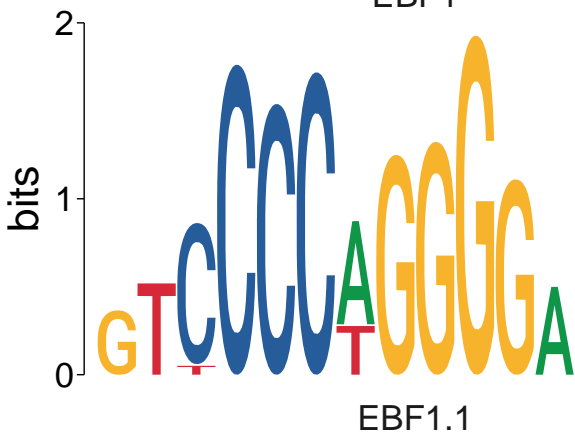
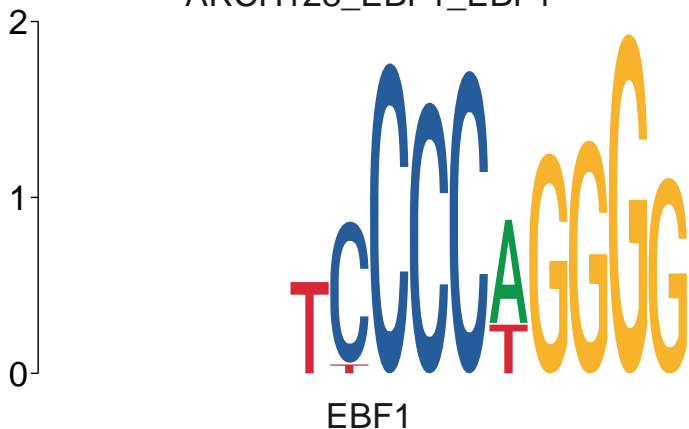
AP-2gamma



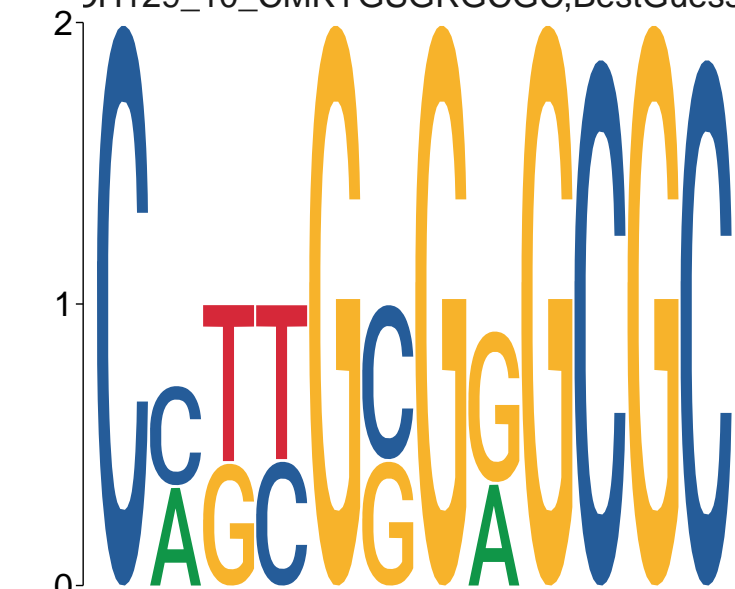
AP-2alpha



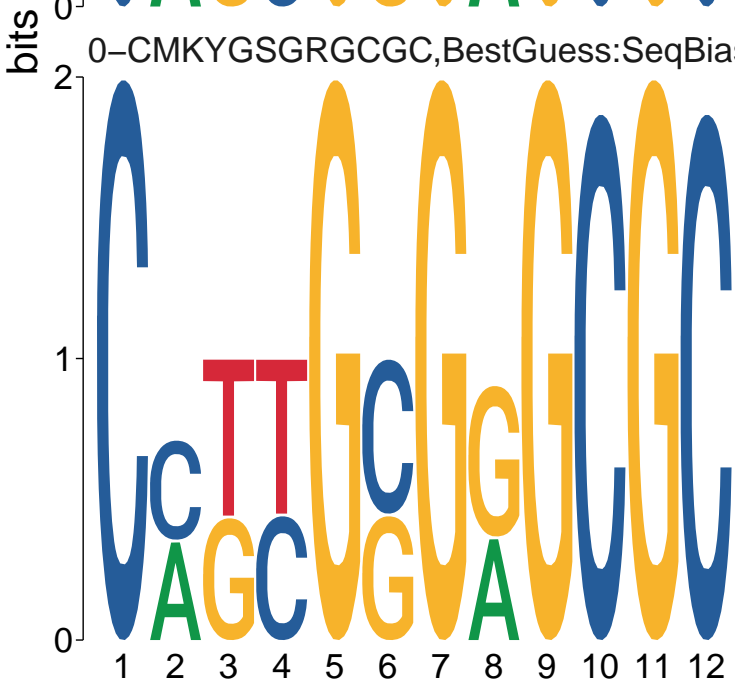
ARCH128_EBF1_EBF1



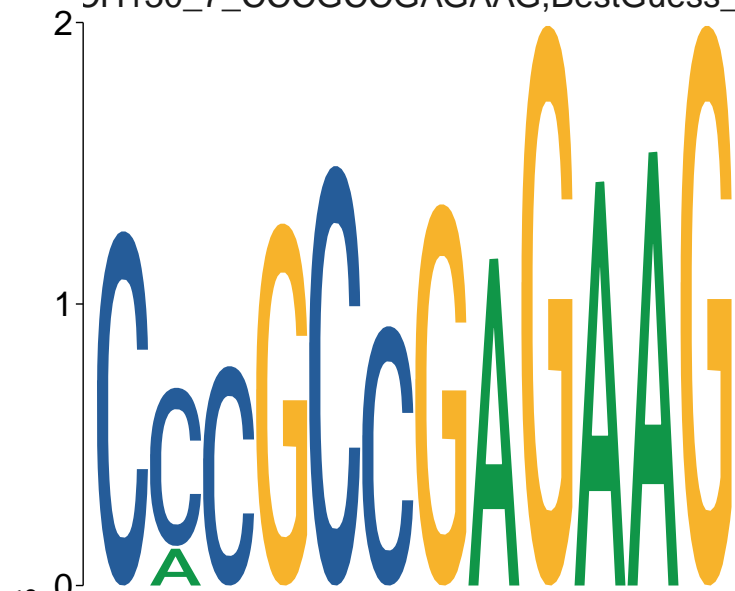
CH129_10_CMKYGSGRGCGC,BestGuess.



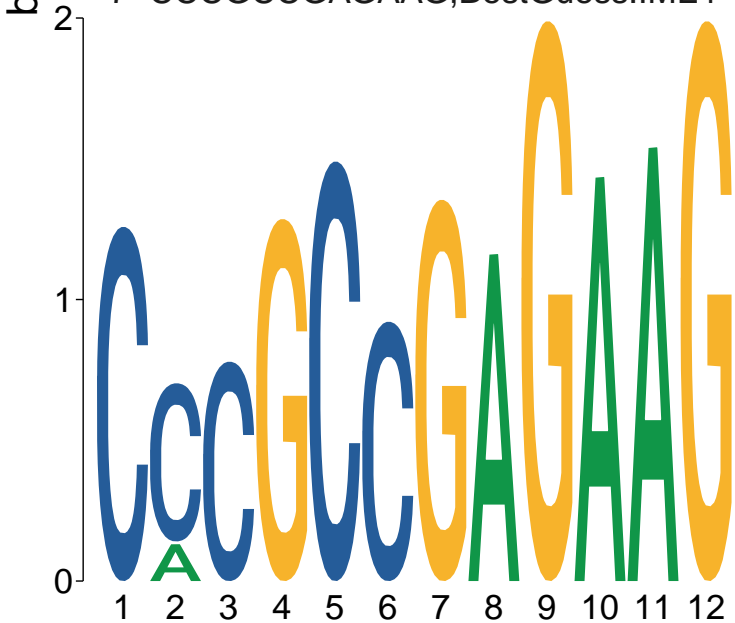
0-CMKYGSGRGCGC,BestGuess:SeqBias



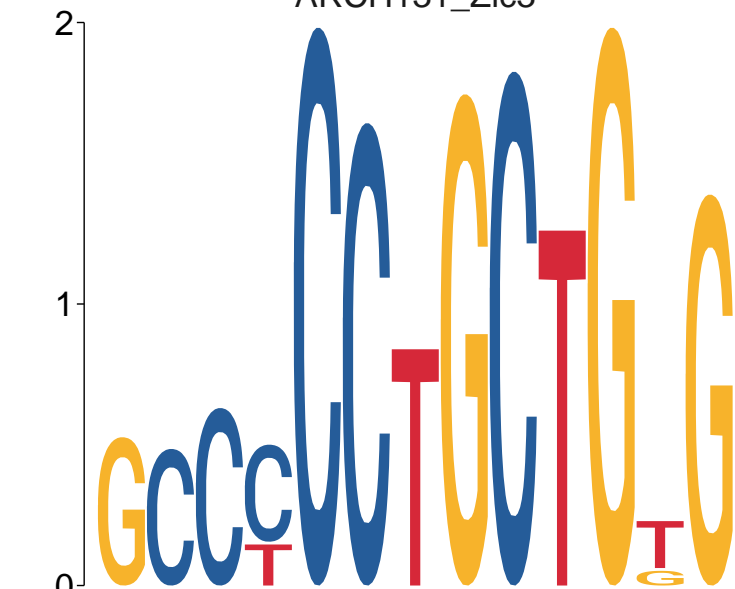
CH130_7_CCCGCCGAGAAG,BestGuess_I



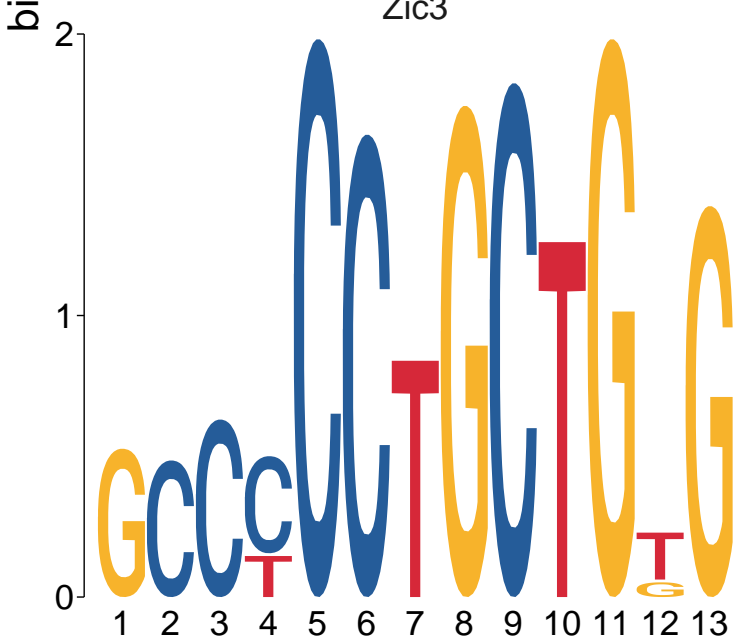
7-CCCGCCGAGAAG,BestGuess:IME1



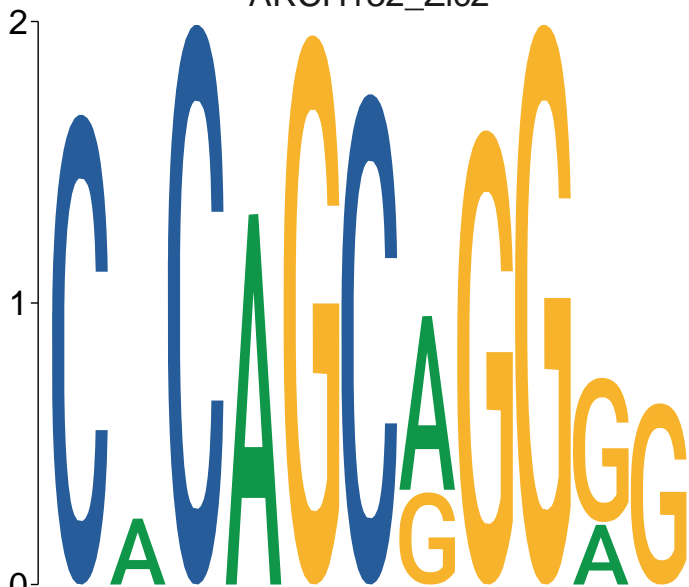
ARCH131_Zic3



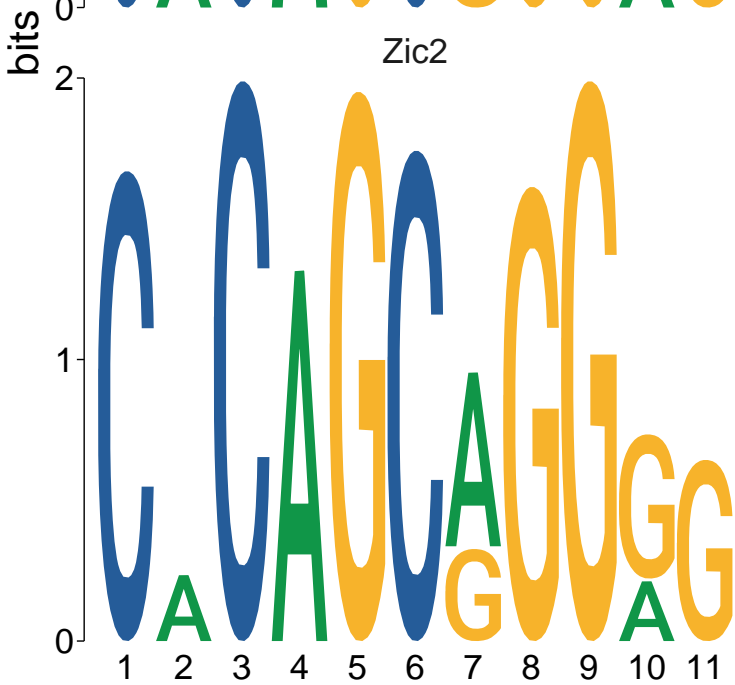
Zic3



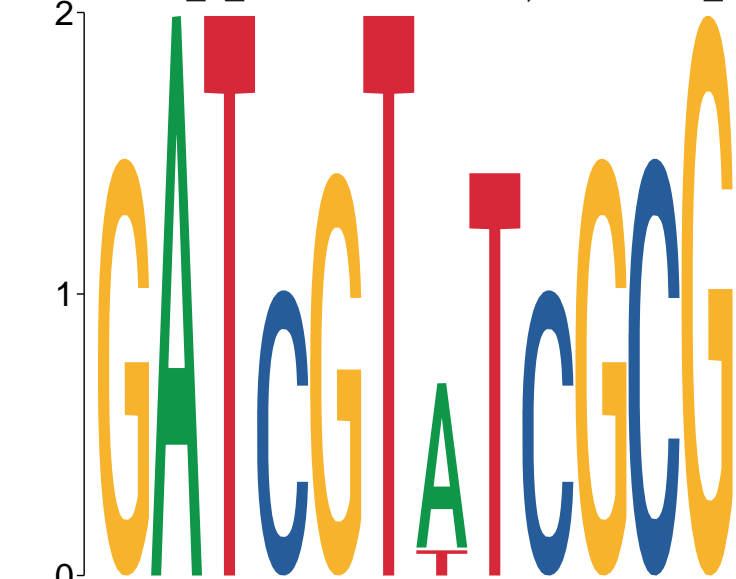
ARCH132_Zic2



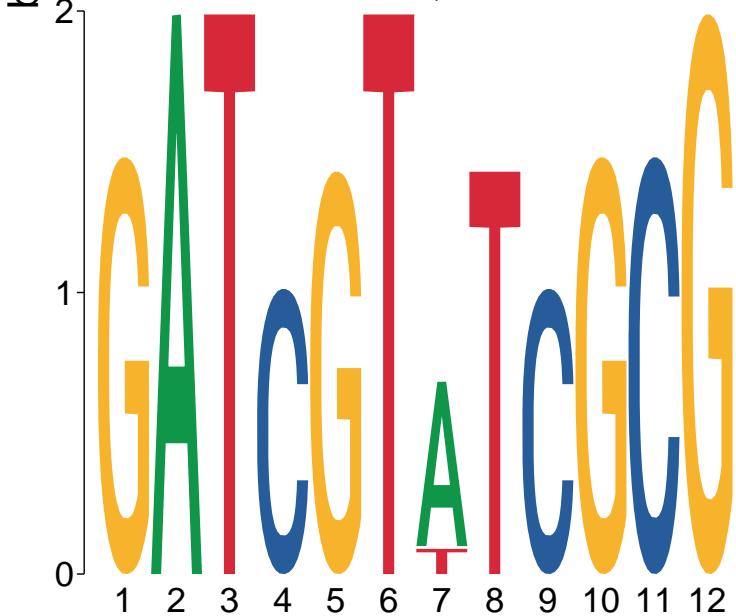
Zic2



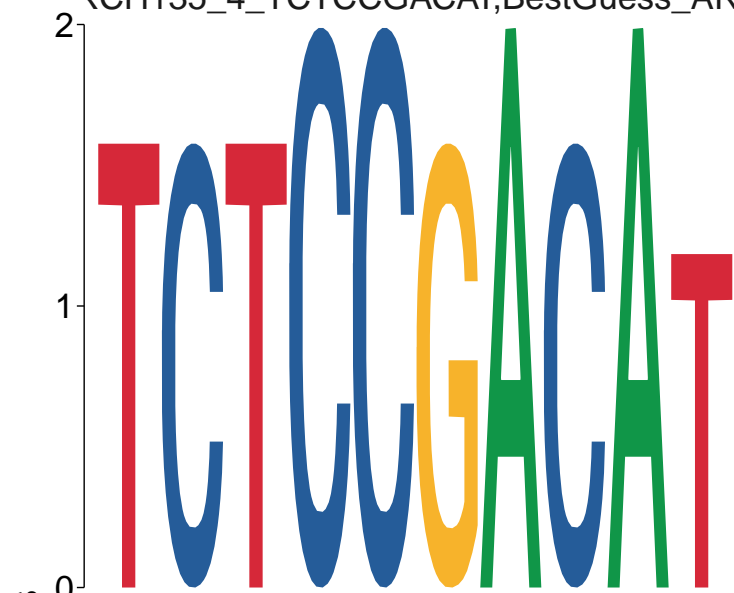
CH134_5_GATCGTATCGCG,BestGuess_R



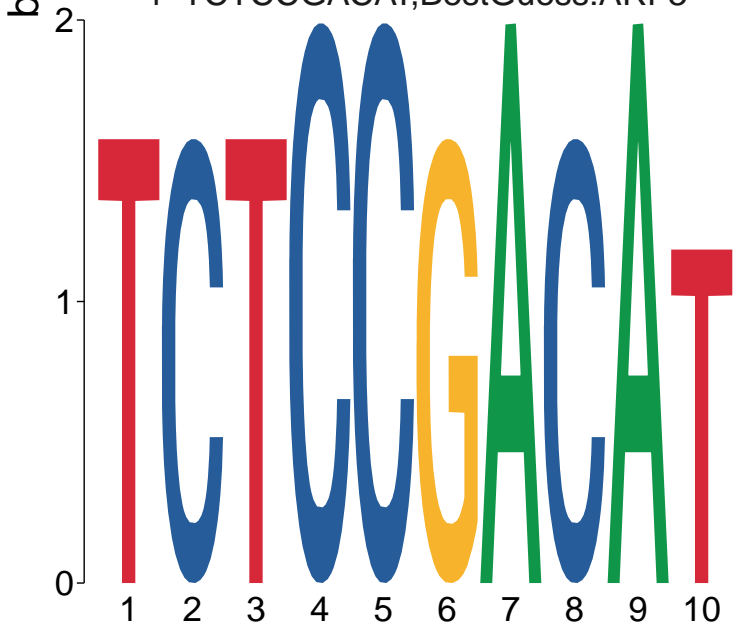
5-GATCGTATCGCG,BestGuess:RBM3



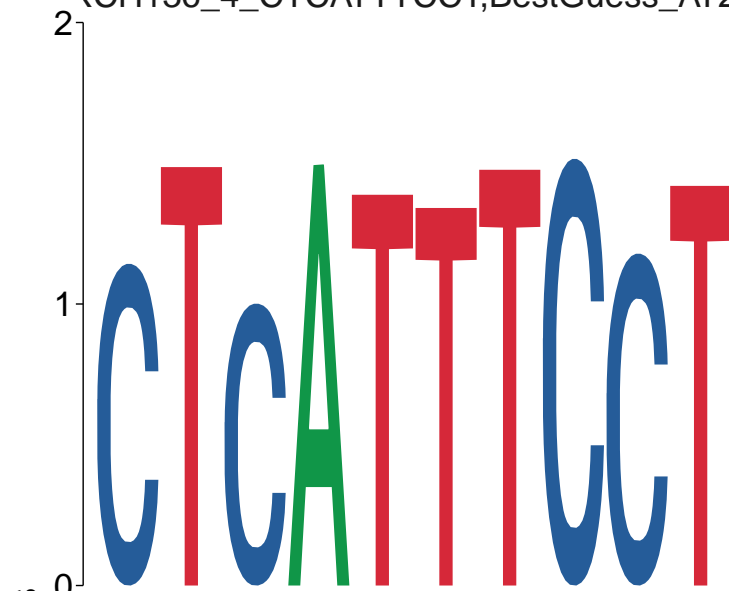
RCH135_4_TCTCCGACAT,BestGuess_ARF



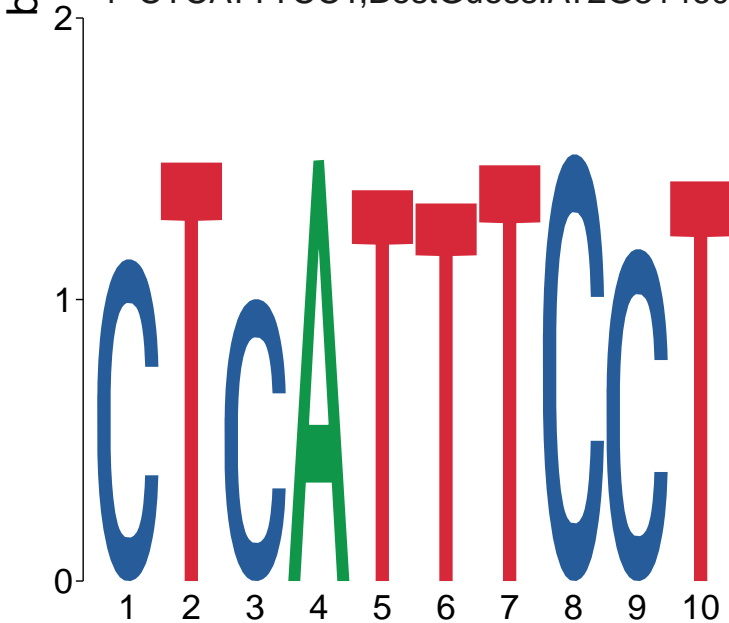
4-TCTCCGACAT,BestGuess:ARF3



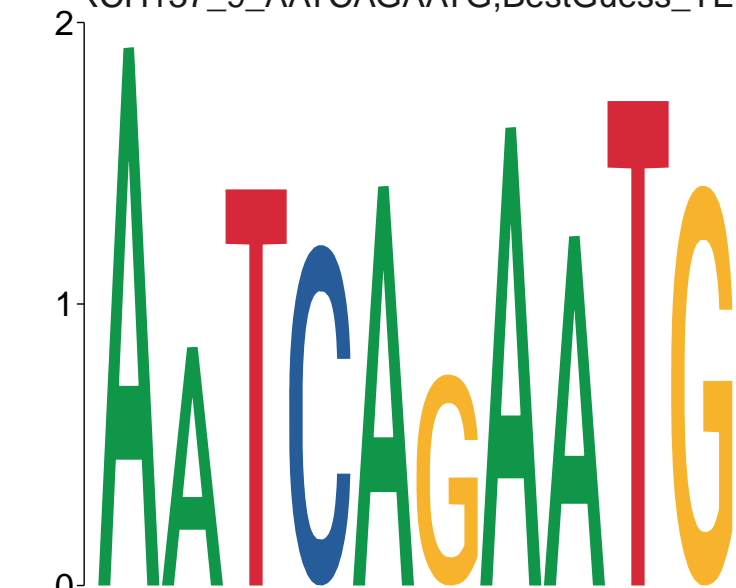
CH136_4 CTCATTTCCT, BestGuess_AT20



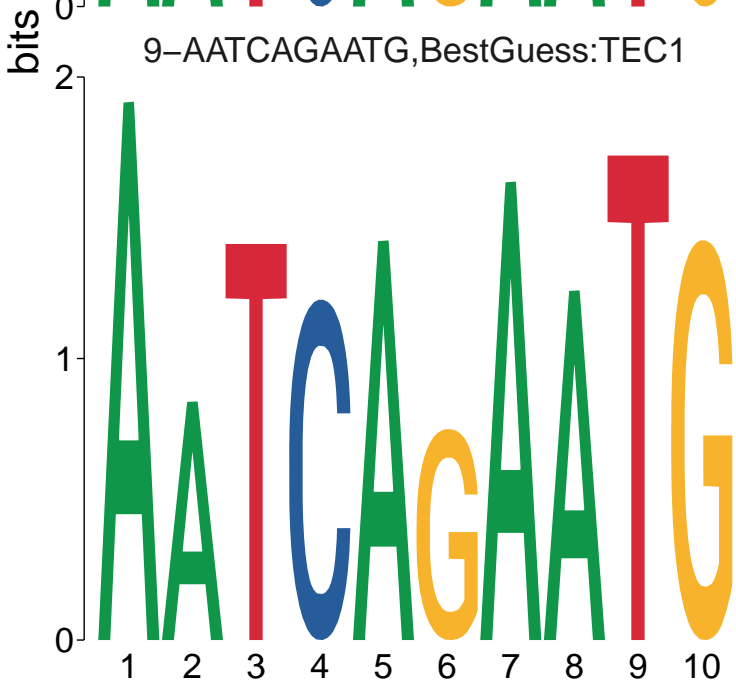
4-CTCATTTCCT, BestGuess:AT2G31460



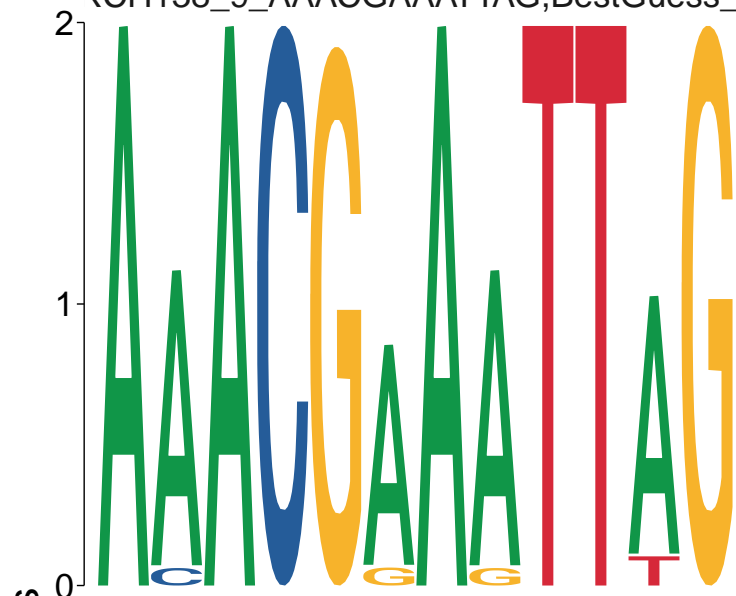
RCH137_9_AATCAGAATG,BestGuess_TEC



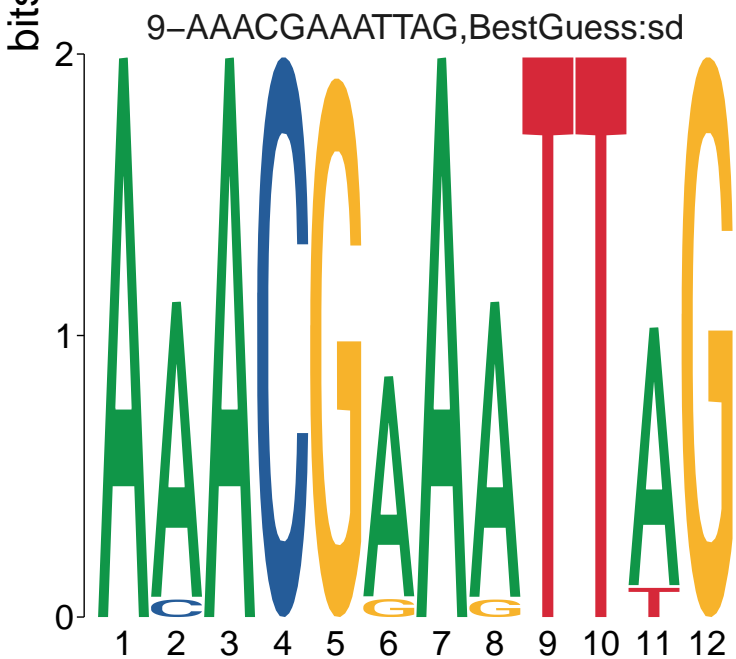
9-AATCAGAATG,BestGuess:TEC1



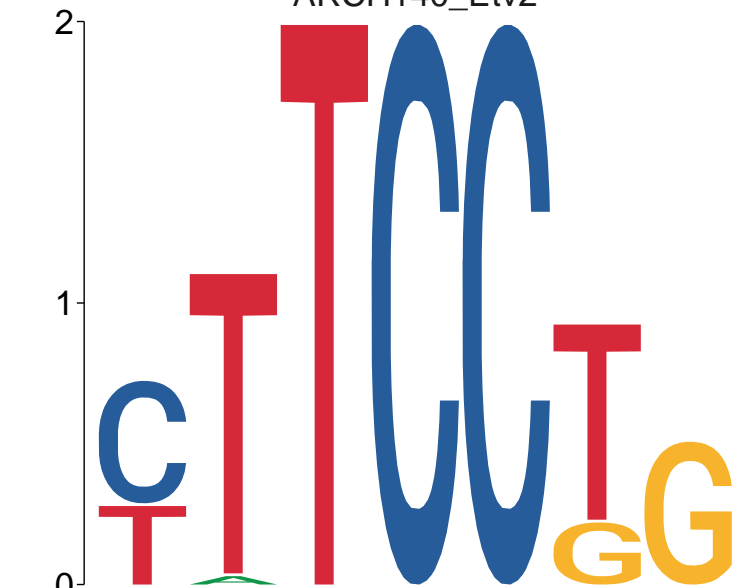
RCH138_9_AAACGAAATTAG,BestGuess_9



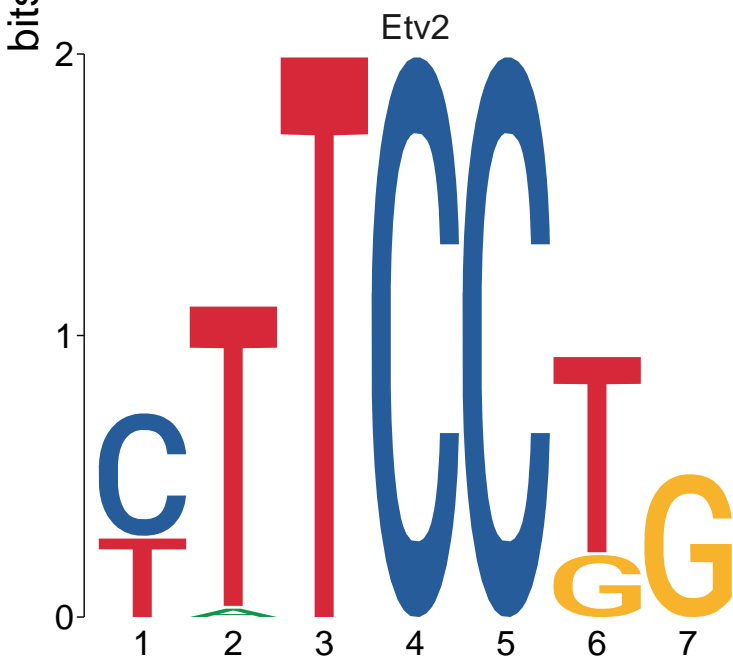
9-AAACGAAATTAG,BestGuess:sd



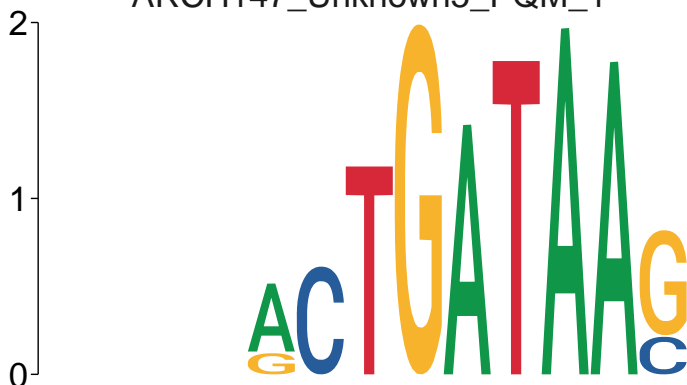
ARCH140_Etv2



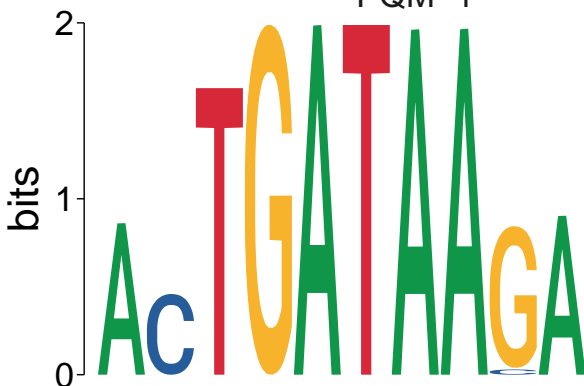
Etv2



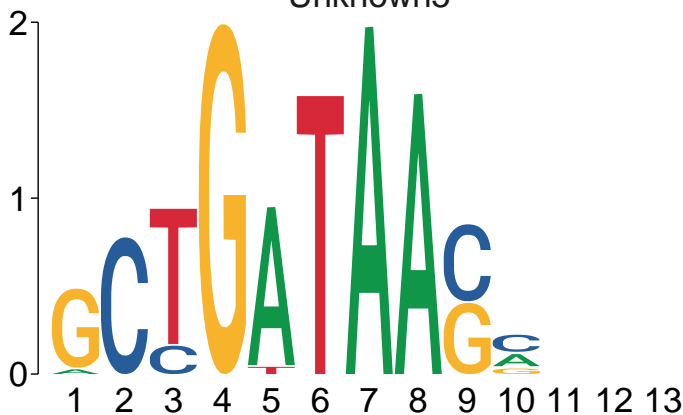
ARCH147_Unknown5_PQM_1



PQM-1



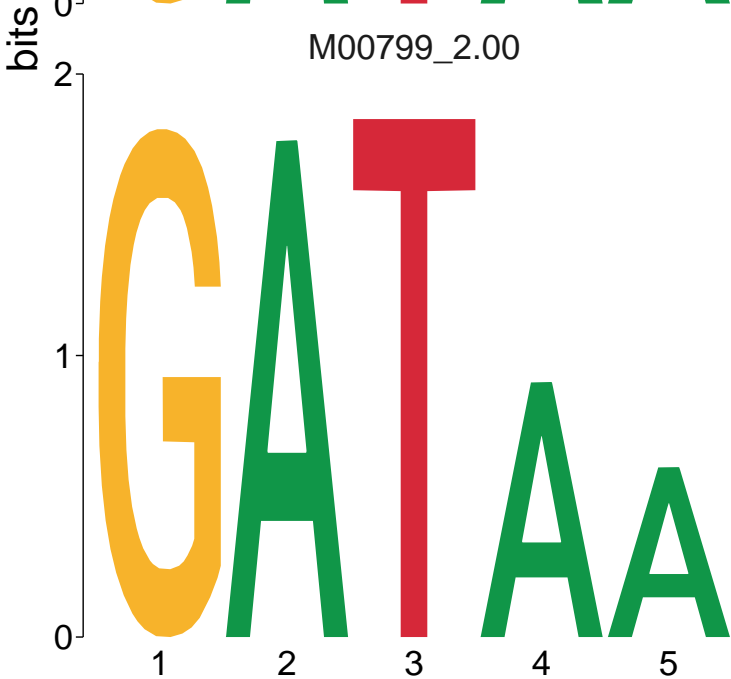
Unknown5



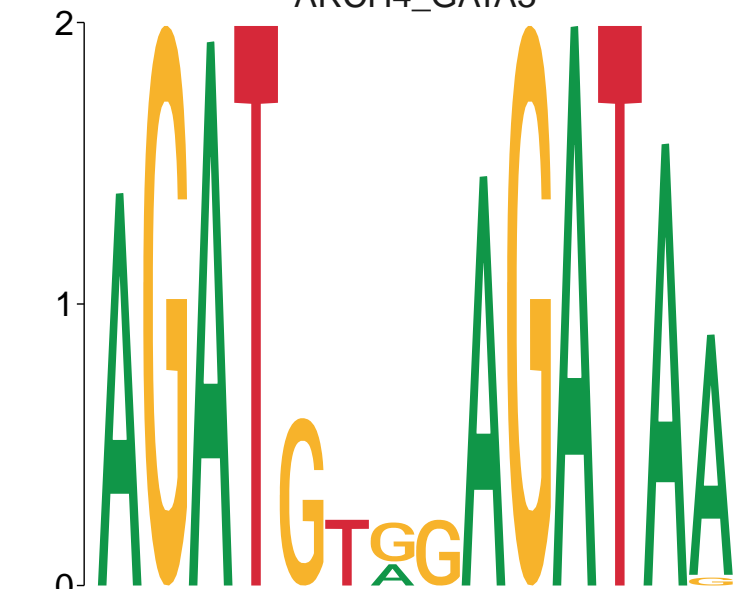
ARCH148_GATA2



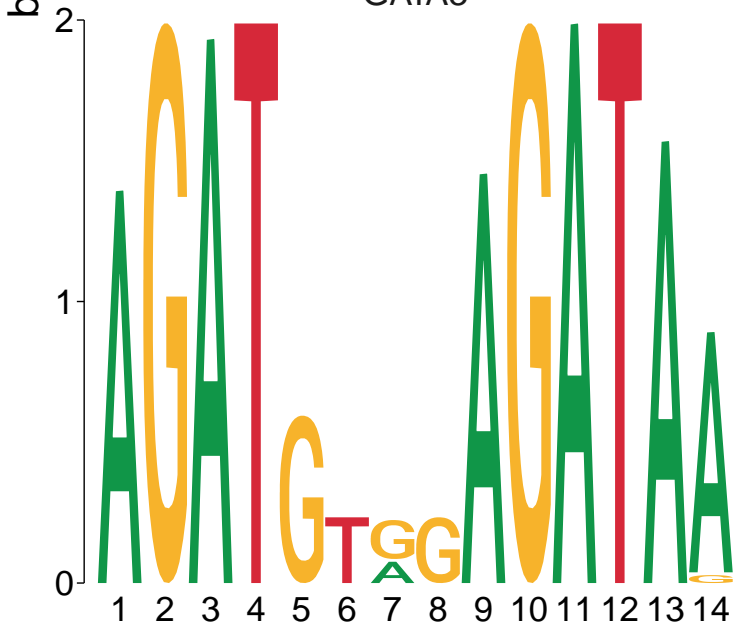
M00799_2.00



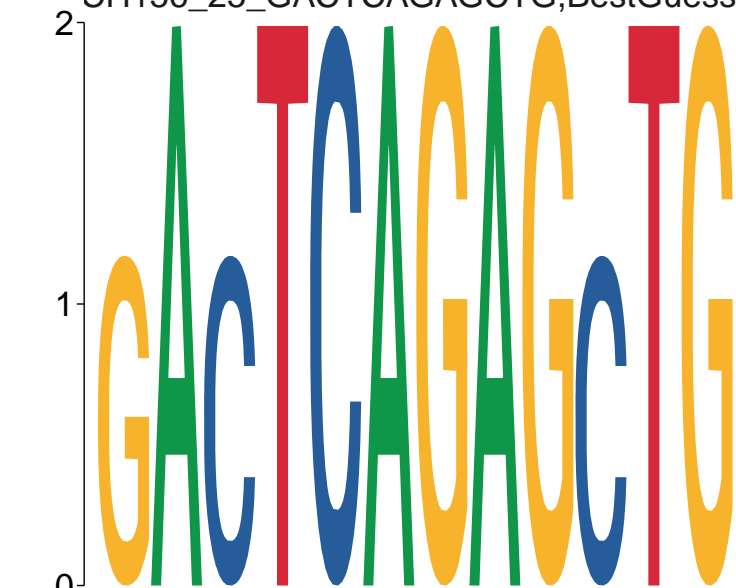
ARCH4_GATA3



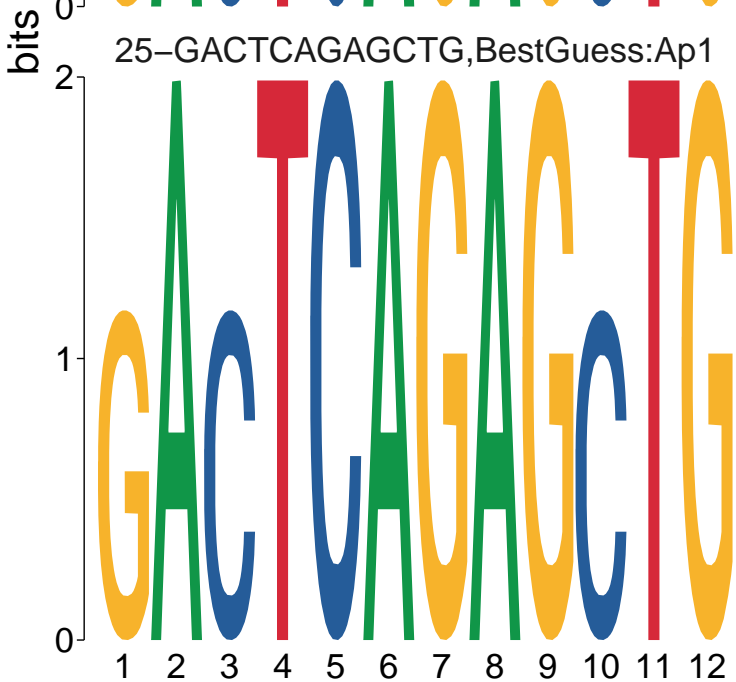
GATA3



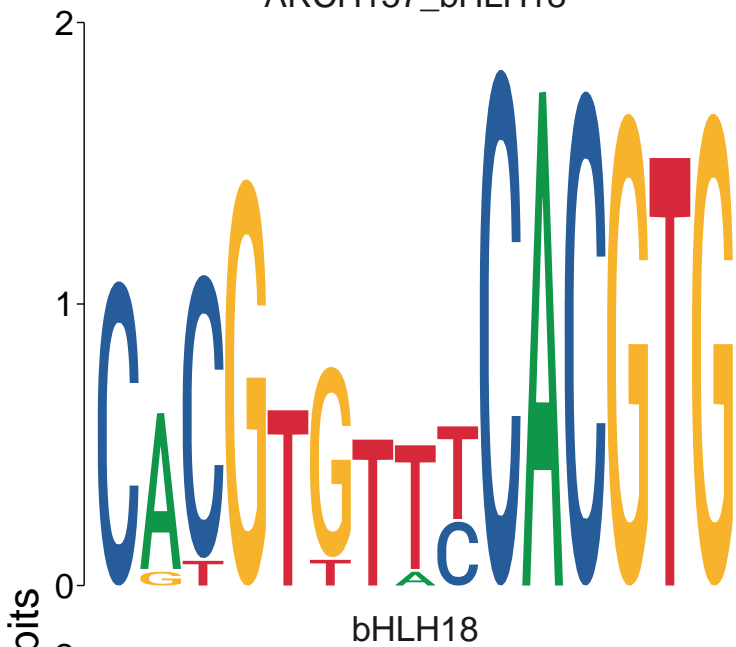
CH156_25_GACTCAGAGCTG,BestGuess_



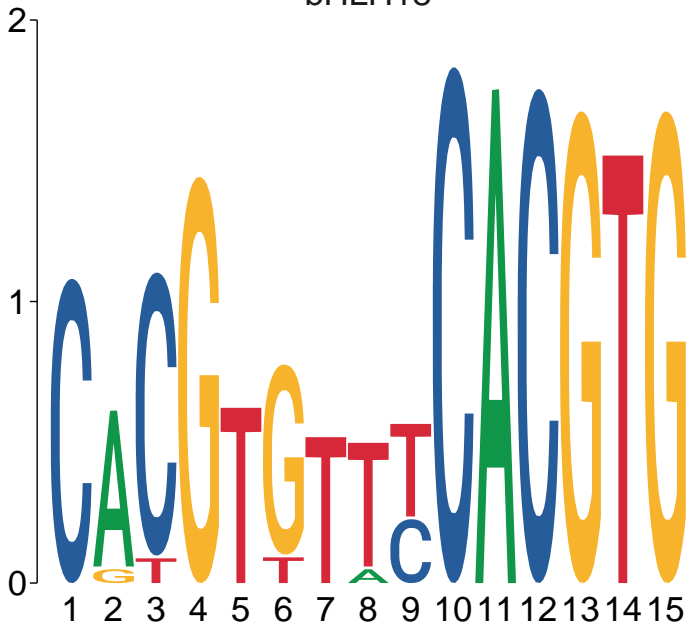
25-GACTCAGAGCTG,BestGuess:Ap1



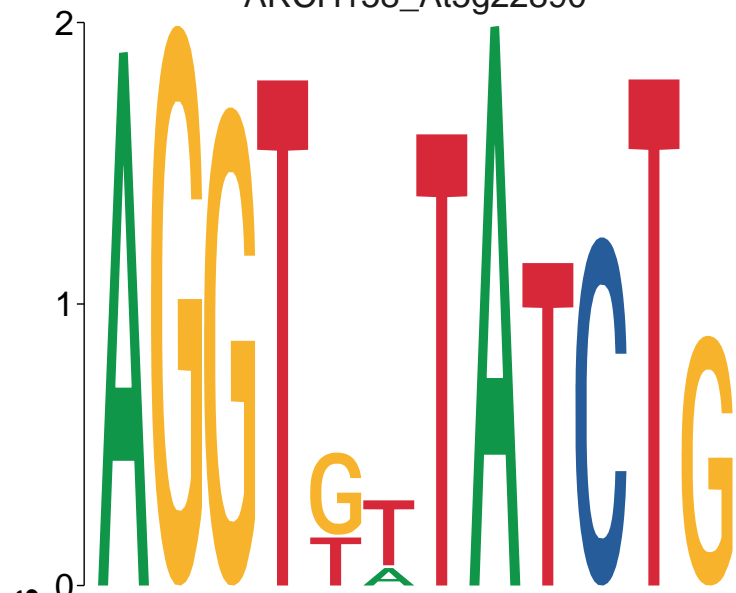
ARCH157_bHLH18



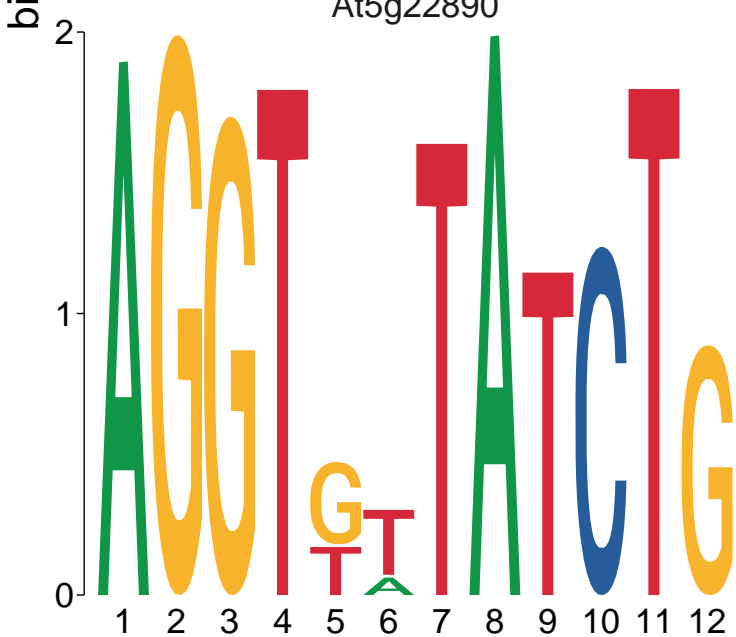
bHLH18



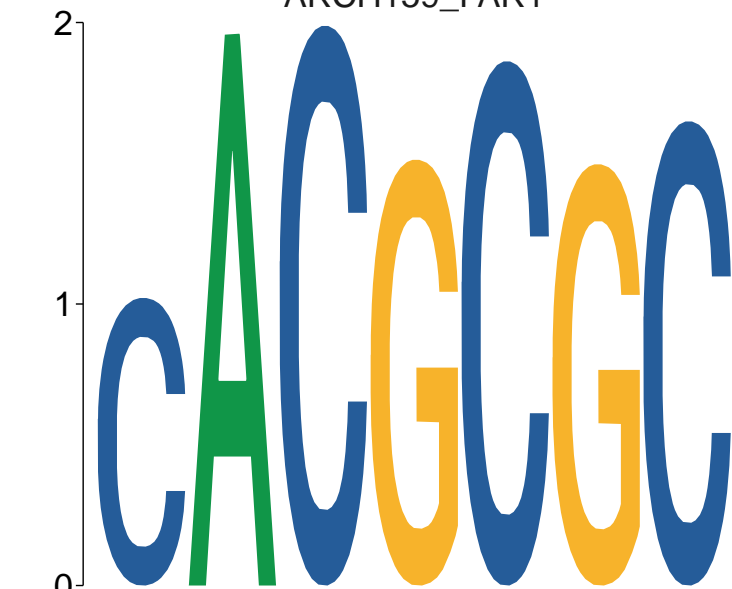
ARCH158_At5g22890



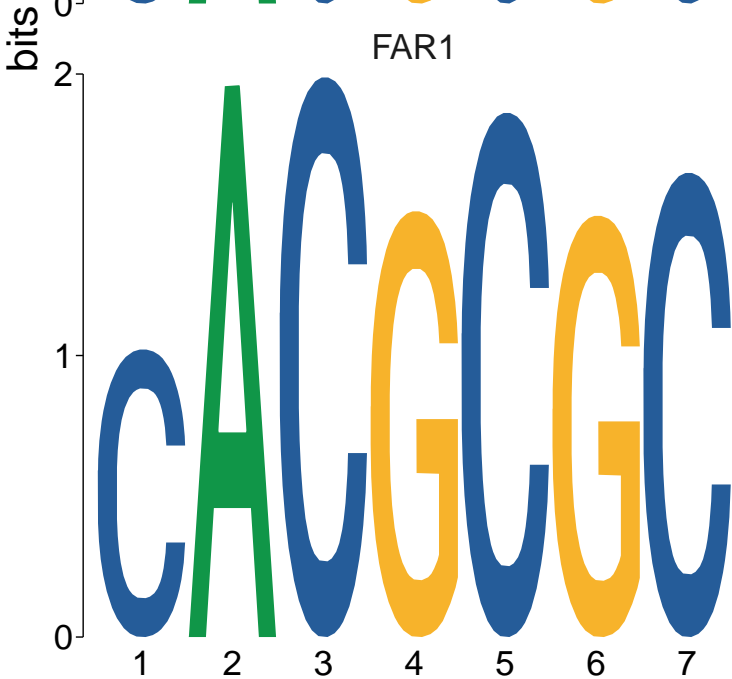
At5g22890



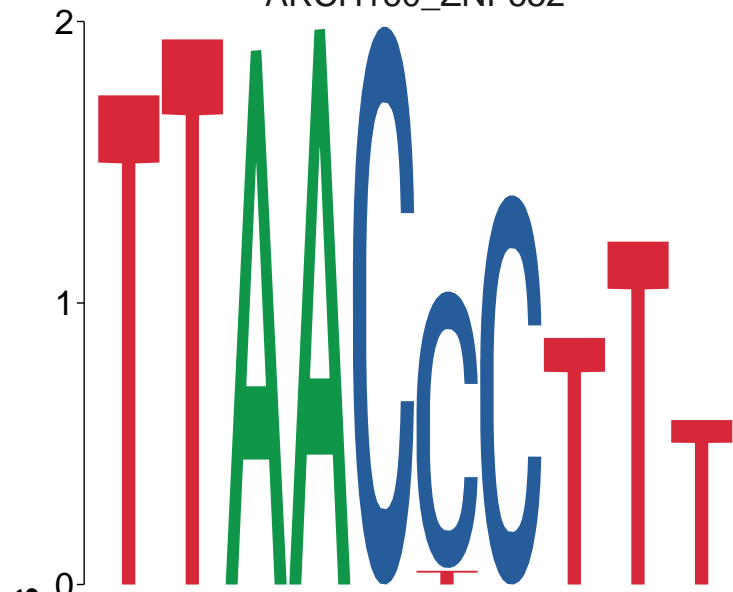
ARCH159_FAR1



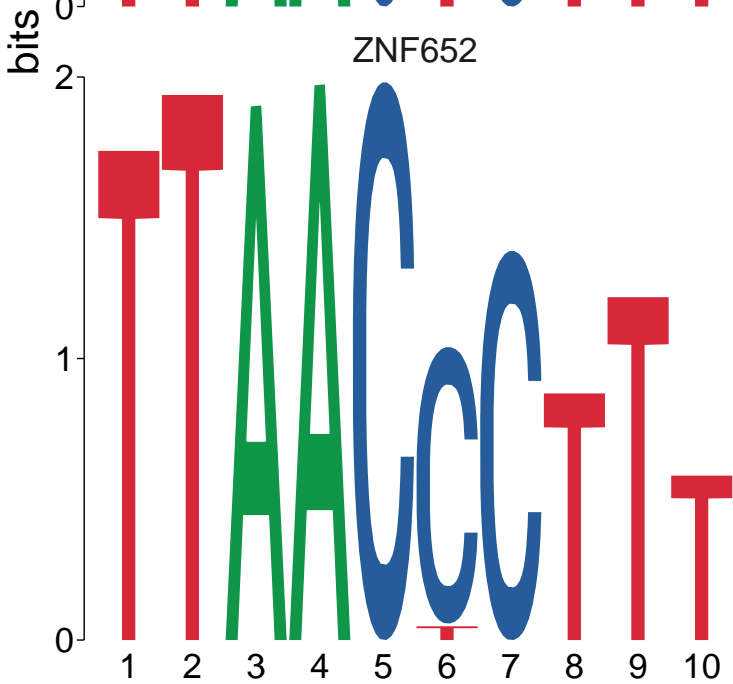
FAR1



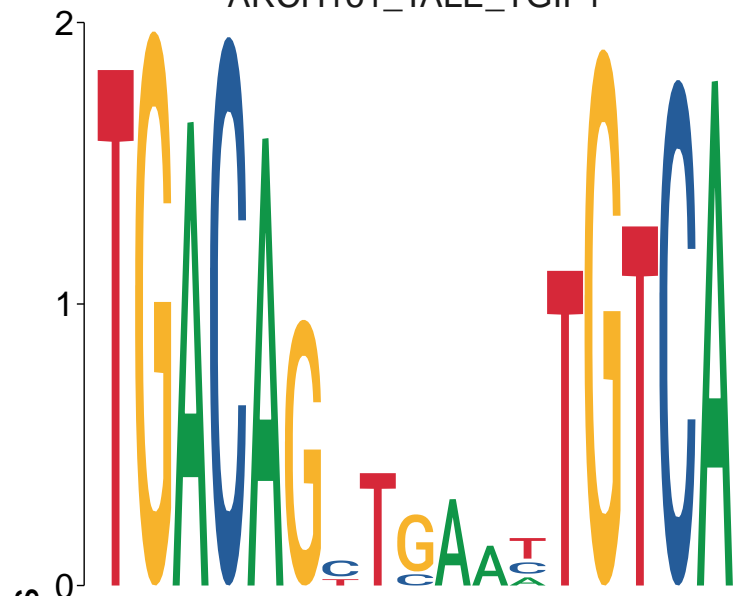
ARCH160_ZNF652



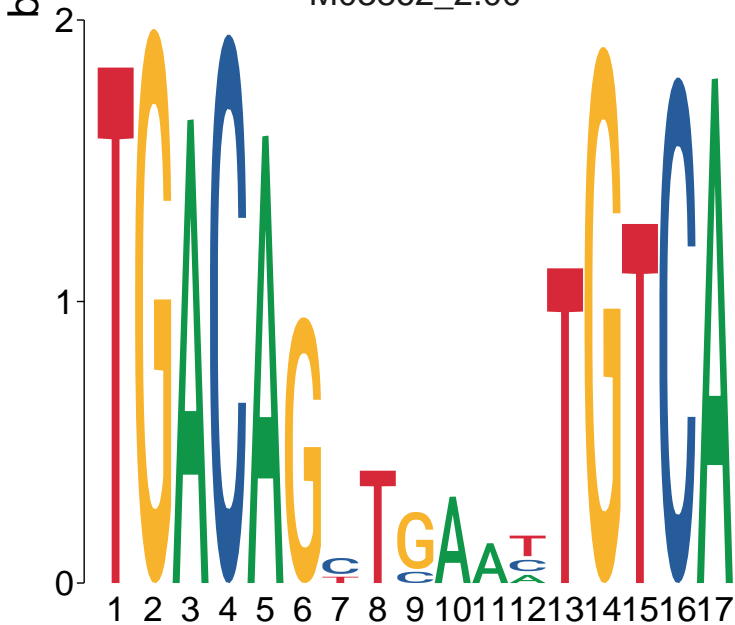
ZNF652



ARCH161_TALE_TGIF1



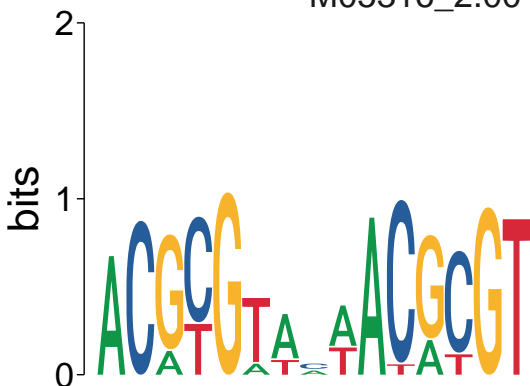
M03862_2.00



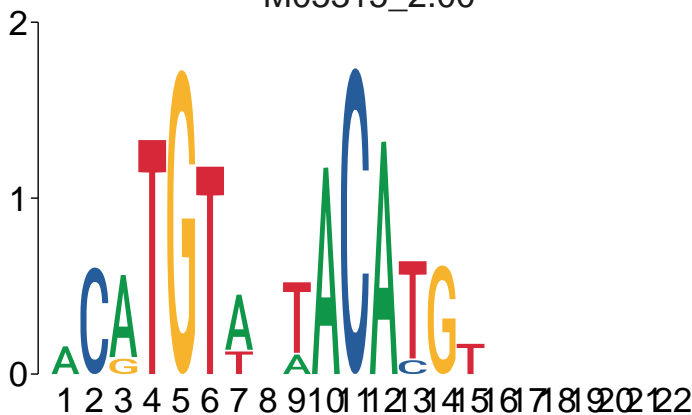
ARCH162_islike_TALE_IRX1_TALE_IRX2



M05316_2.00



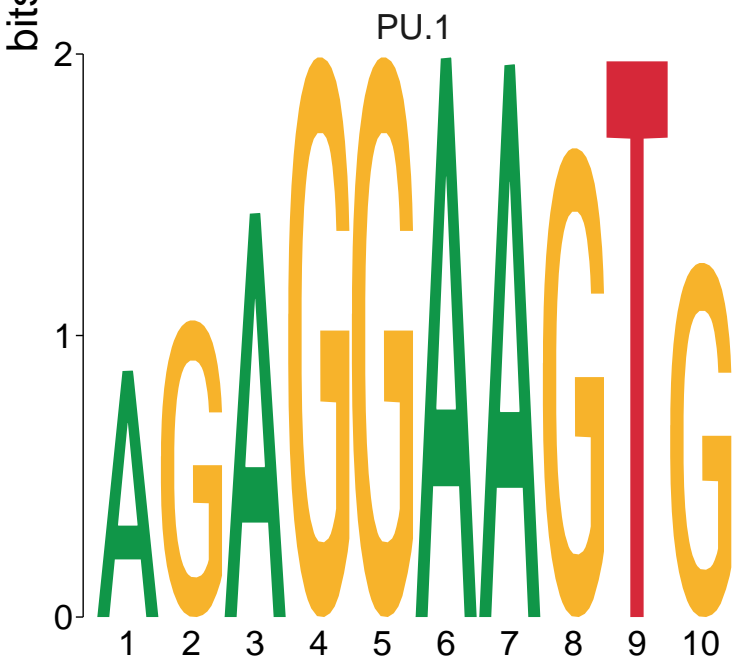
M05315_2.00



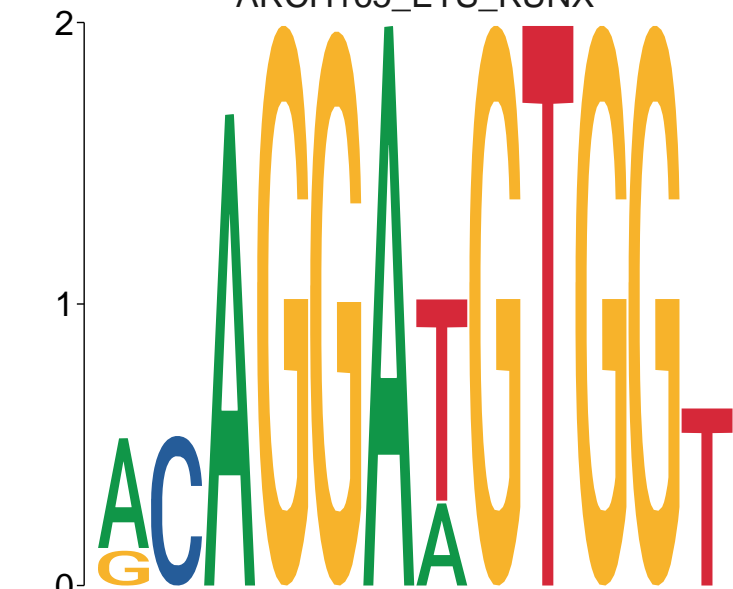
ARCH164_PU.1



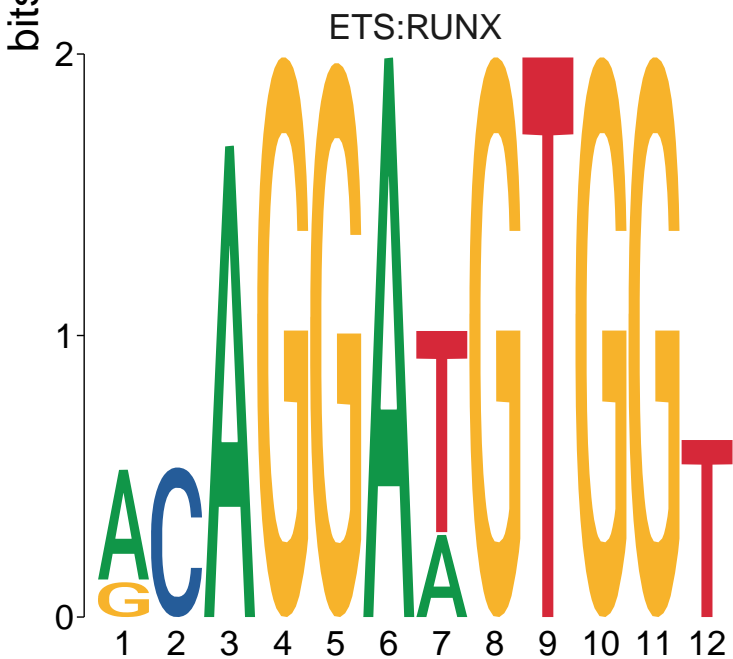
PU.1



ARCH165_ETS_RUNX



ETS:RUNX



RCH169_EWS_FLI1_fusion_EWS_ERG_fu



Ets1-distal



Ets1-distal.1



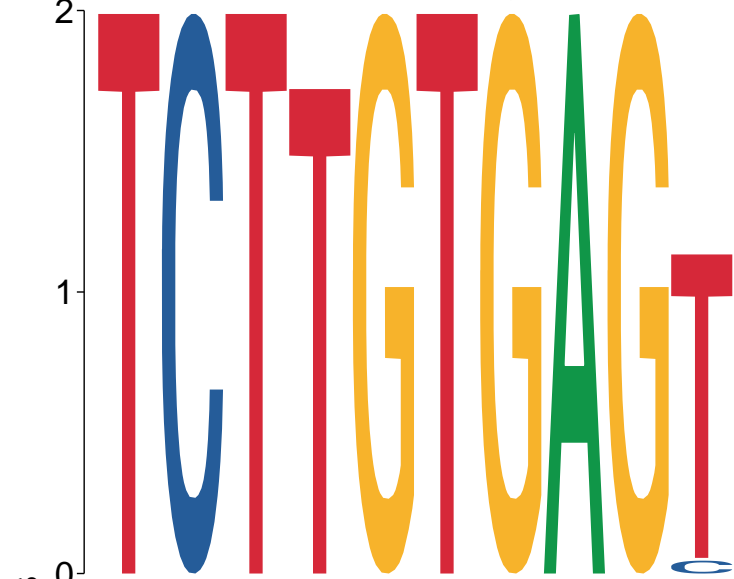
EWS:ERG-fusion



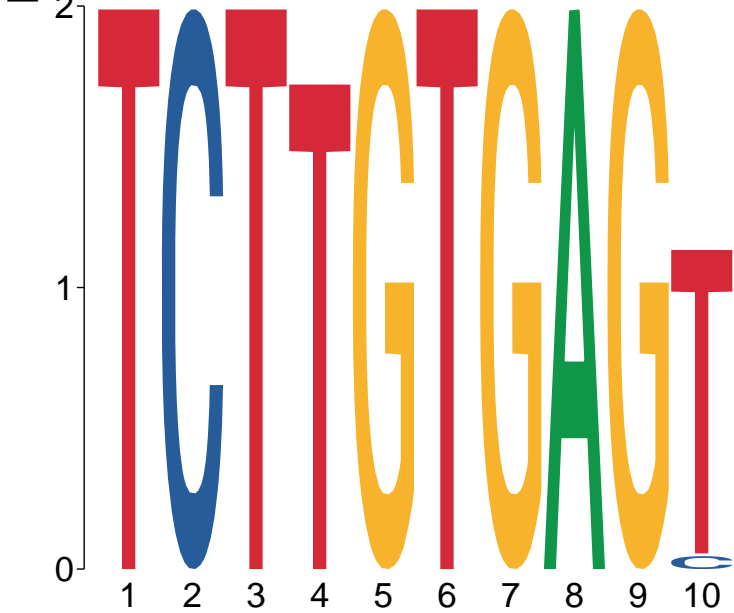
EWS:FLI1-fusion



CH170_20_TCTTGTGAGT,BestGuess_MI'



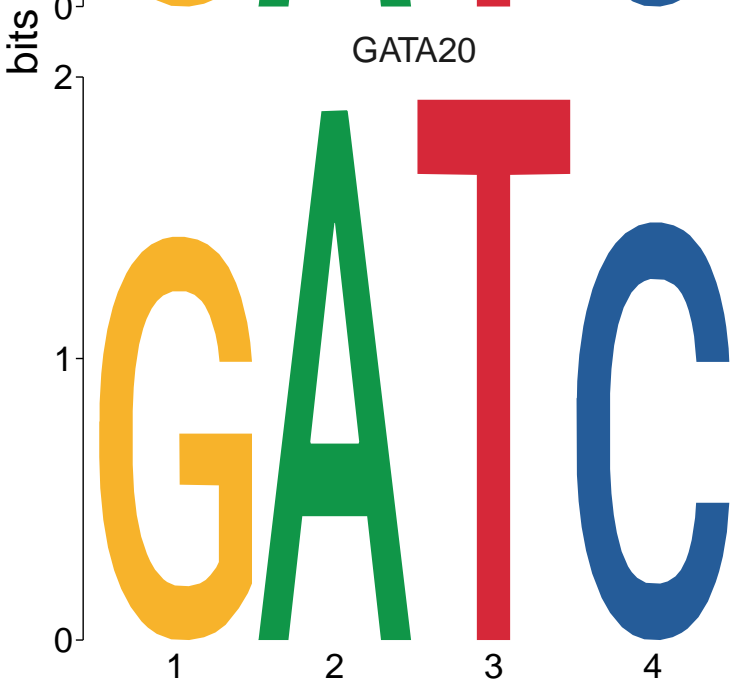
20-TCTTGTGAGT,BestGuess:MITF



ARCH171_GATA20



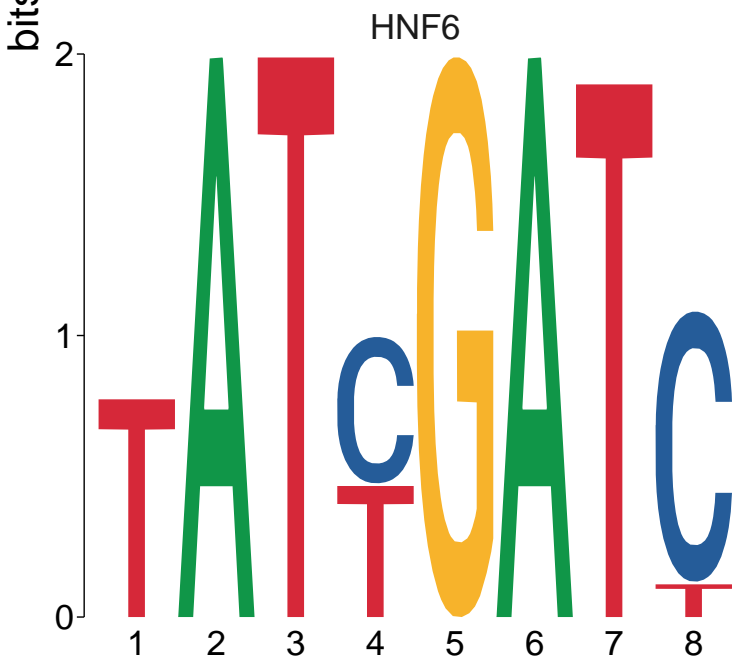
GATA20



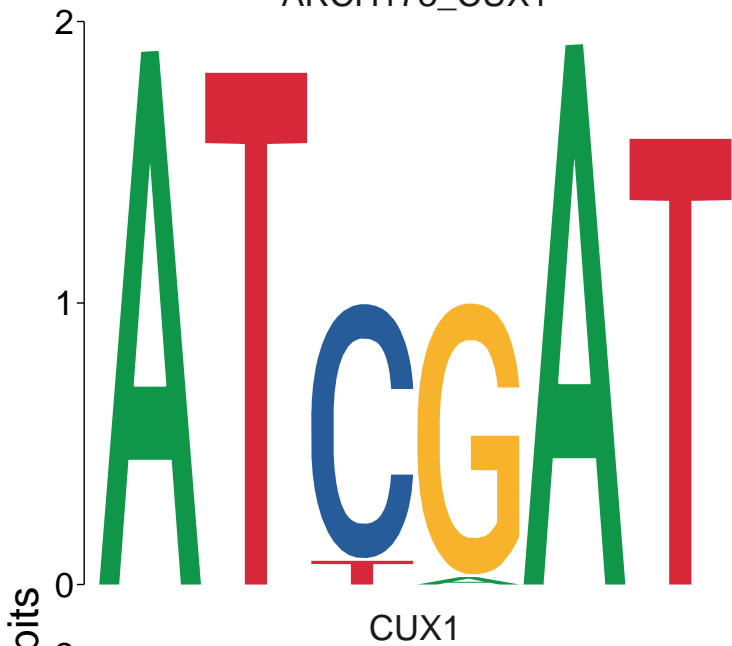
ARCH172_HNF6



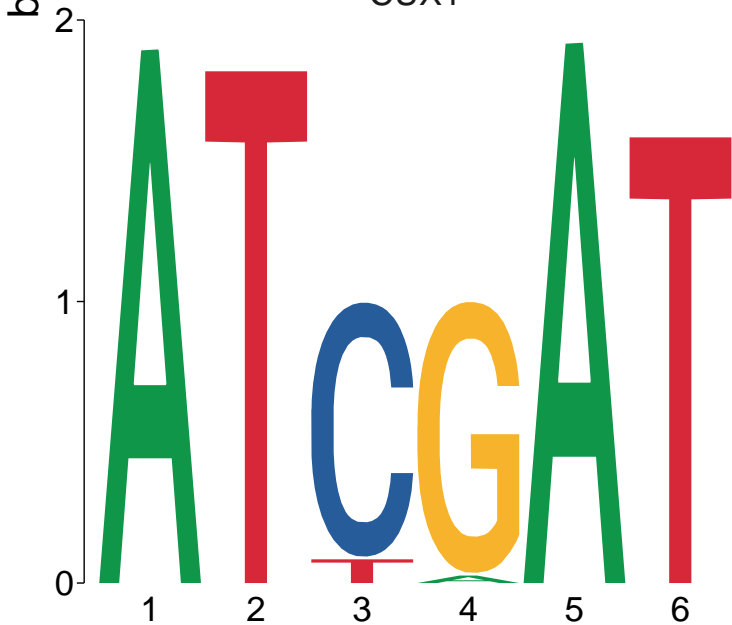
HNF6



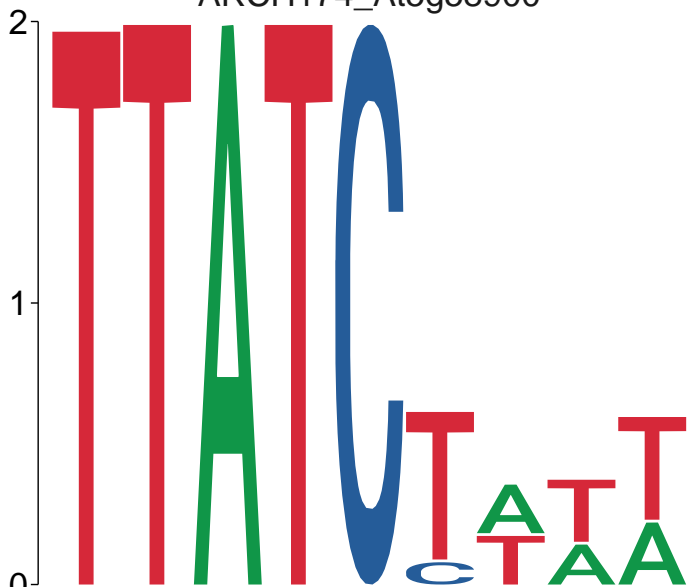
ARCH173_CUX1



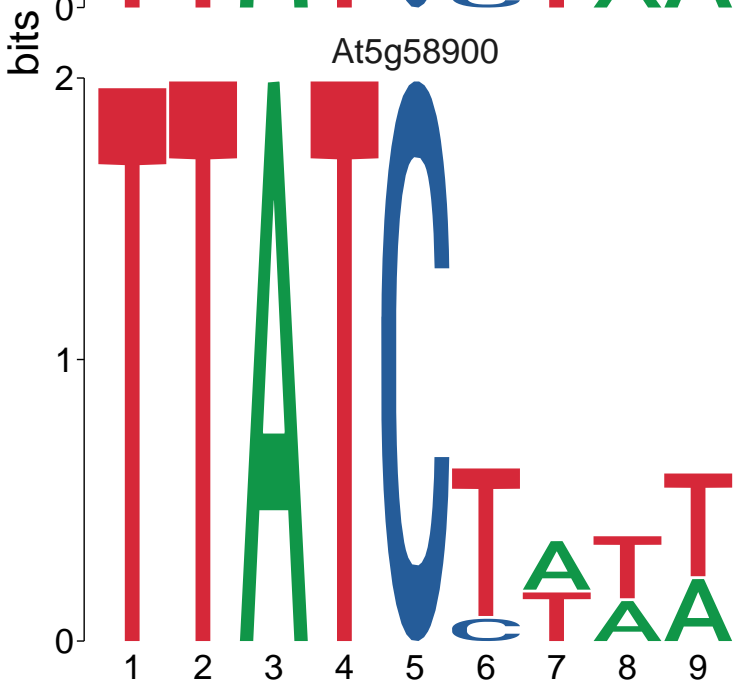
CUX1



ARCH174_At5g58900



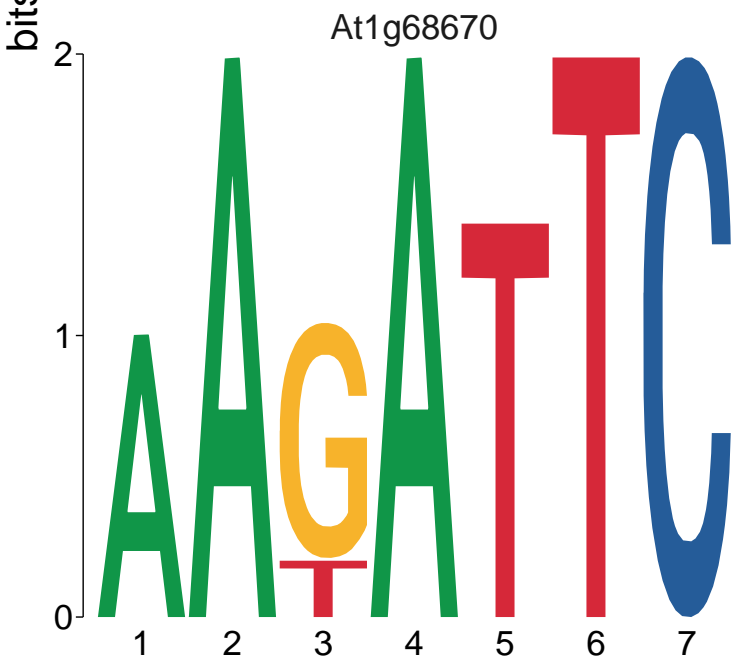
At5g58900



ARCH175_At1g68670



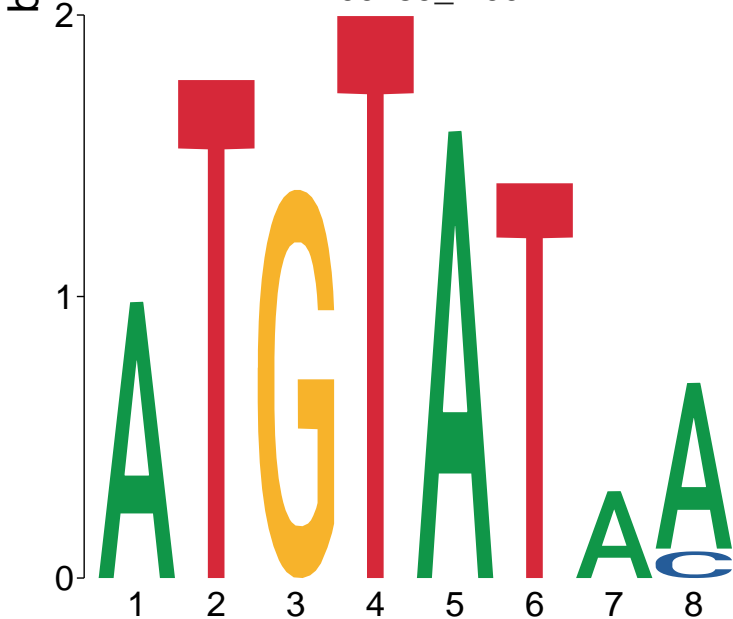
At1g68670



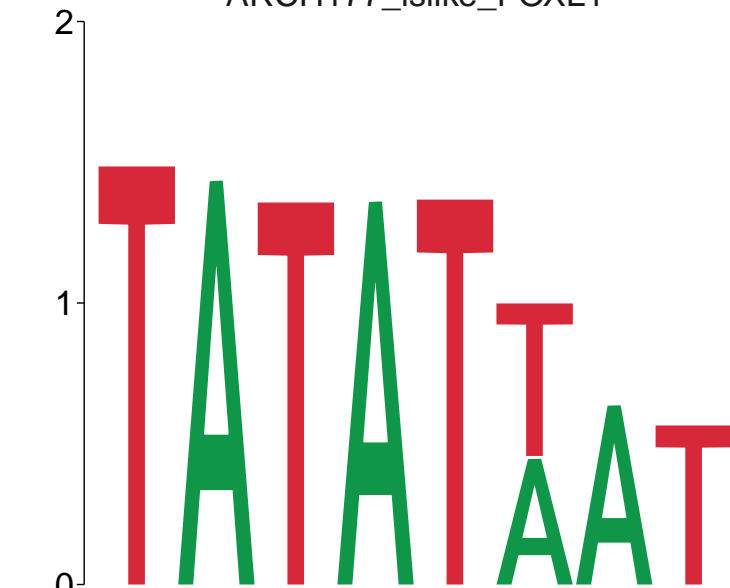
ARCH176_DMRT1



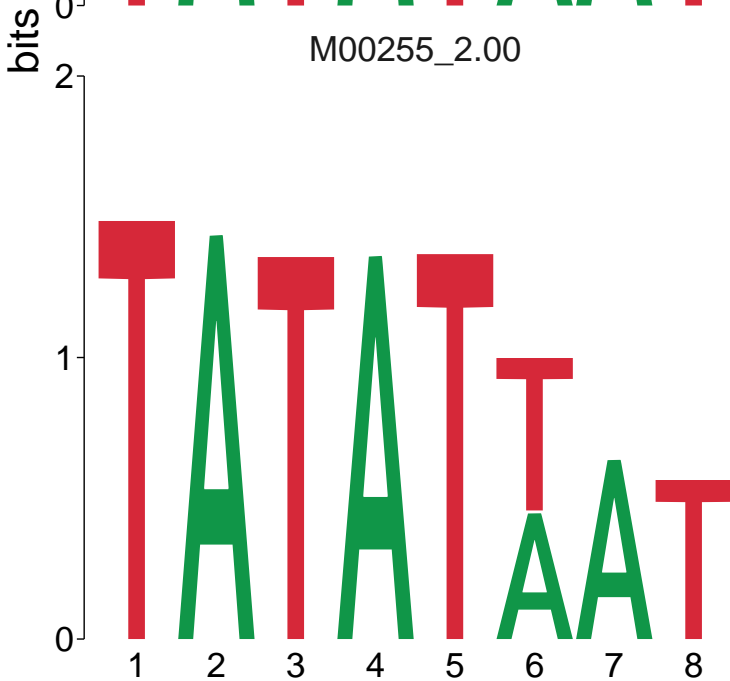
M00789_2.00



ARCH177_islike_FOXL1



M00255_2.00



ARCH178_ANTP_HOXA2_ANTP_HOXA3

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M00285_2.00

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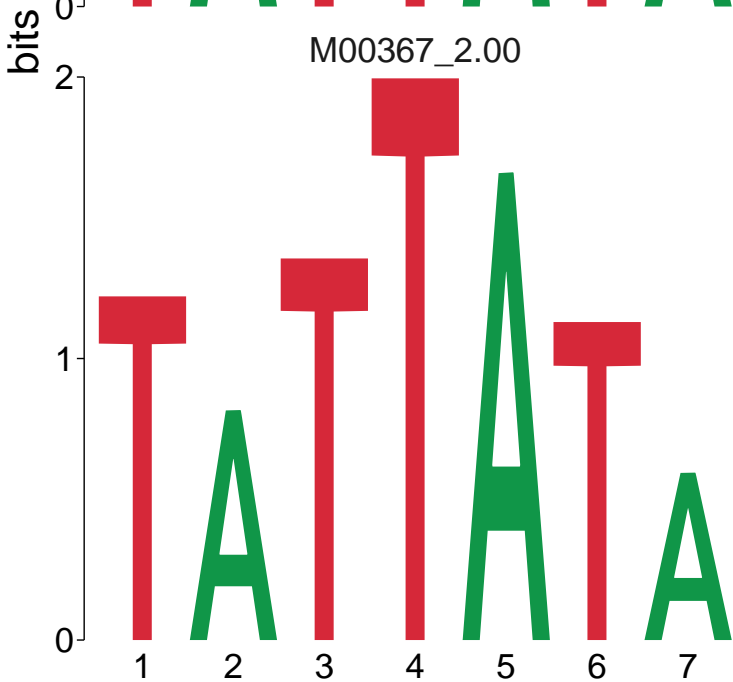
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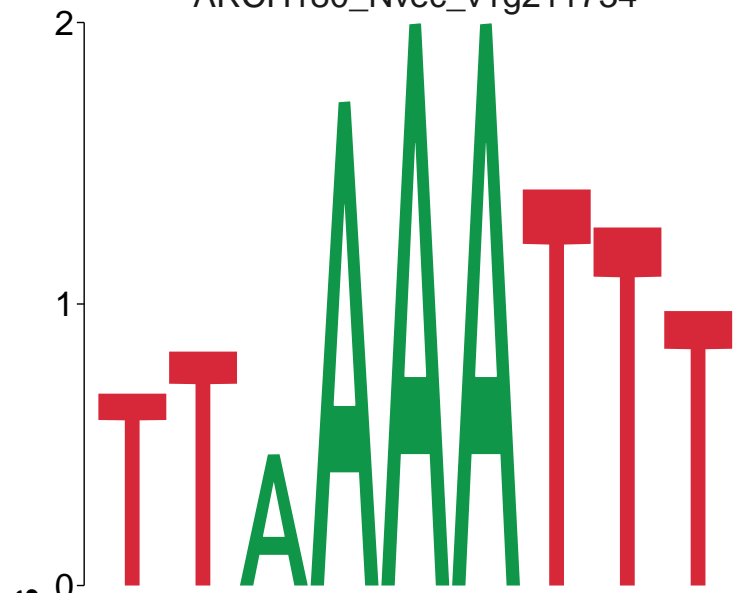
ARCH179_zf-C4_Nuclear_receptors



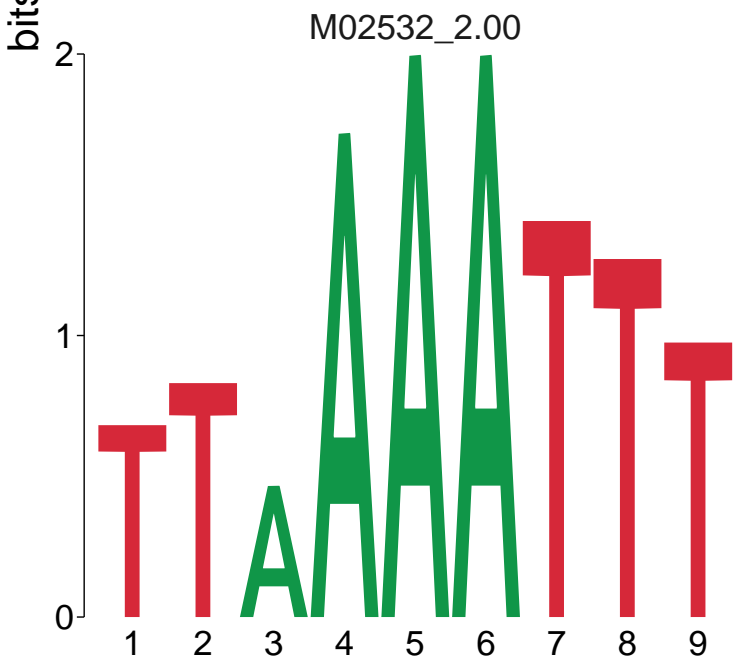
M00367_2.00



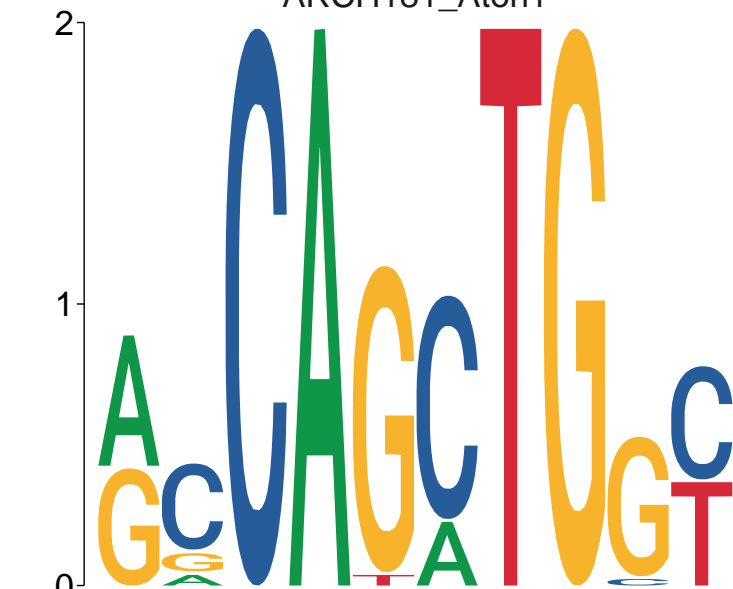
ARCH180_Nvec_v1g211734



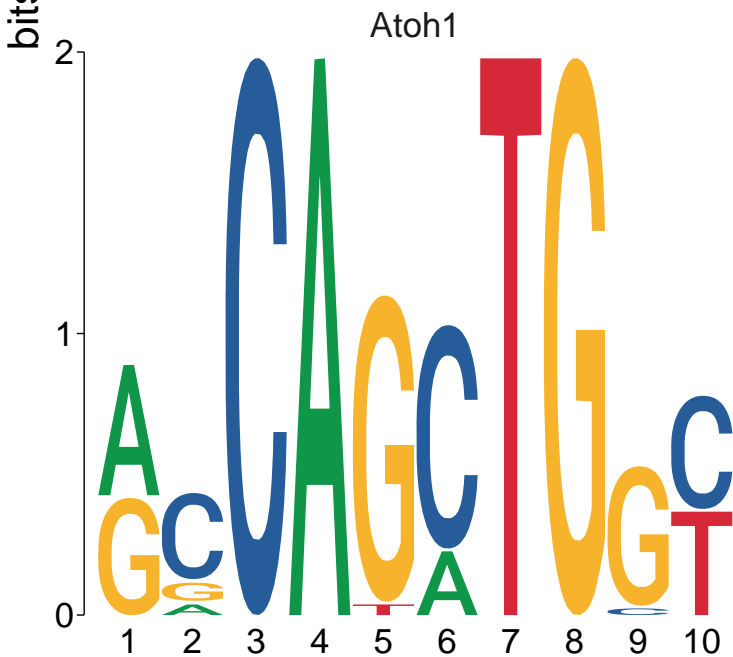
M02532_2.00



ARCH181_Atoh1



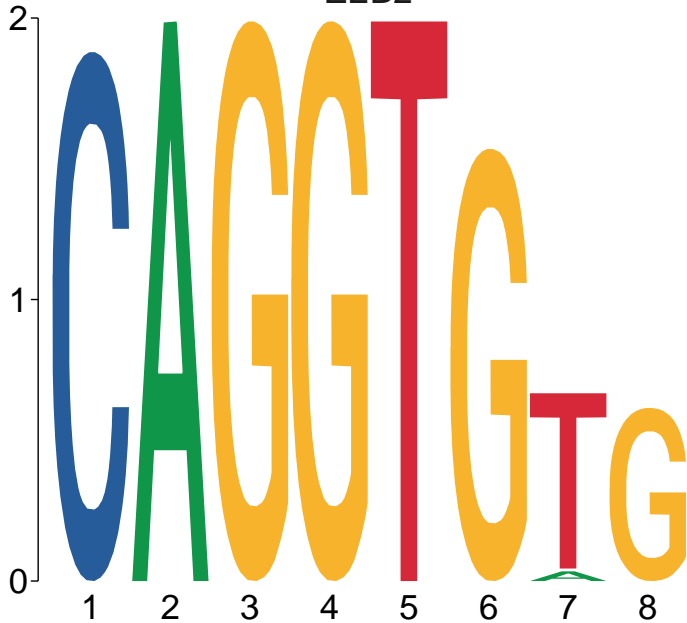
Atoh1



ARCH182_ZEB2



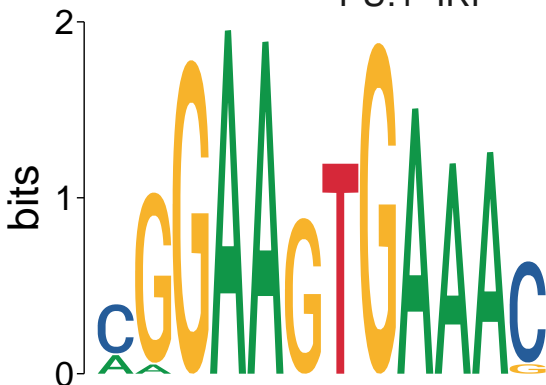
ZEB2



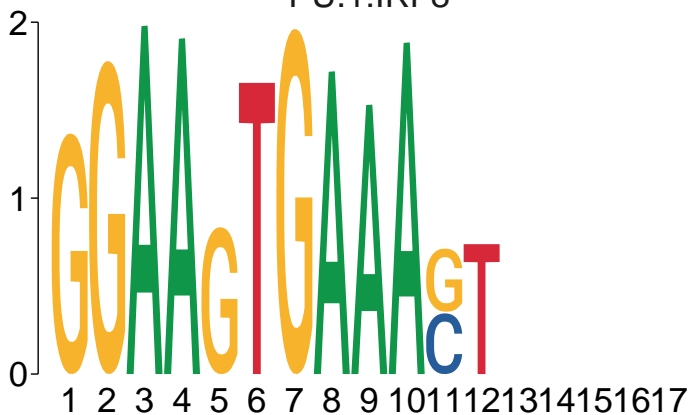
ARCH183_PU.1_IRF8_PU.1_IRF



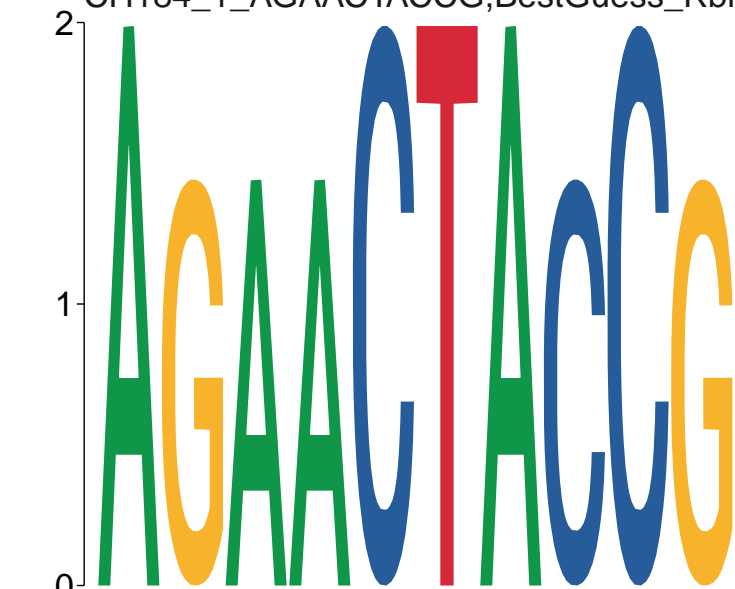
PU.1-IRF



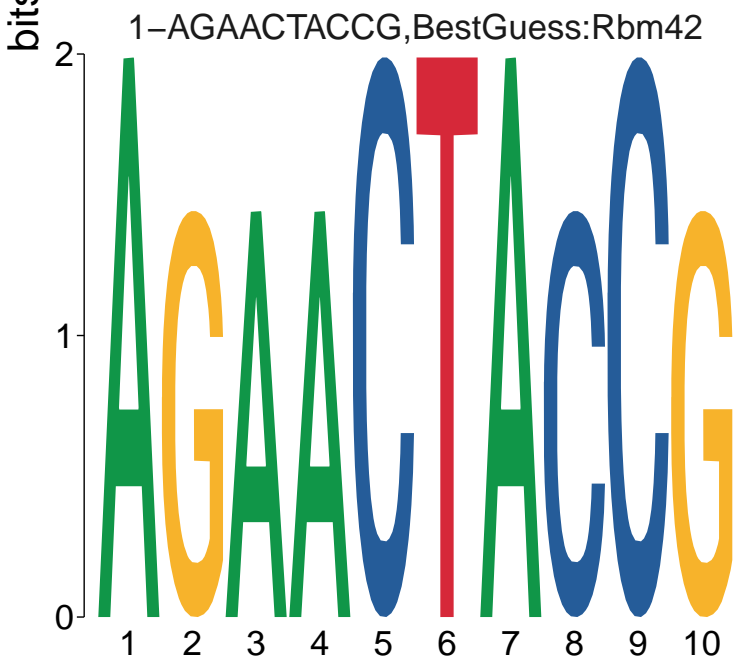
PU.1:IRF8



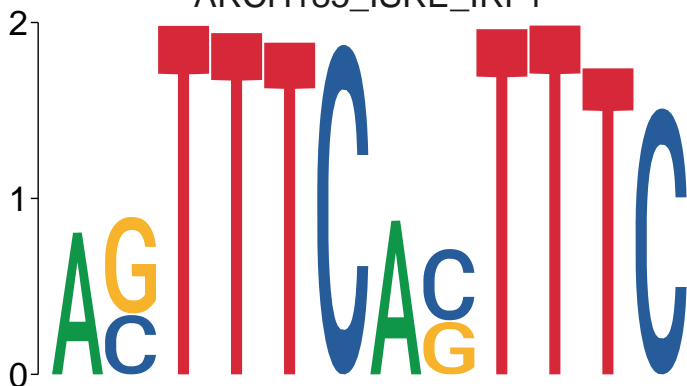
CH184_1_AGAACCTACCG,BestGuess_Rbr



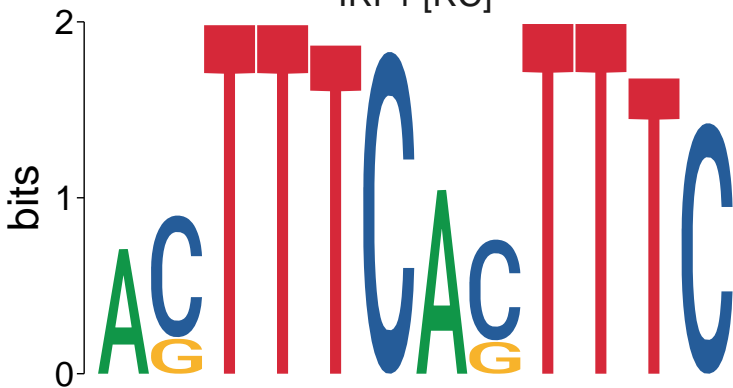
1-AGAACCTACCG,BestGuess:Rbm42



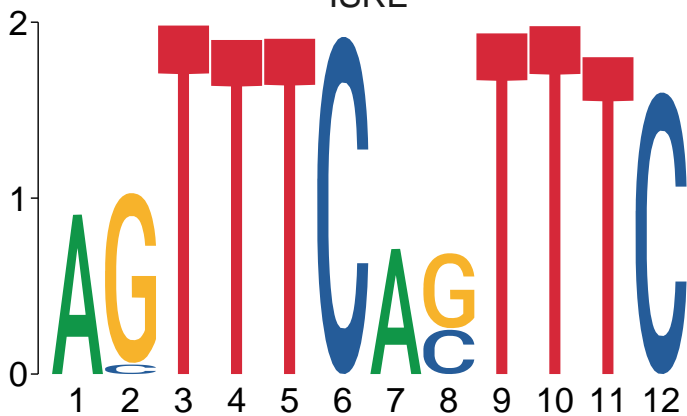
ARCH185_ISRE_IRF1



IRF1 [RC]



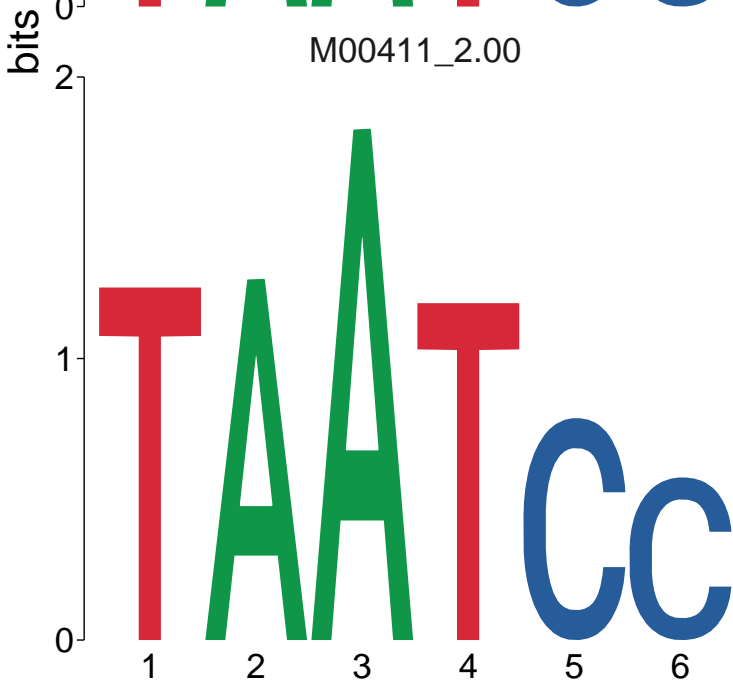
ISRE



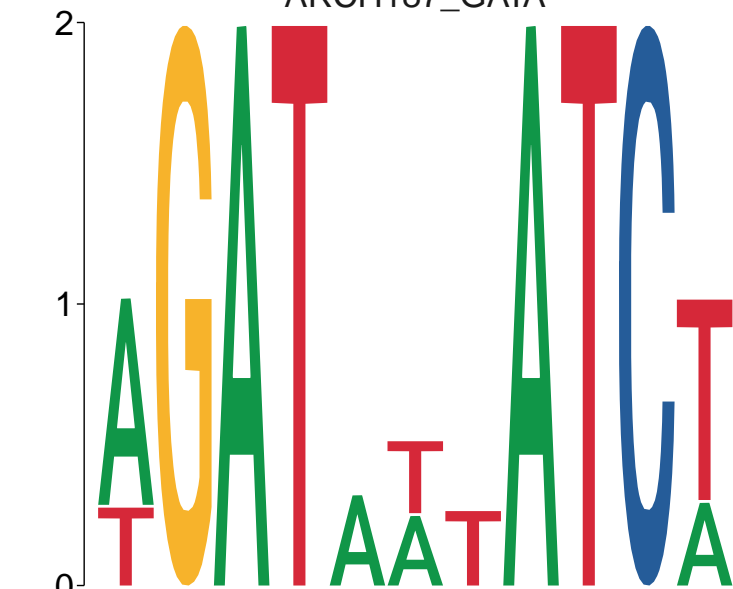
ARCH186_PRD_DMBX1



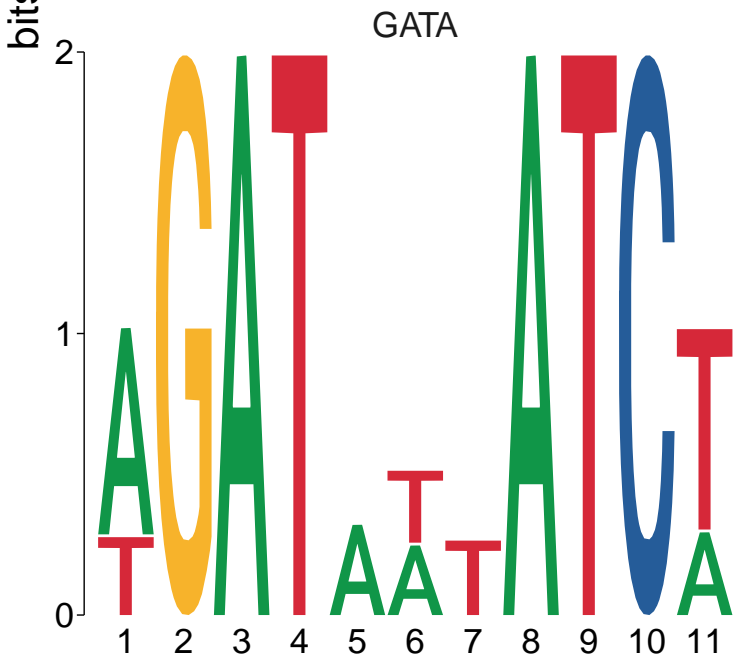
M00411_2.00



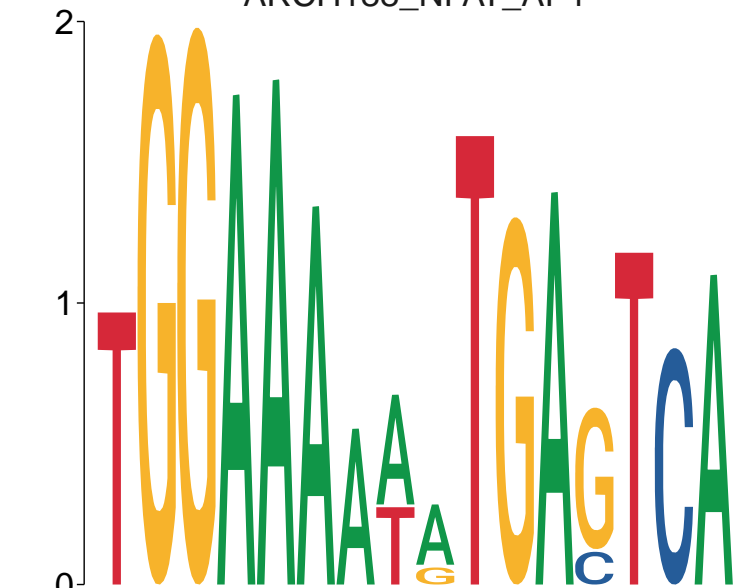
ARCH187_GATA



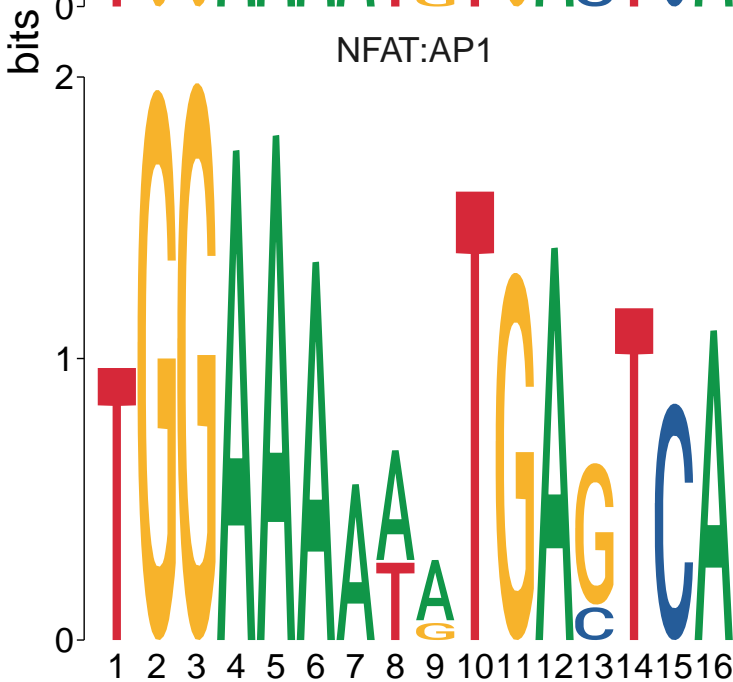
GATA



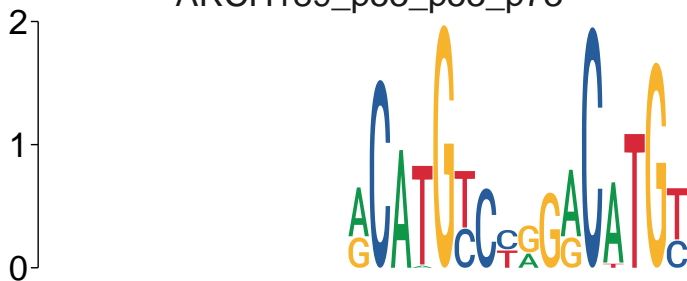
ARCH188_NFAT_AP1



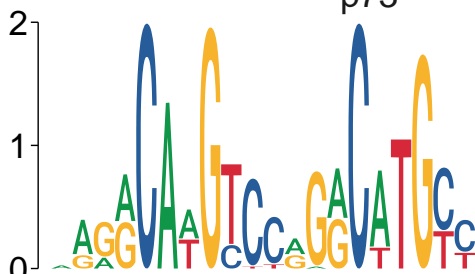
NFAT:AP1



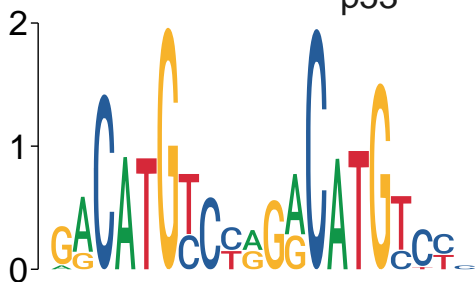
ARCH189_p53_p53_p73



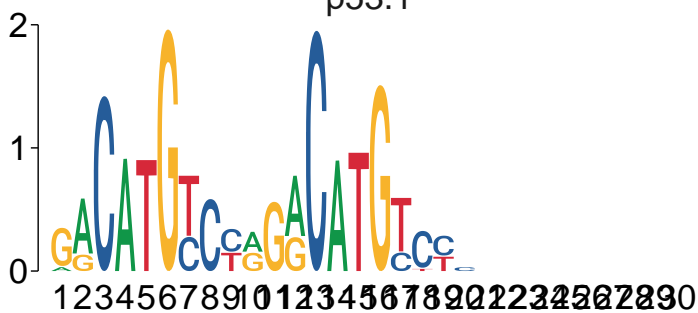
p73



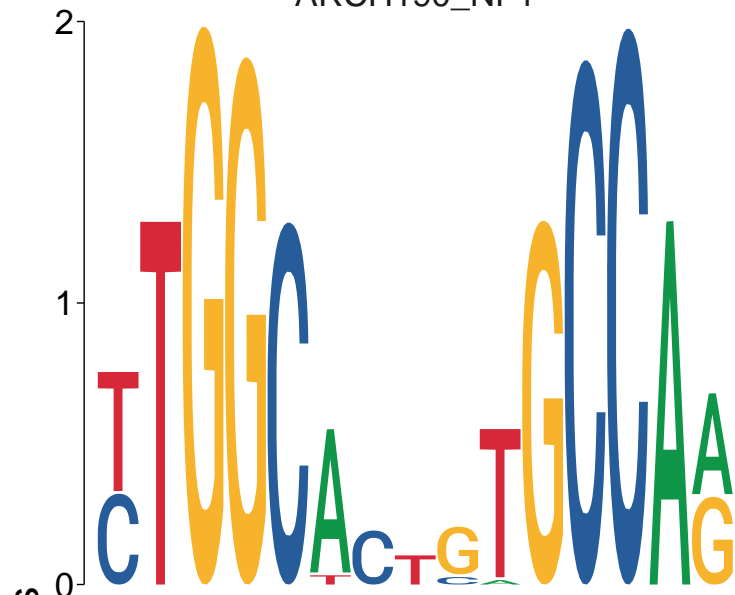
p53



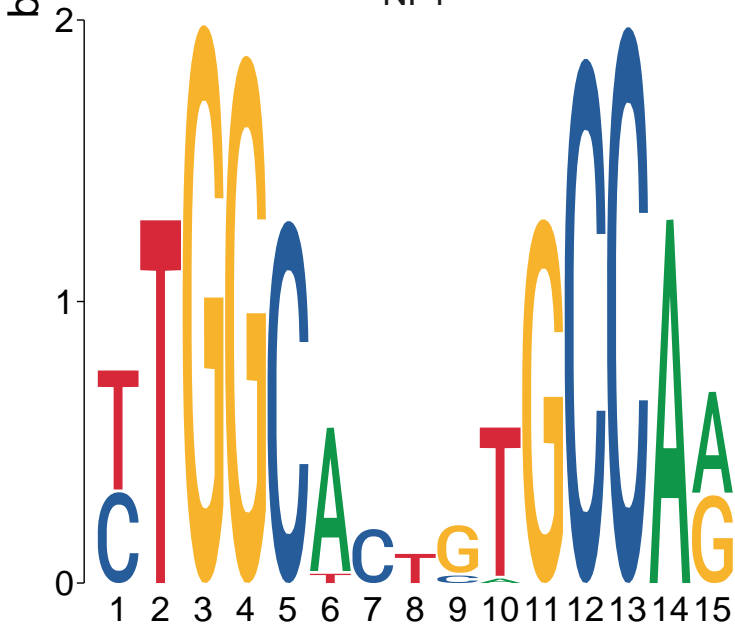
p53.1



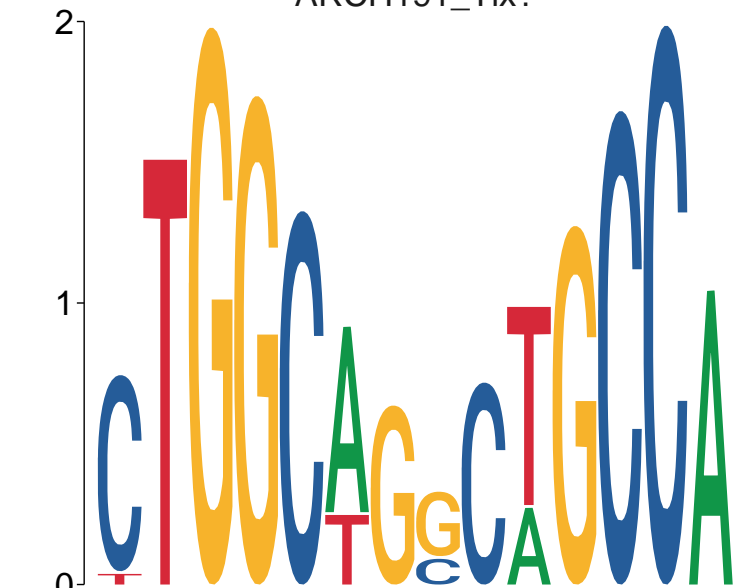
ARCH190_NF1



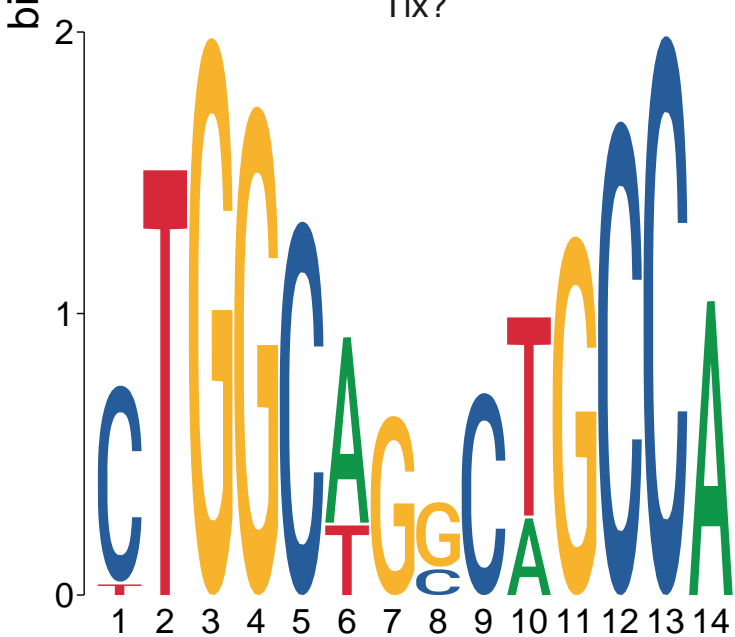
NF1



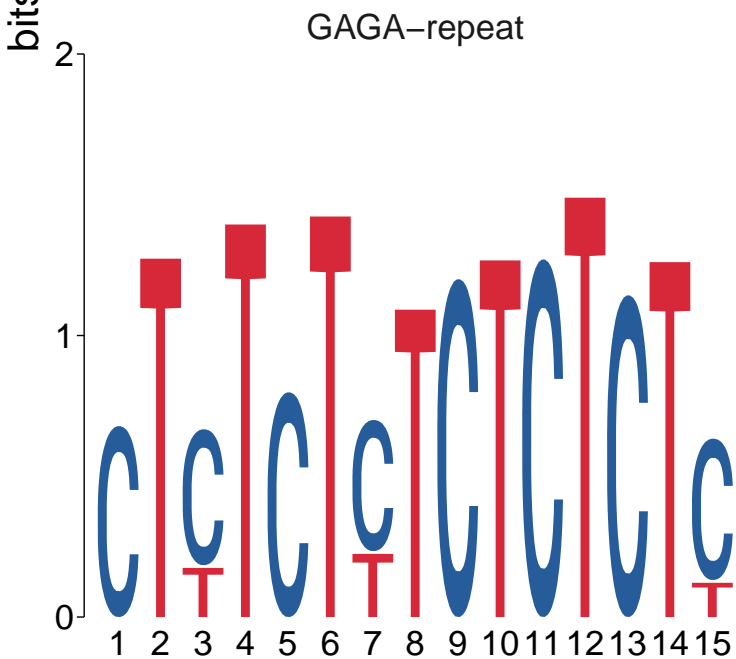
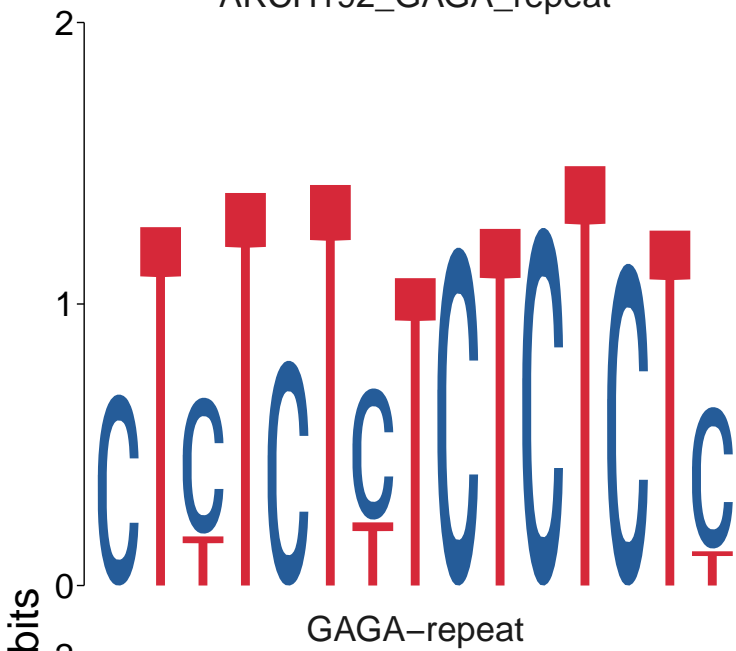
ARCH191_Tlx?



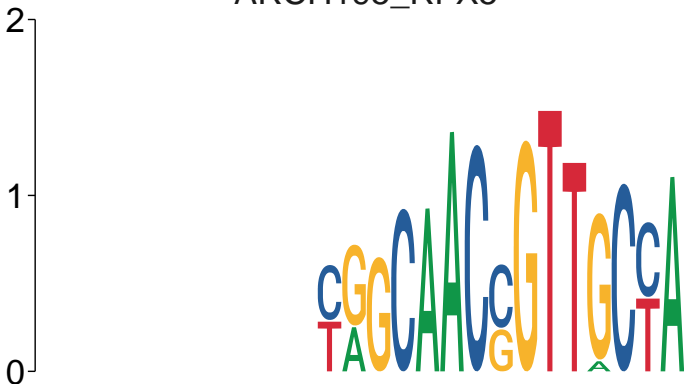
Tlx?



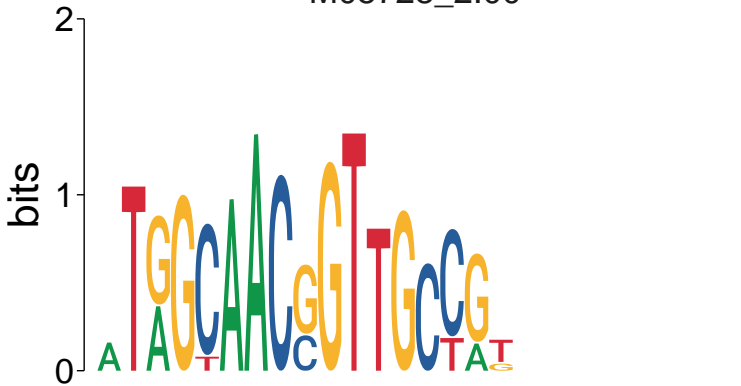
ARCH192_GAGA_repeat



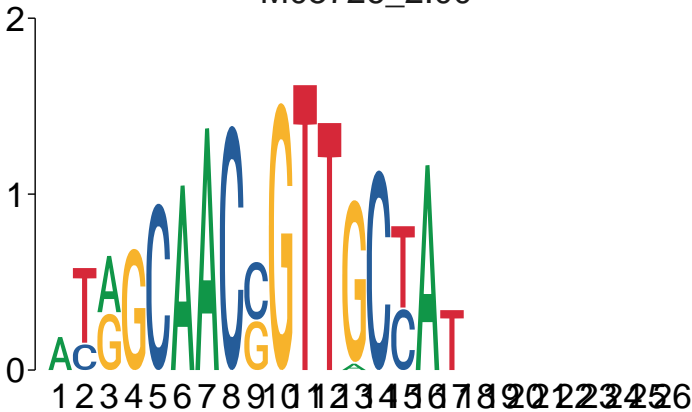
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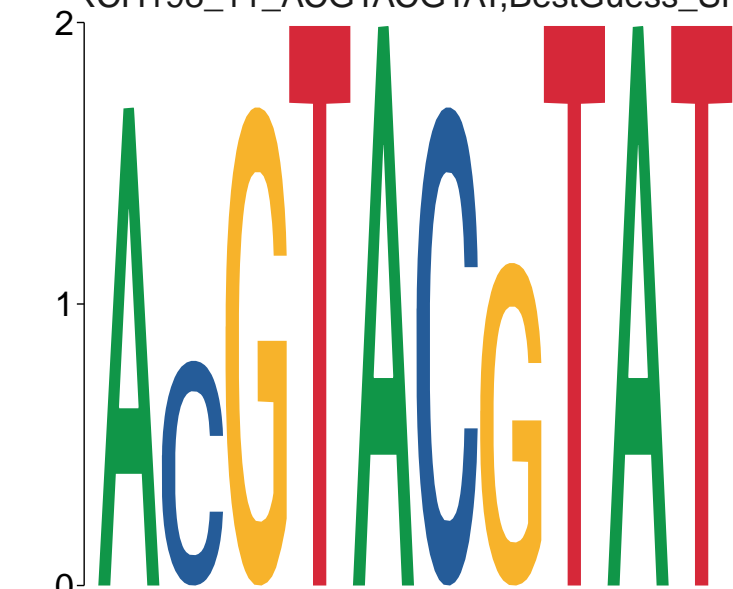
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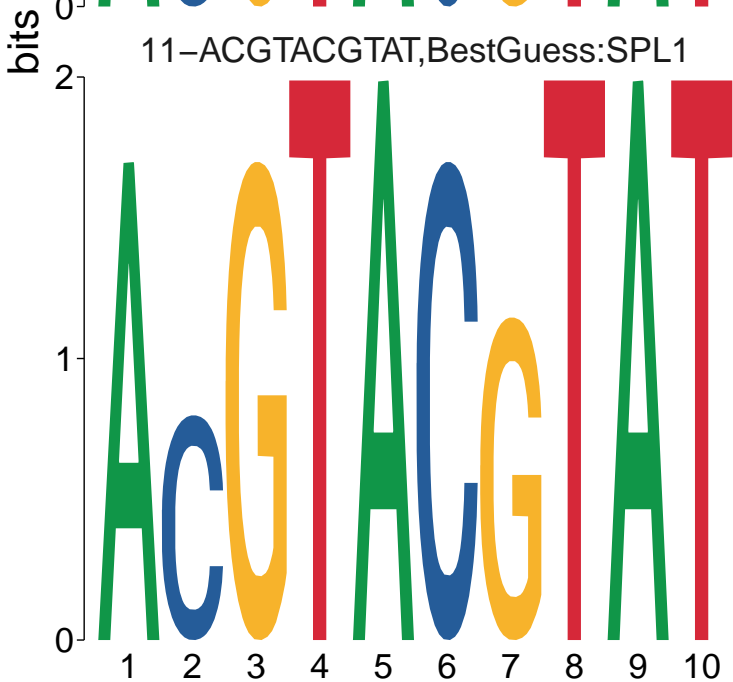
2



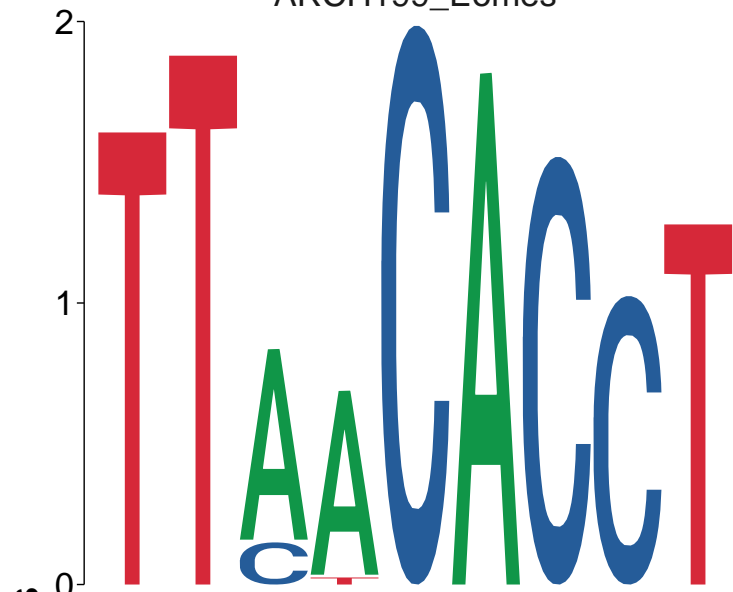
RCH198_11_ACGTACGTAT,BestGuess_SP



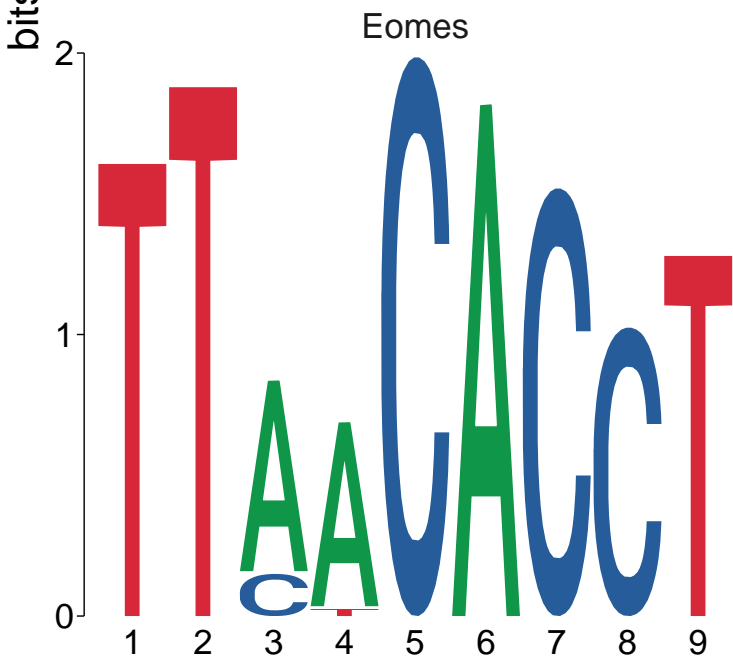
11-ACGTACGTAT,BestGuess:SPL1



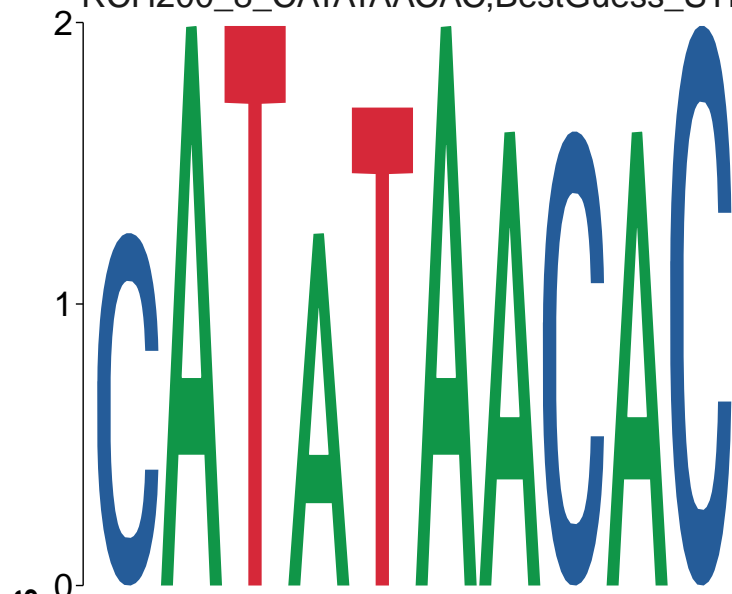
ARCH199_Eomes



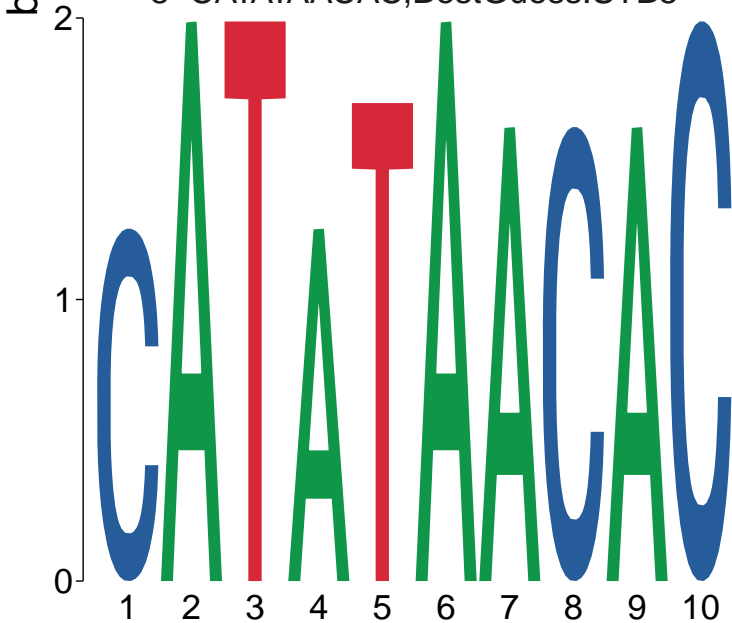
Eomes



RCH200_8_CATATAACAC,BestGuess_STB



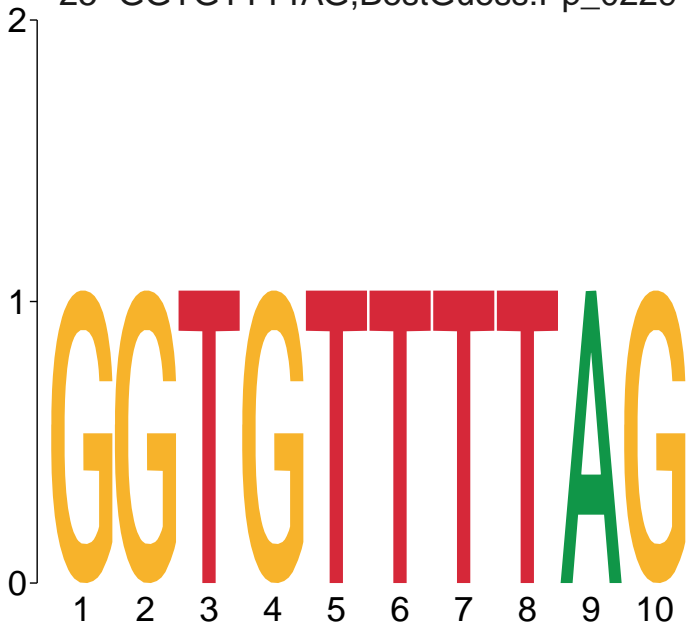
8-CATATAACAC,BestGuess:STB5



CH201_25_GGTGTTTTAG,BestGuess_Pp



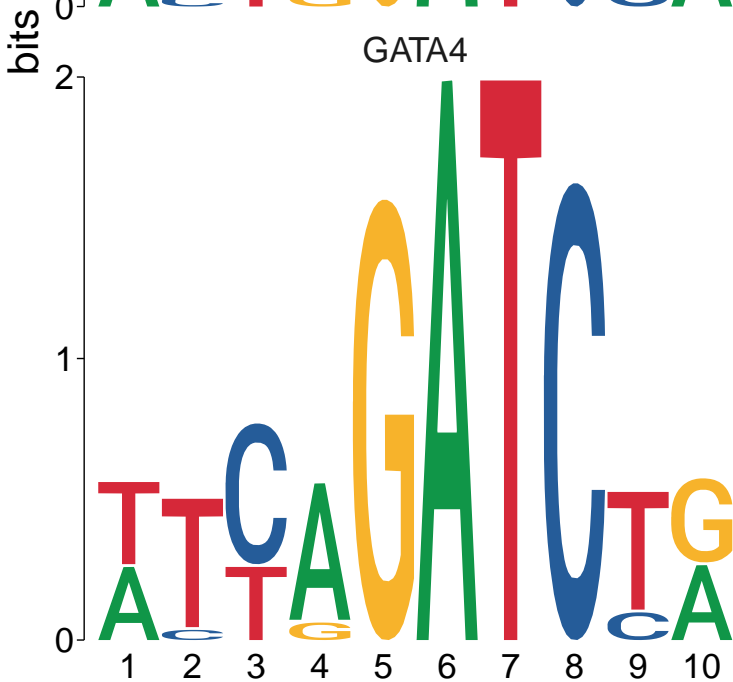
25-GGTGTTTTAG,BestGuess:Pp_0229



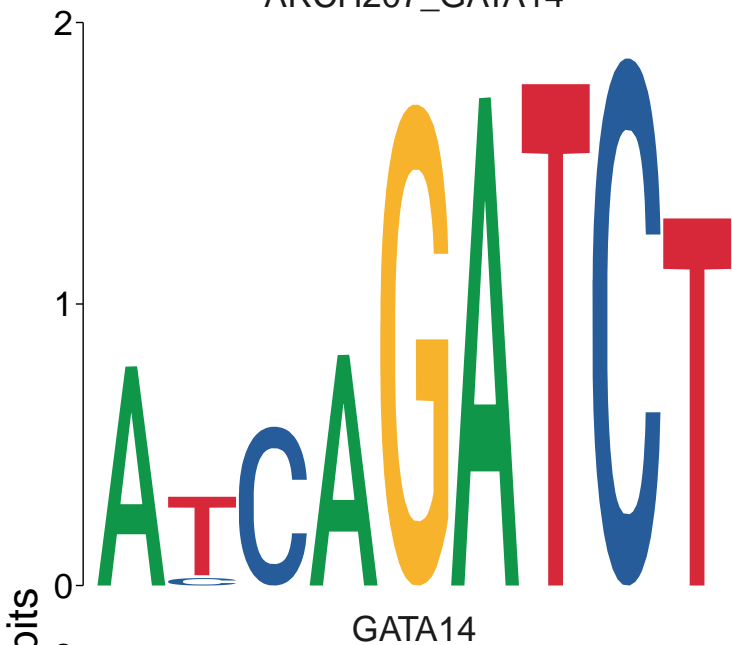
ARCH206_GATA4



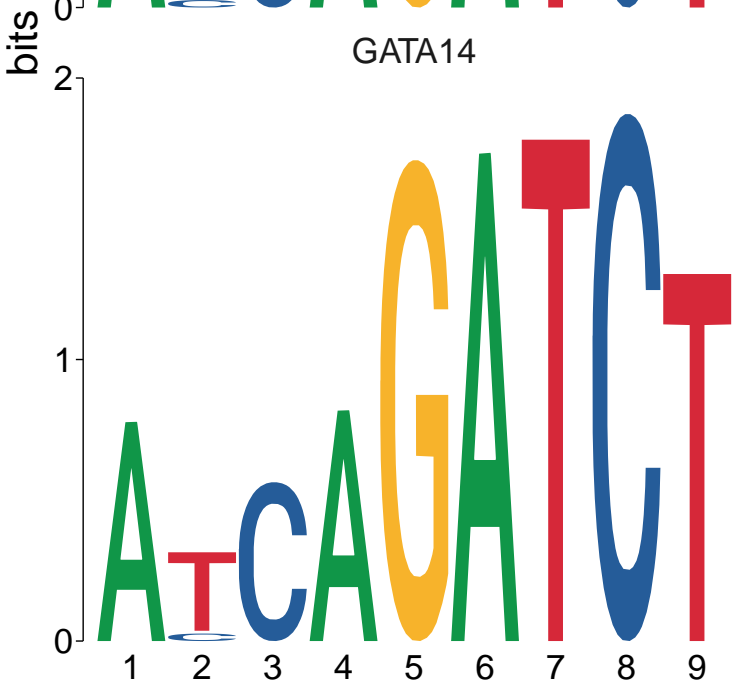
GATA4



ARCH207_GATA14



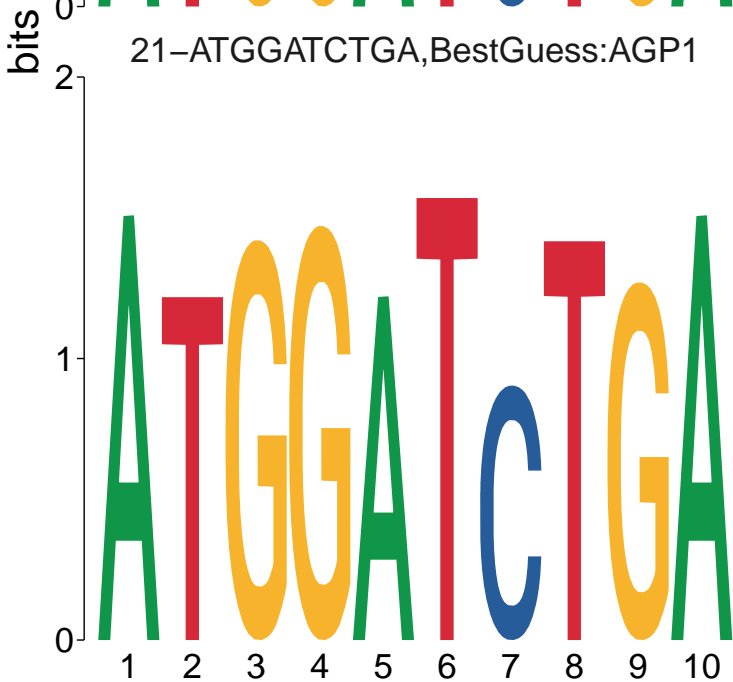
GATA14



CH208_21_ATGGATCTGA,BestGuess_AG



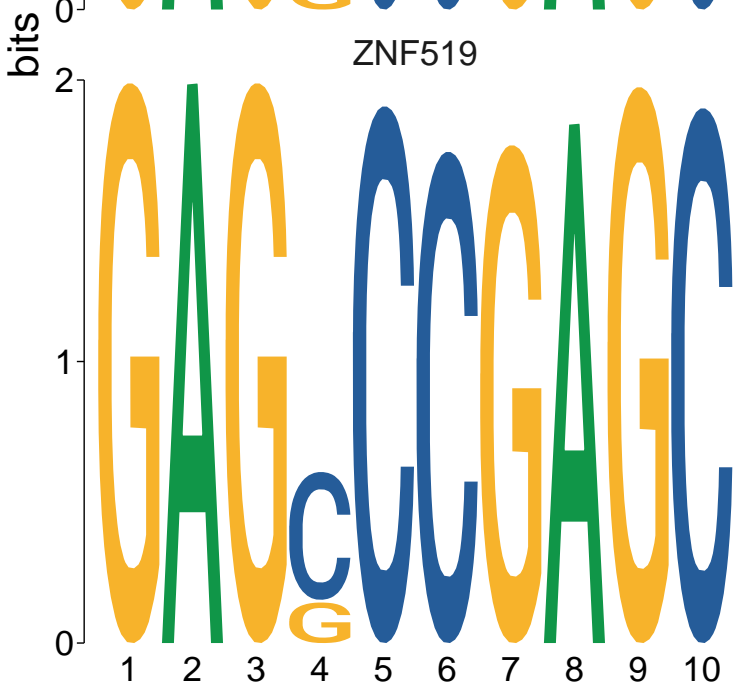
21-ATGGATCTGA,BestGuess:AGP1



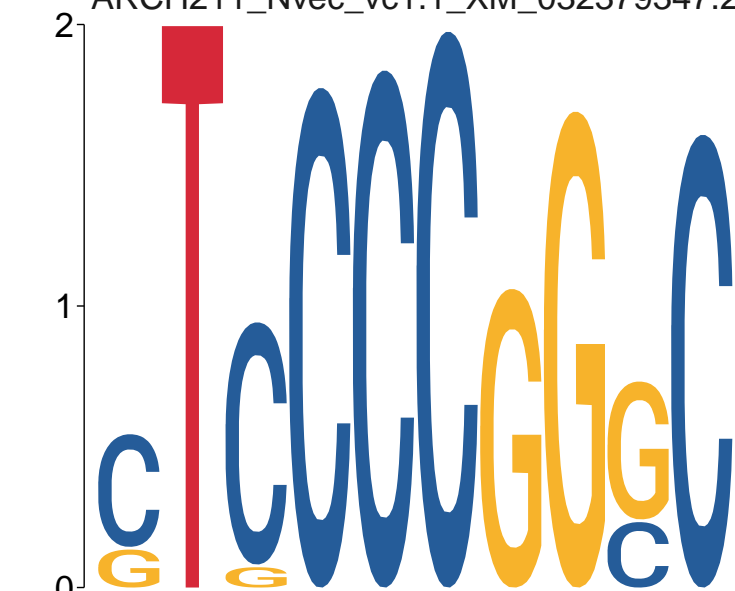
ARCH209_ZNF519



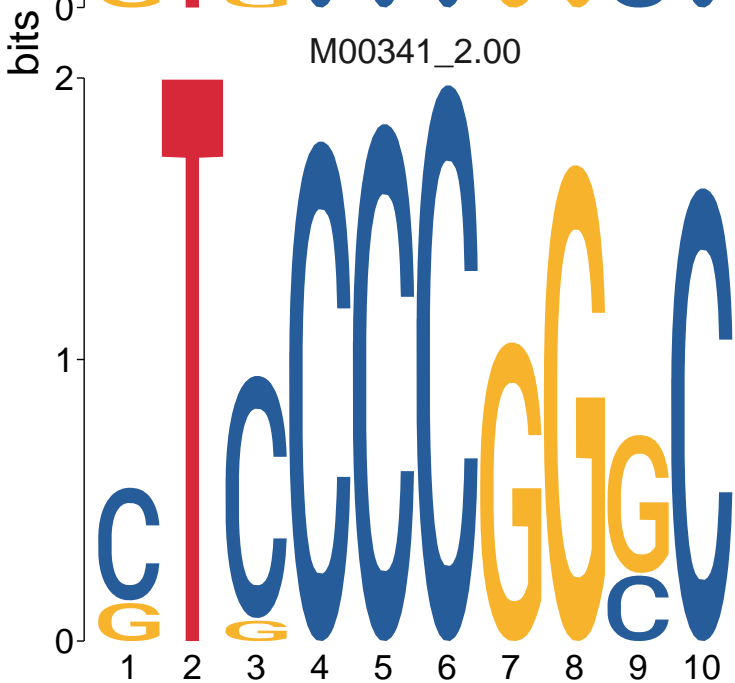
ZNF519



ARCH211_Nvec_vc1.1_XM_032379347.2



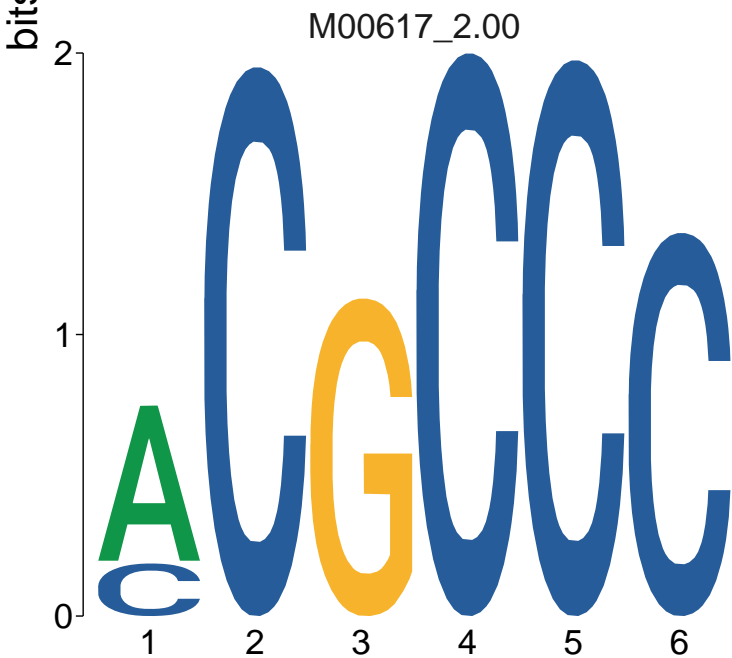
M00341_2.00



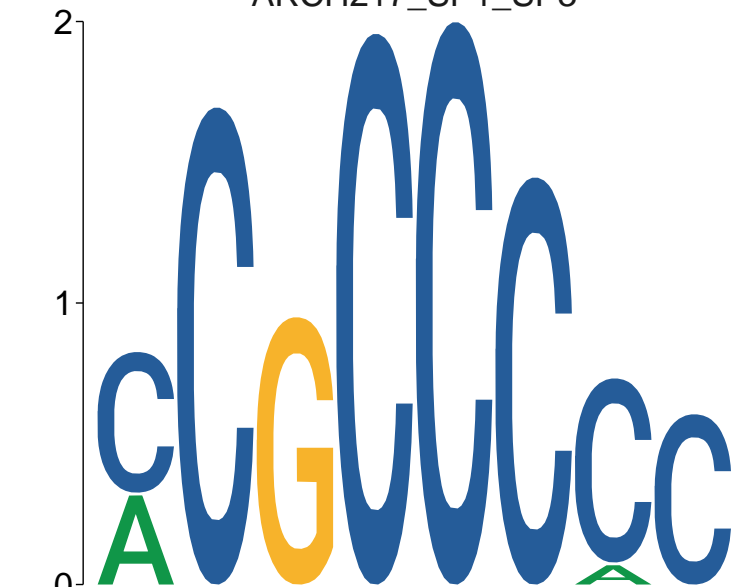
ARCH216_KLF6



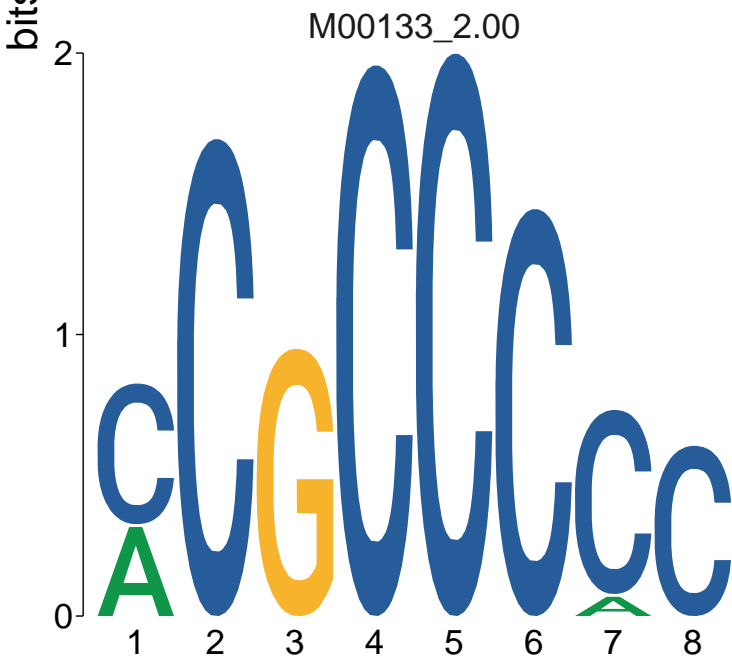
M00617_2.00



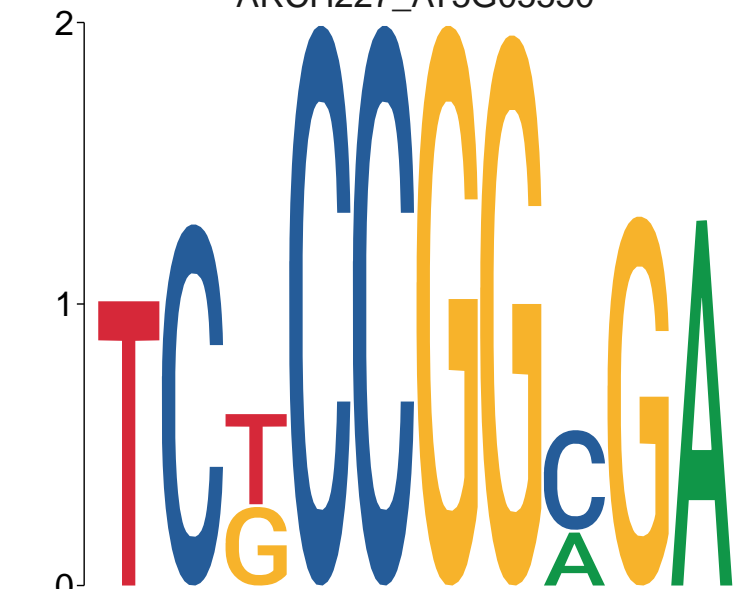
ARCH217_SP1_SP3



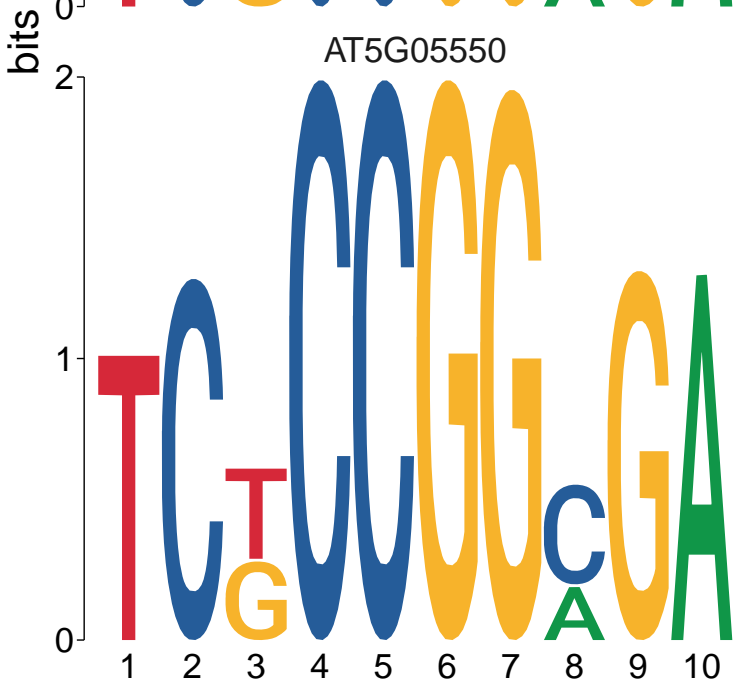
M00133_2.00



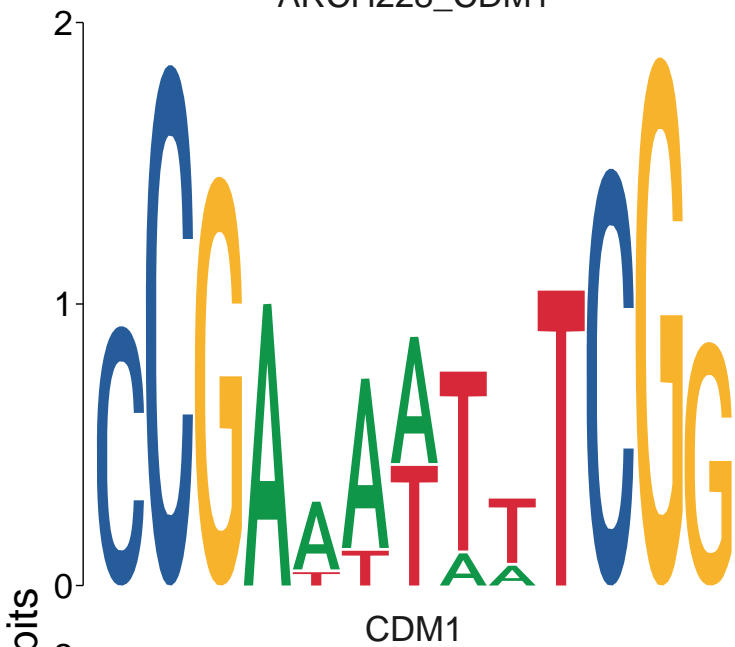
ARCH227_AT5G05550



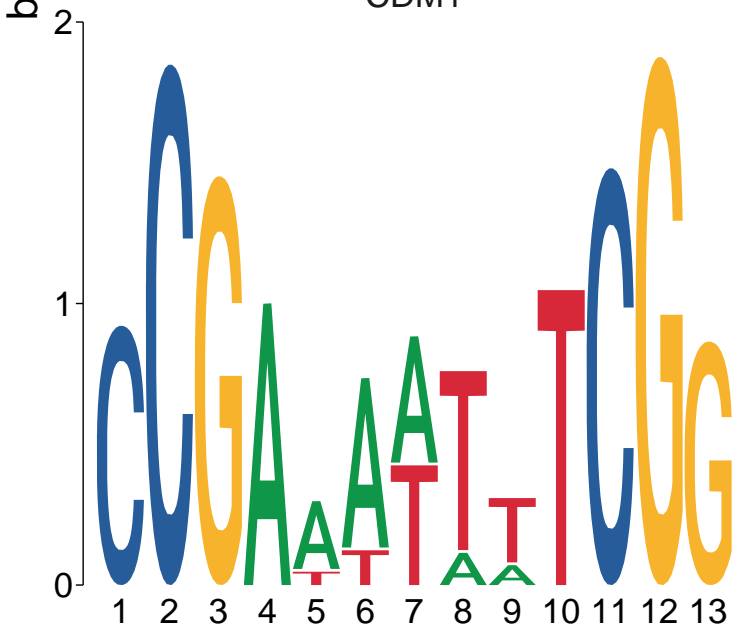
AT5G05550



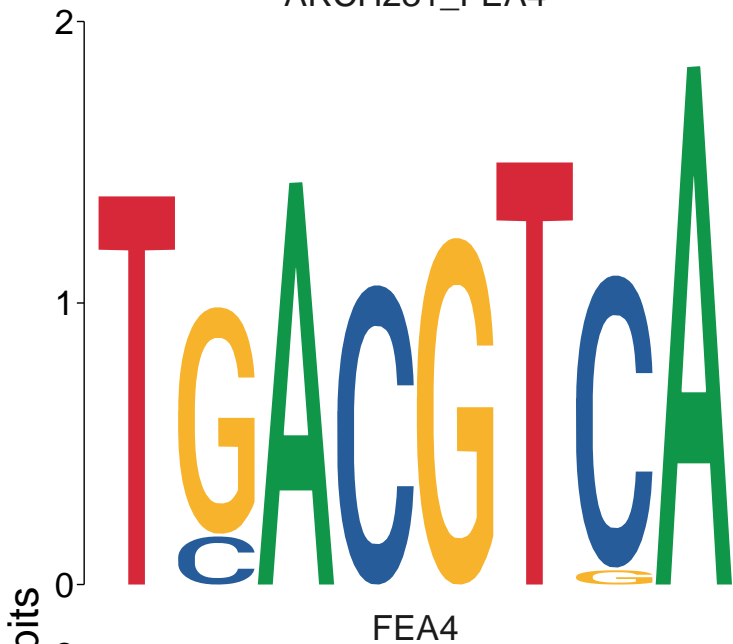
ARCH228_CDM1



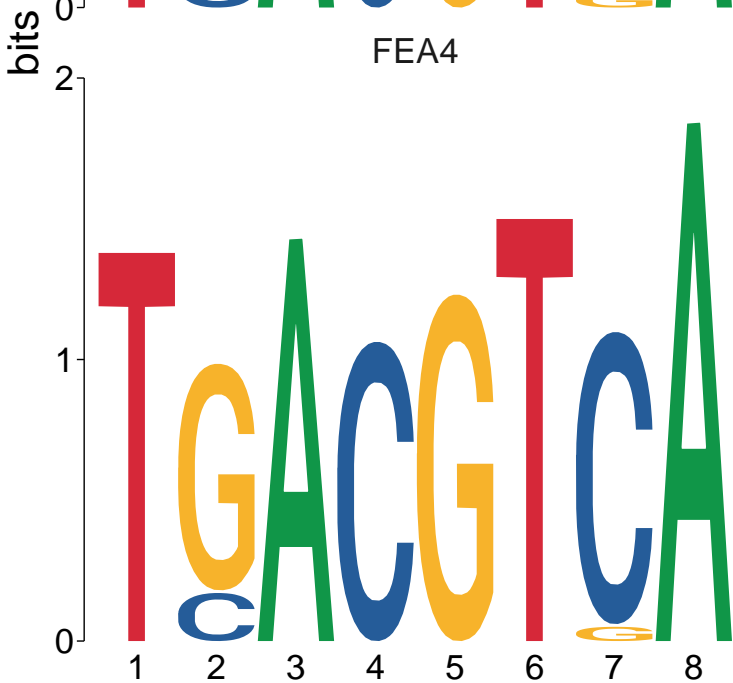
CDM1



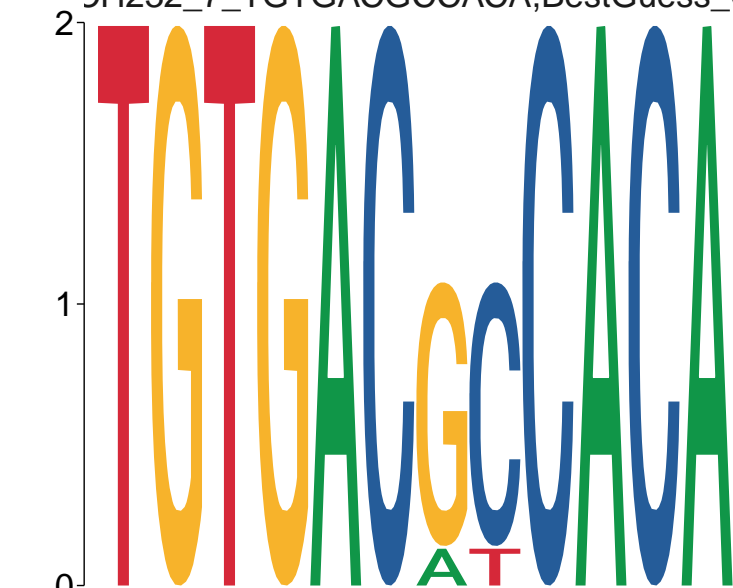
ARCH231_FEA4



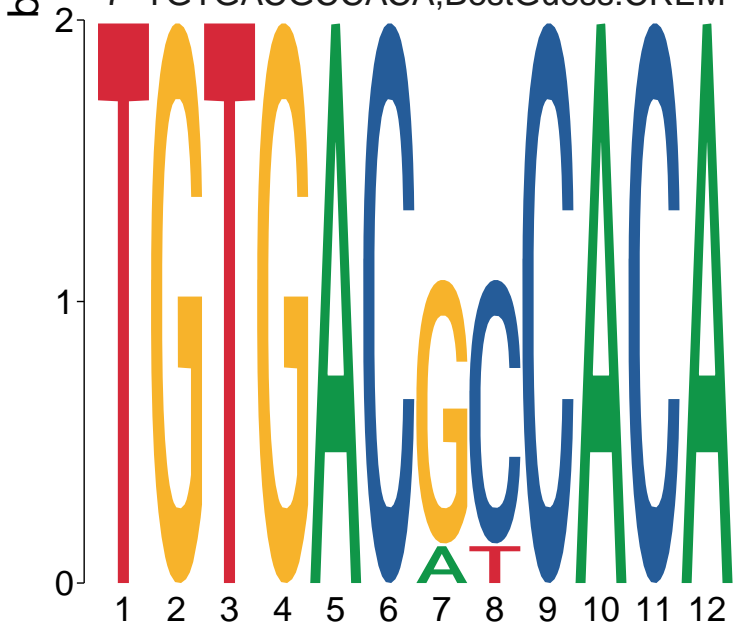
FEA4



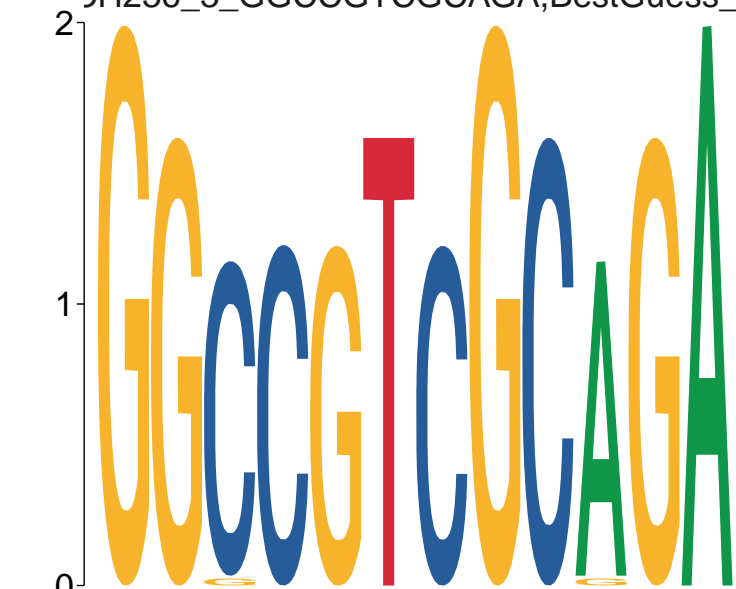
CH232_7_TGTGACGCCACA,BestGuess_C



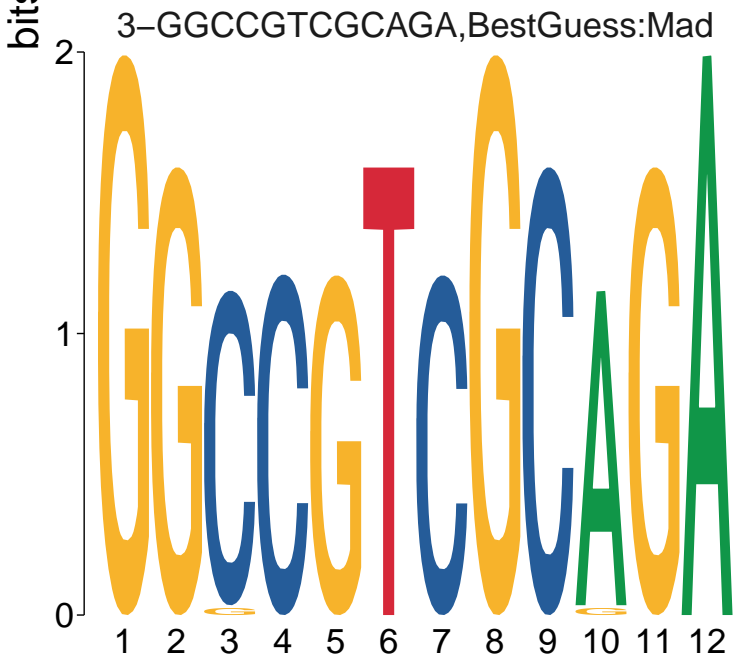
7-TGTGACGCCACA,BestGuess:CREM



CH236_3_GGCCGTCGCAGA,BestGuess_1



3-GGCCGTCGCAGA,BestGuess:Mad



ARCH238_ANTP_HOXA2_ANTP_HOXA3

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M03874_2.00

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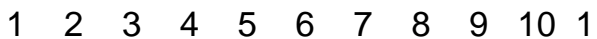
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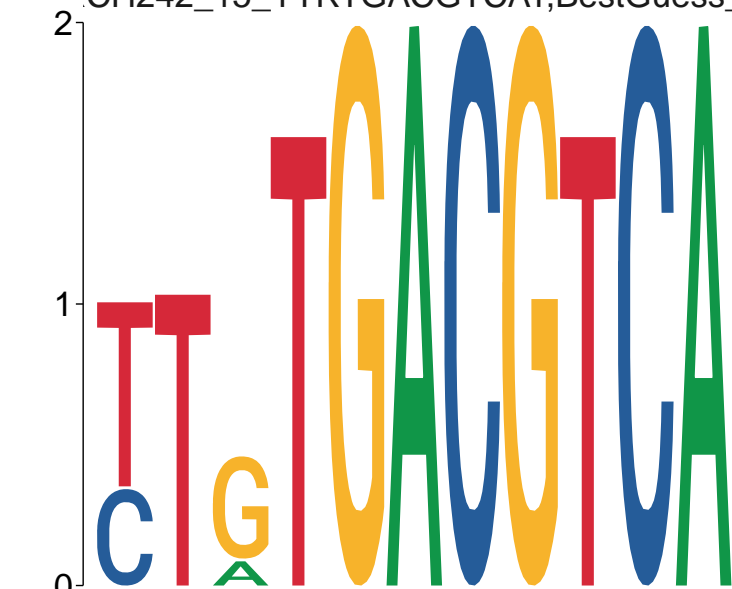
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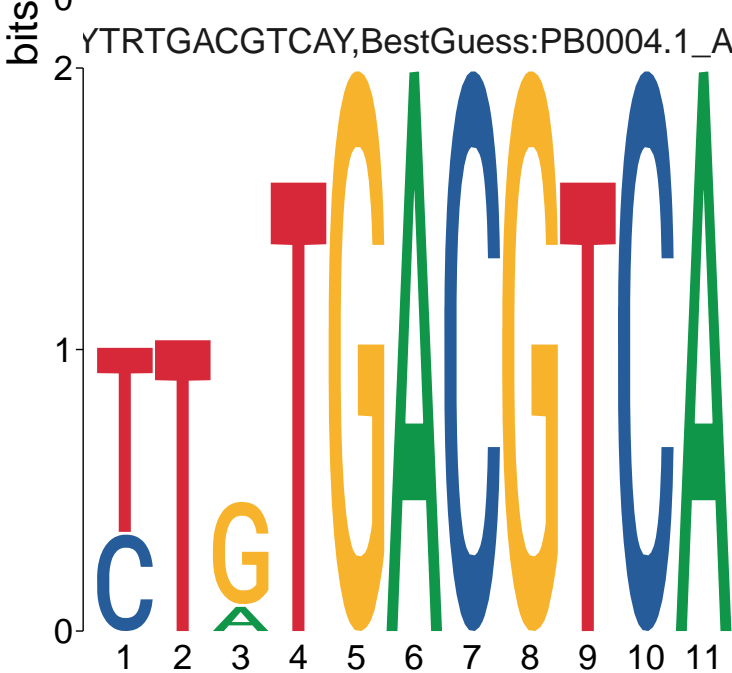
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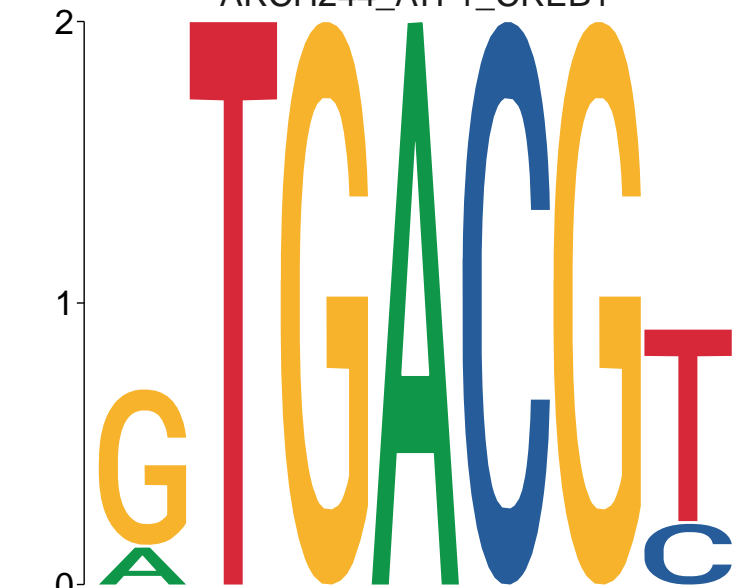
CH242_15_YTRTGACGTCAY,BestGuess_



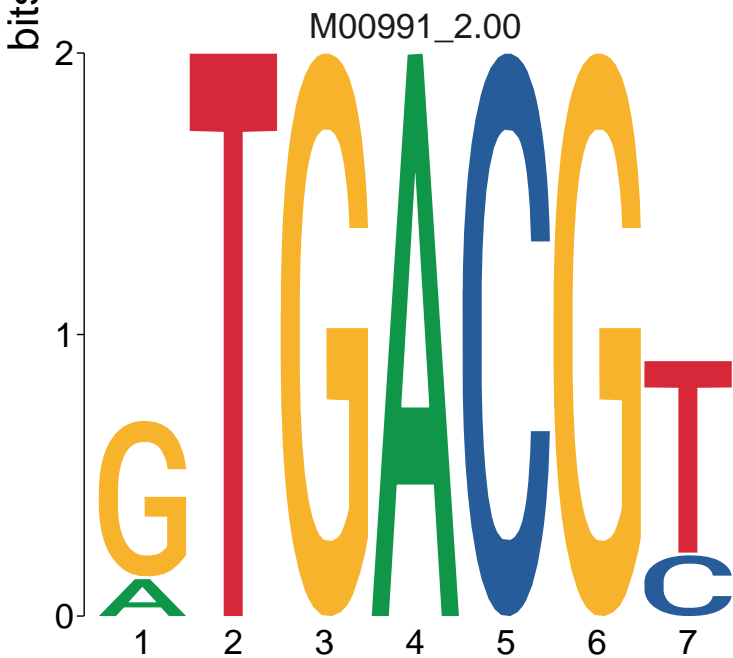
YTRTGACGTCAY,BestGuess:PB0004.1_At



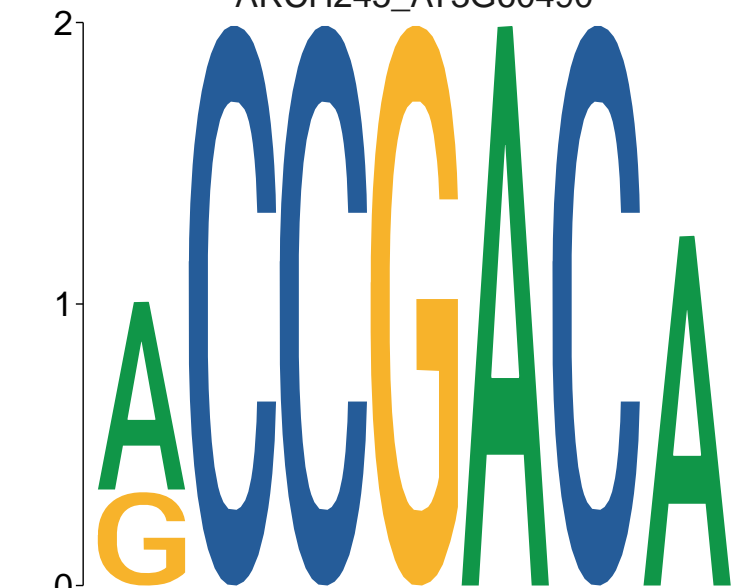
ARCH244_ATF1_CREB1



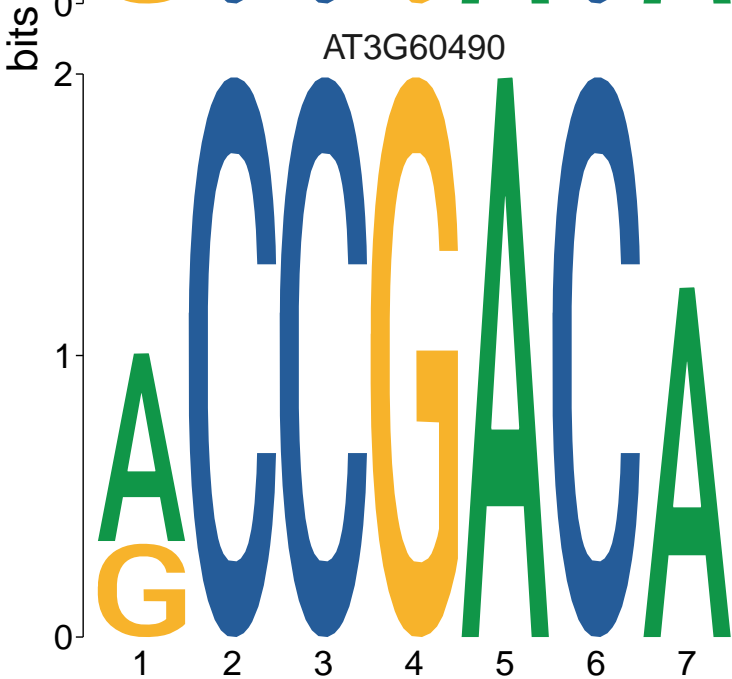
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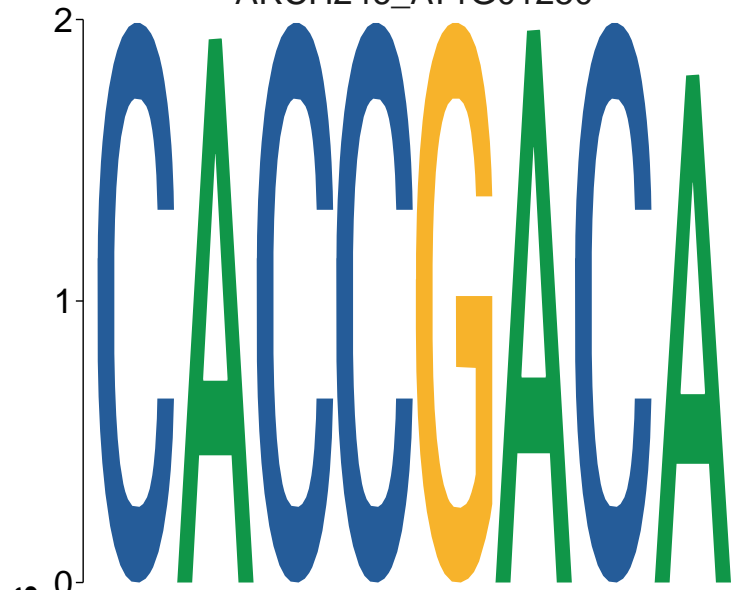
ARCH245_AT3G60490



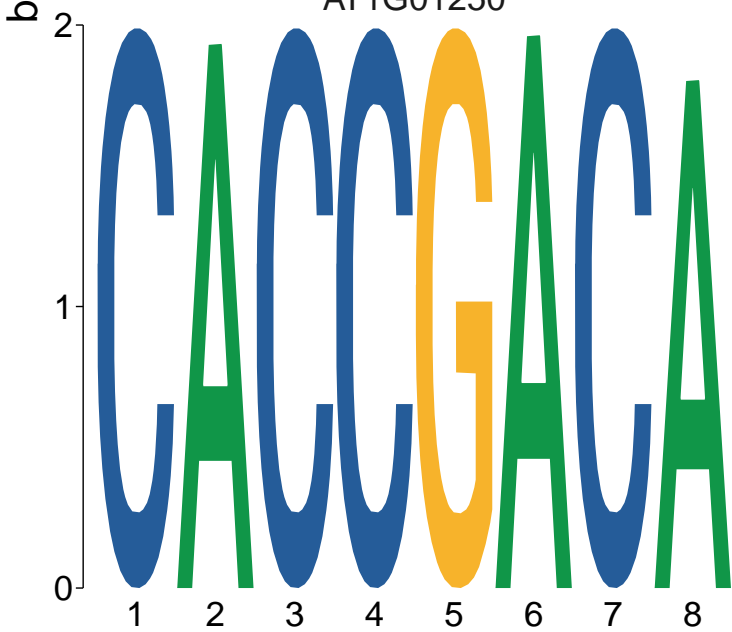
AT3G60490



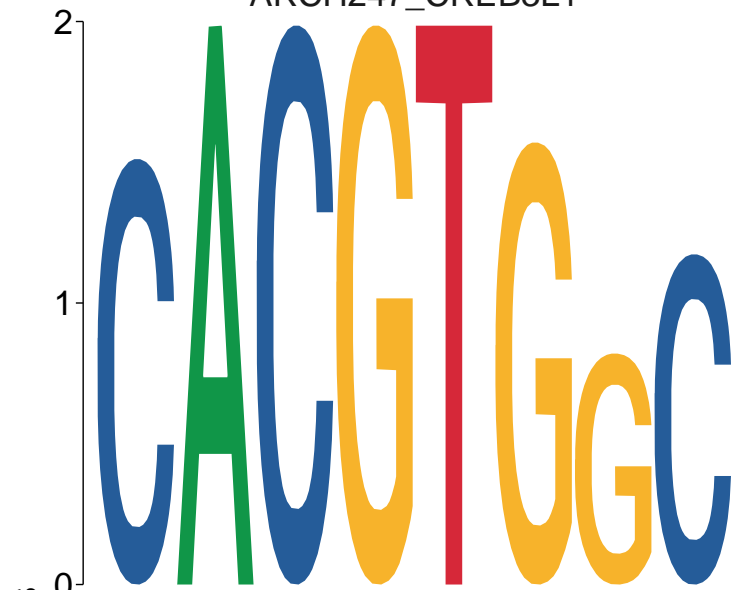
ARCH246_AT1G01250



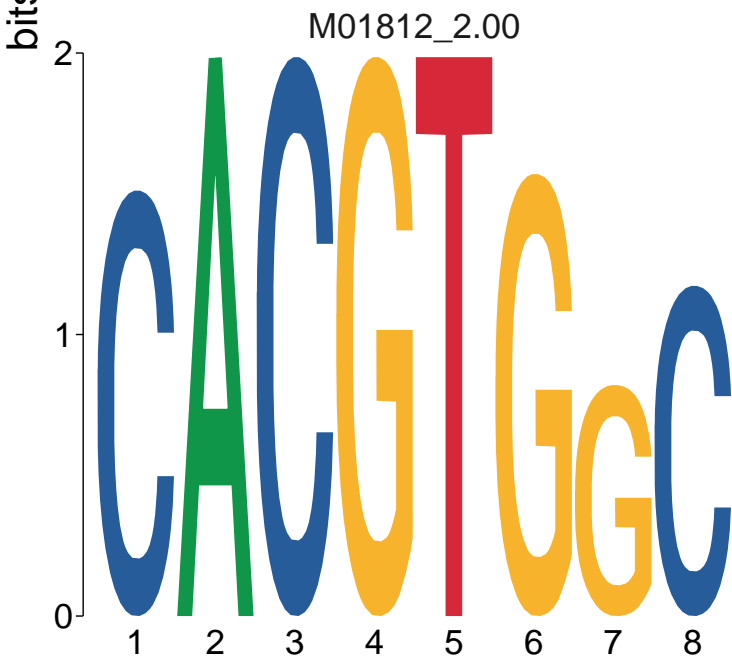
AT1G01250



ARCH247_CREB3L1



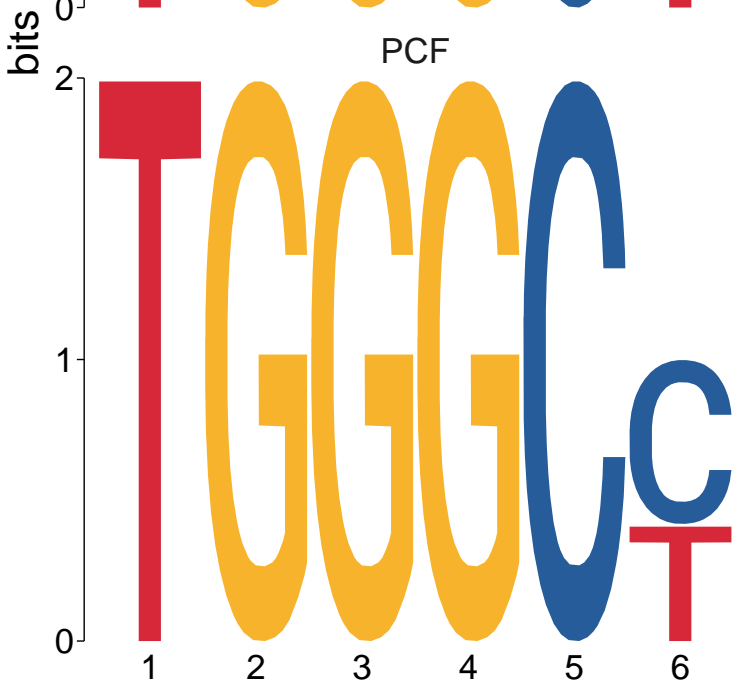
M01812_2.00



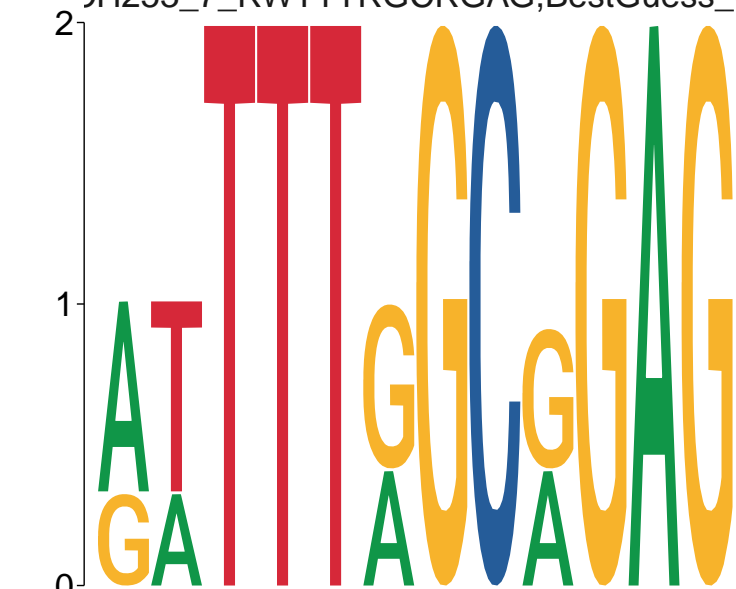
ARCH250_PCF



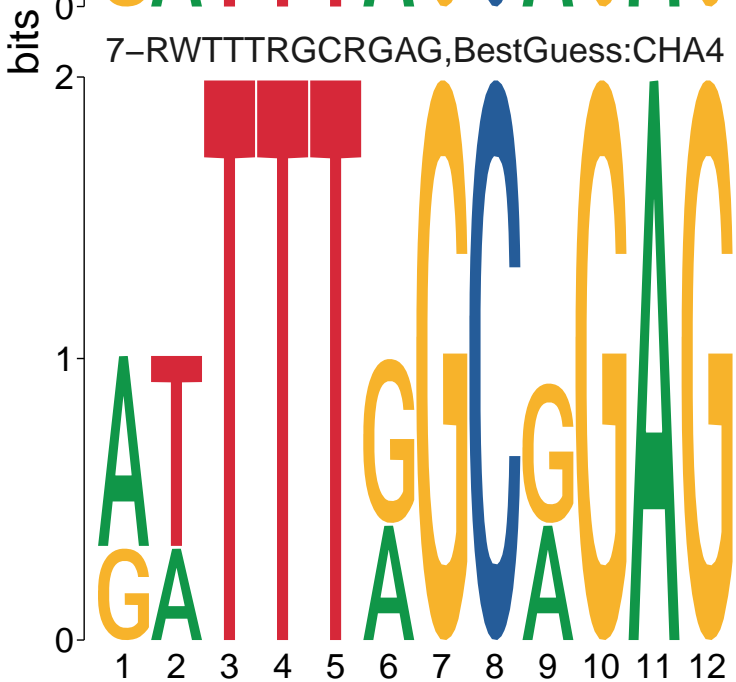
PCF



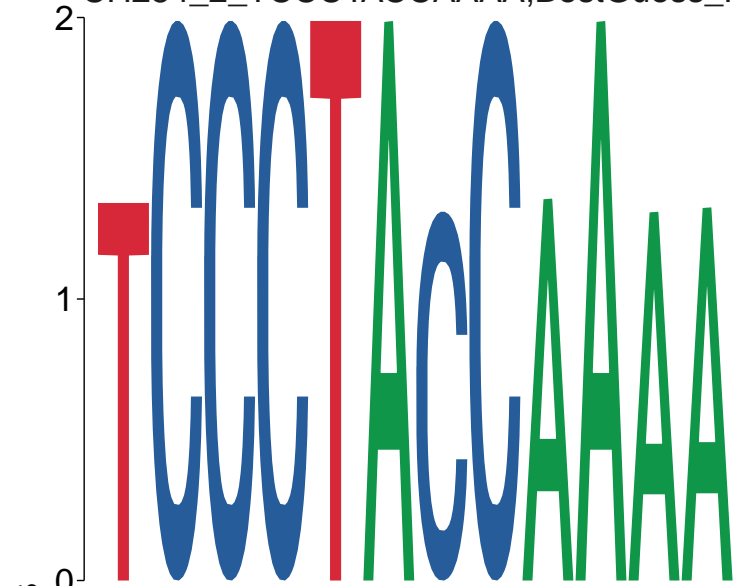
CH253_7_RWTTTTRGCRGAG,BestGuess_C



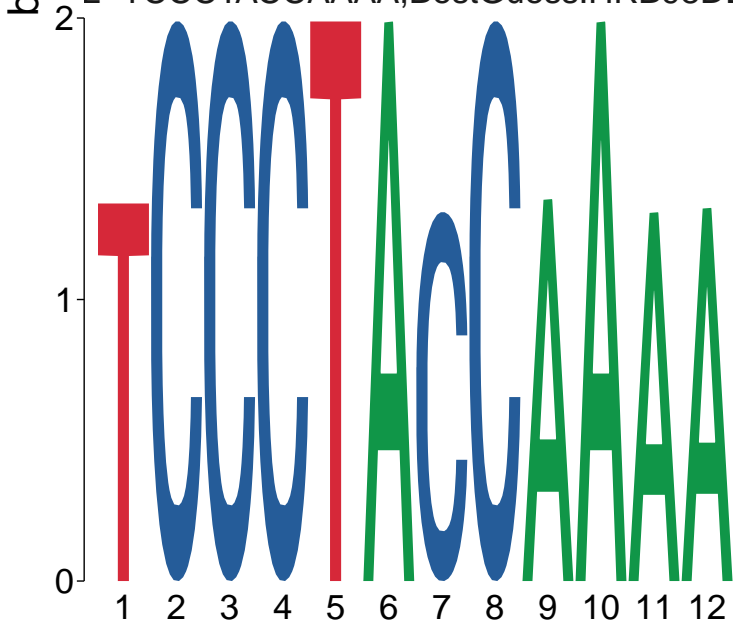
7-RWTTTTRGCRGAG,BestGuess:CHA4



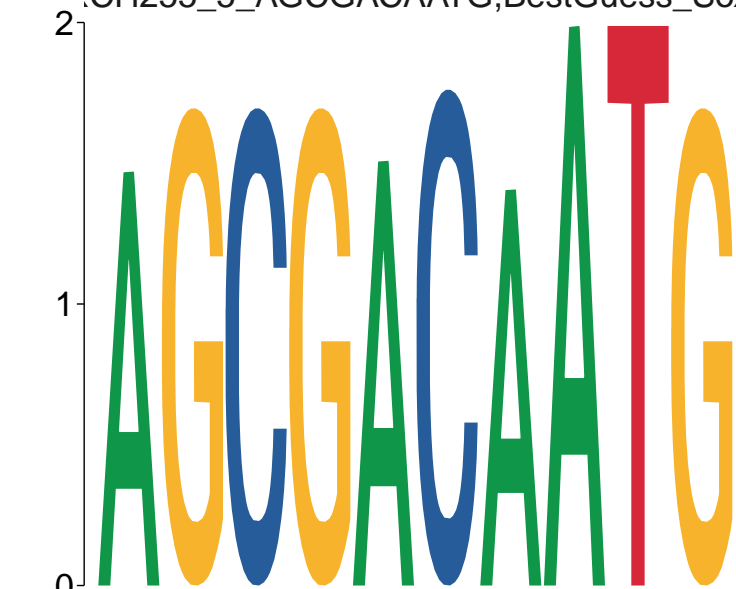
CH254_2_TCCCTACCAAAA,BestGuess_H



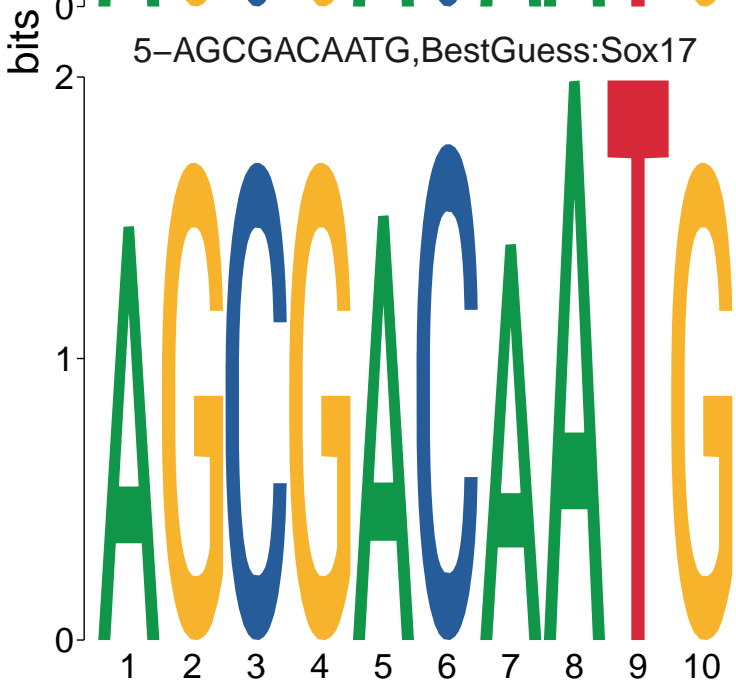
2-TCCCTACCAAAA,BestGuess:HRB98DE



CH255_5_AGCGACAATG,BestGuess_Sox



5-AGCGACAATG,BestGuess:Sox17



CH256_6_TMKCGCCATW,BestGuess_E2

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6-TMKCGCCATW,BestGuess:E2FA

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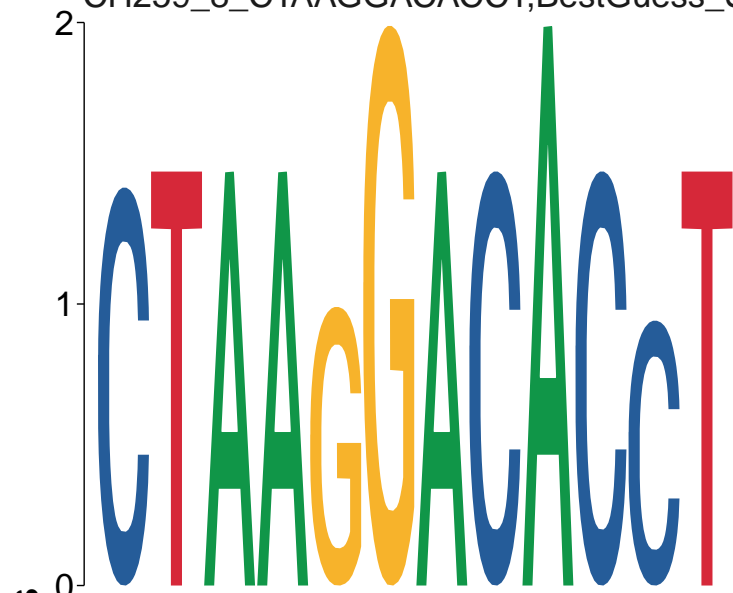
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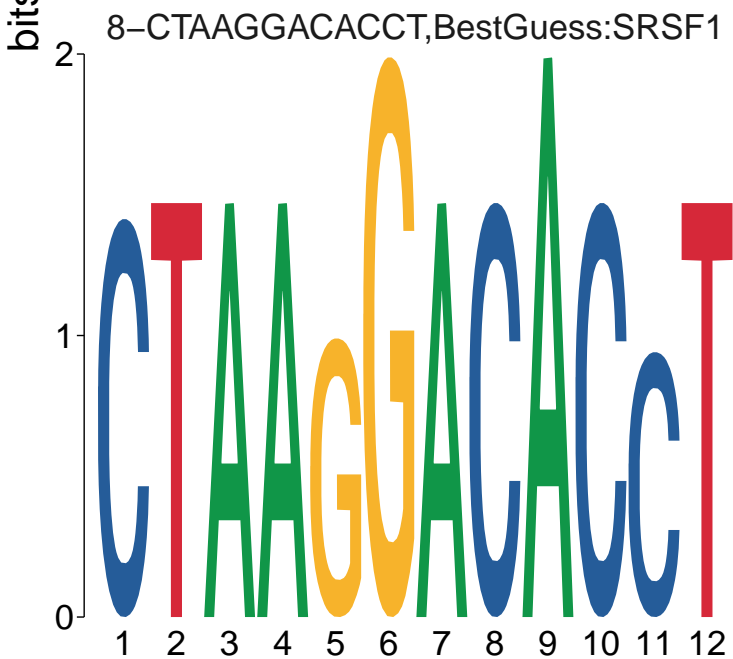


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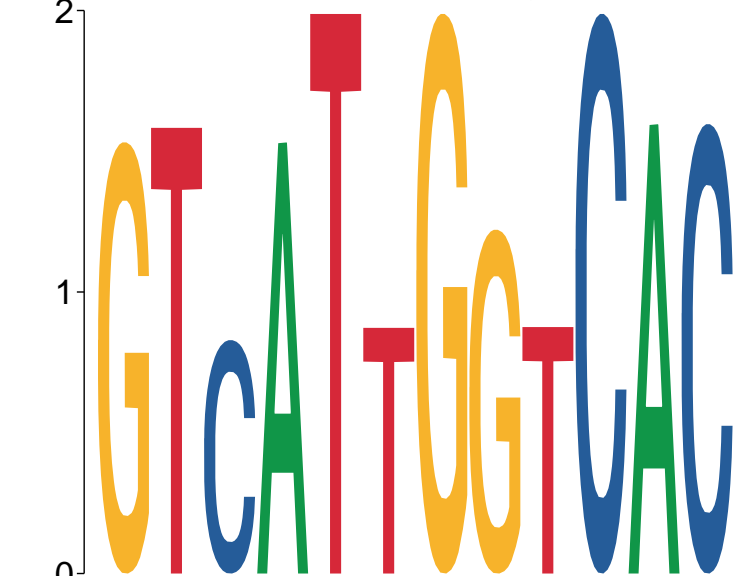
CH259_8_CTAAGGACACCT,BestGuess_S



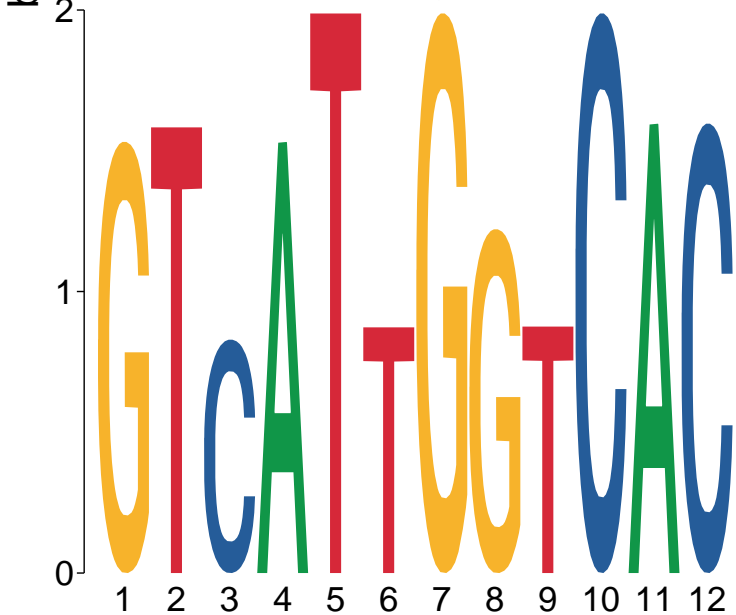
8-CTAAGGACACCT,BestGuess:SRSF1



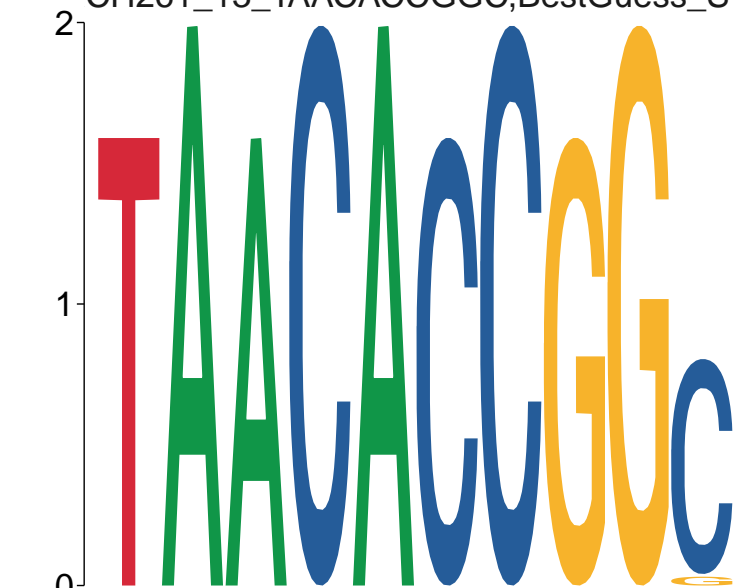
CH260_1_GTCATTGGTCAC,BestGuess_R



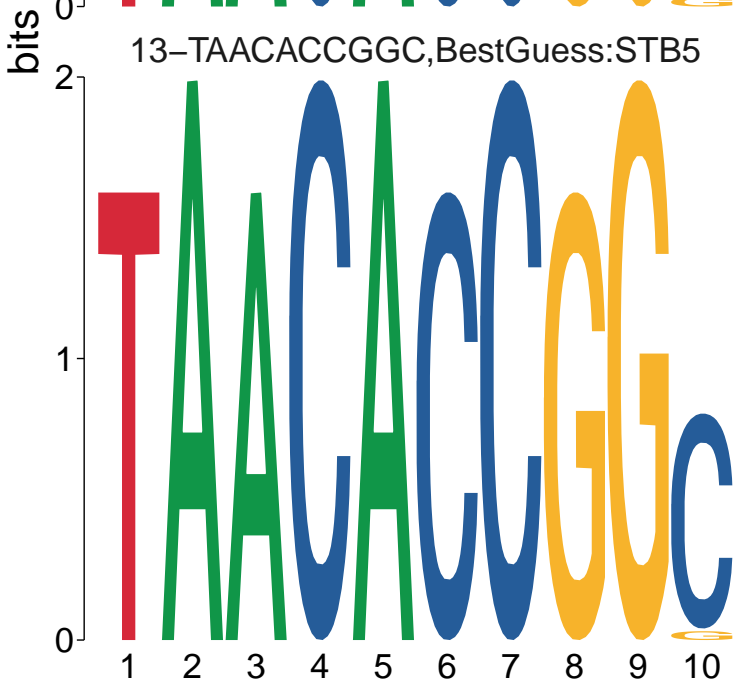
1-GTCATTGGTCAC,BestGuess:RTG3



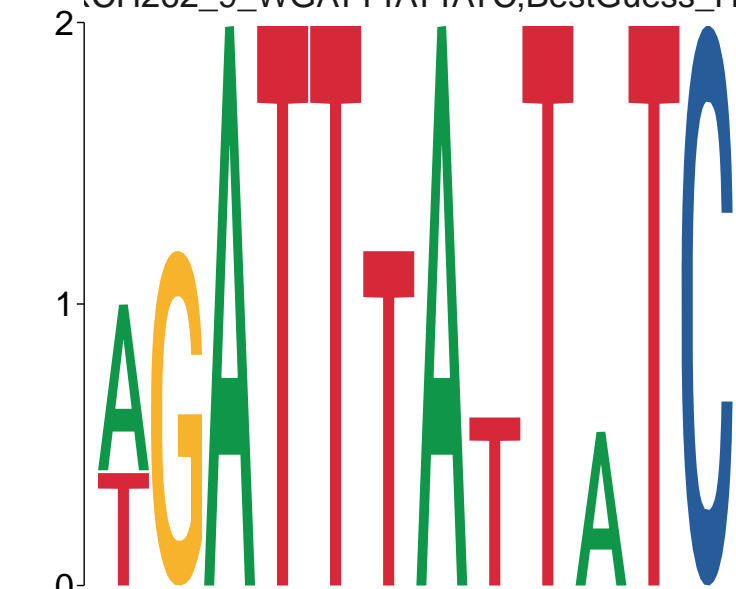
CH261_13_TAACACCGGC,BestGuess_ST



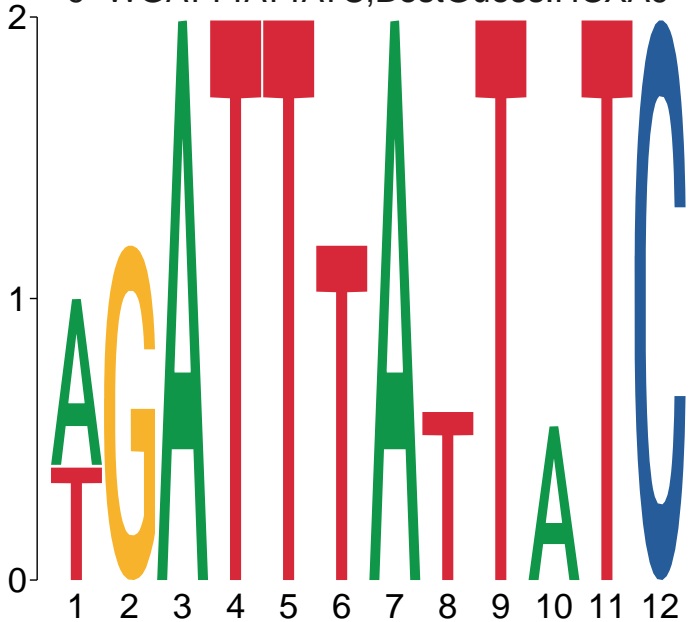
13-TAACACCGGC,BestGuess:STB5



CH262_9_WGATTATTATC,BestGuess_HOXA9



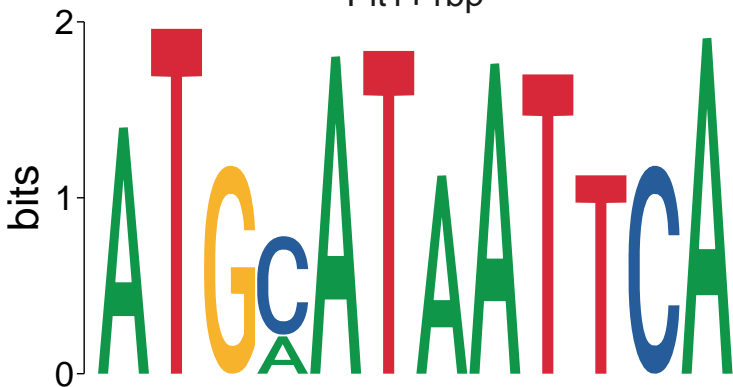
9-WGATTATTATC,BestGuess:HOXA9



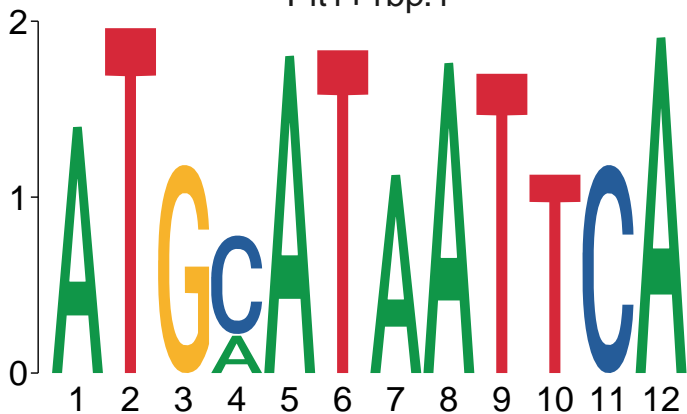
ARCH263_Pit1+1bp_Pit1+1bp



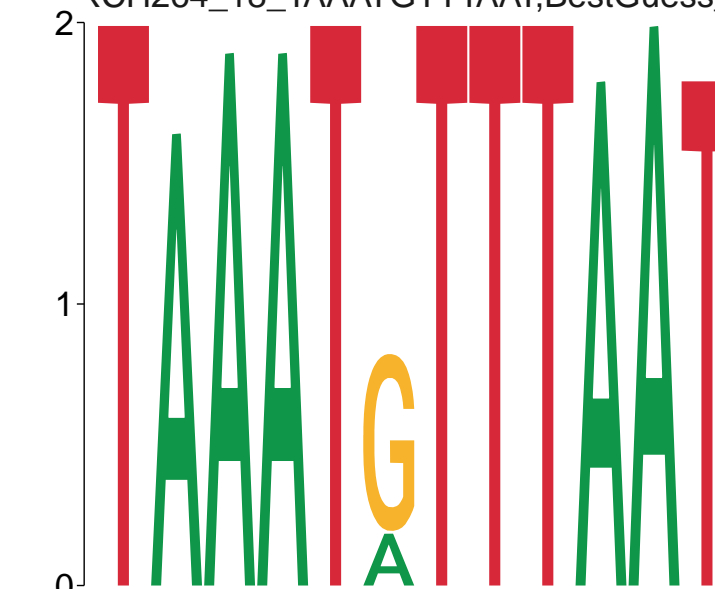
Pit1+1bp



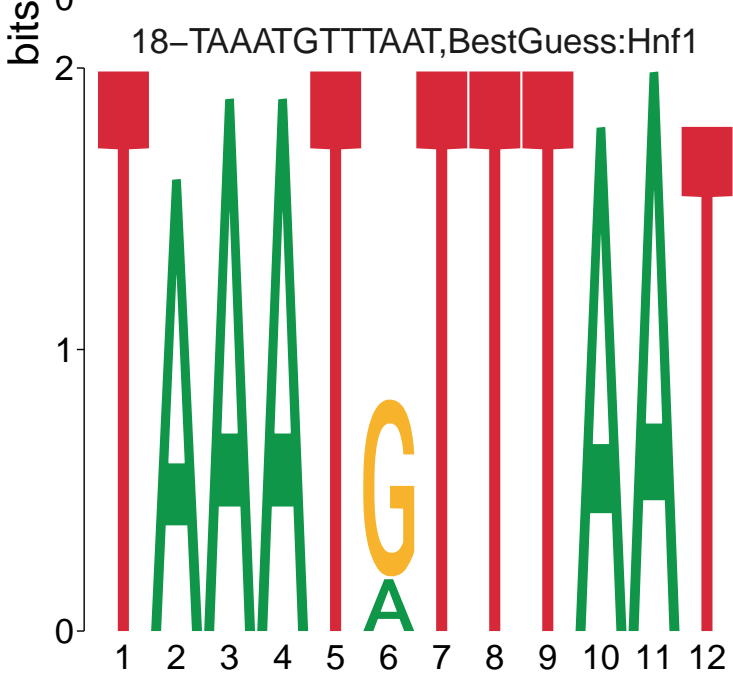
Pit1+1bp.1



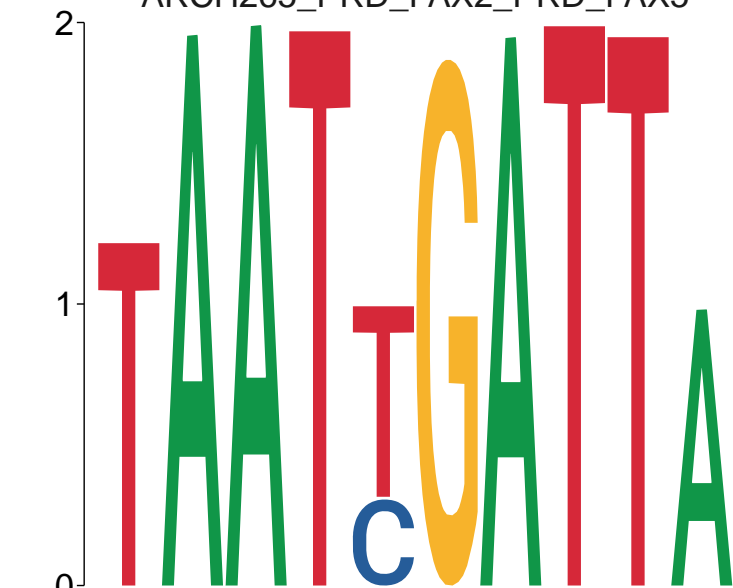
RCH264_18_TAAATGTTTAAT,BestGuess_1



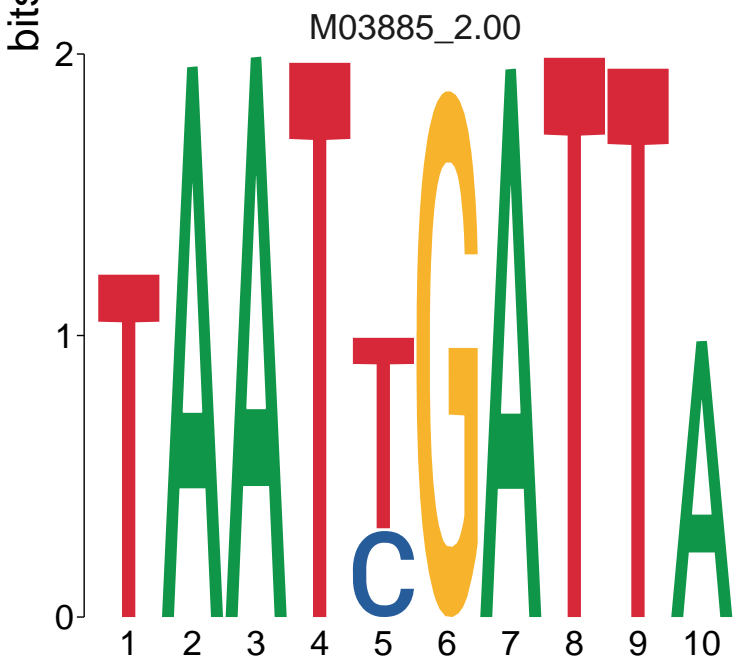
18-TAAATGTTTAAT,BestGuess:Hnf1



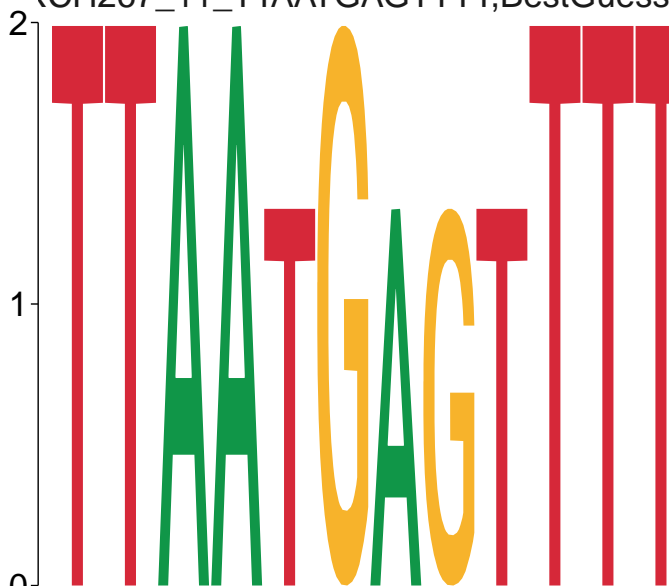
ARCH265_PRD_PAX2_PRD_PAX5



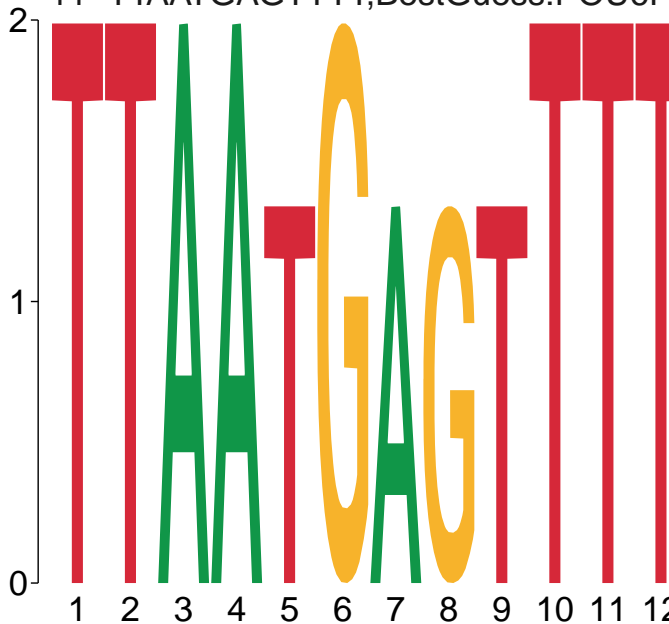
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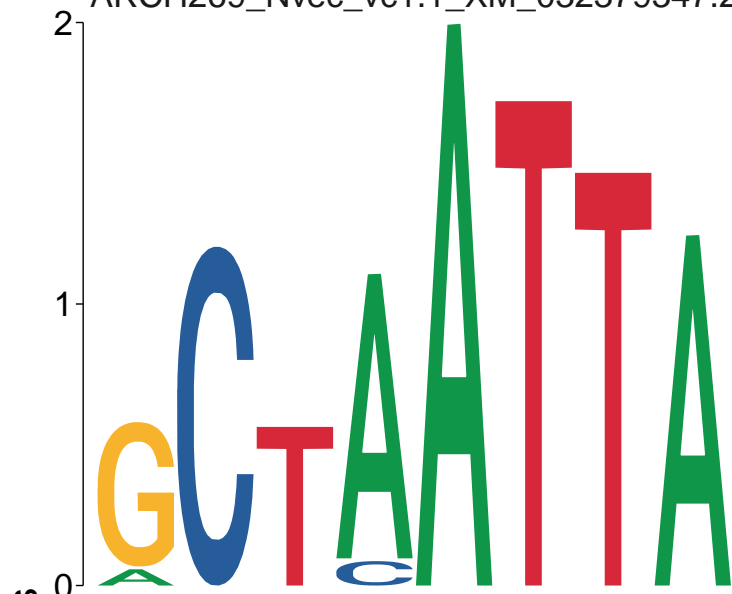
2CH267_11_TTAATGAGTTTT, BestGuess_F



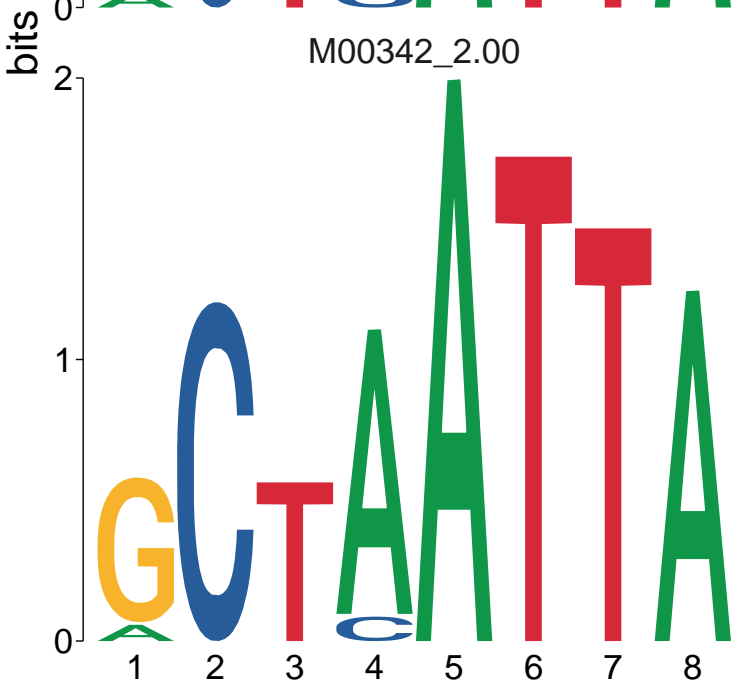
11-TTAATGAGTTTT, BestGuess:POU6F2



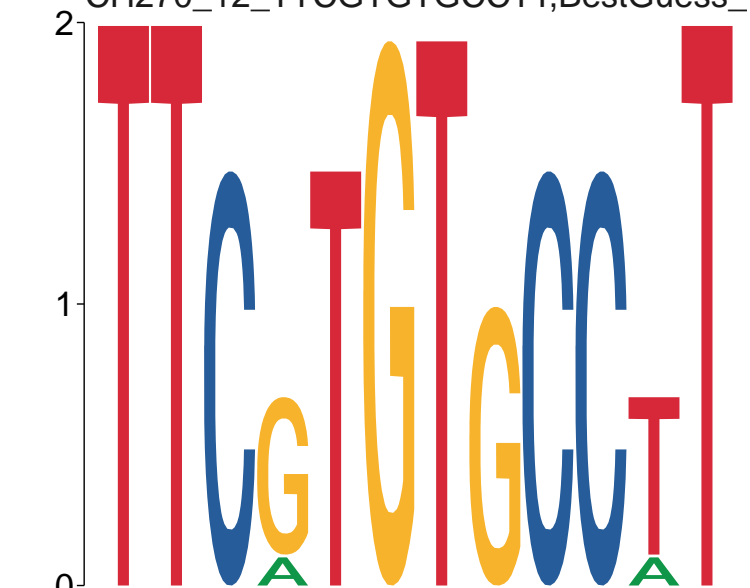
ARCH269_Nvec_vc1.1_XM_032379347.2



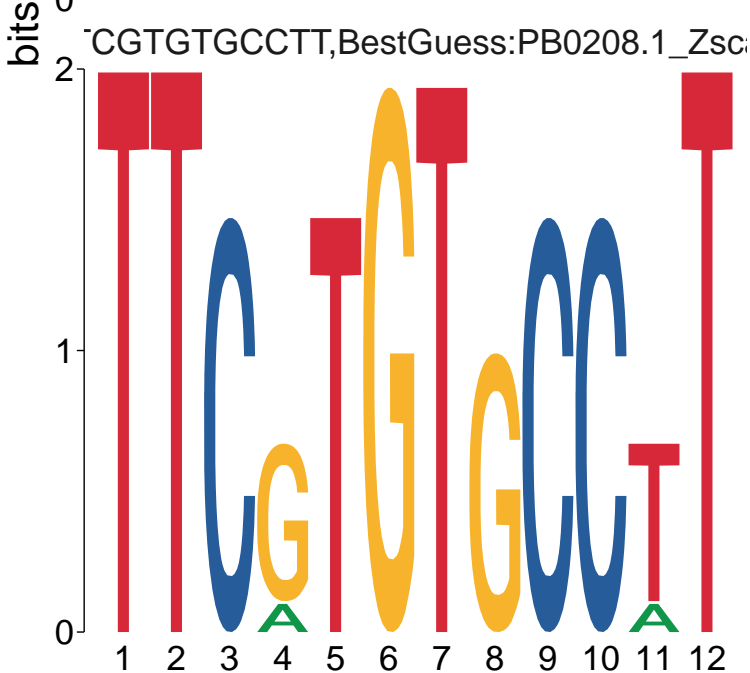
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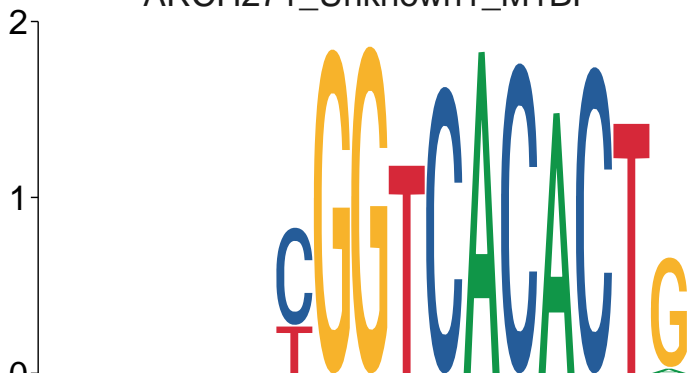
CH270_12_TTCGTGTGCCTT,BestGuess_



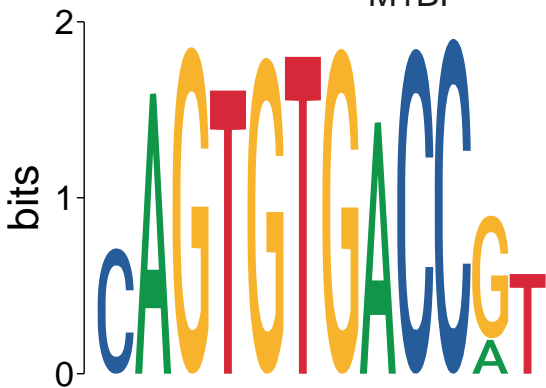
CGTGTGTGCCTT,BestGuess:PB0208.1_Zsc



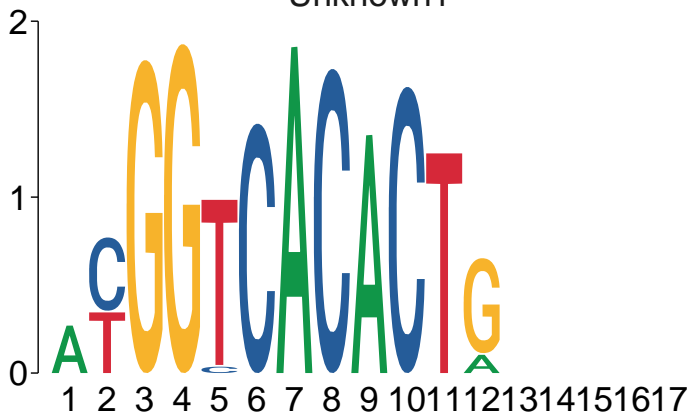
ARCH271_Unknown1_M1BP



M1BP

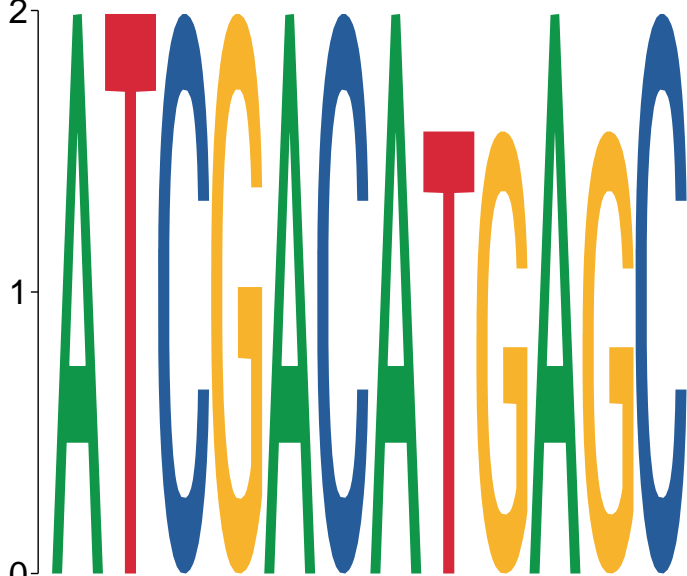


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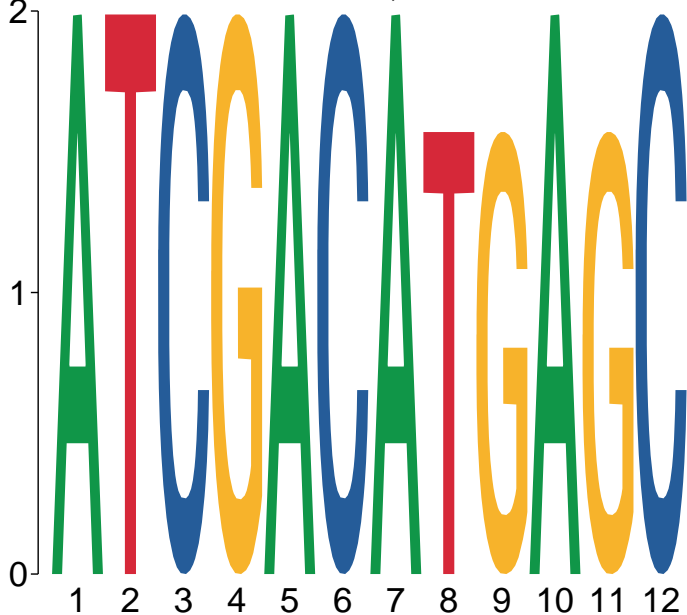


CH272_5_ATCGACATGAGC,BestGuess_D

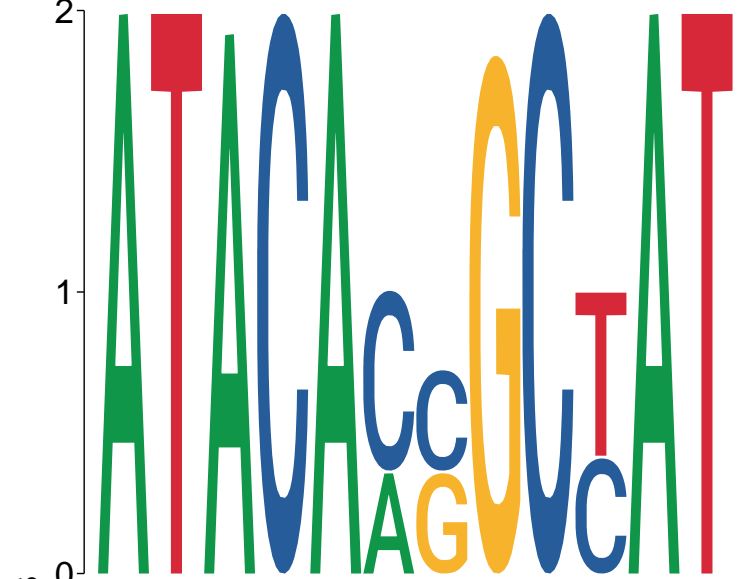
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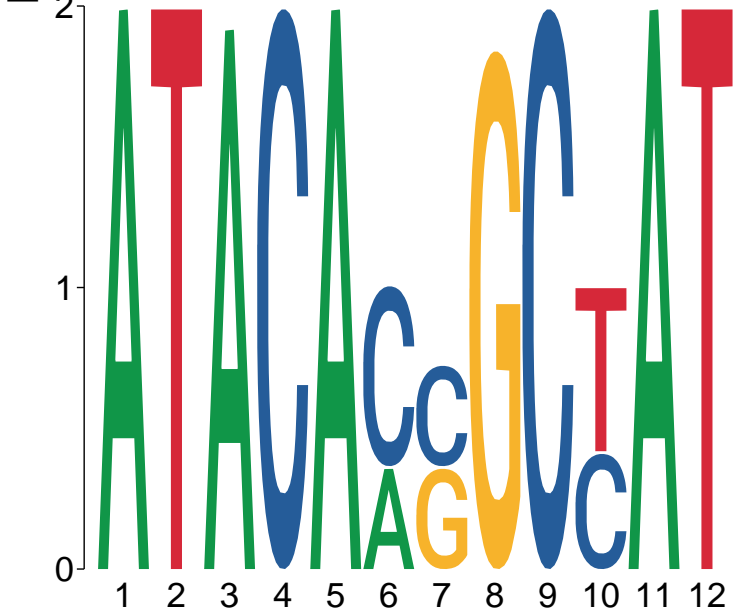
5-ATCGACATGAGC,BestGuess:DDF2



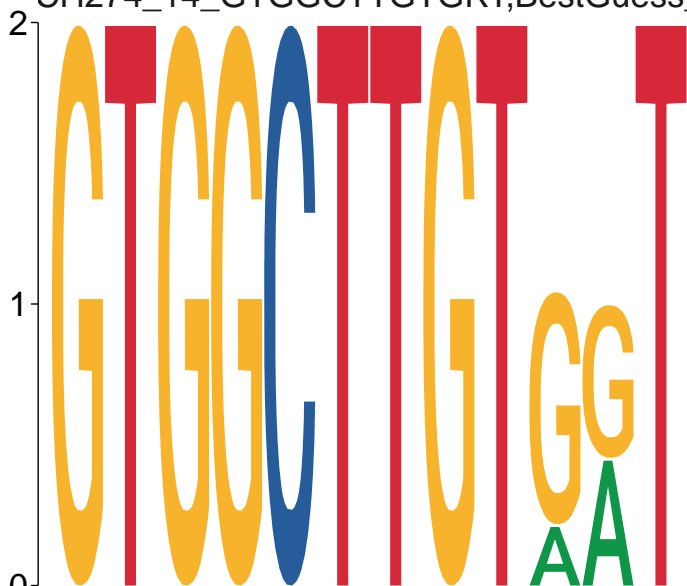
CH273_7_ATACAMSGCYAT,BestGuess_M



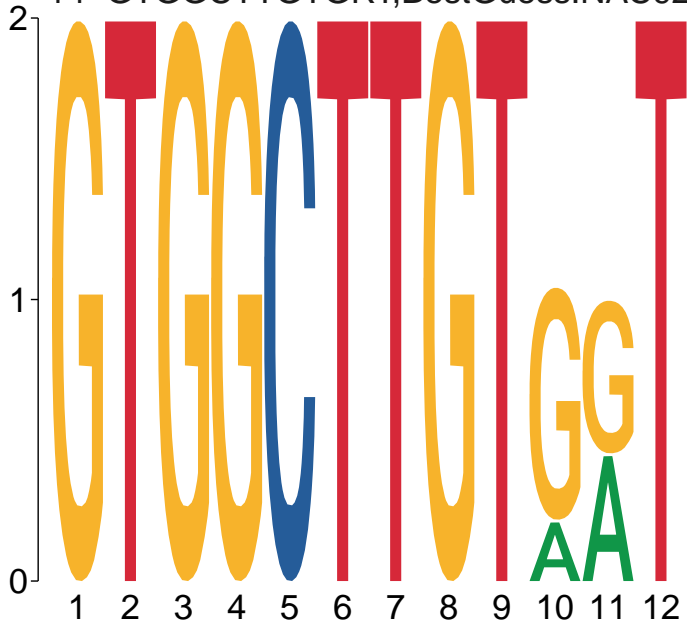
-ATACAMSGCYAT,BestGuess:MAL13P1.3!



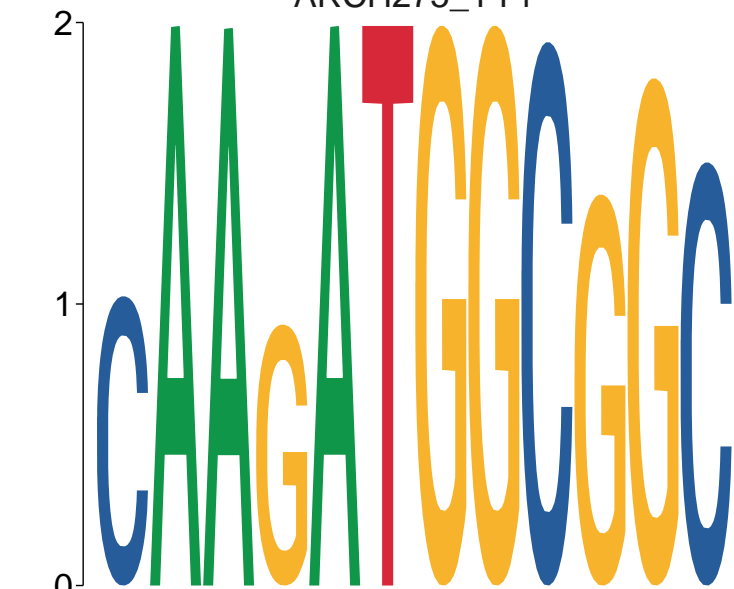
CH274_14_GTGGCTTGTGRT,BestGuess_



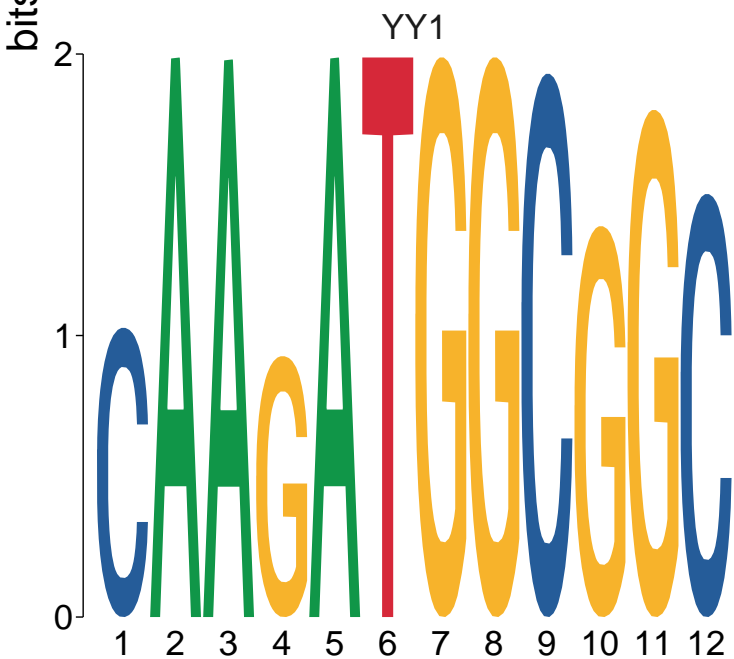
14-GTGGCTTGTGRT,BestGuess:NAC92



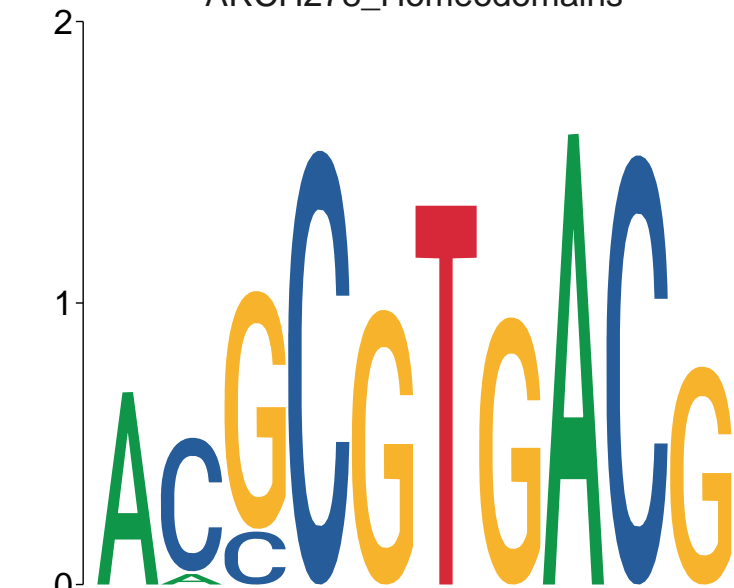
ARCH275_YY1



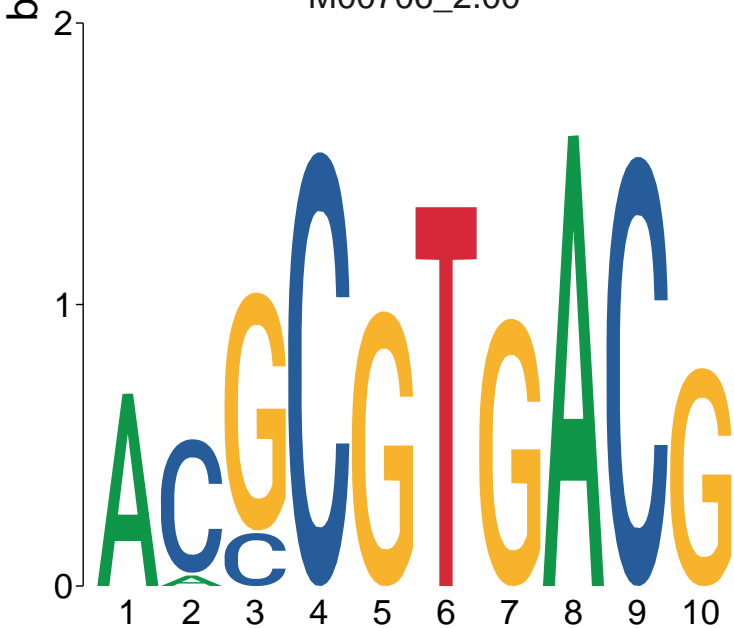
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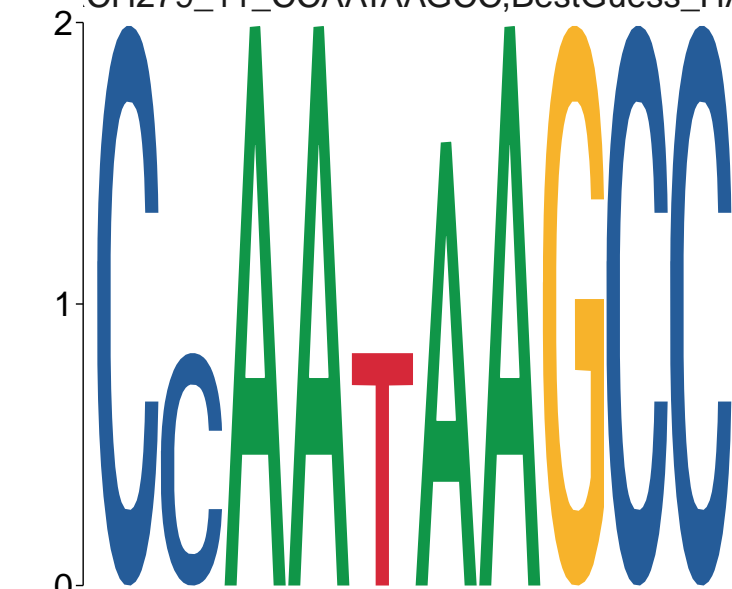
ARCH278_Homeodomains



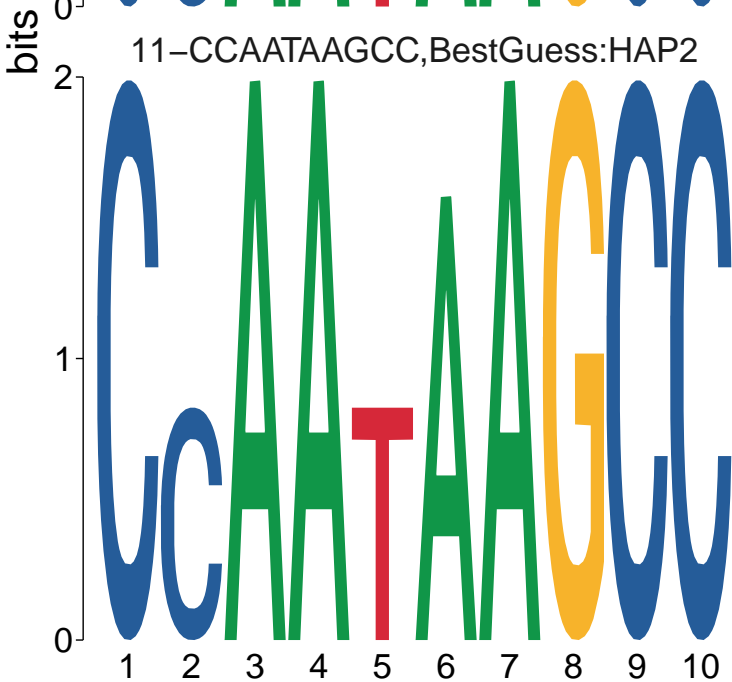
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CH279_11_CCAATAAGCC,BestGuess_HA



11-CCAATAAGCC,BestGuess:HAP2



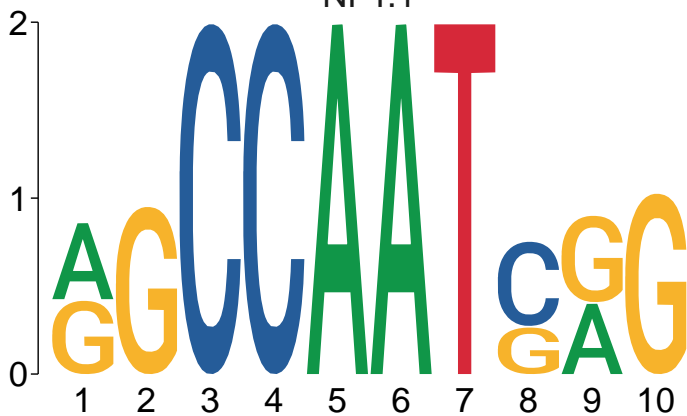
ARCH280_NFY_NFY



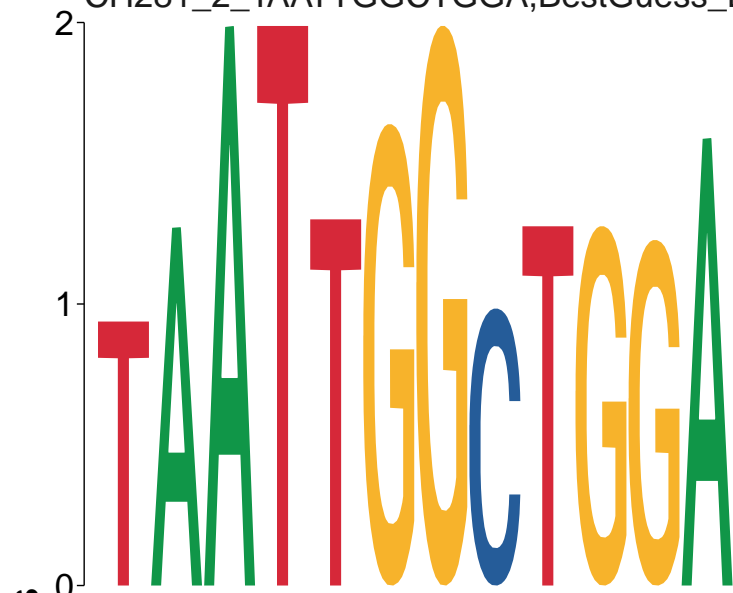
NFY



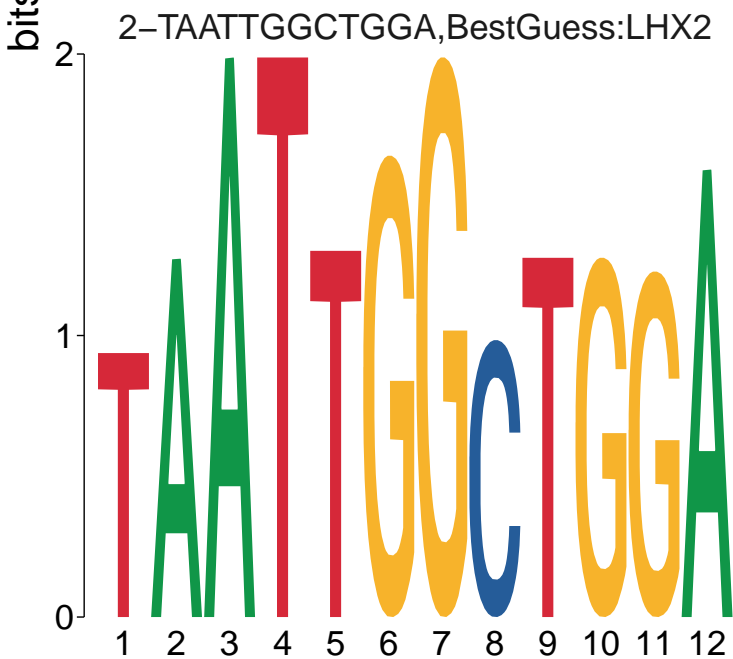
NFY.1



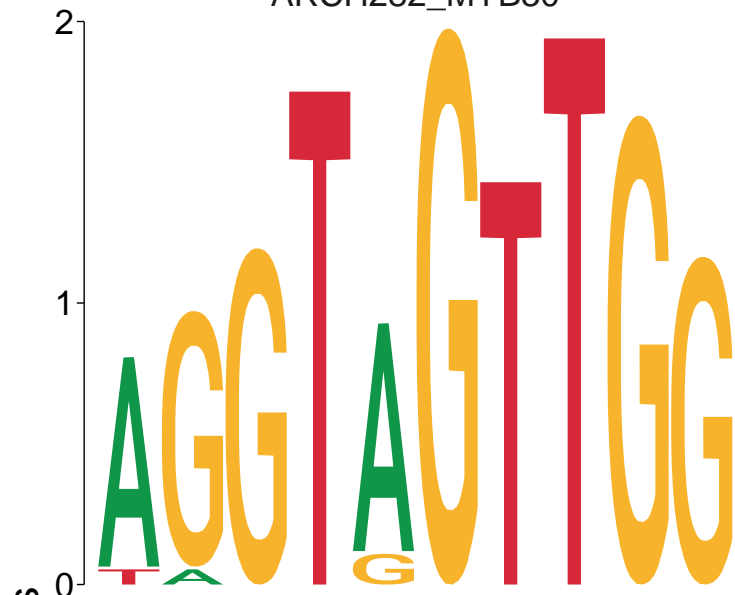
CH281_2_TAATTGGCTGGA,BestGuess_L



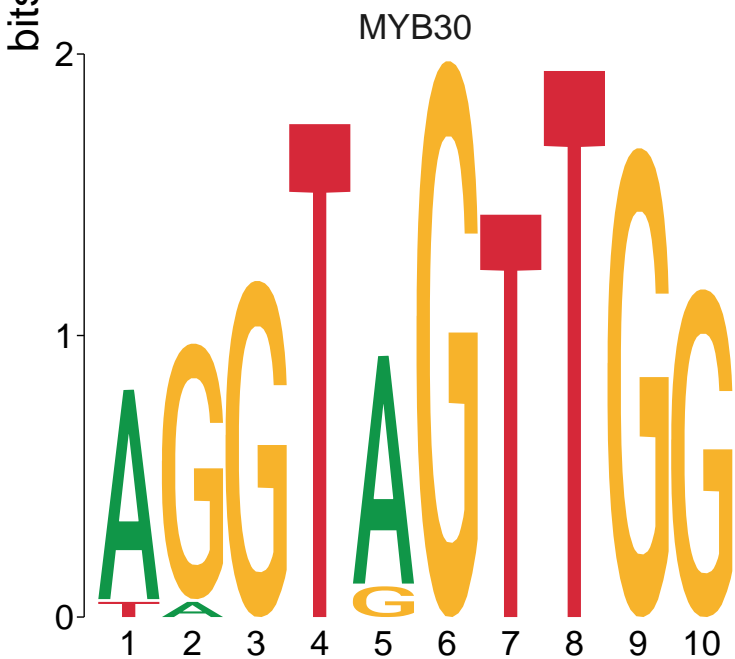
2-TAATTGGCTGGA,BestGuess:LHX2



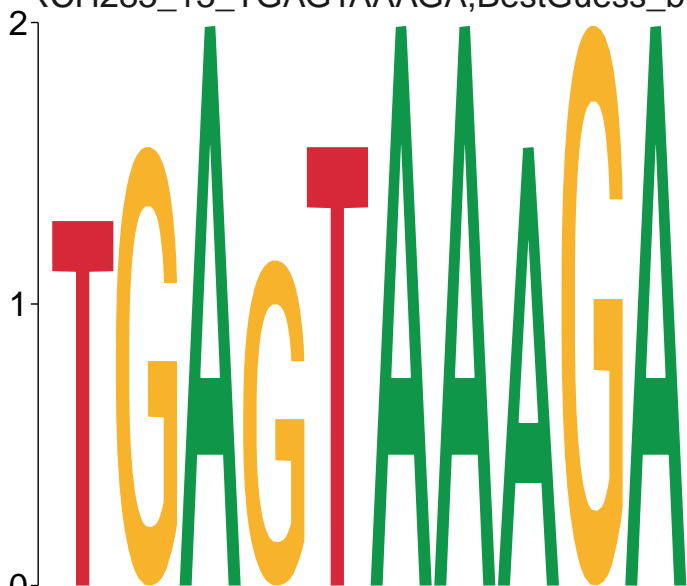
ARCH282_MYB30



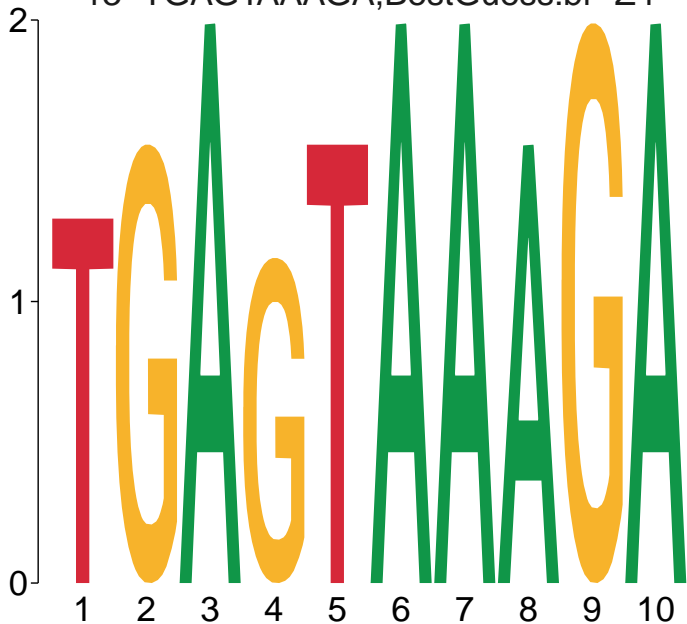
MYB30



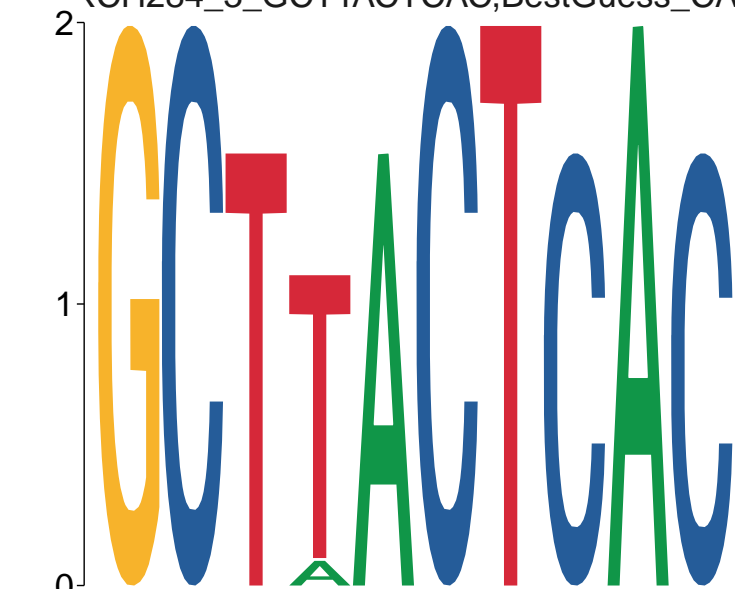
RCH283_13_TGAGTAAAGA,BestGuess_br.



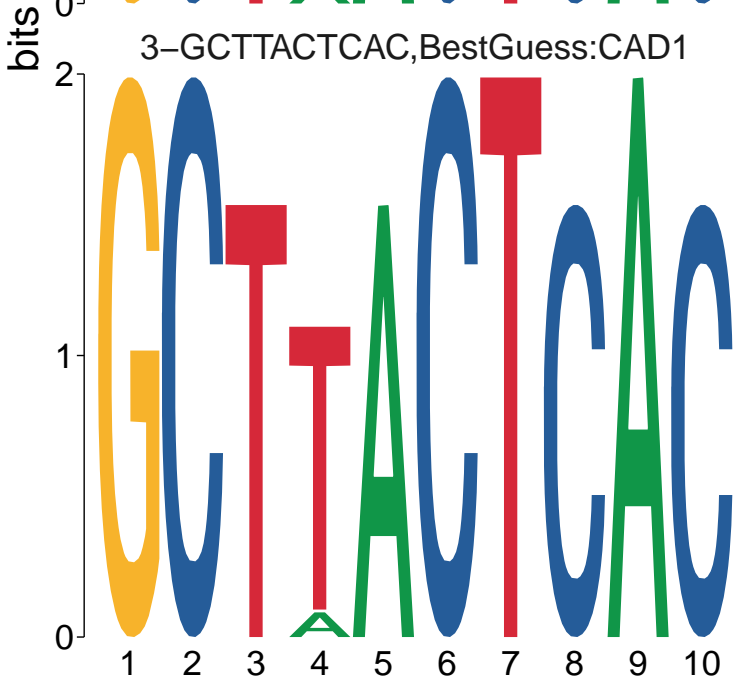
bits 13-TGAGTAAAGA,BestGuess:br-Z4



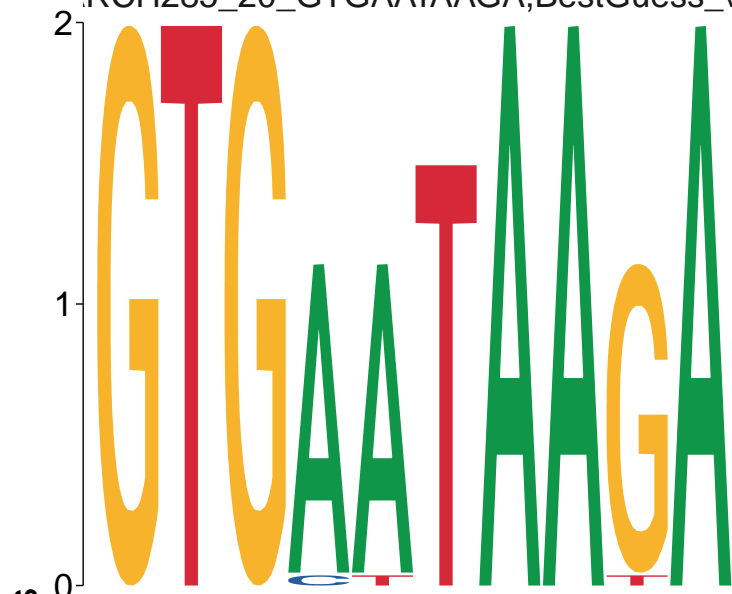
RCH284_3_GCTTACTCAC, BestGuess_CAI



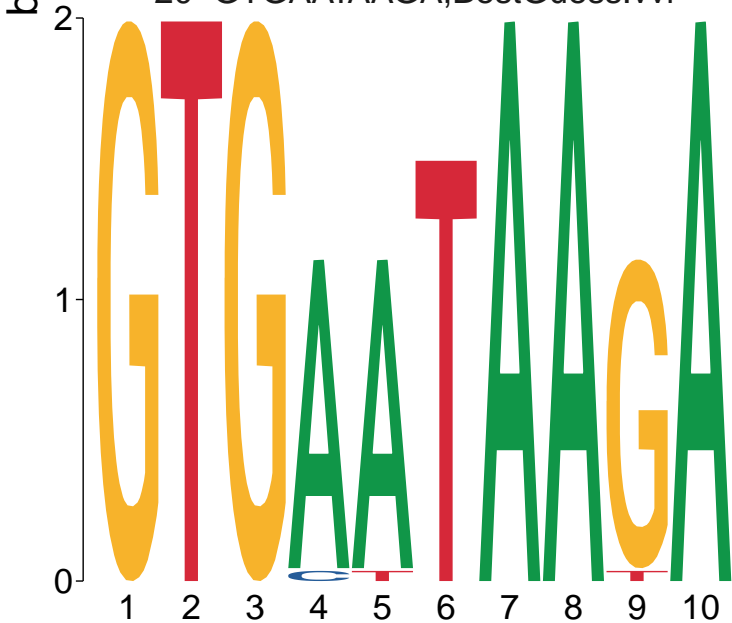
3-GCTTACTCAC, BestGuess:CAD1



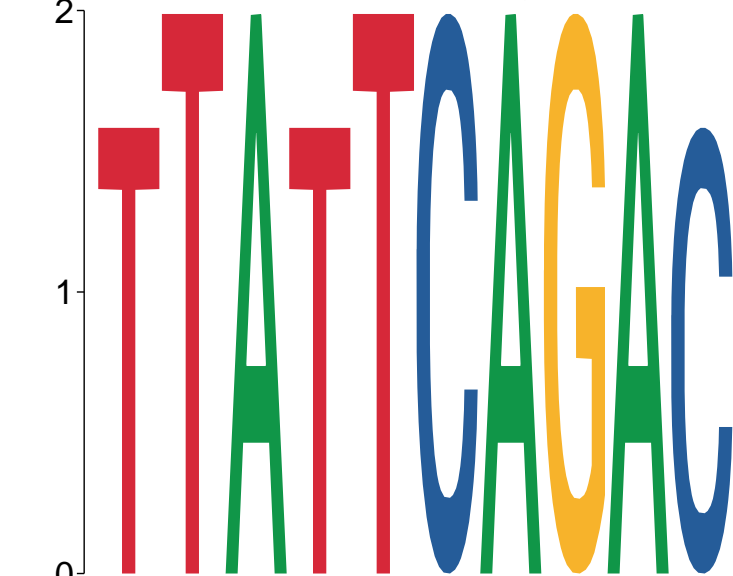
RCH285_20_GTGAATAAGA,BestGuess_v1



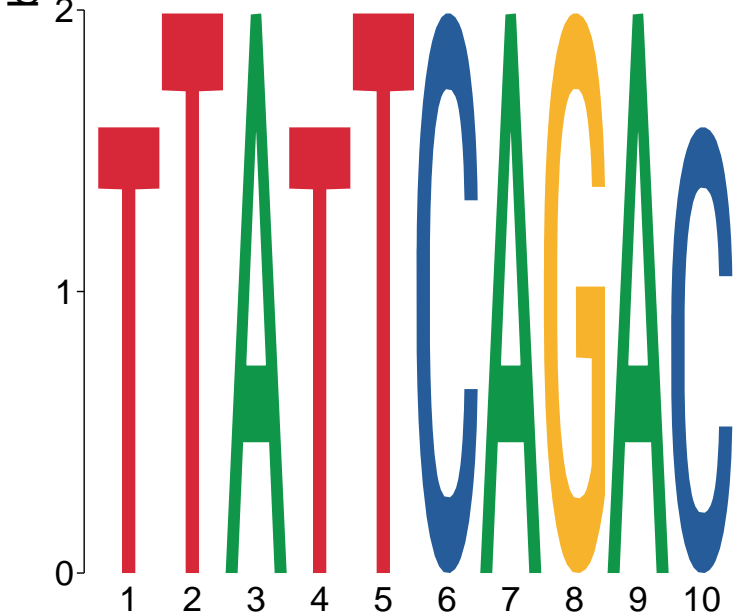
20-GTGAATAAGA,BestGuess:v1



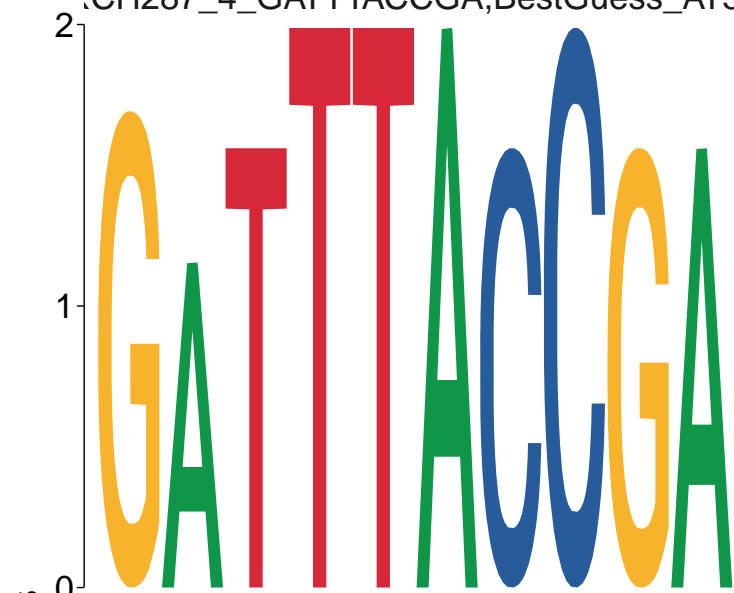
ARCH286_22_TTATTCAGAC,BestGuess_v1



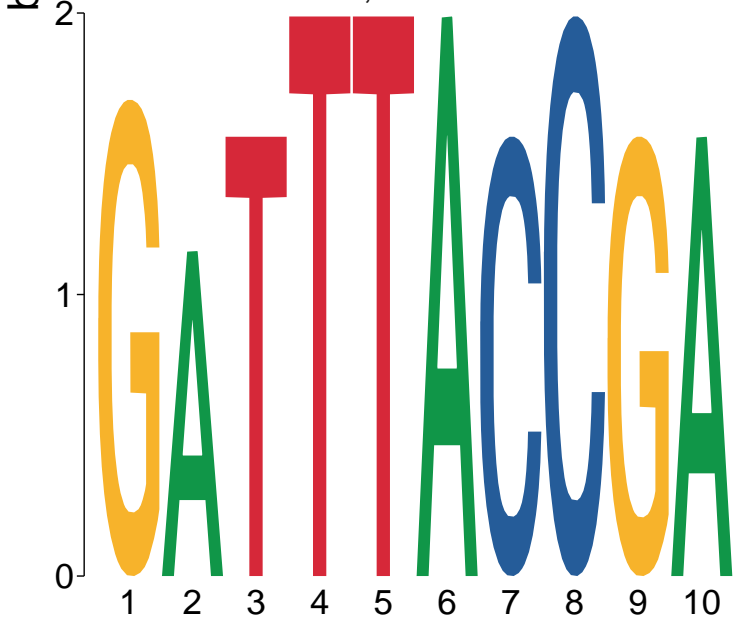
22-TTATTCAGAC,BestGuess:v1



CH287_4_GATTACCGA,BestGuess_AT50

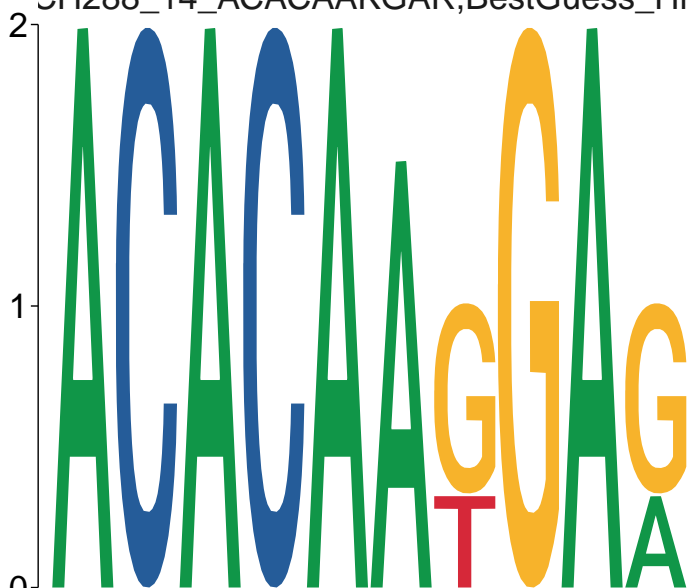


4-GATTTACCGA,BestGuess:AT5G47660

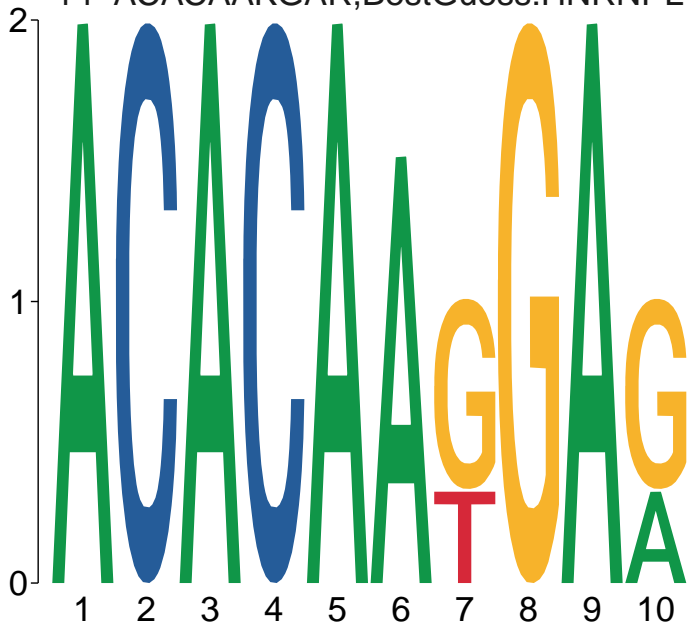


CH288_14_ACACAAKGAR,BestGuess_HN

bits



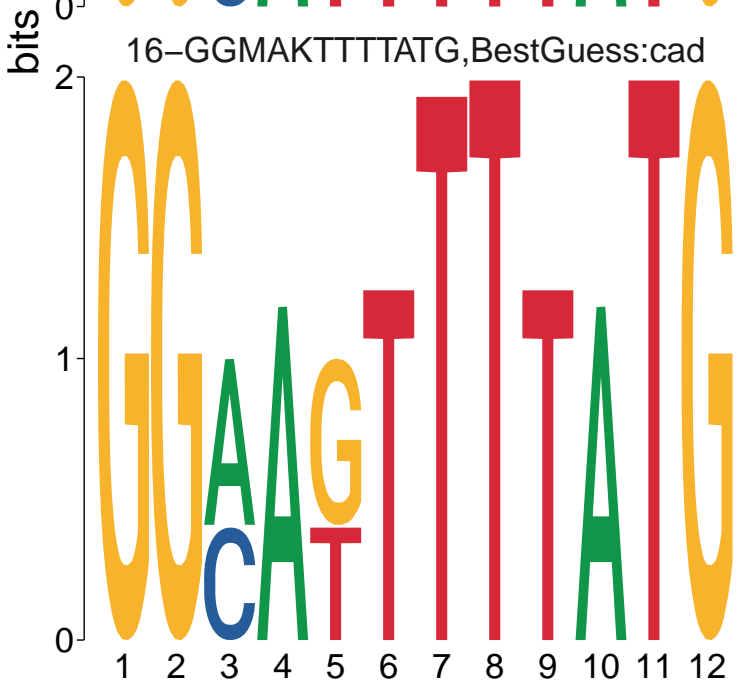
14-ACACAAKGAR,BestGuess:HNRNPL



CH290_16_GGMAKTTTTATG,BestGuess_



16-GGMAKTTTTATG,BestGuess:cad



CH291_12_ACAACGAAAAAT,BestGuess_

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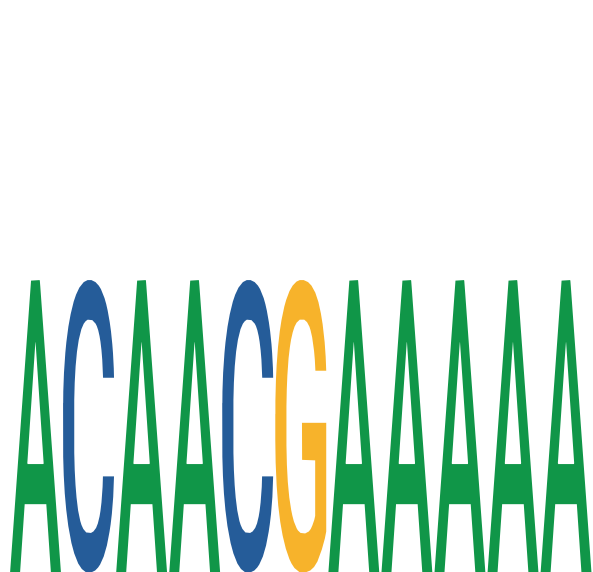
ACGAAAAAT,BestGuess:PB0032.1_IRC9C

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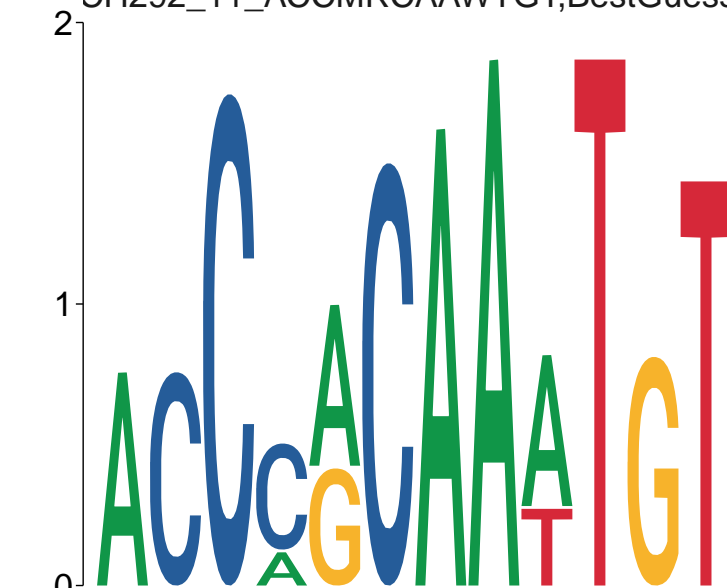
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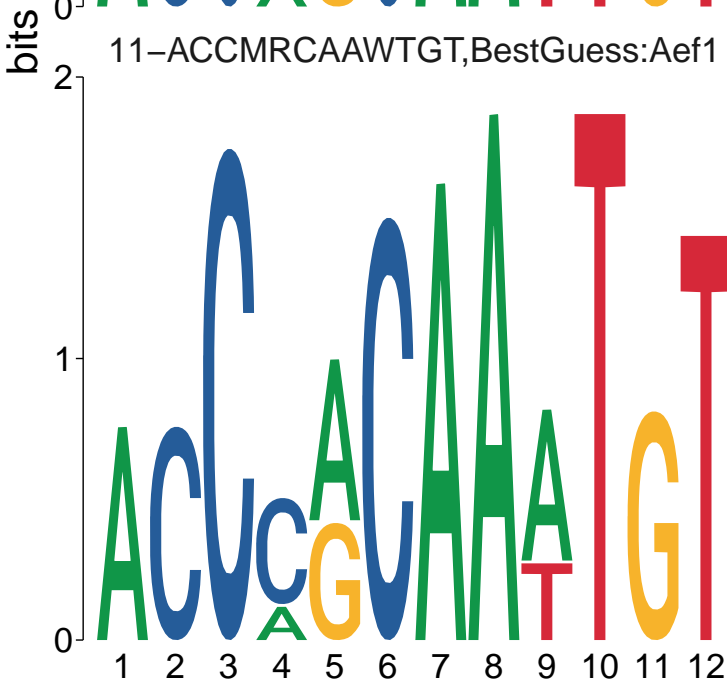
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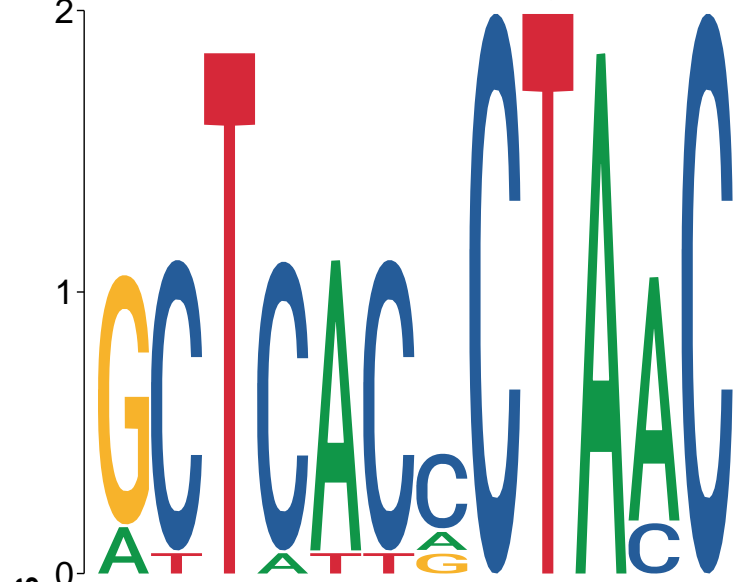
CH292_11_ACCMRCAAWTGT, BestGuess_



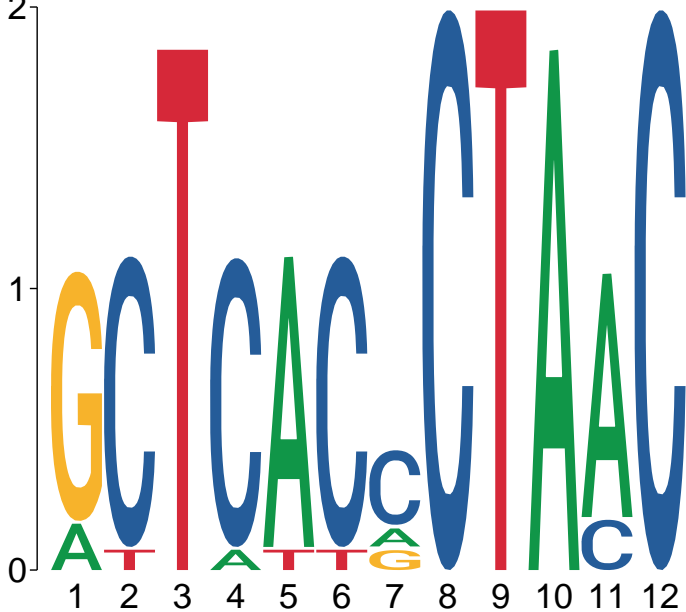
11-ACCMRCAAWTGT, BestGuess:Aef1



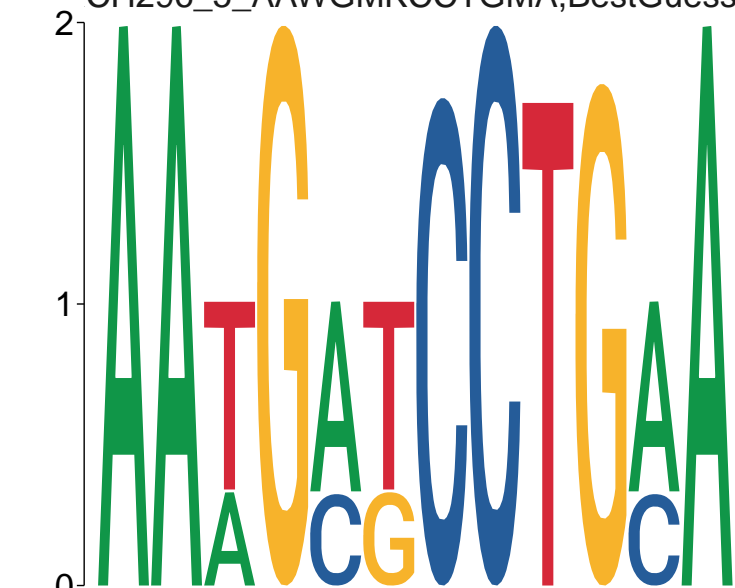
CH295_19_GCTCACVCTAAC,BestGuess_



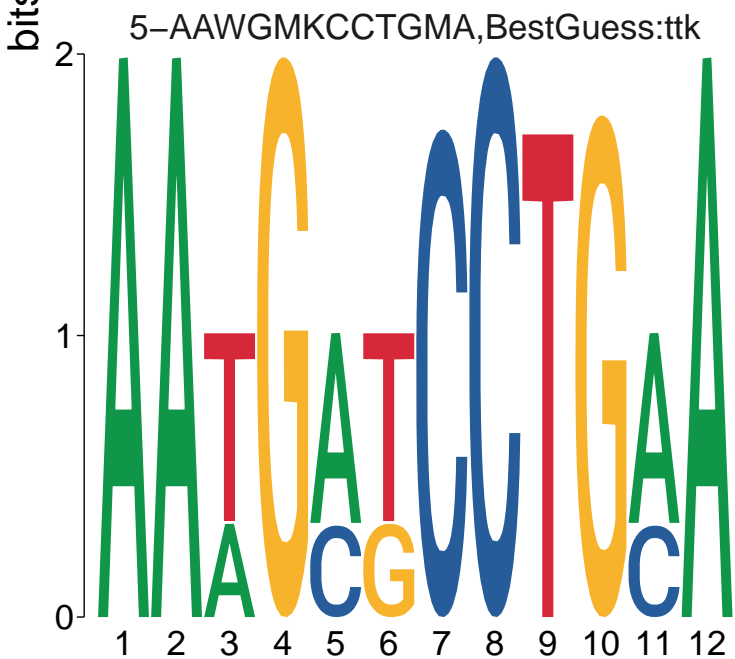
9-GCTCACVCTAAC,BestGuess:HRB98DE



CH296_5_AAWGMKCCTGMA,BestGuess_



5-AAWGMKCCTGMA,BestGuess:ttk



CH297_18_AACTCATGCTGT,BestGuess_

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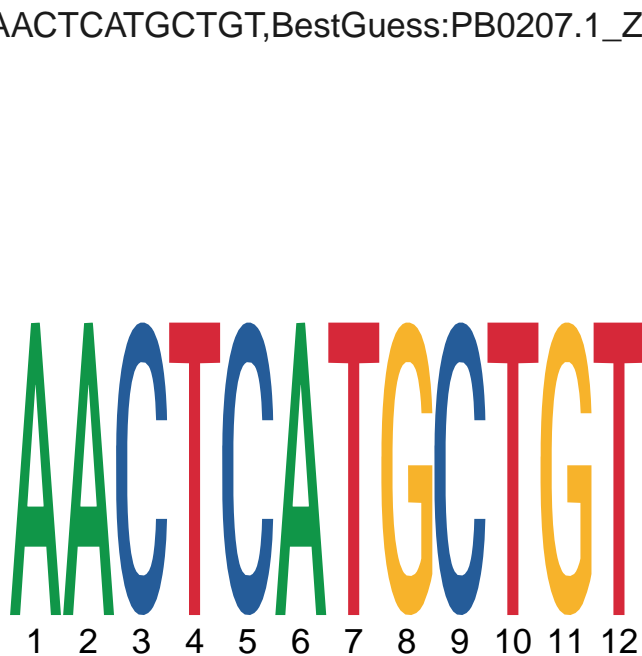
AACTCATGCTGT,BestGuess:PB0207.1_Zic

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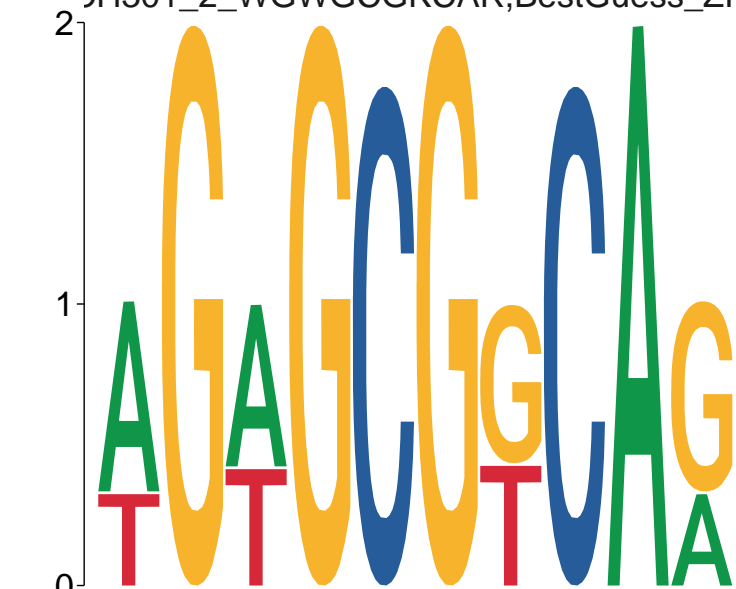
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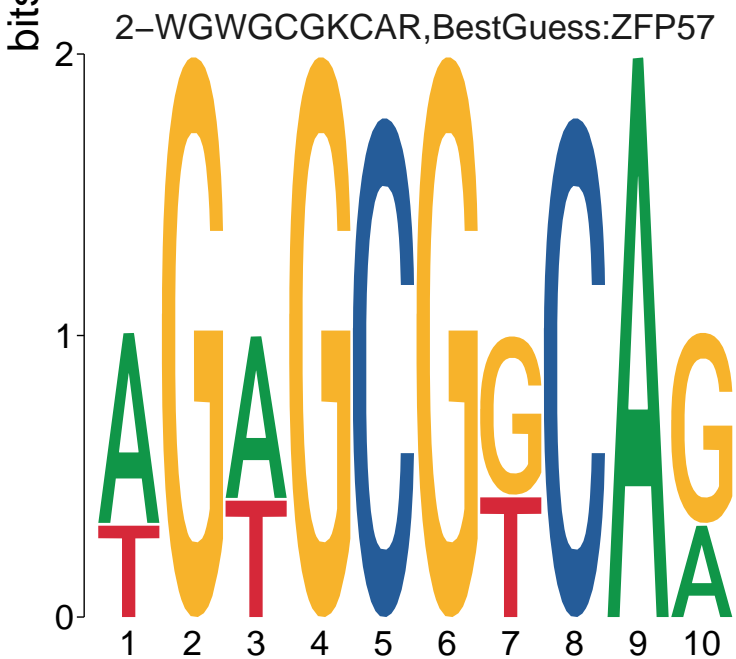
1 2 3 4 5 6 7 8 9 10 11 12



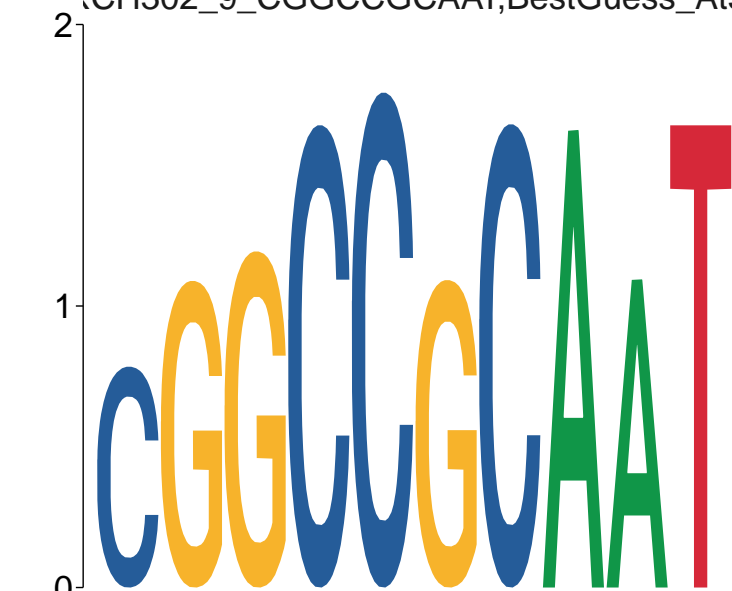
CH301_2_WGWGCGKCAR,BestGuess_ZF



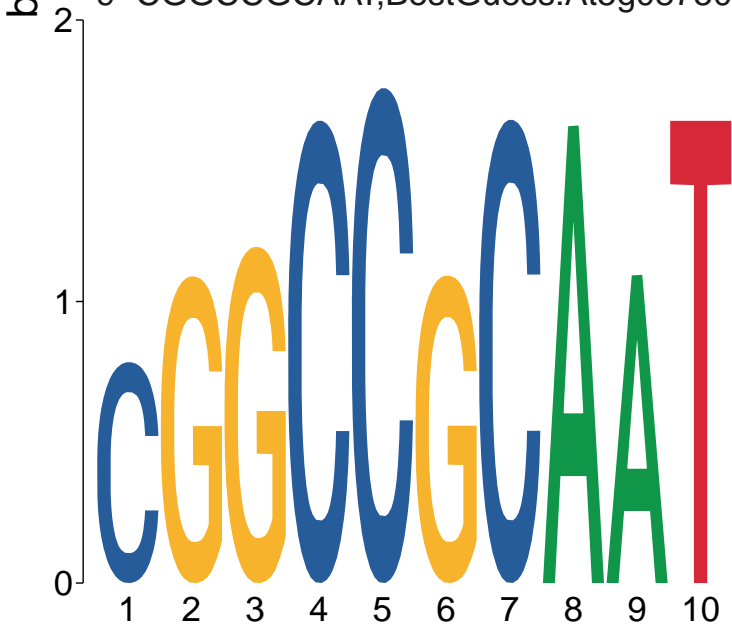
2-WGWGCGKCAR,BestGuess:ZFP57



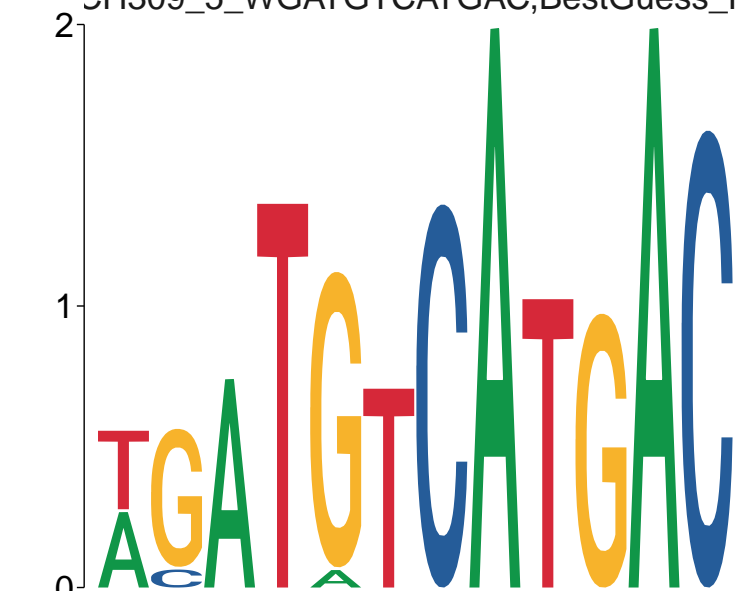
CH302_9_CGGCCGCAAT,BestGuess_At5



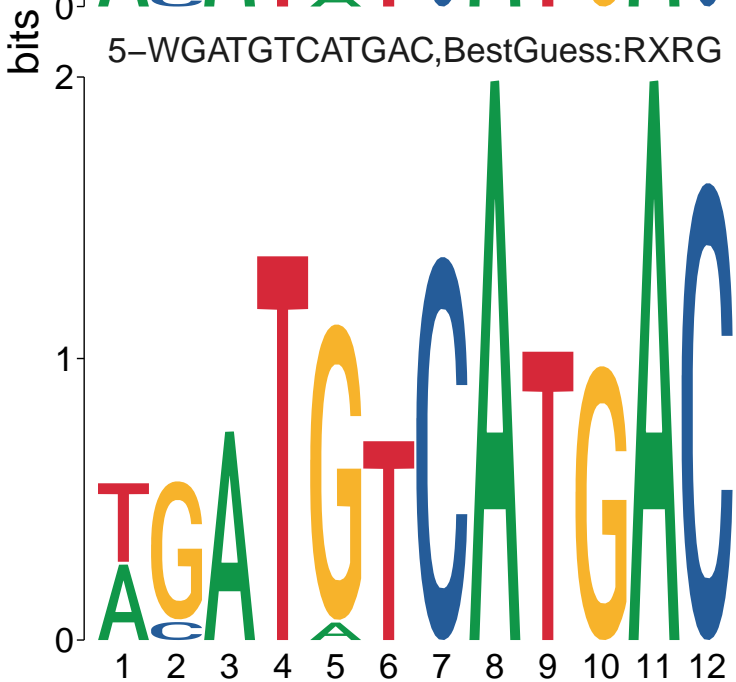
9-CGGCCGCAAT,BestGuess:At5g08750



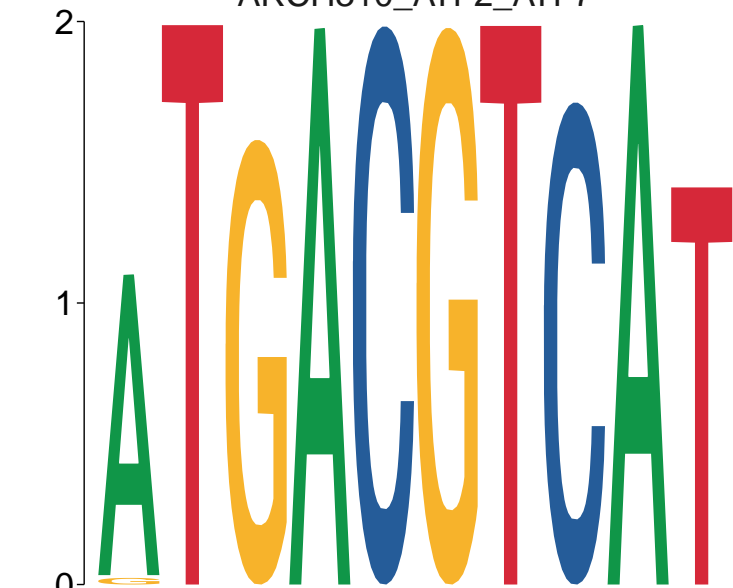
CH309_5_WGATGTCATGAC,BestGuess_R



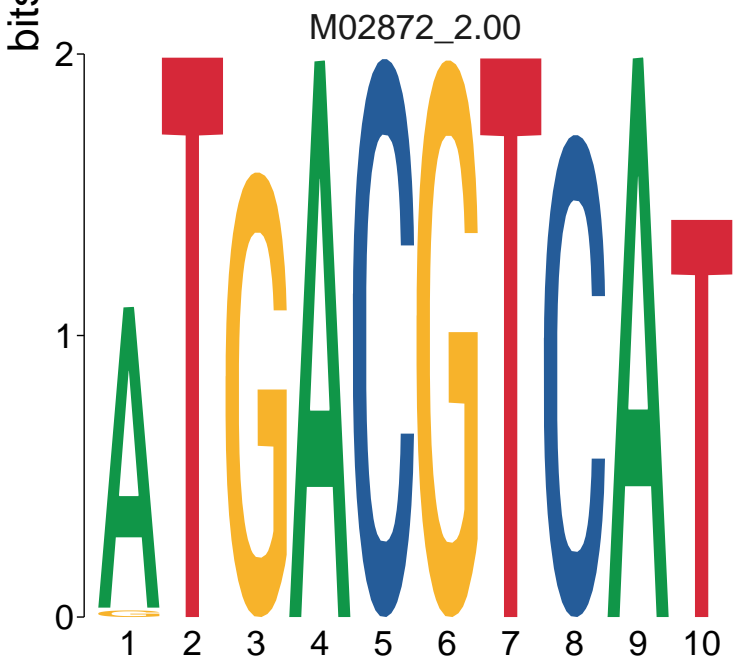
5-WGATGTCATGAC,BestGuess:RXRG



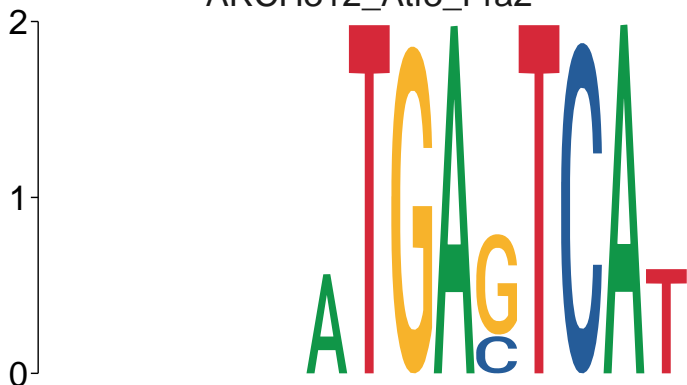
ARCH310_ATF2_ATF7



M02872_2.00



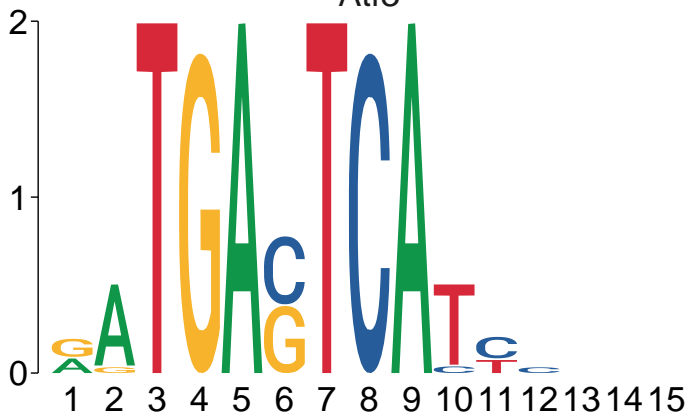
ARCH312_At3_Fra2



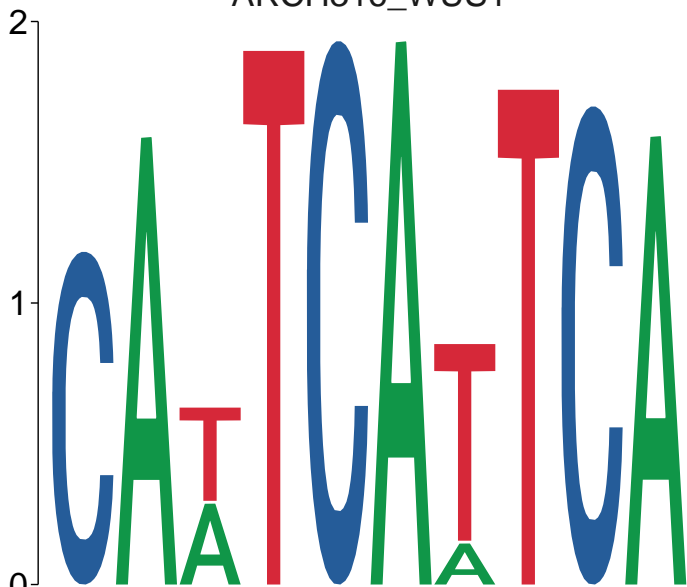
Fra2



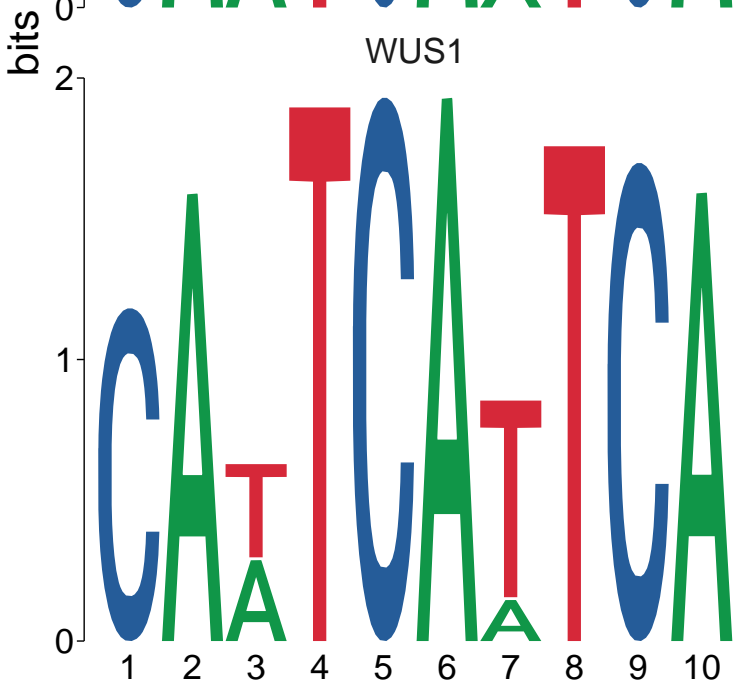
Atf3



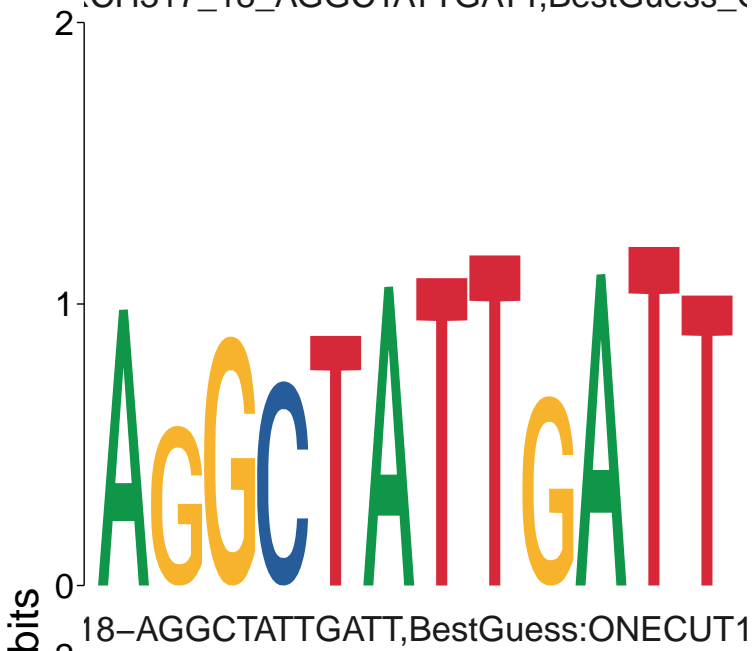
ARCH316_WUS1



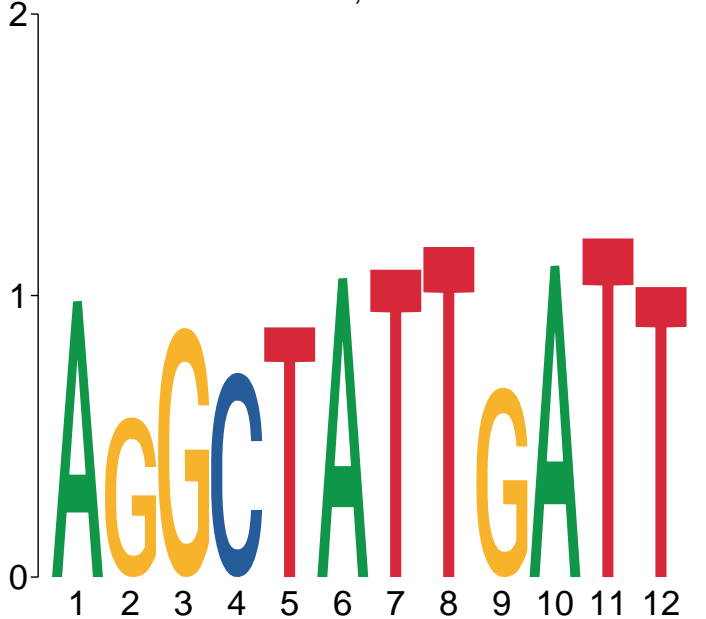
WUS1



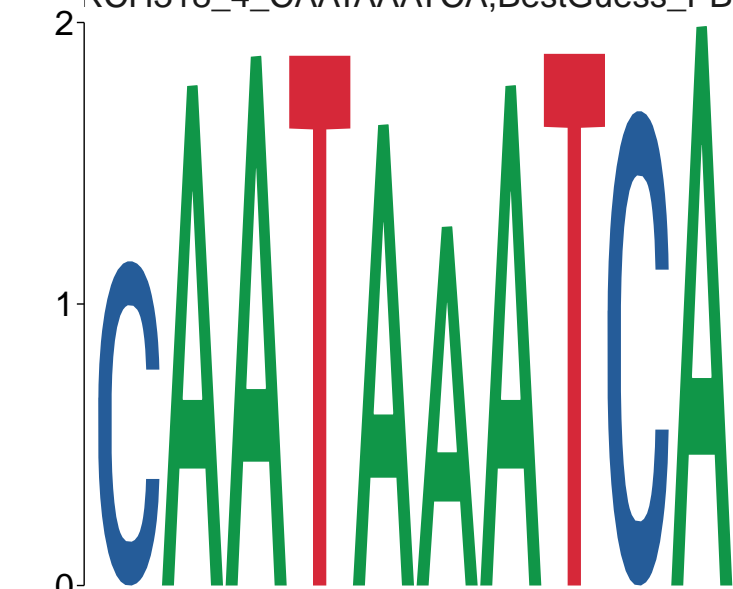
CH317_18_AGGCTATTGATT,BestGuess_C



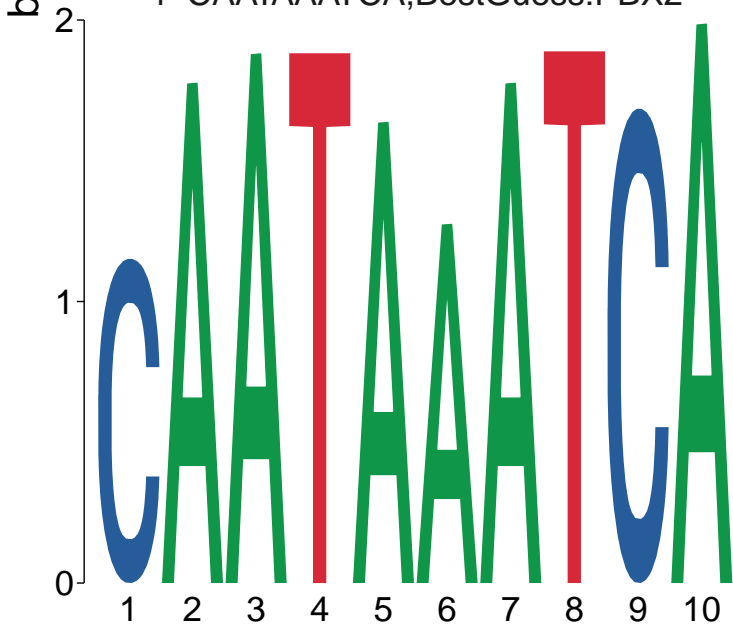
18-AGGCTATTGATT,BestGuess:ONECUT1



RCH318_4_CAATAAATCA,BestGuess_PBX



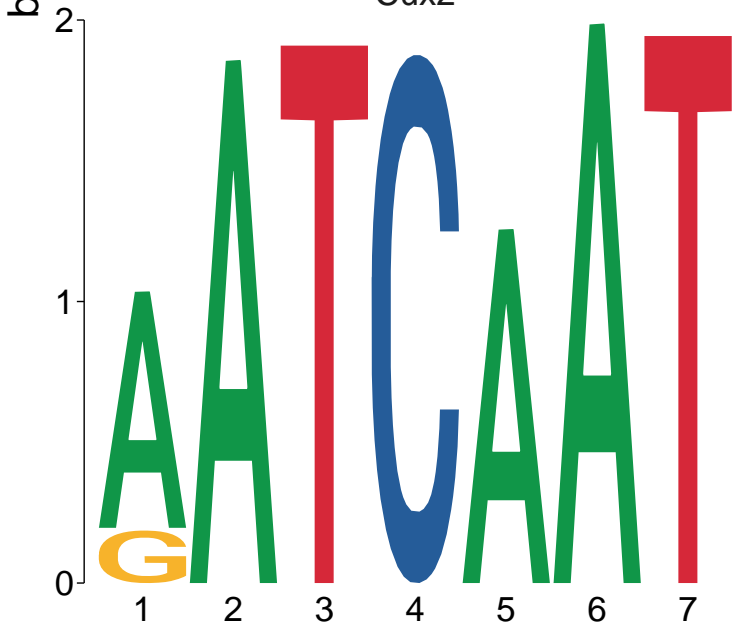
4-CAATAAATCA,BestGuess:PBX2



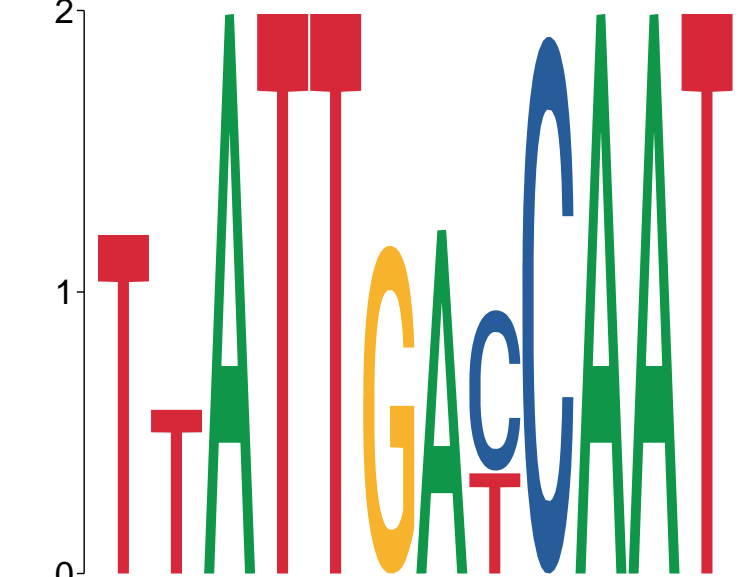
ARCH319_Cux2



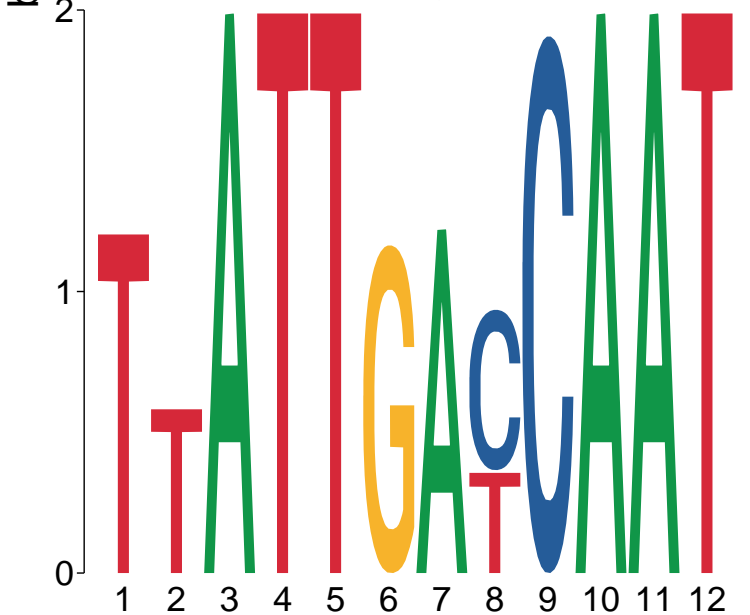
Cux2



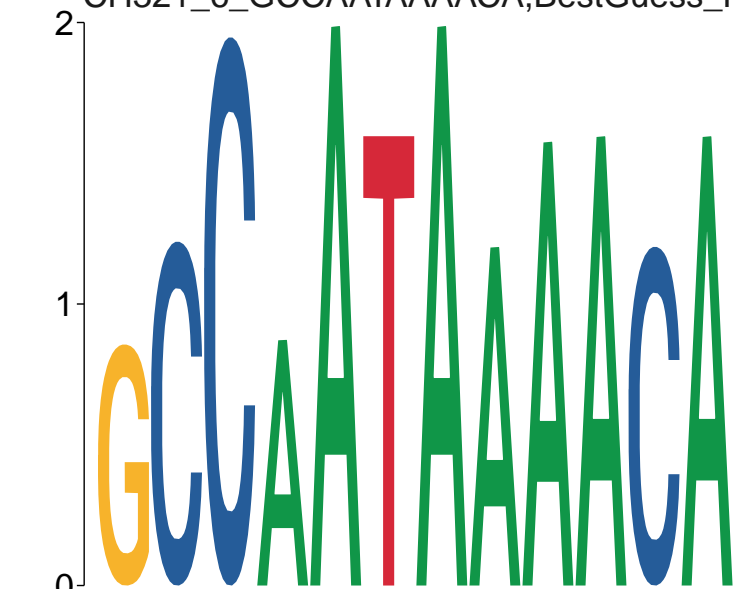
RCH320_1_TTATTGAYCAAT,BestGuess_sl



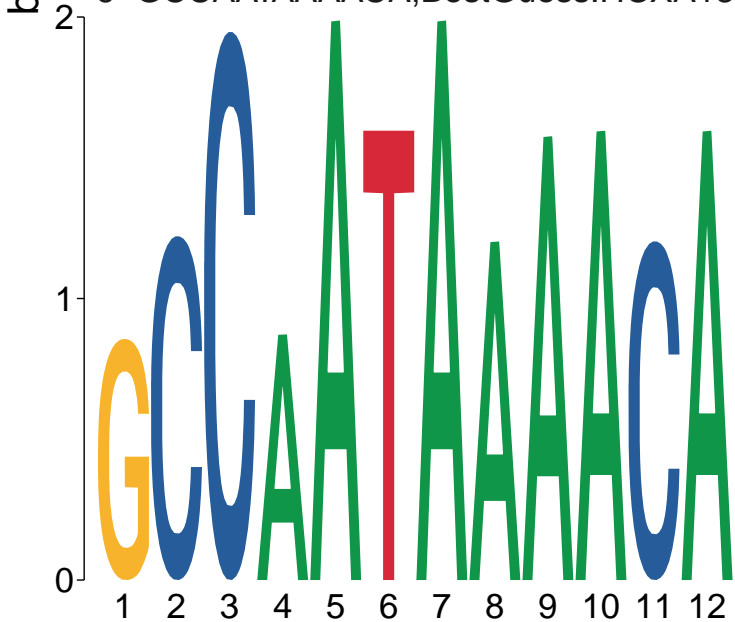
1-TTATTGAYCAAT,BestGuess:slbo



CH321_6_GCCAATAAAACA, BestGuess_H



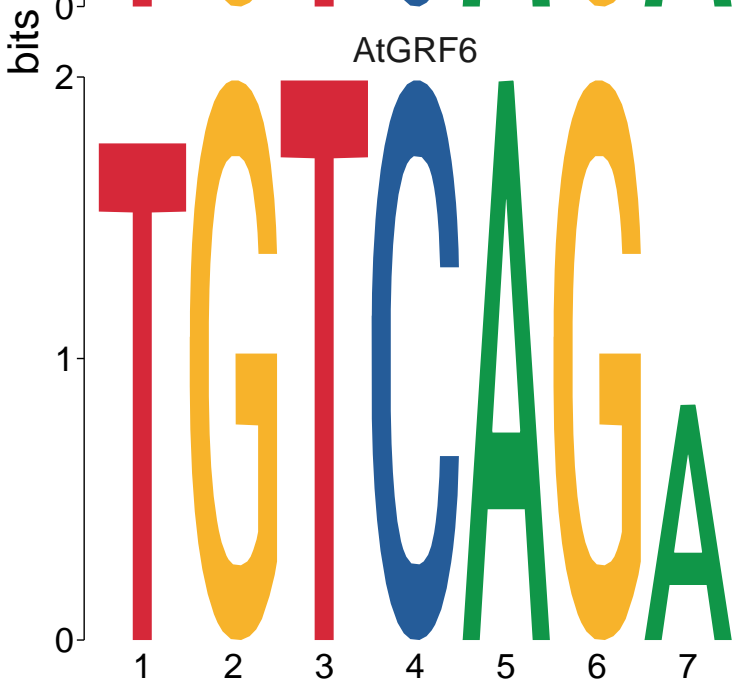
6-GCCAATAAAACA, BestGuess:HOXA13



ARCH323_AtGRF6



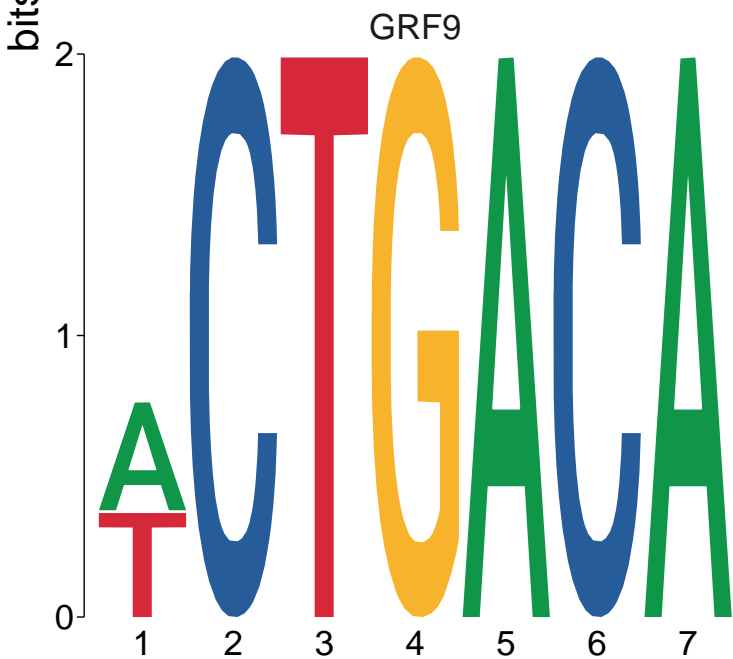
AtGRF6



ARCH324_GRF9



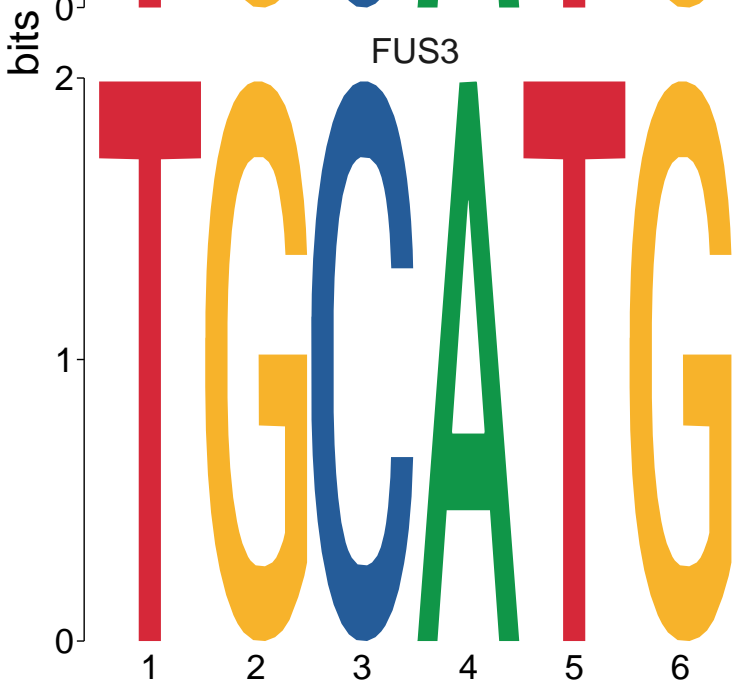
GRF9



ARCH326_FUS3



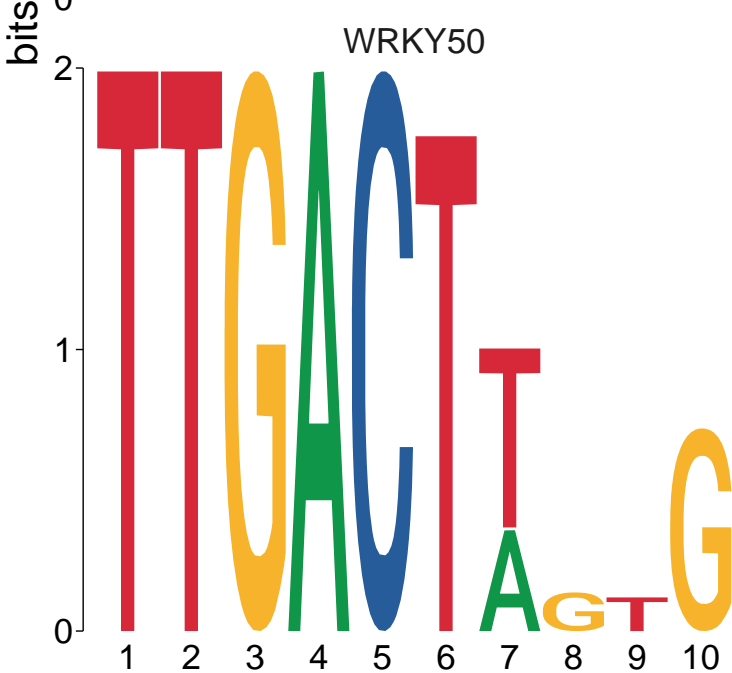
FUS3



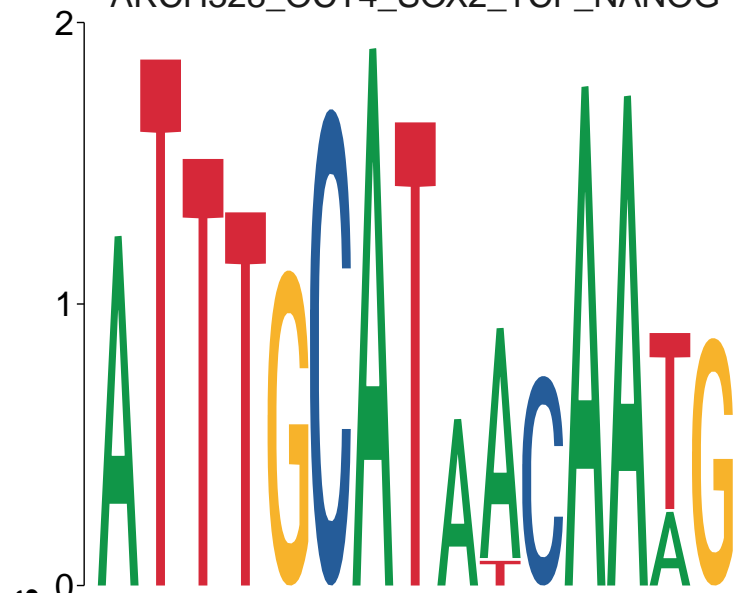
ARCH327_WRKY50



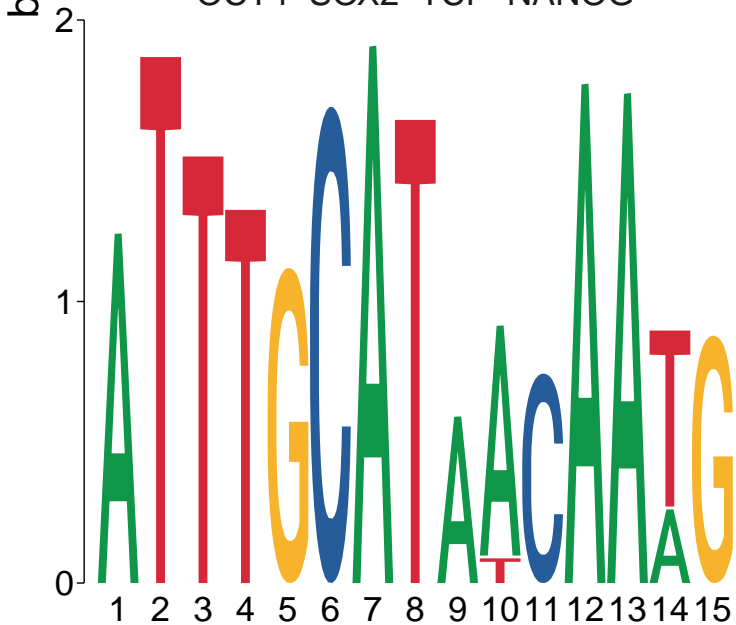
WRKY50



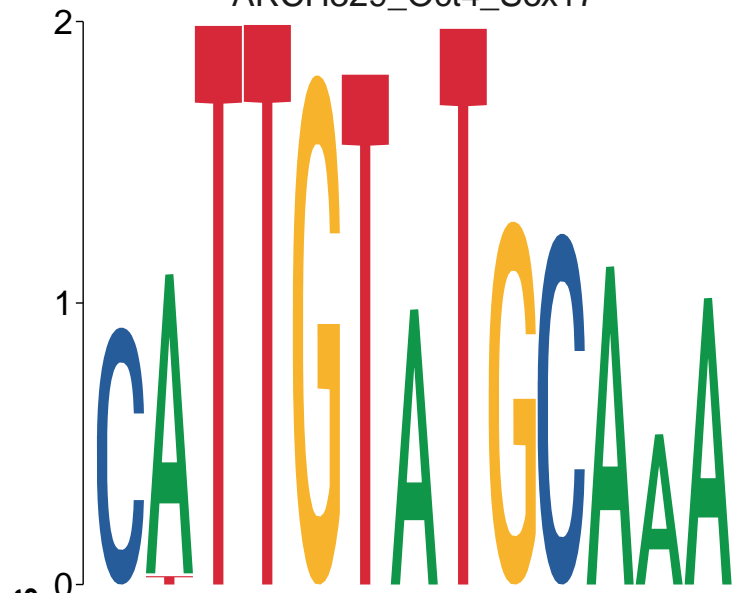
ARCH328_OCT4_SOX2_TCF_NANOG



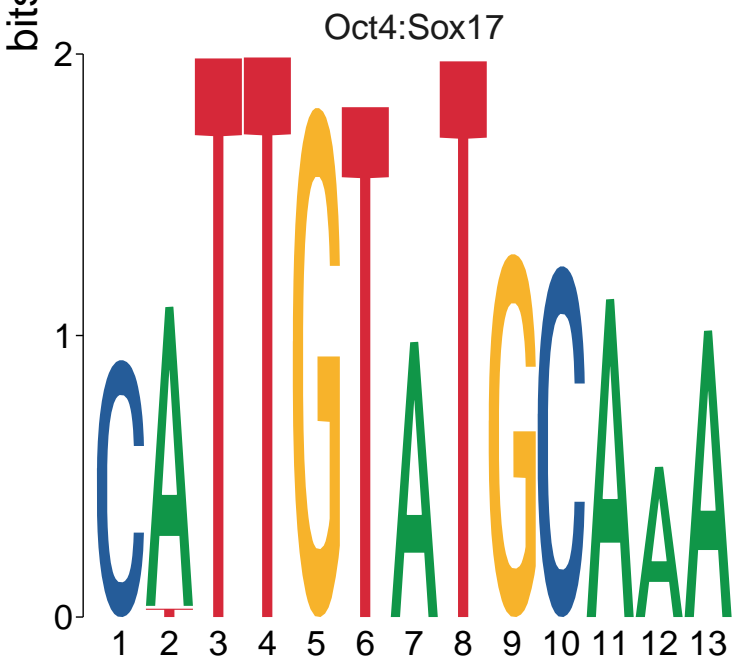
OCT4-SOX2-TCF-NANOG



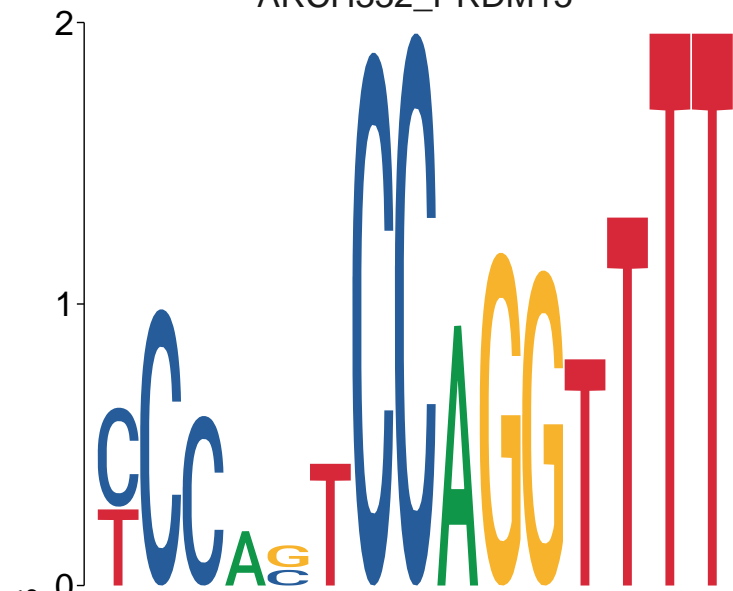
ARCH329_Oct4_Sox17



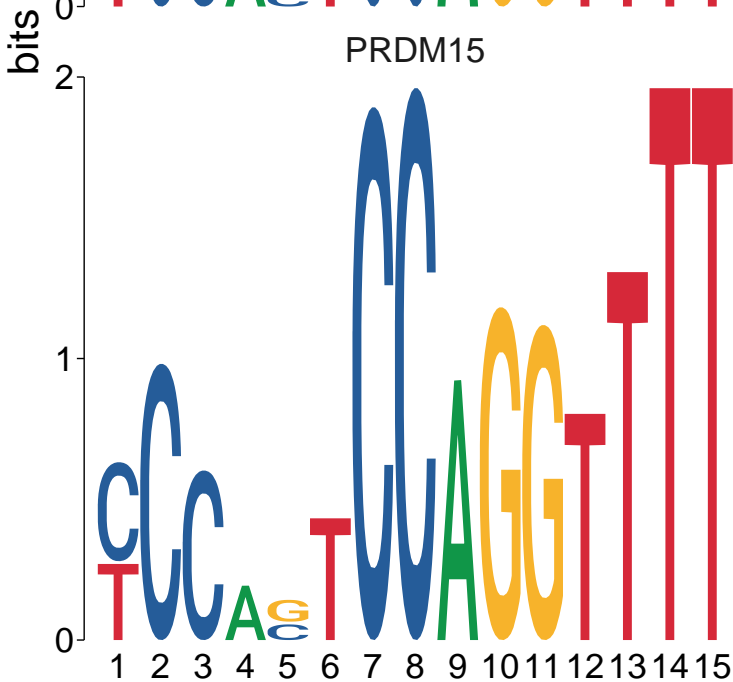
Oct4:Sox17



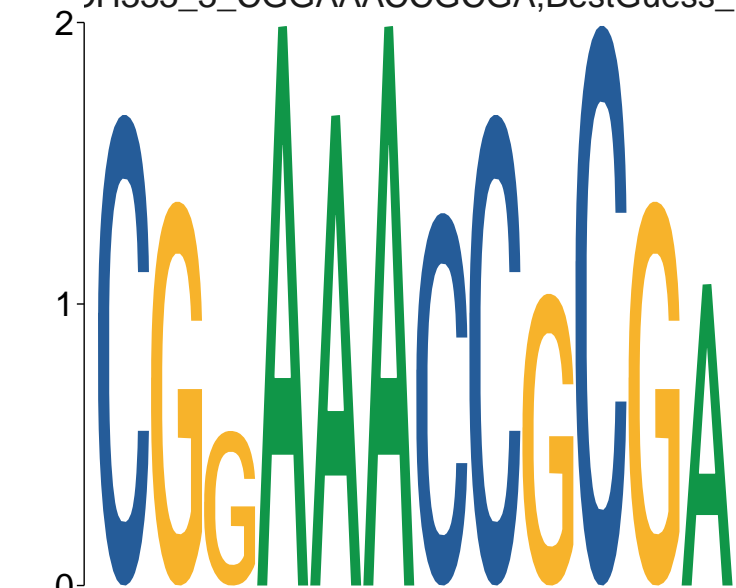
ARCH332_PRDM15



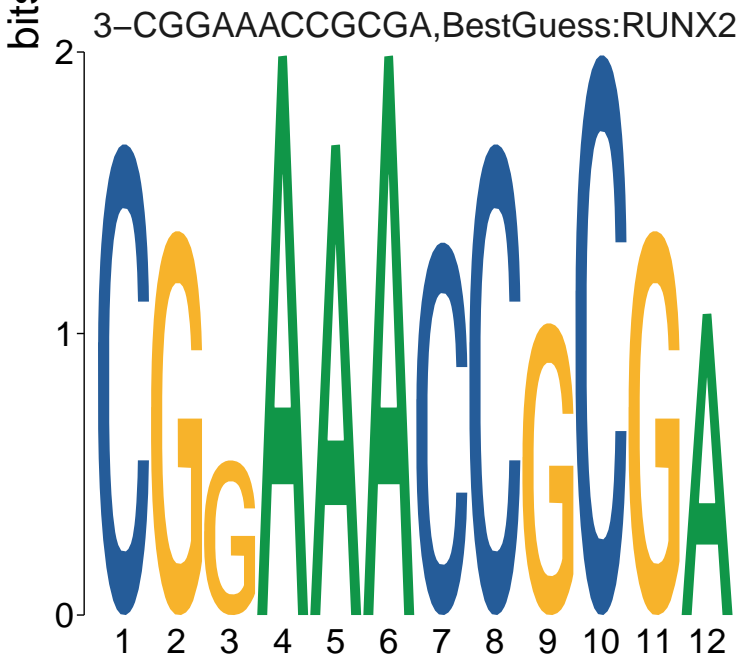
PRDM15



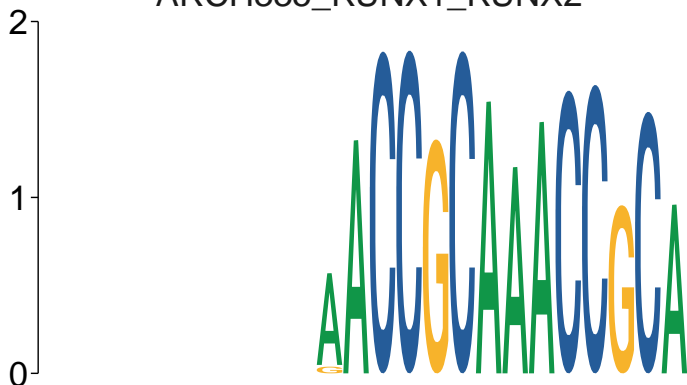
CH333_3_CGGAAACCGCGA,BestGuess_F



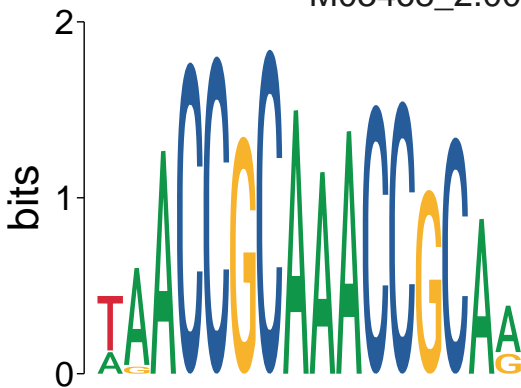
3-CGGAAACCGCGA,BestGuess:RUNX2



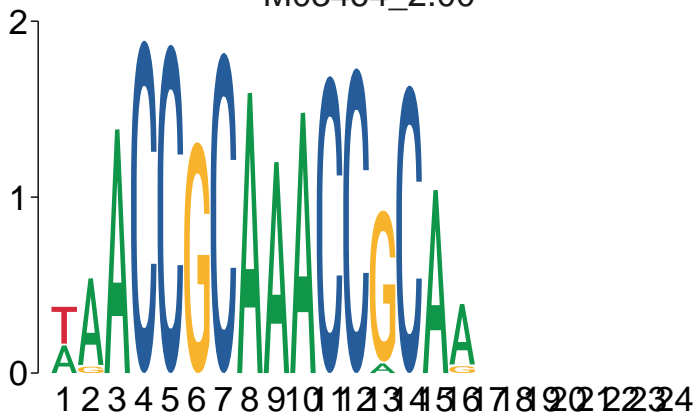
ARCH335_RUNX1_RUNX2



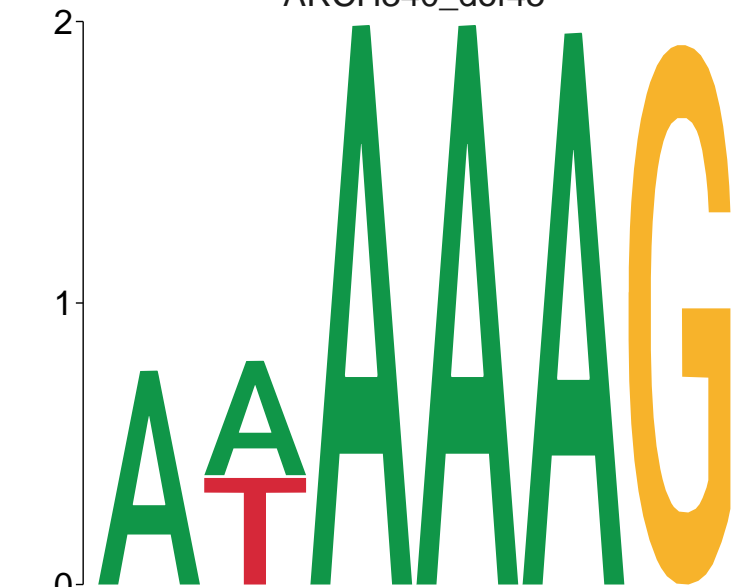
M03468_2.00



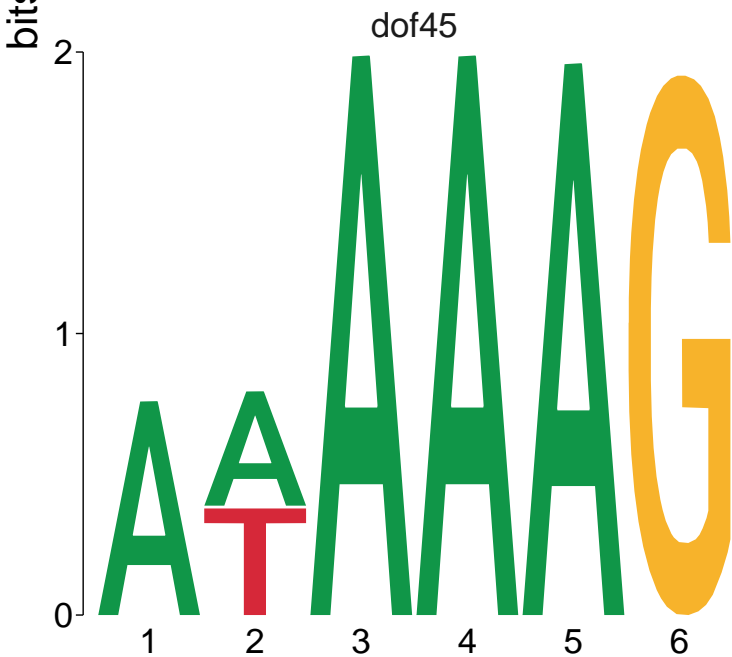
M03464_2.00



ARCH340_dof45



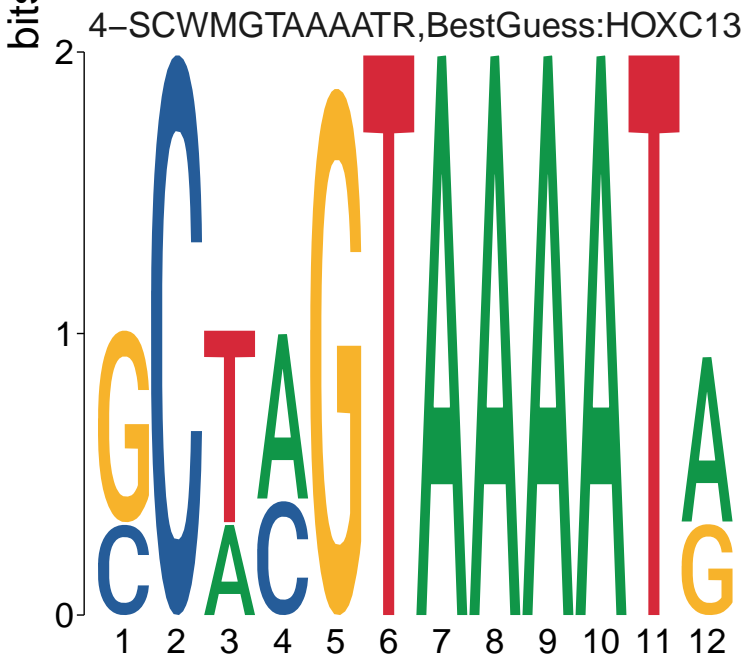
dof45



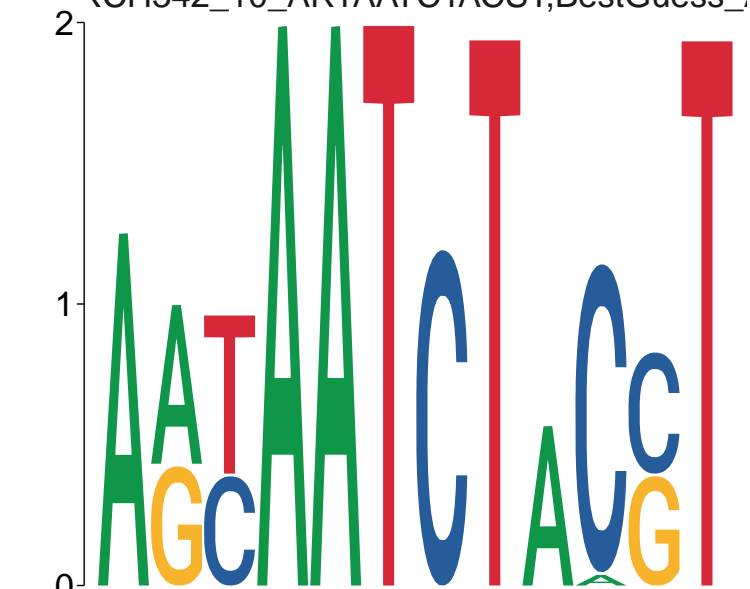
CH341_4_SCWMGTAAAATR,BestGuess_1



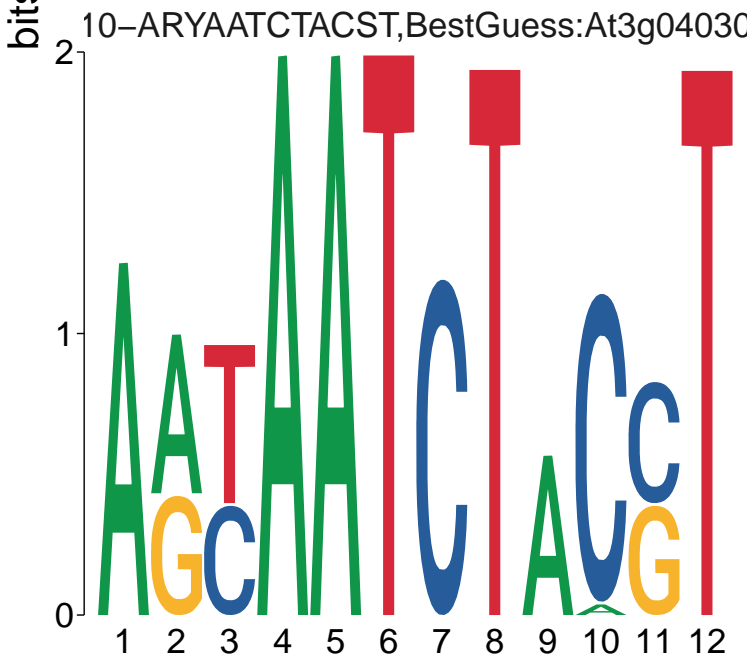
4-SCWMGTAAAATR,BestGuess:HOXC13



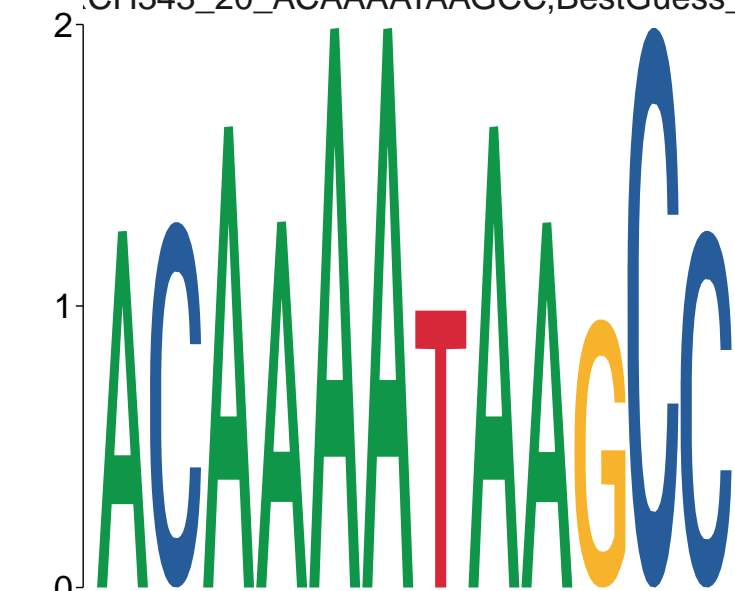
RCH342_10_ARYAATCTACST,BestGuess_



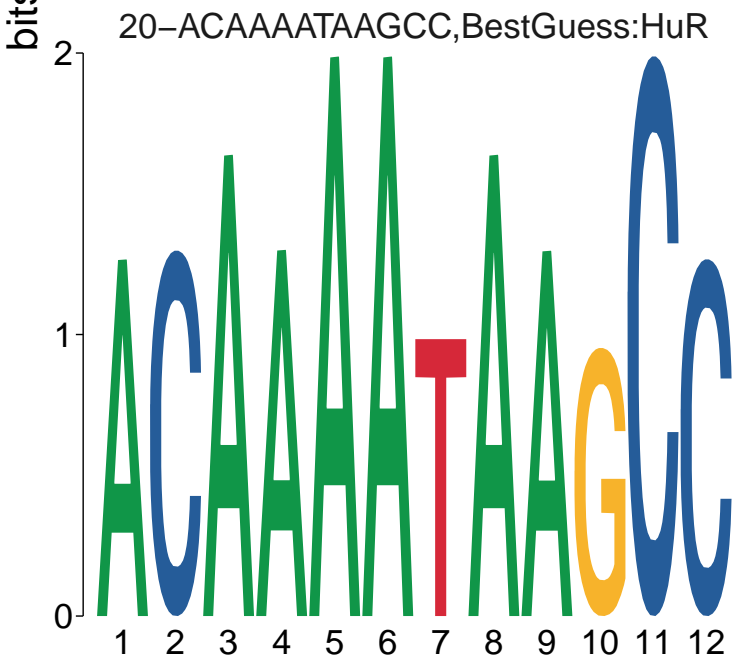
10-ARYAATCTACST,BestGuess:At3g0403C



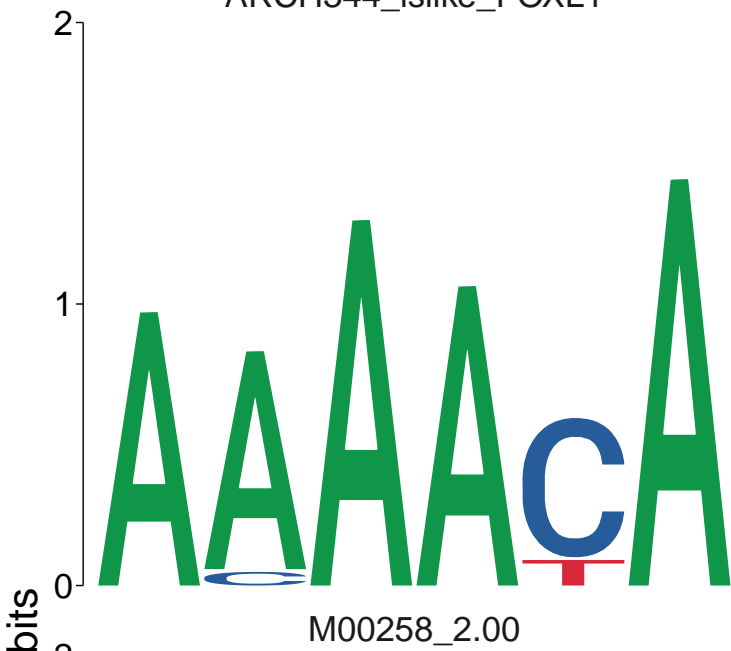
CH343_20_ACAAAATAAGCC,BestGuess_



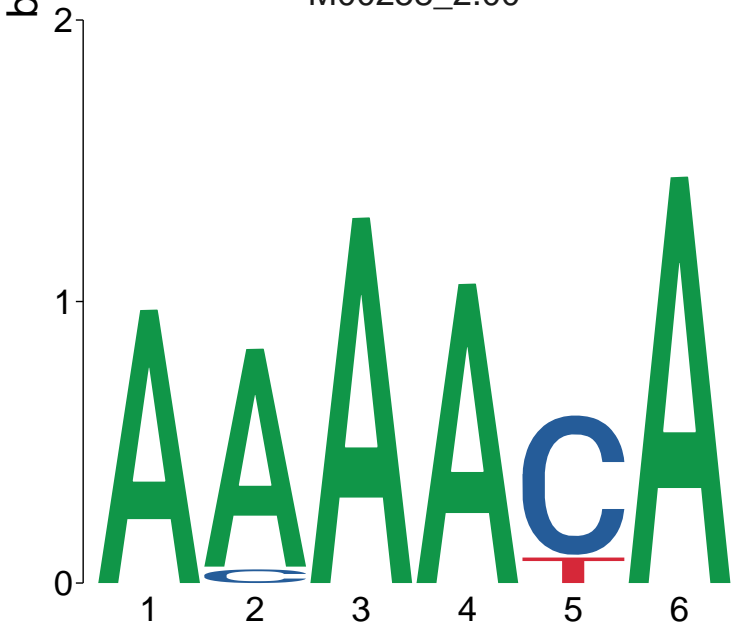
20-ACAAAATAAGCC,BestGuess:HuR



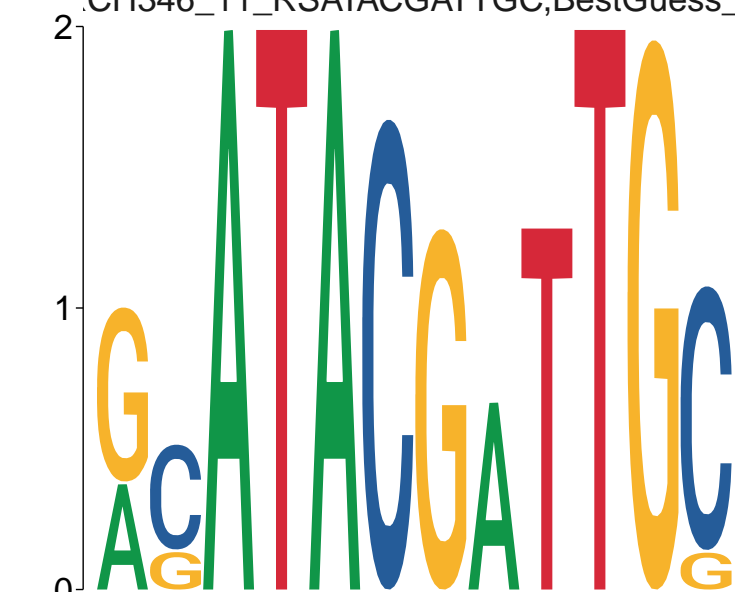
ARCH344_islike_FOXL1



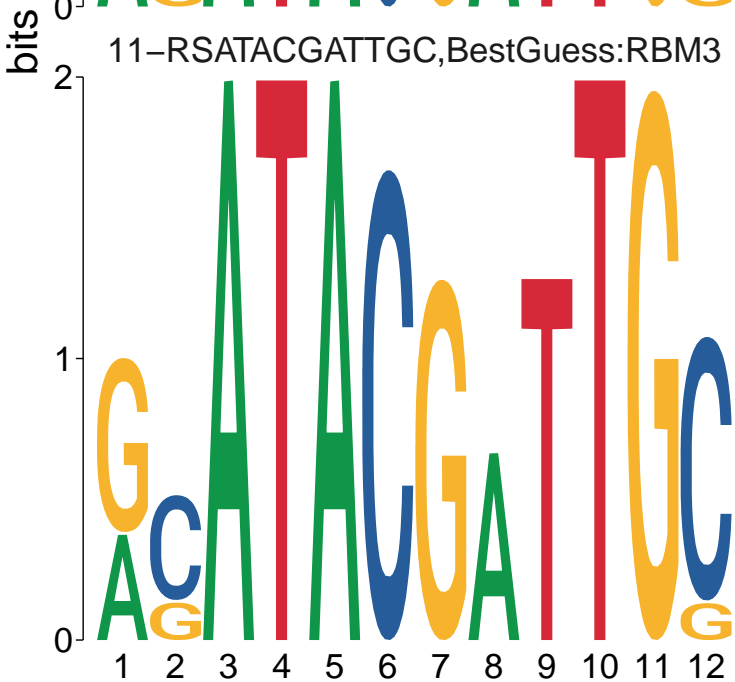
M00258_2.00



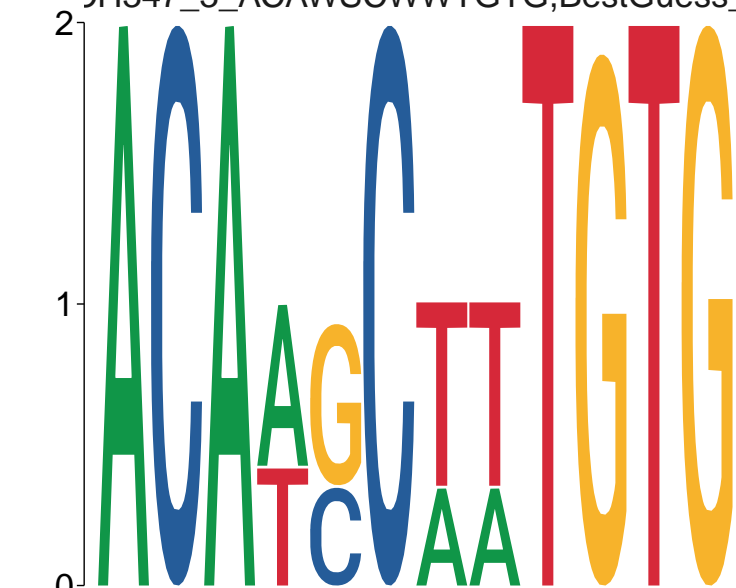
CH346_11_RSATACGATTGC,BestGuess_1



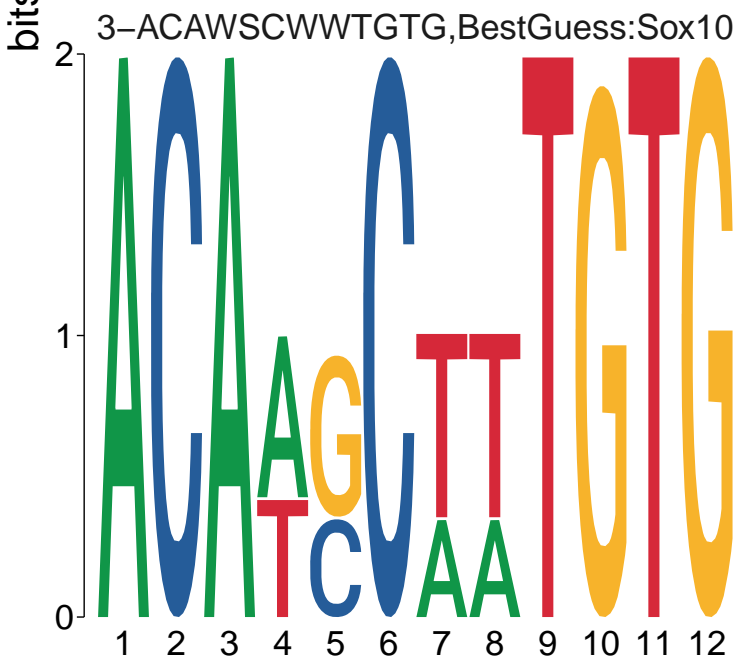
11-RSATACGATTGC,BestGuess:RBM3



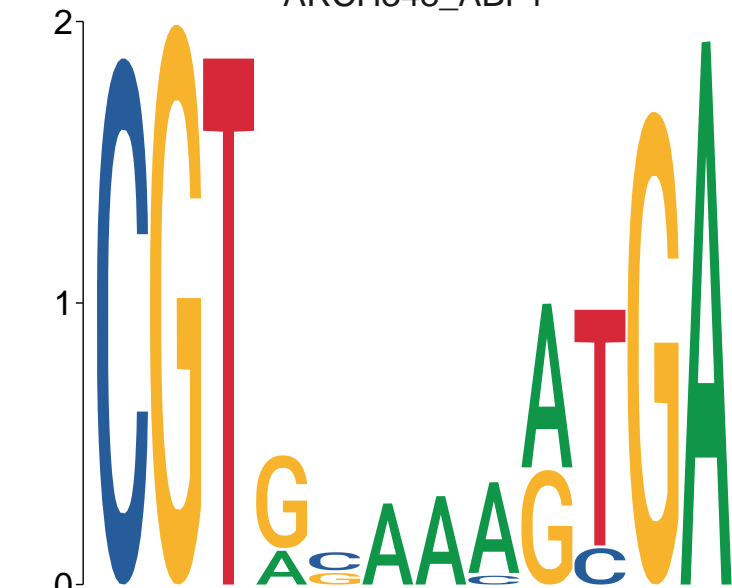
CH347_3_ACAWSCWWTGTG,BestGuess_



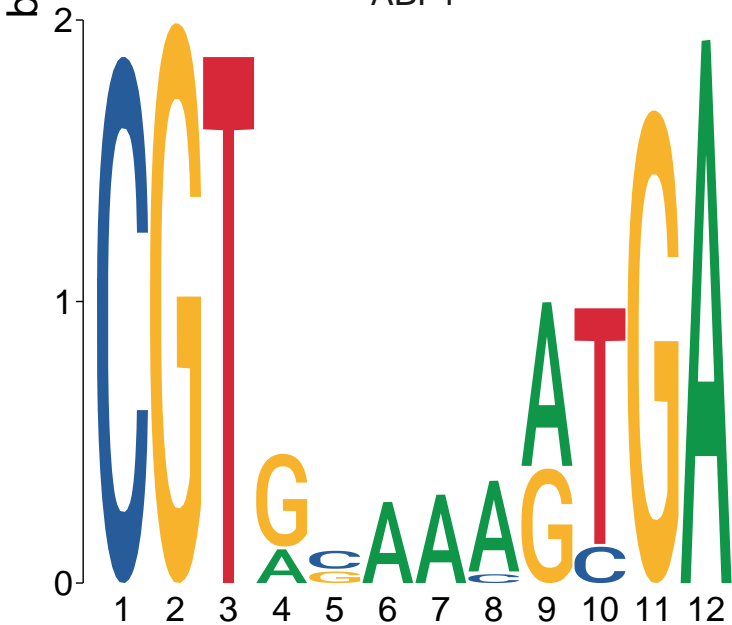
3-ACAWSCWWTGTG,BestGuess:Sox10



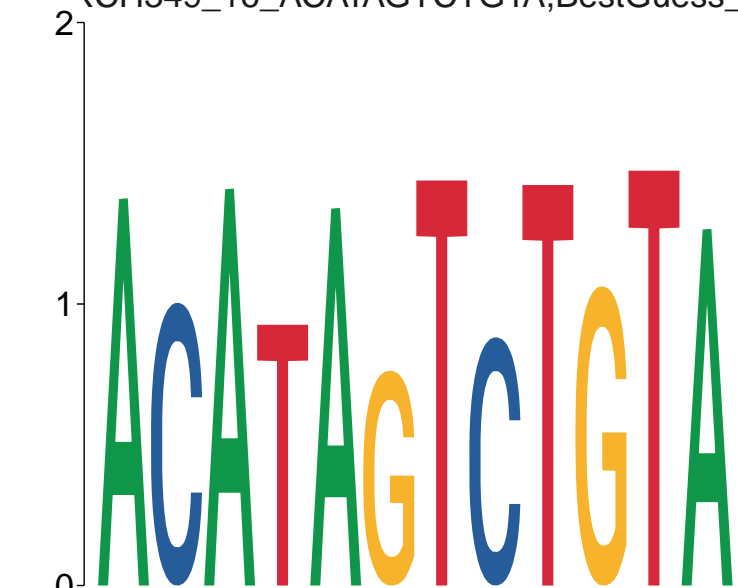
ARCH348_ABF1



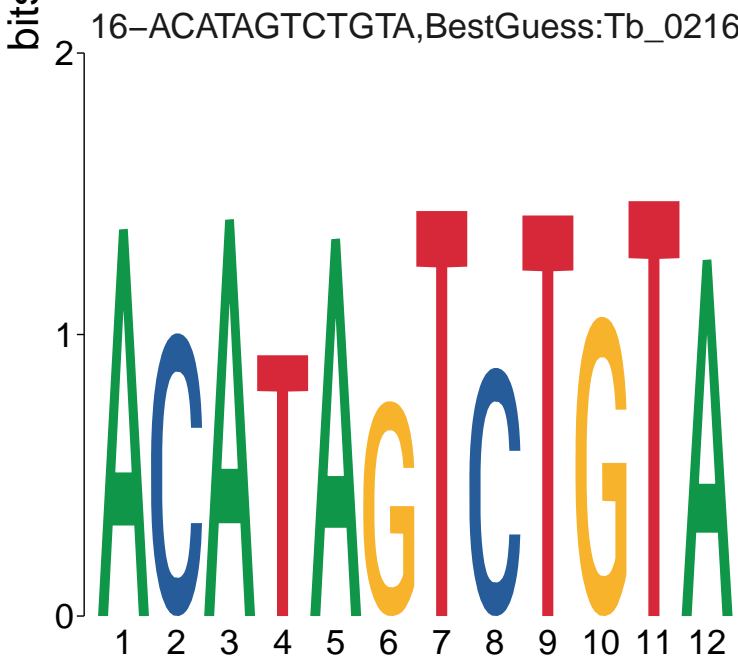
ABF1



16-ACATAGTCTGTA,BestGuess_



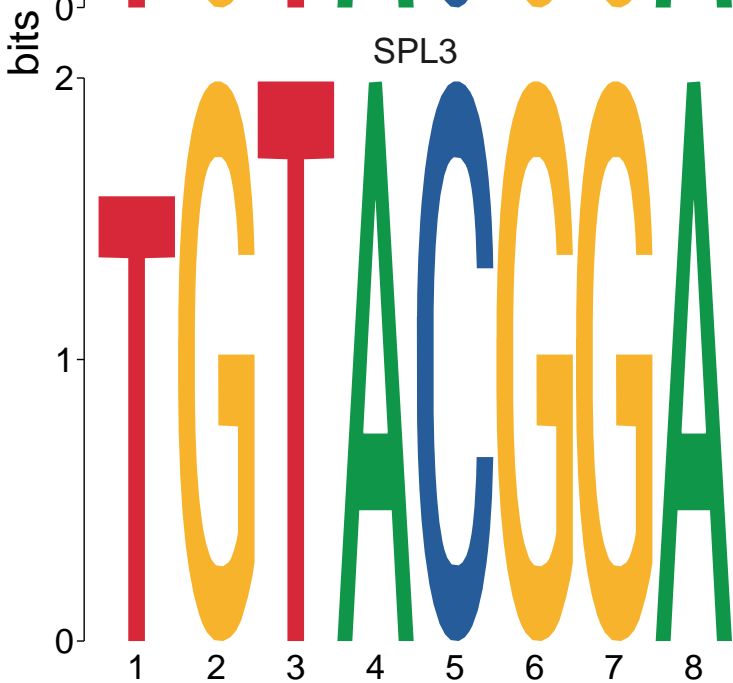
16-ACATAGTCTGTA,BestGuess:Tb_0216



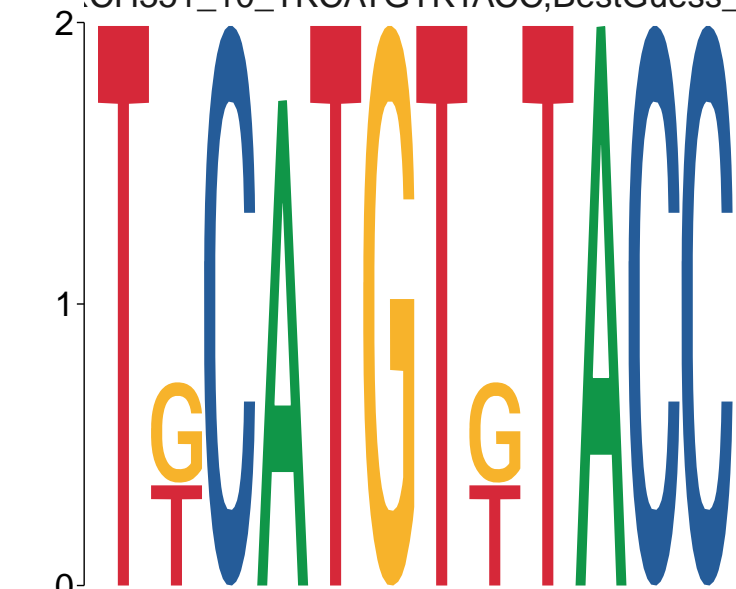
ARCH350_SPL3



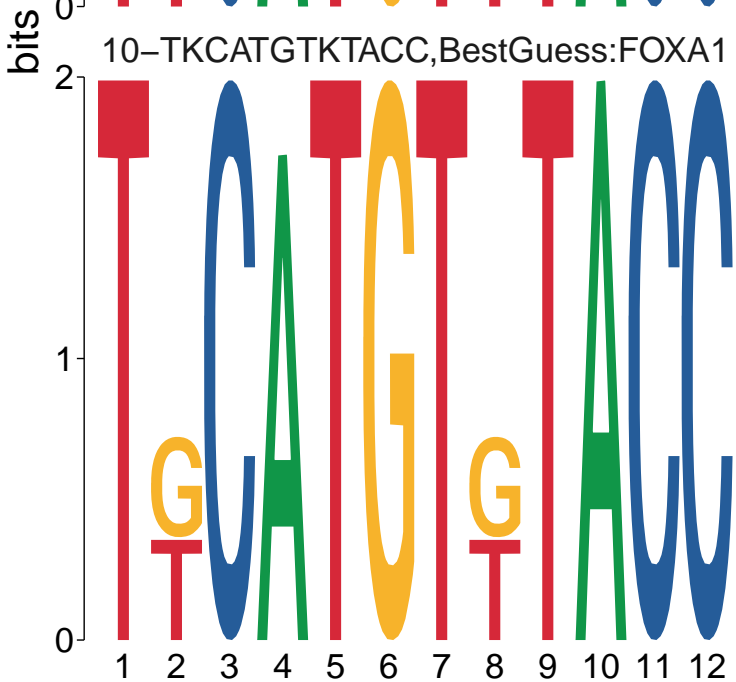
SPL3



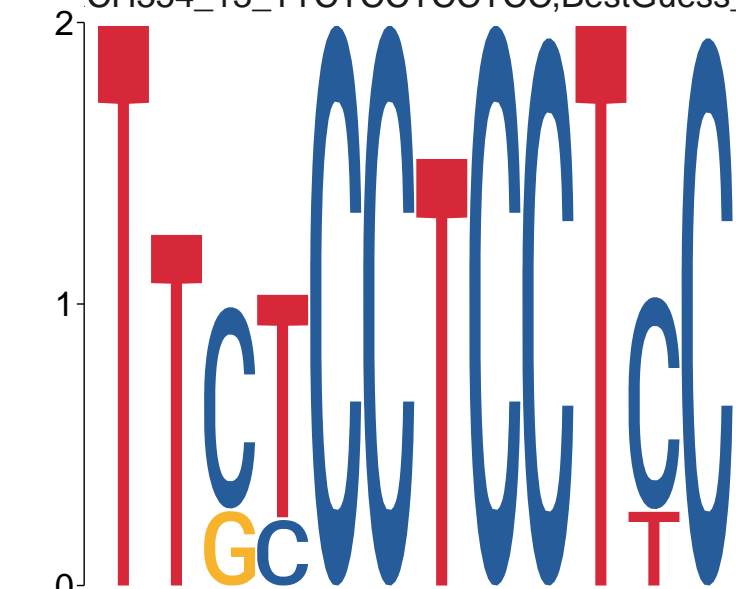
CH351_10_TKCATGTKTACC,BestGuess_I



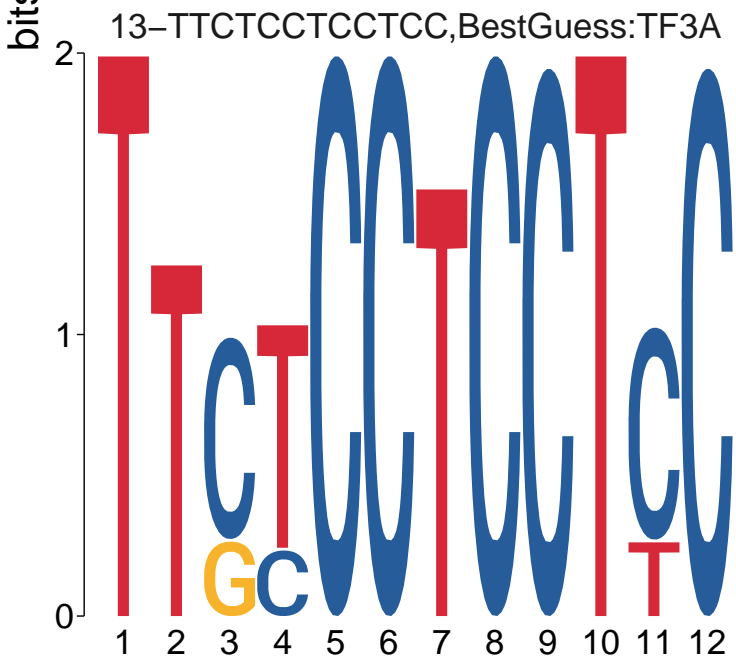
10-TKCATGTKTACC,BestGuess:FOXA1



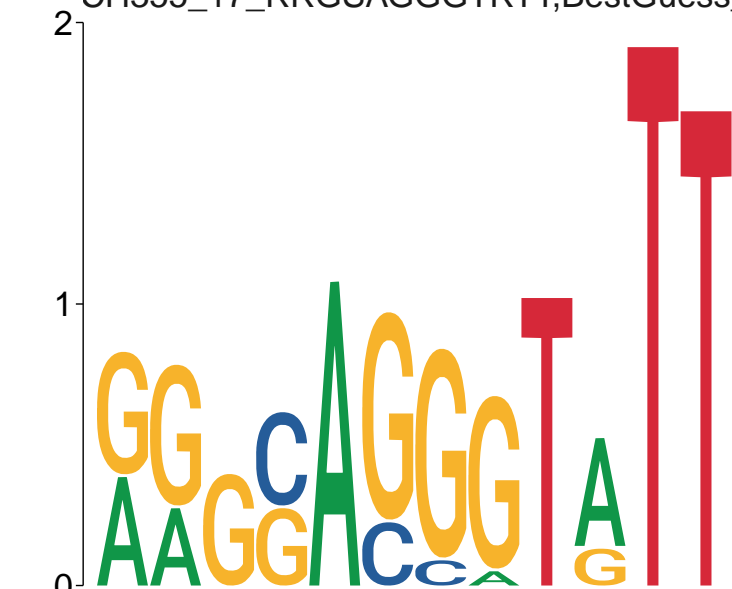
CH354_13_TTCTCCTCCTCC,BestGuess_



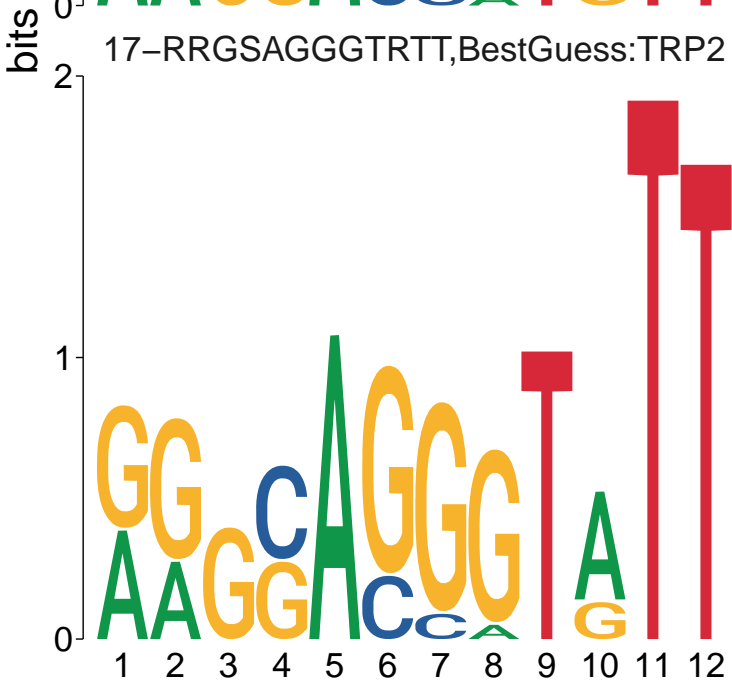
13-TTCTCCTCCTCC,BestGuess:TF3A



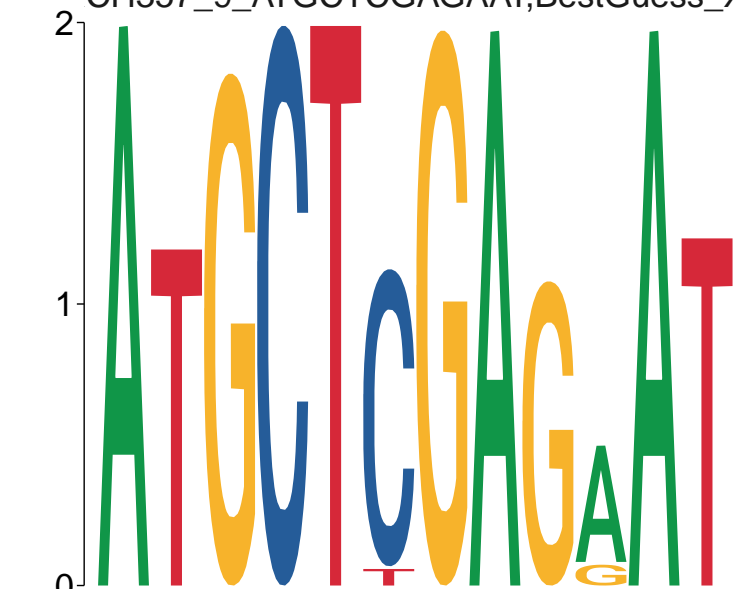
CH355_17_RRGSAAGGGTRTT,BestGuess_



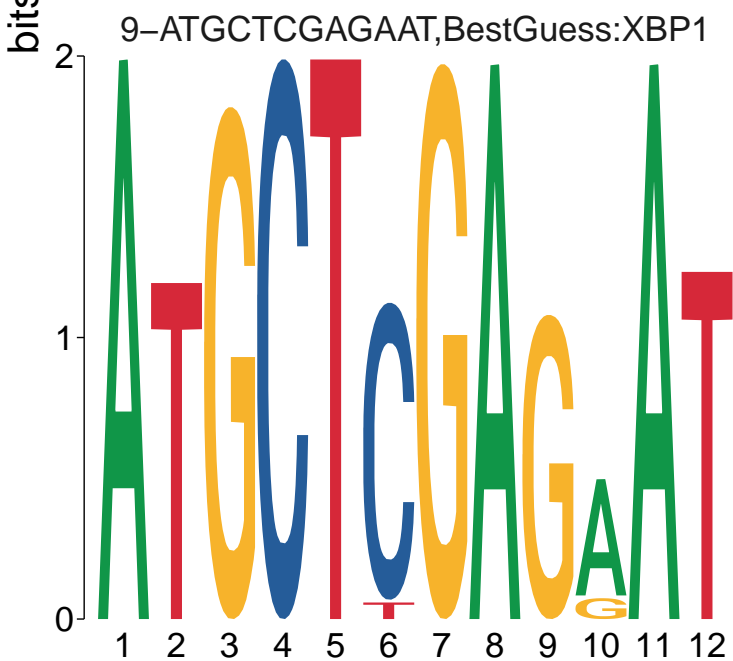
17-RRGSAGGGTRTT,BestGuess:TRP2



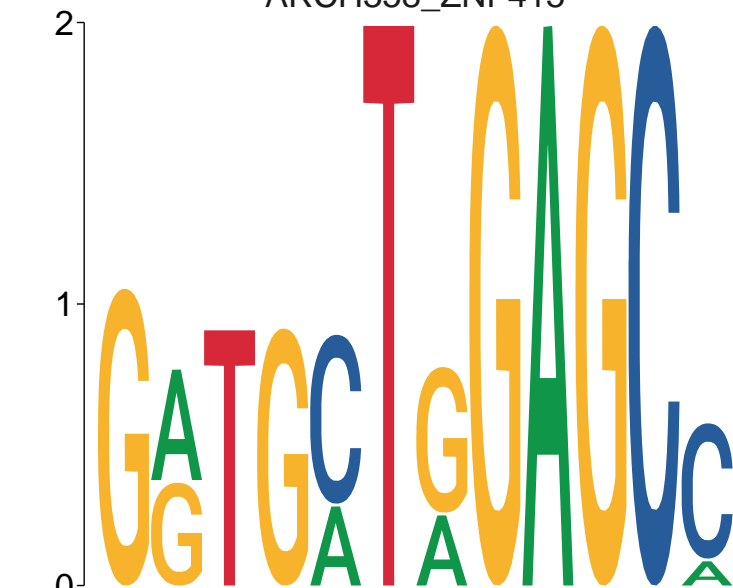
CH357_9_ATGCTCGAGAAT,BestGuess_X



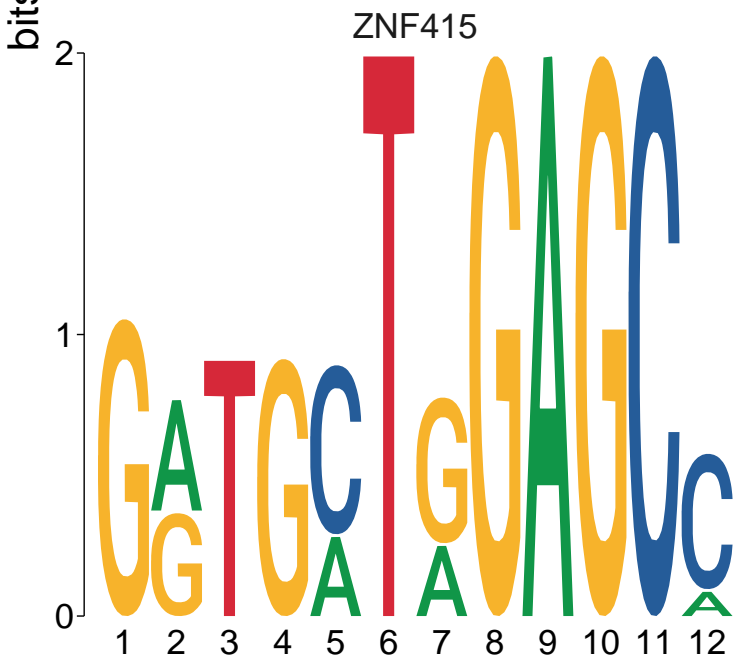
9-ATGCTCGAGAAT,BestGuess:XBP1



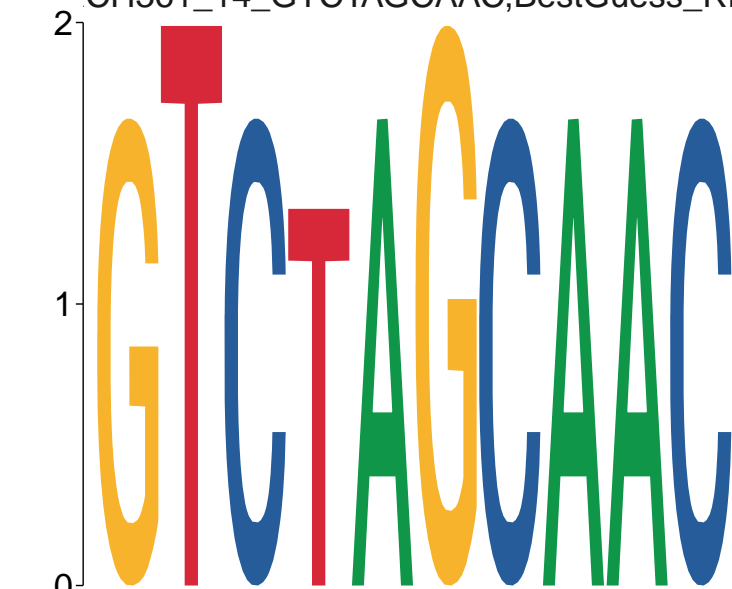
ARCH358_ZNF415



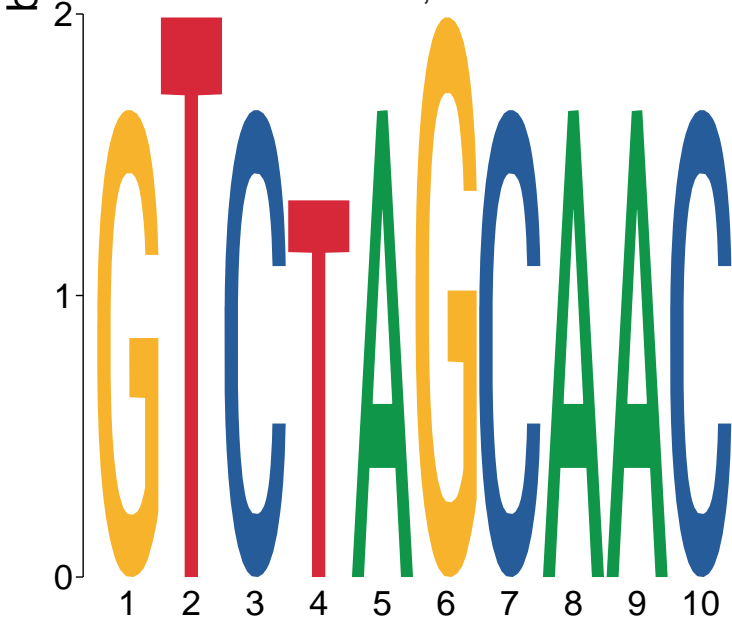
ZNF415



CH361_14_GTCTAGCAAC,BestGuess_RF



14-GTCTAGCAAC,BestGuess:RFX1



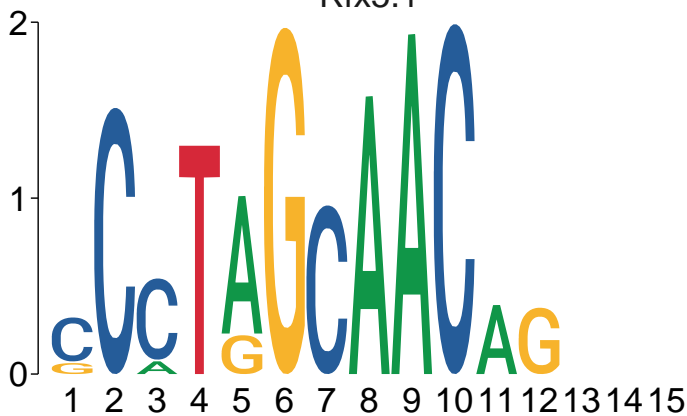
ARCH362_Rfx5_Rfx5



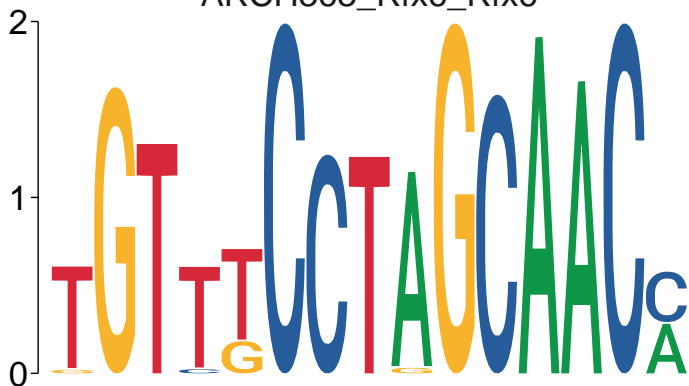
Rfx5



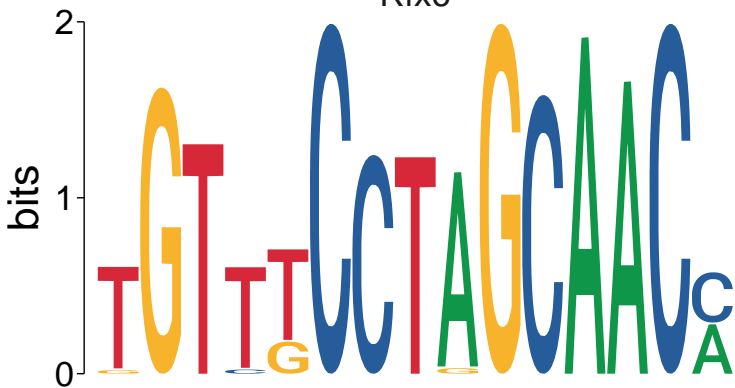
Rfx5.1



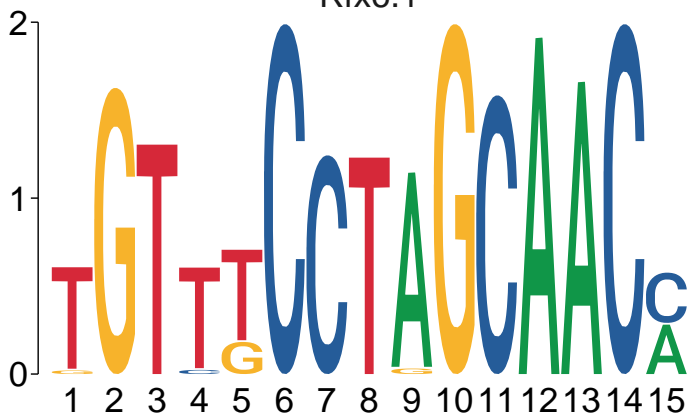
ARCH363_Rfx6_Rfx6



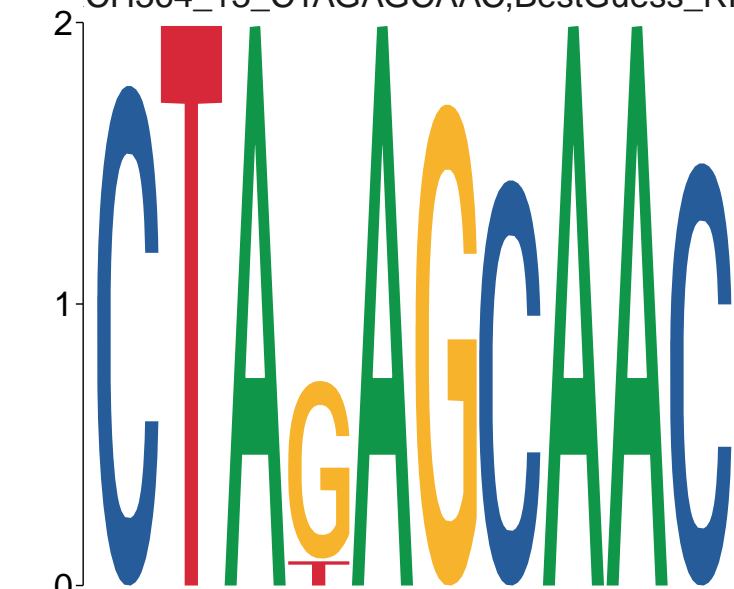
Rfx6



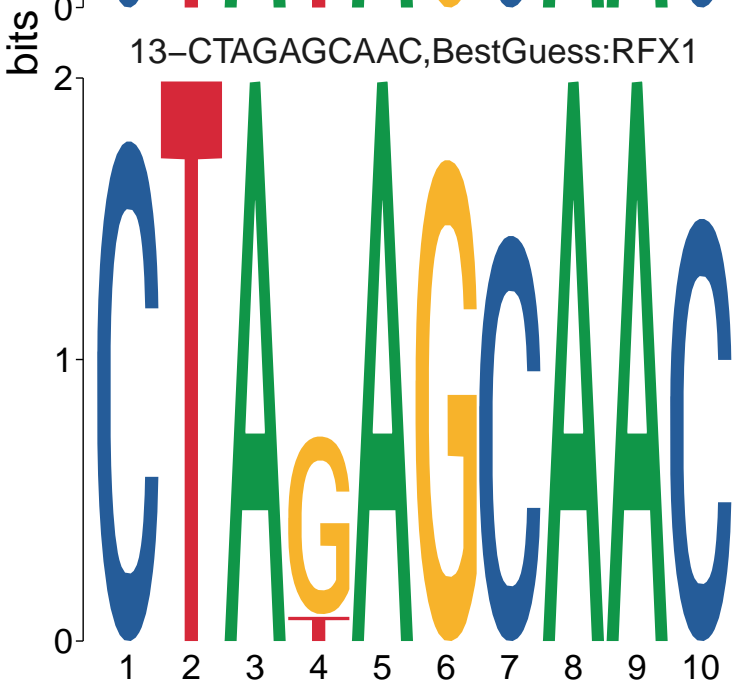
Rfx6.1



CH364_13_CTAGAGCAAC,BestGuess_RF



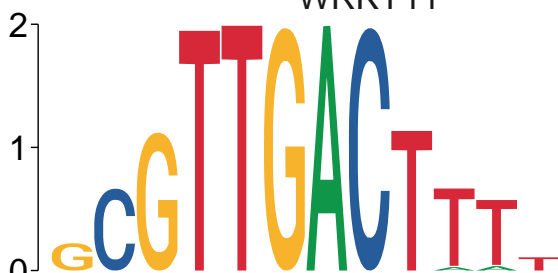
13-CTAGAGCAAC,BestGuess:RFX1



ARCH367_WRKY46_WRKY22_WRKY11



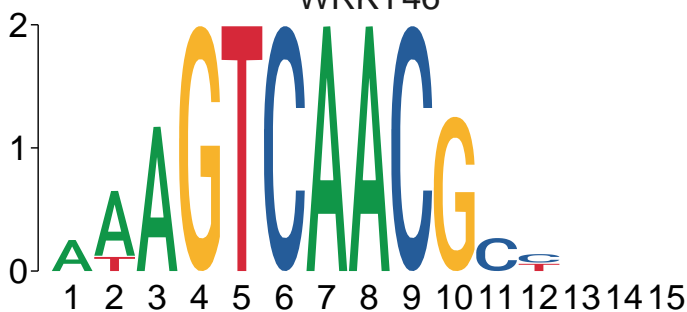
WRKY11



WRKY22



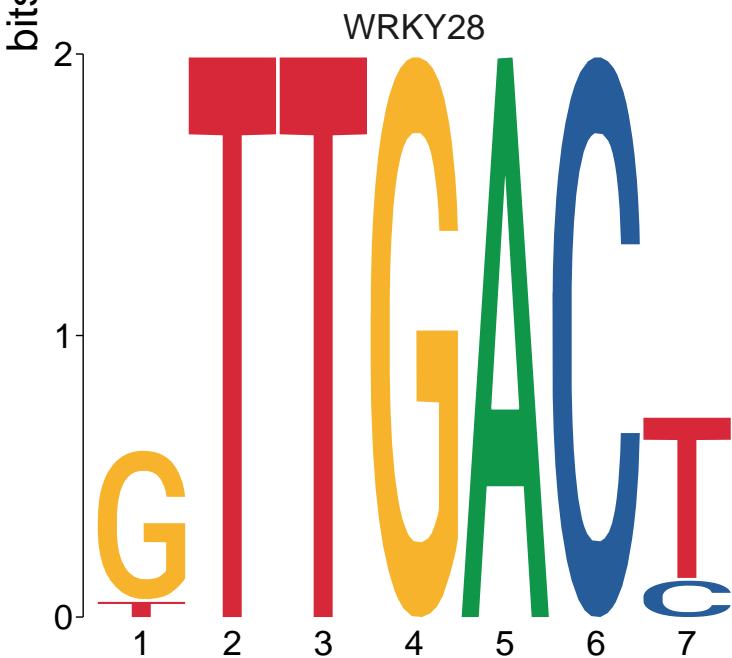
WRKY46



ARCH368_WRKY28



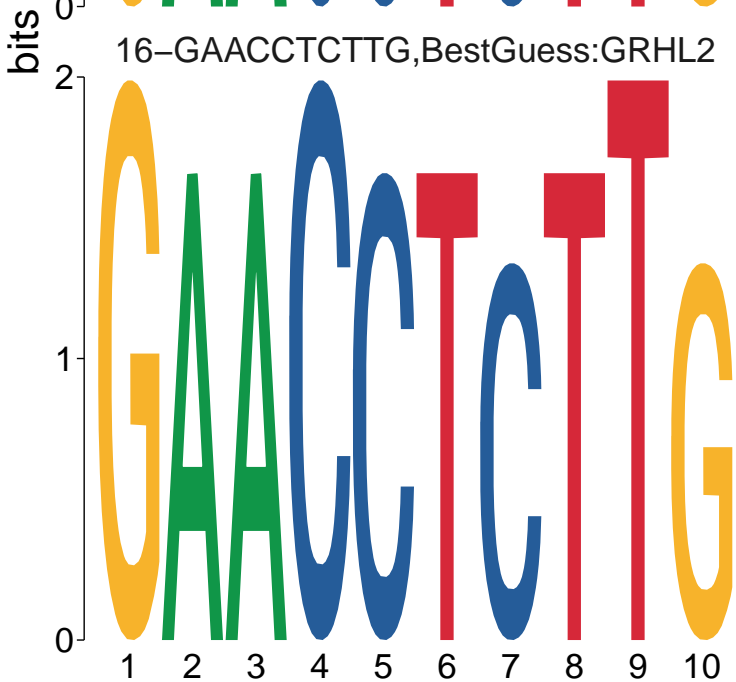
WRKY28



CH369_16_GAACCTCTTG,BestGuess_GR



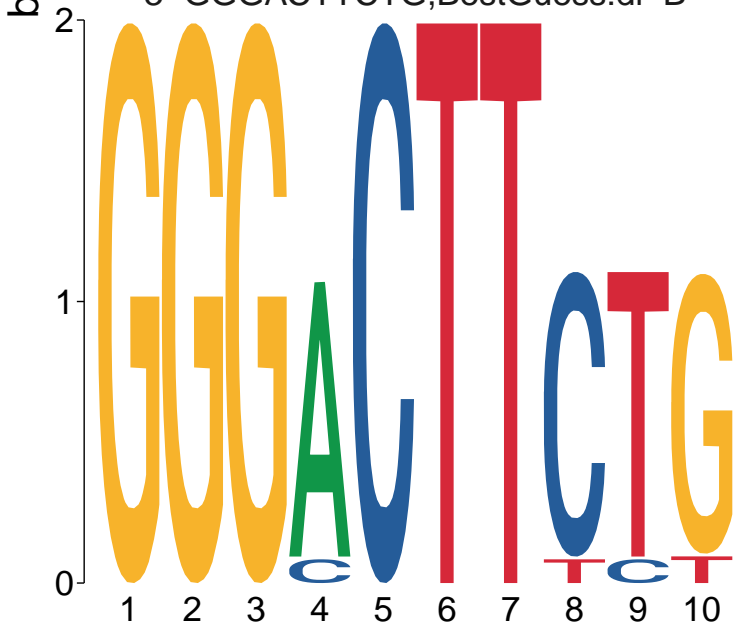
16-GAACCTCTTG,BestGuess:GRHL2



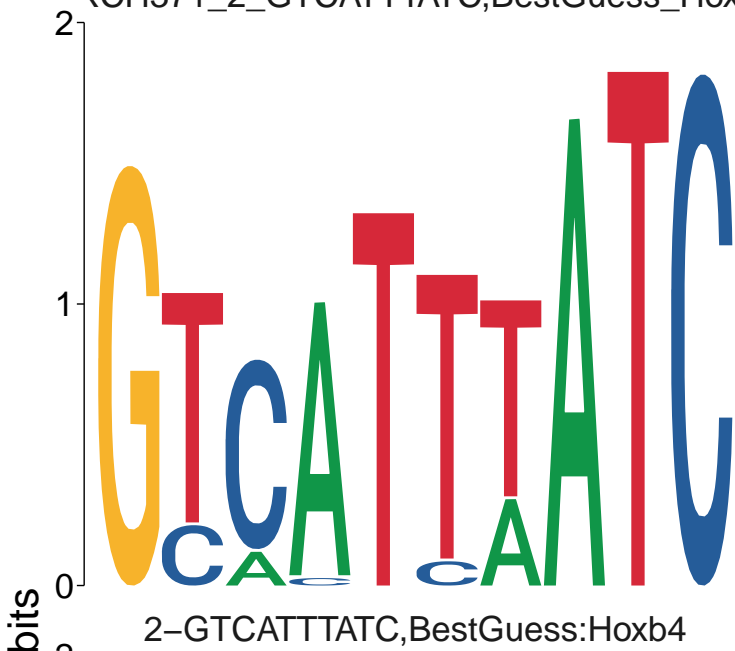
RCH370_3_GGGACTTCTG,BestGuess_dl_



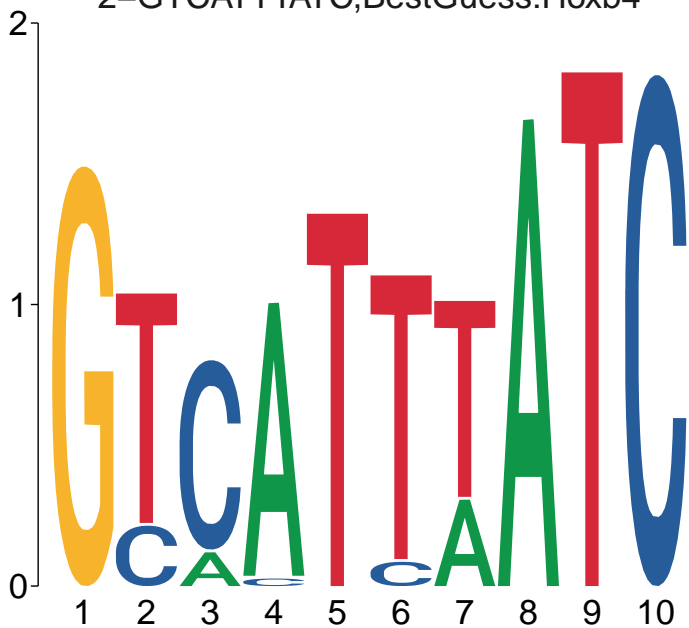
3-GGGACTTCTG,BestGuess:dl-B



RCH371_2_GTCATTTATC,BestGuess_Hoxl



2-GTCATTTATC,BestGuess:Hoxb4



ARCH372_LIN_39_Pdx1_Pdx1_LIN_39



LIN-39



Pdx1 [RC]



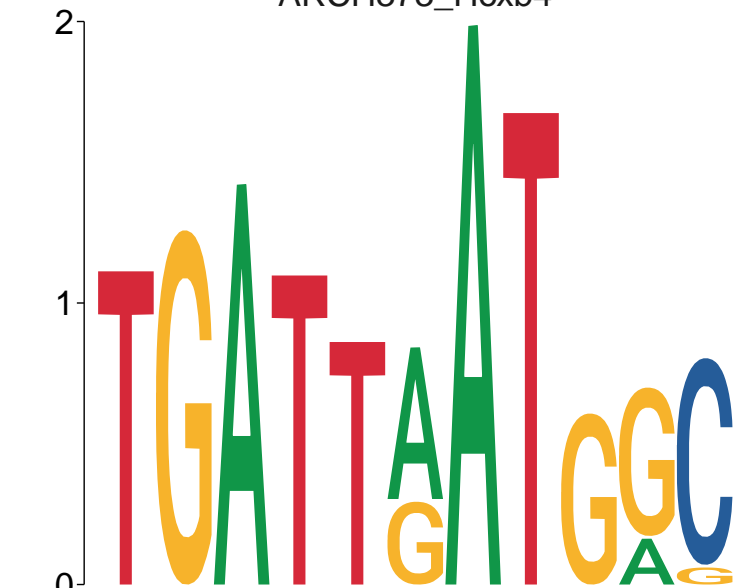
Pdx1.1 [RC]



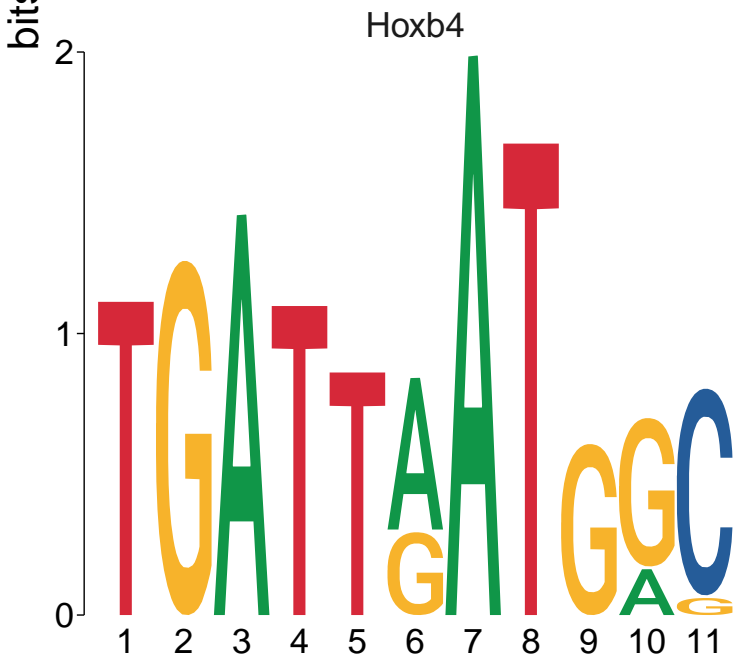
LIN-39.1



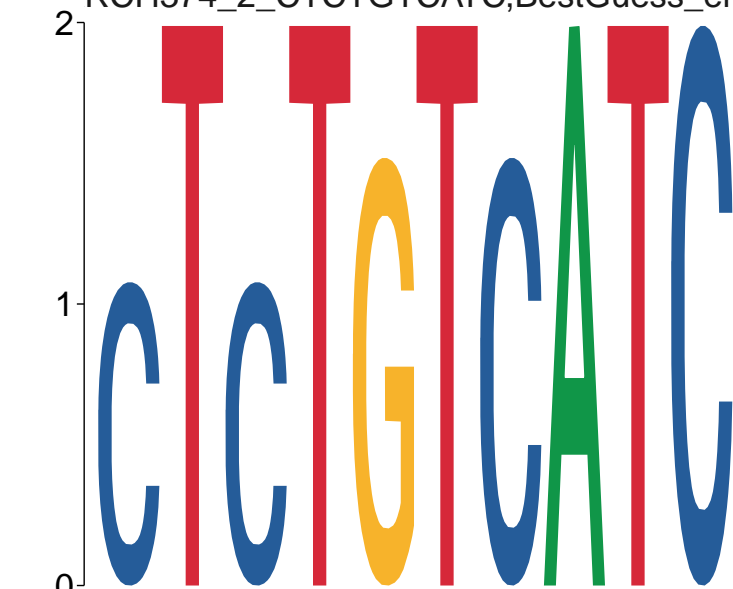
ARCH373_Hoxb4



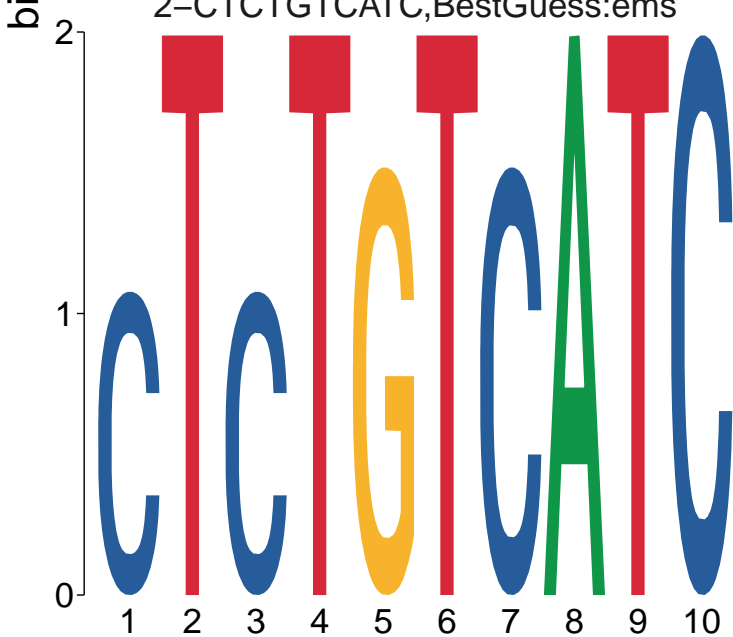
Hoxb4



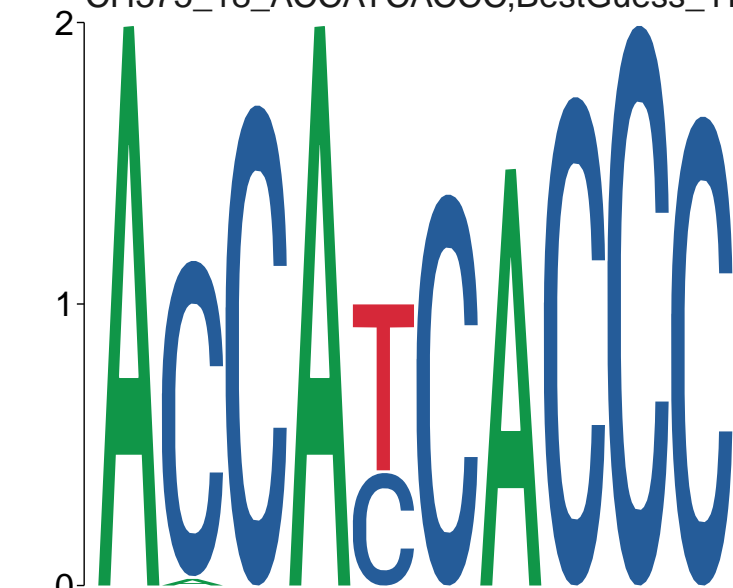
RCH374_2_CTCTGTCATC,BestGuess_em



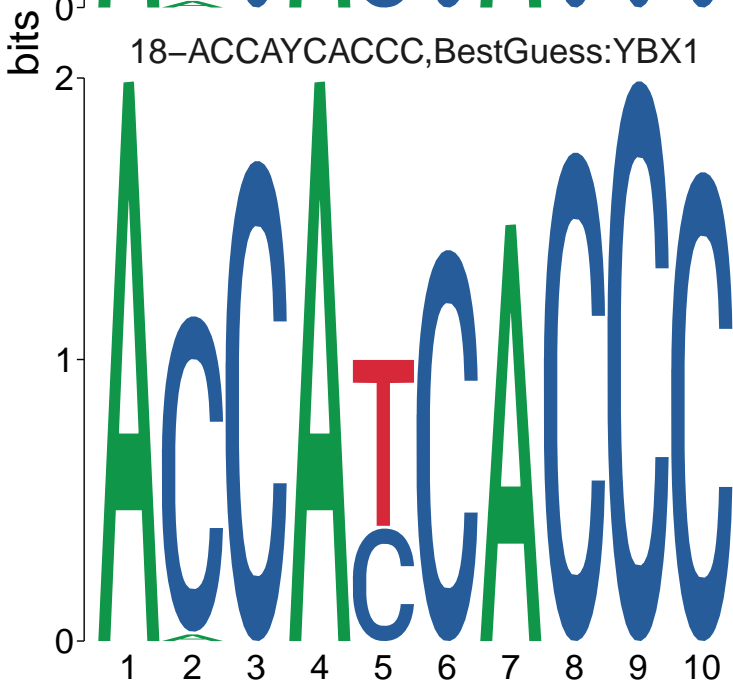
2-CTCTGTCATC,BestGuess:ems



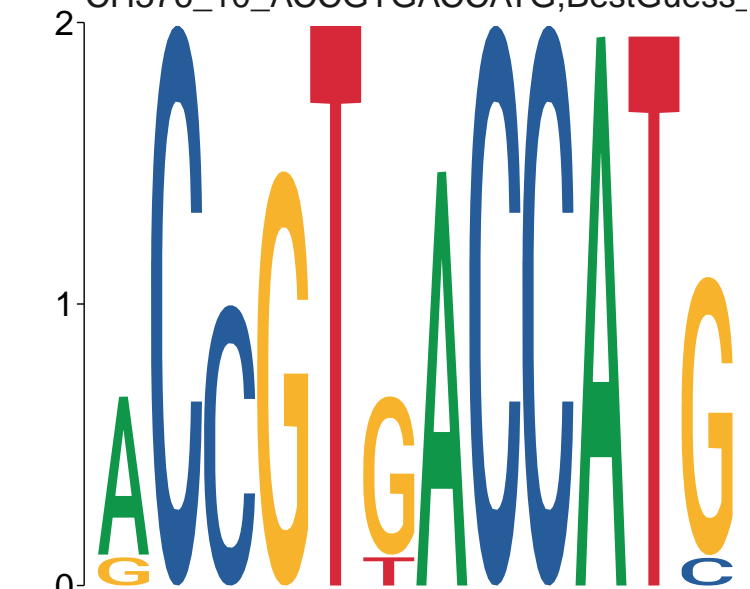
CH375_18_ACCAYCACCC,BestGuess_YB



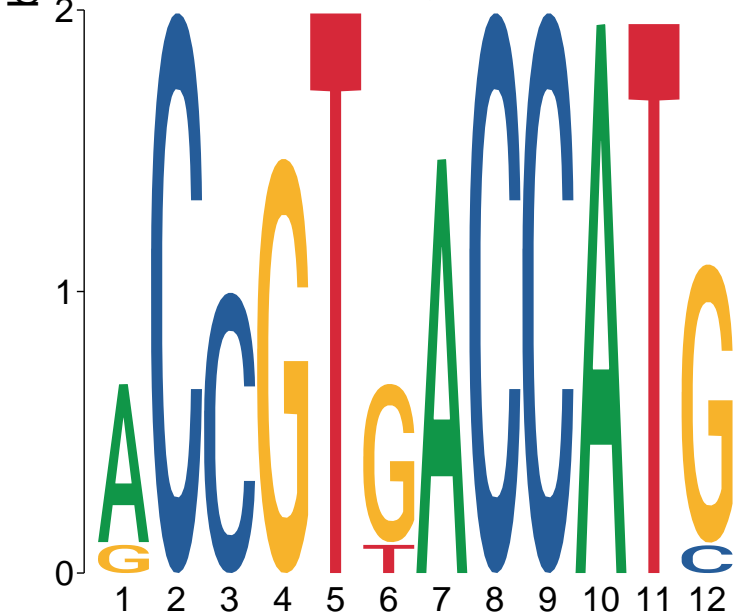
18-ACCAYCACCC,BestGuess:YBX1



CH376_10_ACCGTGACCATG,BestGuess_



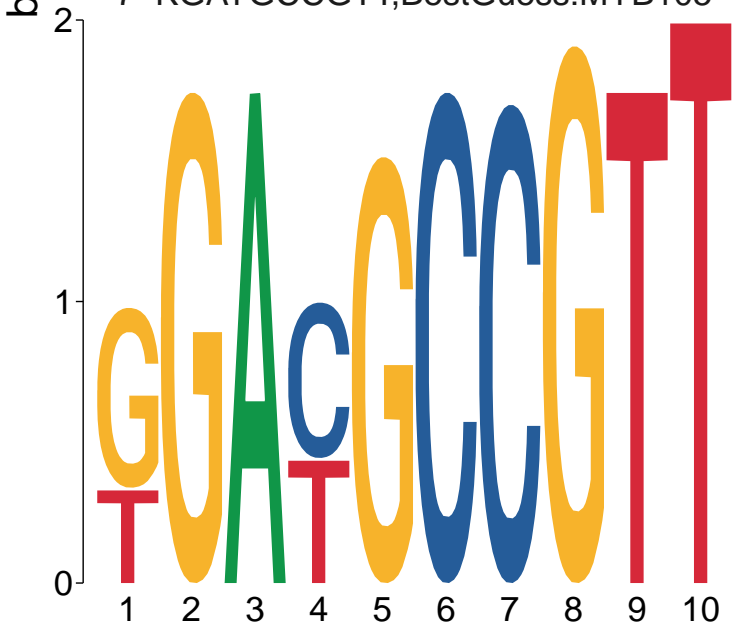
10-ACCGTGACCATG,BestGuess:Tv_0259



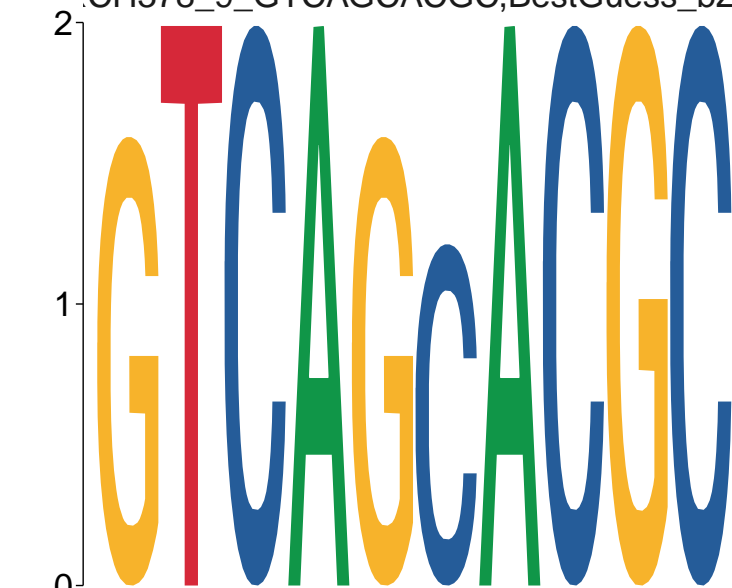
CH377_7_KGAYGCCGTT,BestGuess_MYE



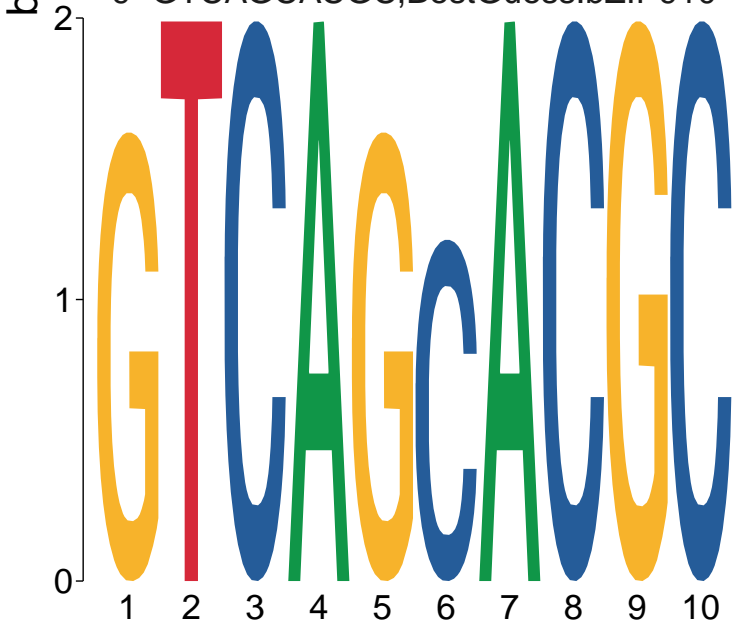
7-KGAYGCCGTT,BestGuess:MYB105



CH378_9_GTCAGCACGC,BestGuess_bZI



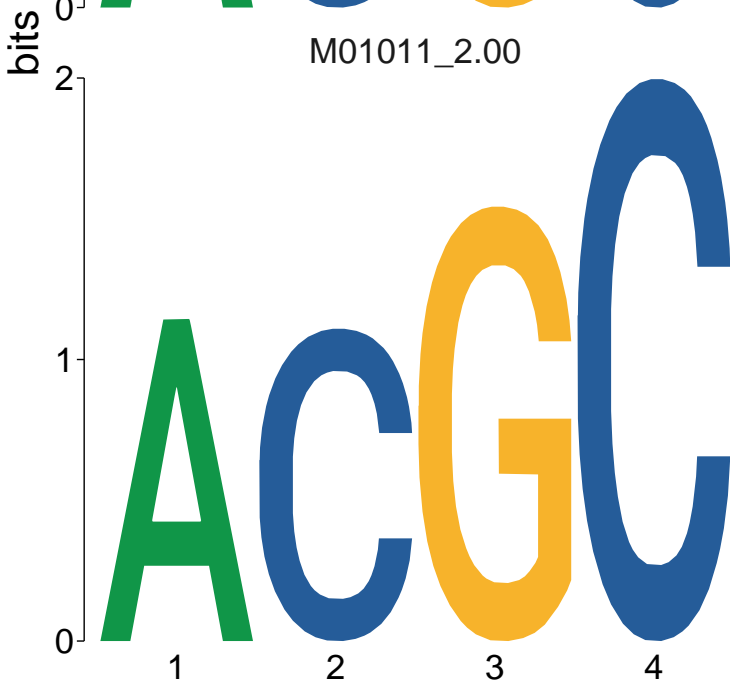
9-GTCAGCACGC,BestGuess:bZIP910



ARCH379_FOXN1_FOXN4_FOXR1



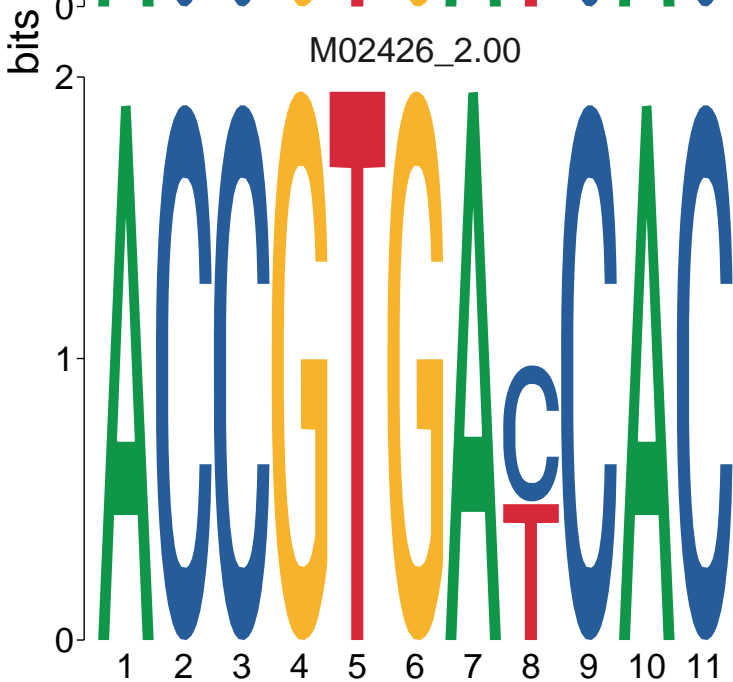
M01011_2.00



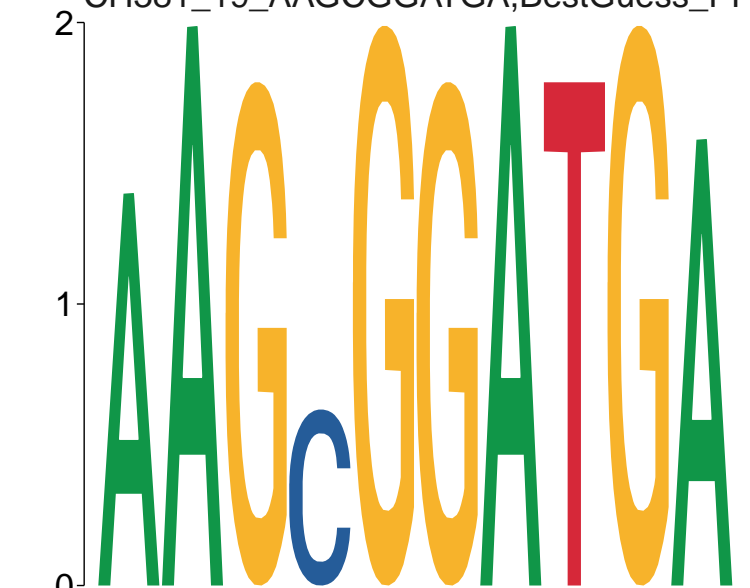
ARCH380_Homeodomains



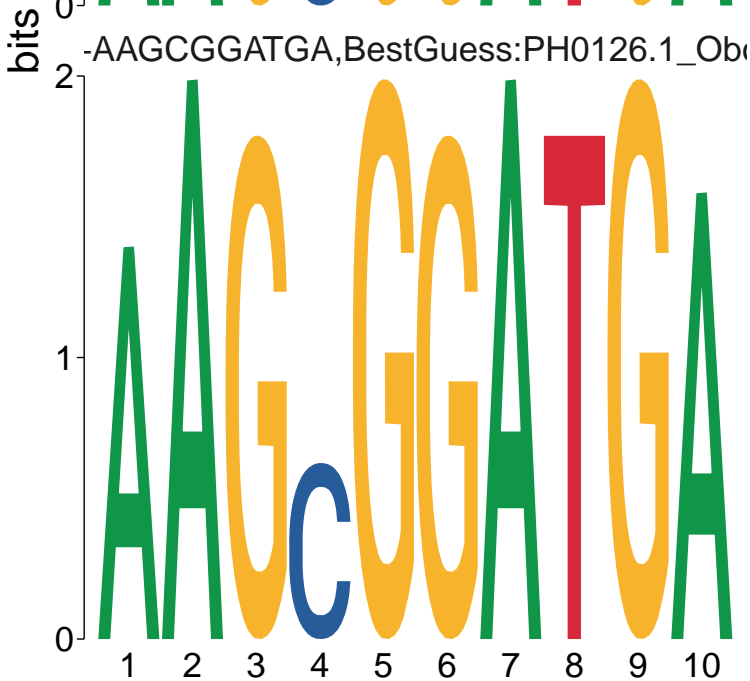
M02426_2.00



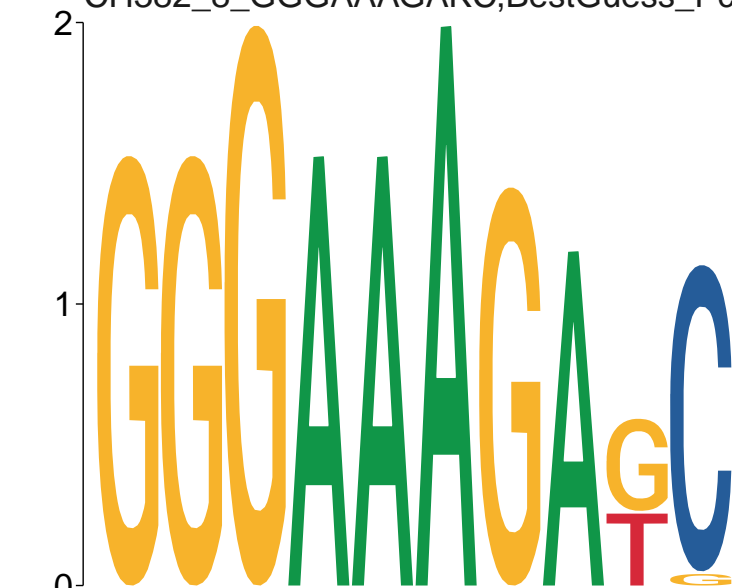
CH381_19_AAGCGGATGA,BestGuess_PT



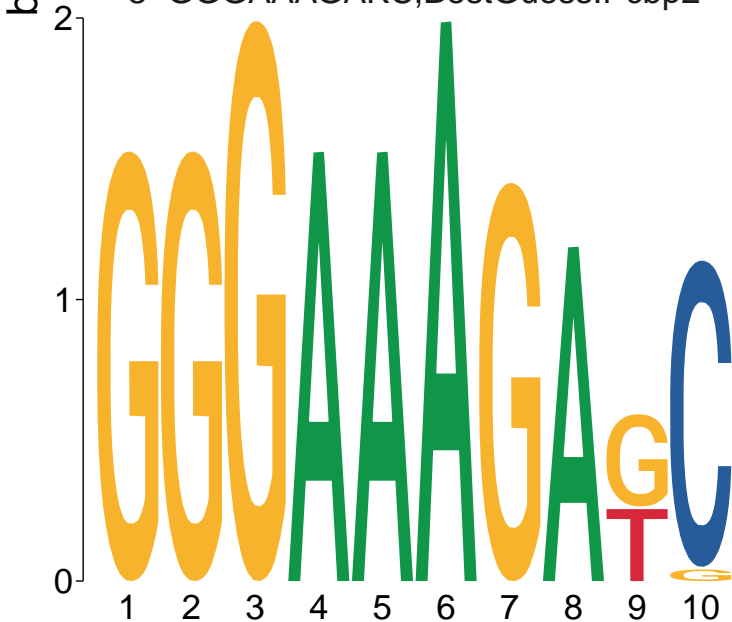
-AAGCGGATGA,BestGuess:PH0126.1_Ob



CH382_8_GGGAAAGAKC,BestGuess_Pck



8-GGGAAAGAKC,BestGuess:Pcbp2



RCH383_1_RAGMSGATTT,BestGuess_Cr

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1-RAGMSGATTT,BestGuess:Crx

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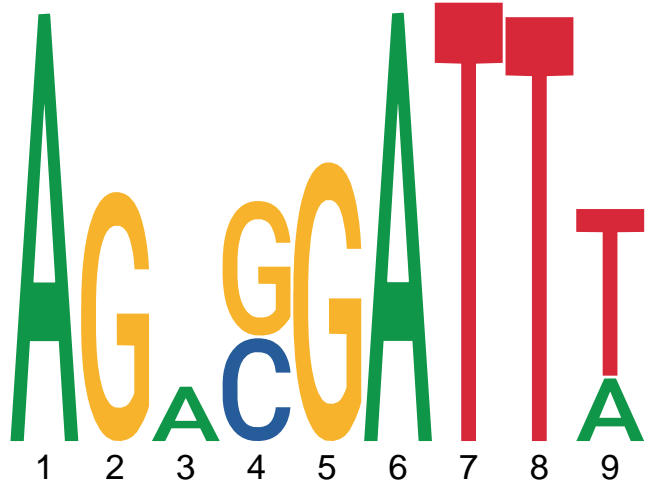
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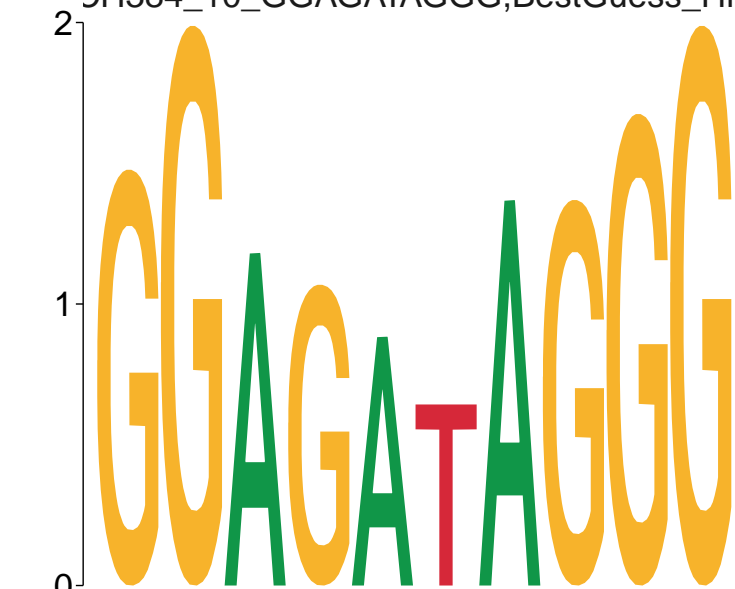
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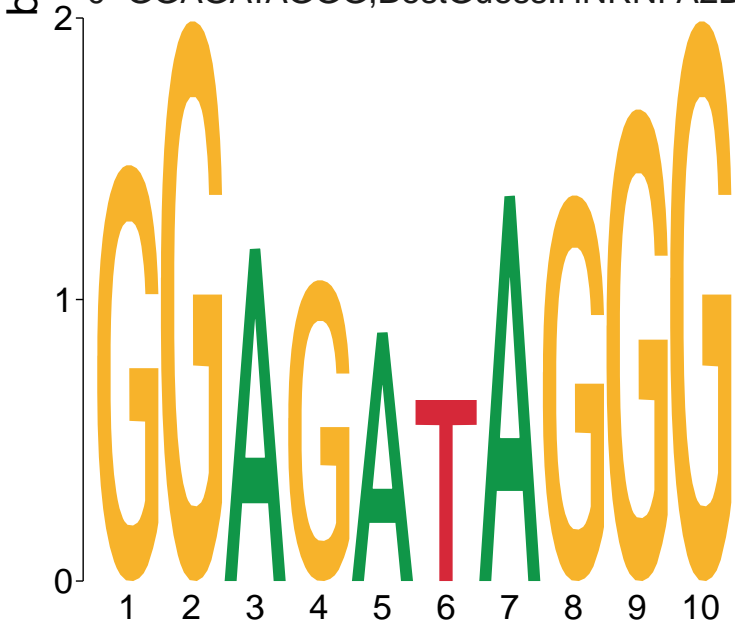
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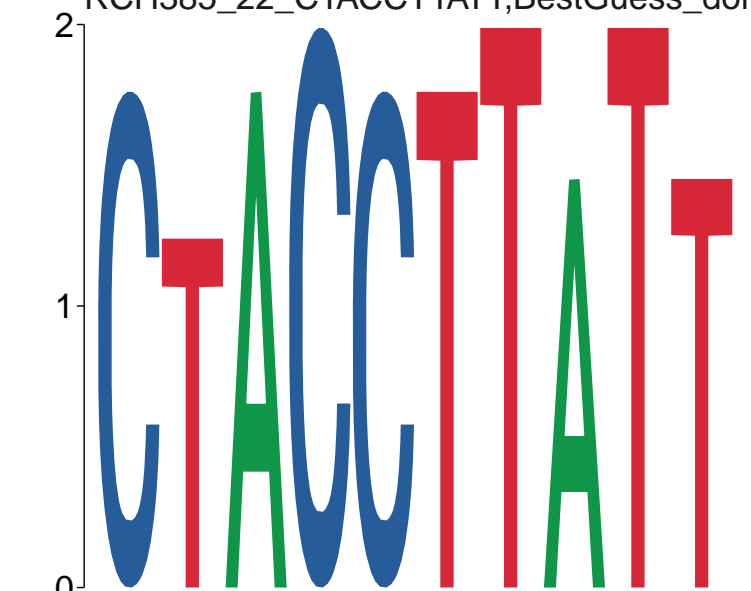
CH384_10_GGAGATAGGG, BestGuess_HN



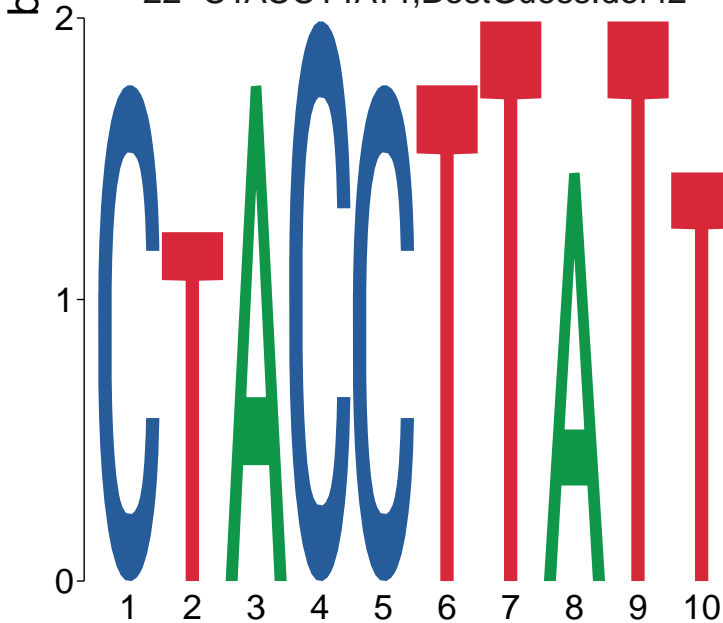
0-GGAGATAGGG, BestGuess: HNRNPA2B



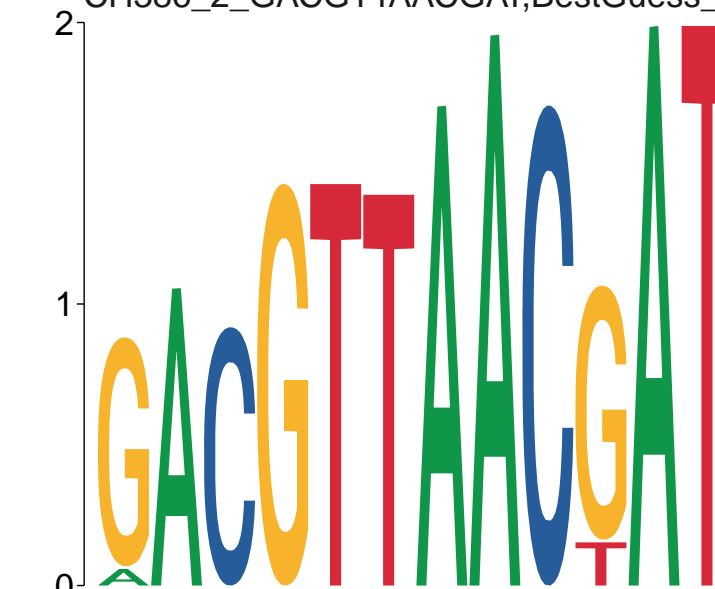
RCH385_22_CTACCTTATT,BestGuess_dof



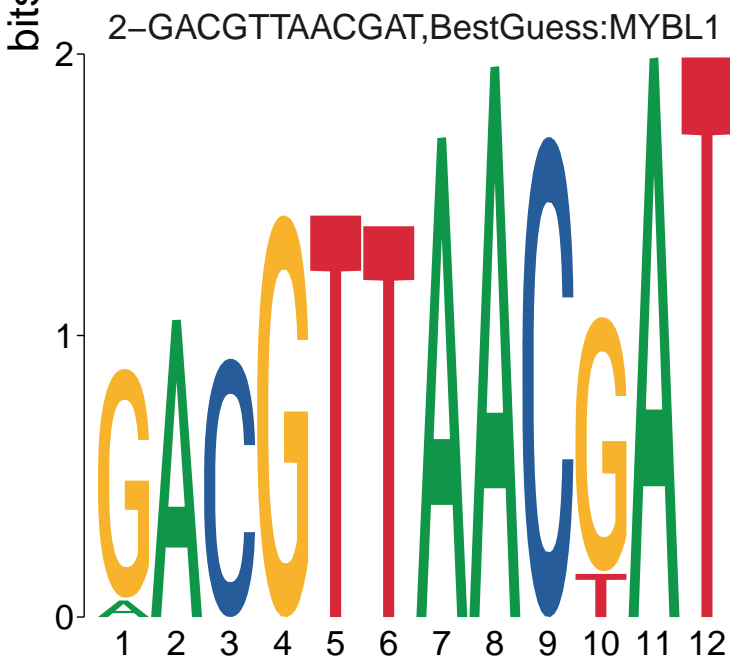
22-CTACCTTATT,BestGuess:dof42



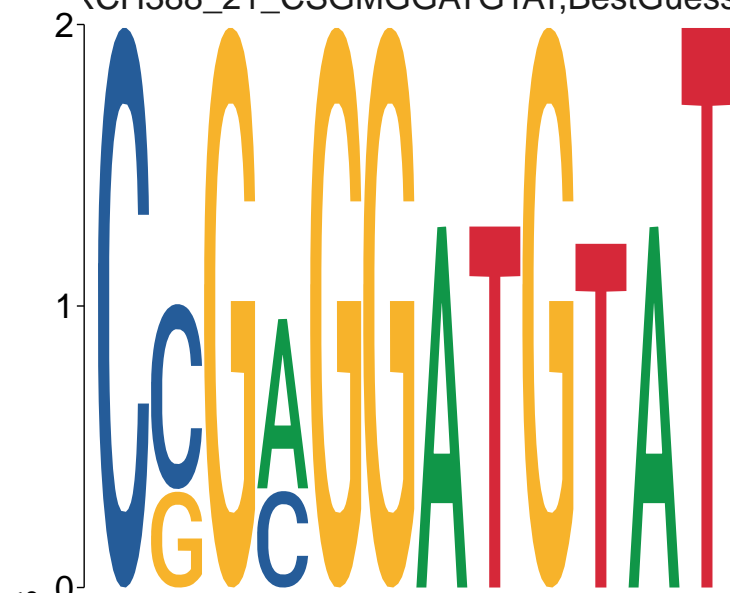
CH386_2_GACGTTAACGAT,BestGuess_M



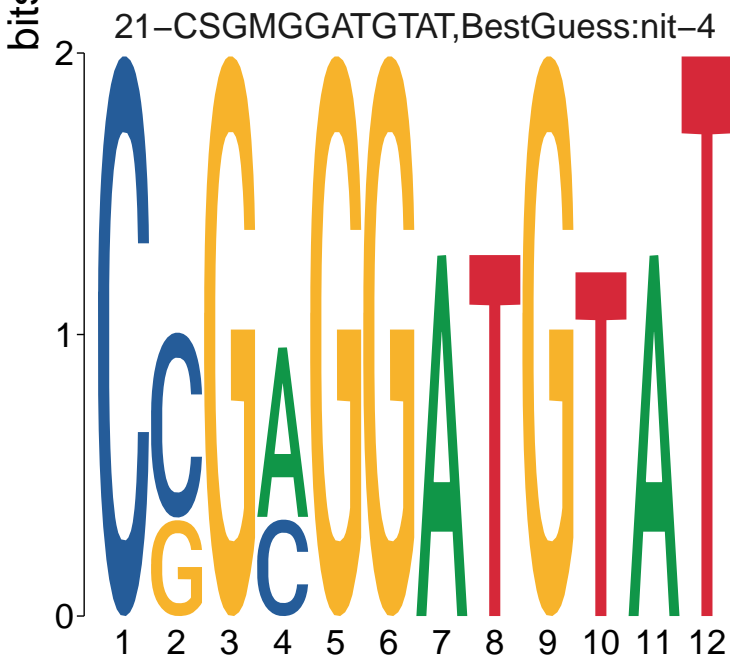
2-GACGTTAACGAT,BestGuess:MYBL1



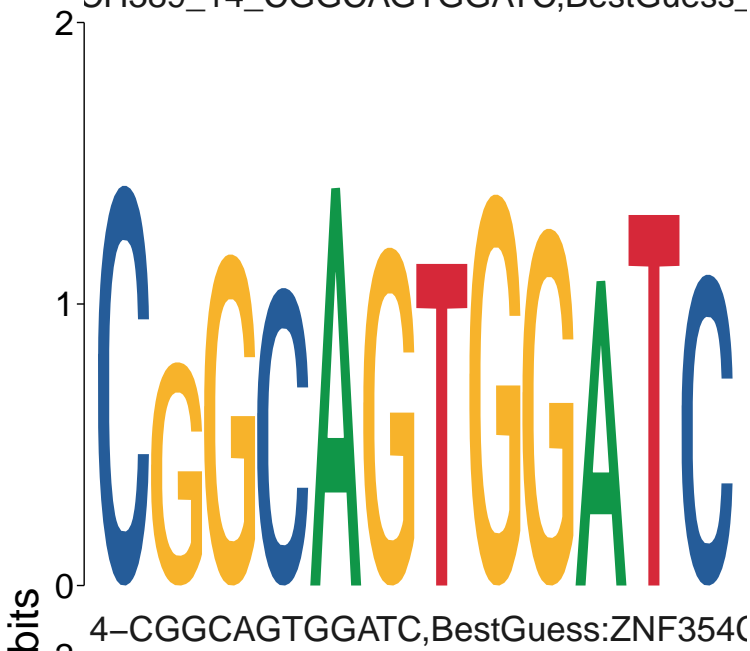
21-CH388_21_CSGMGGATGTAT,BestGuess_



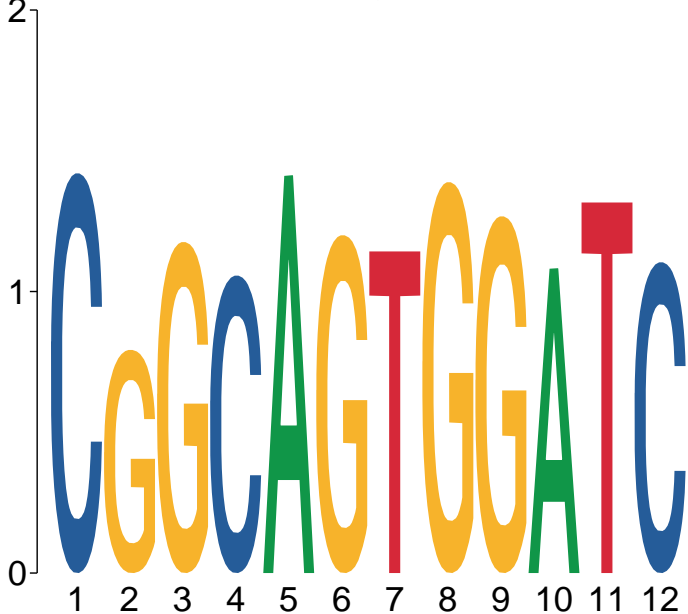
21-CSGMGGATGTAT,BestGuess:nit-4



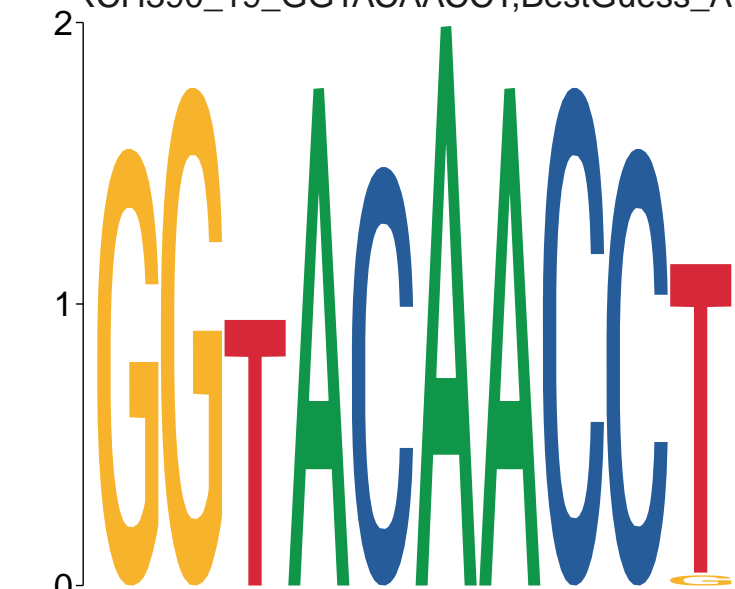
CH389_14_CGGCAGTGGATC,BestGuess_



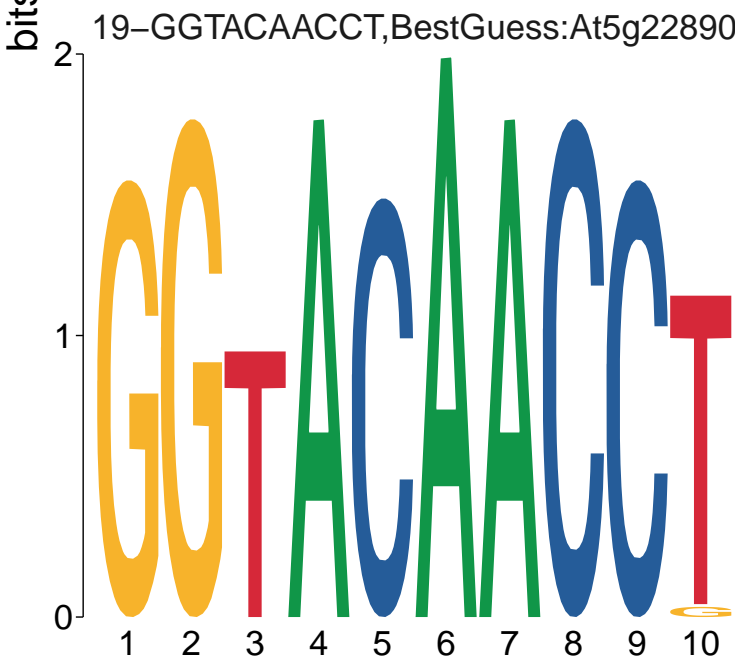
4-CGGCAGTGGATC,BestGuess:ZNF354C



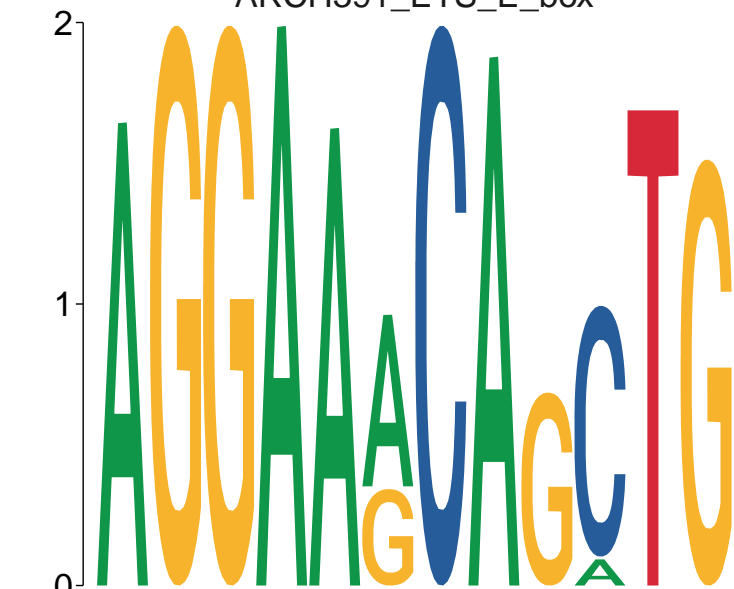
RCH390_19_GGTACAACCT,BestGuess_At5g22890



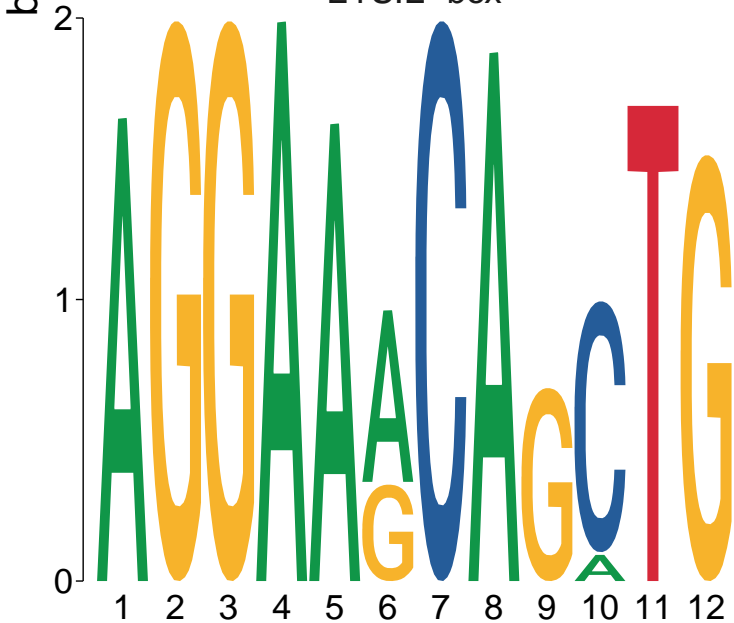
19-GGTACAACCT,BestGuess:At5g22890

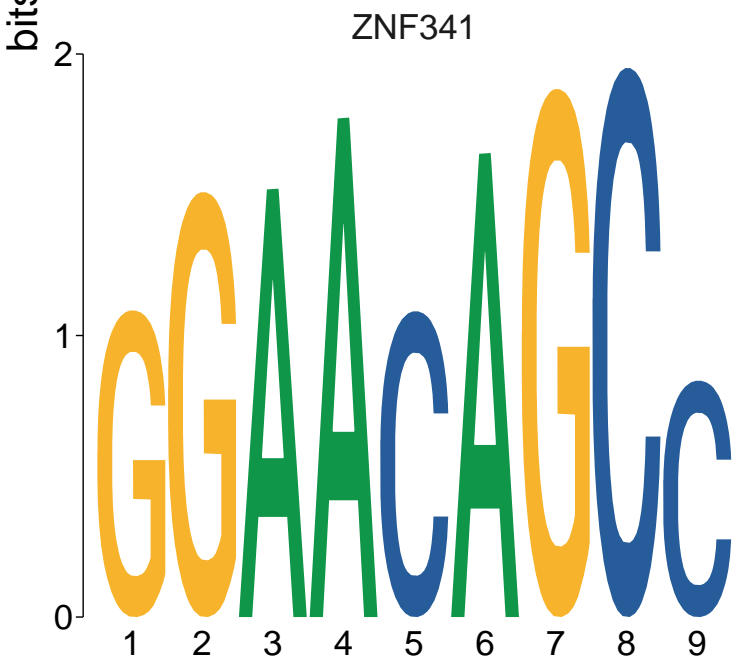
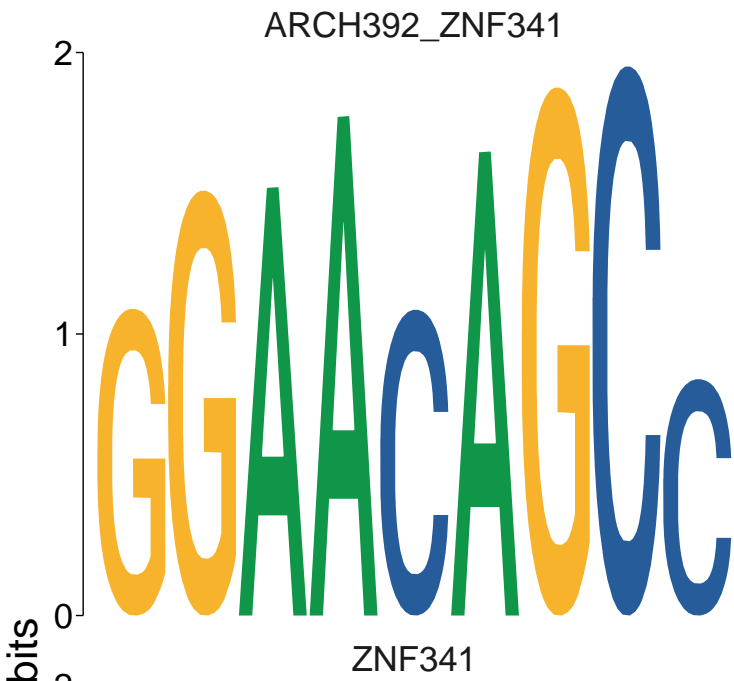


ARCH391_ETS_E_box

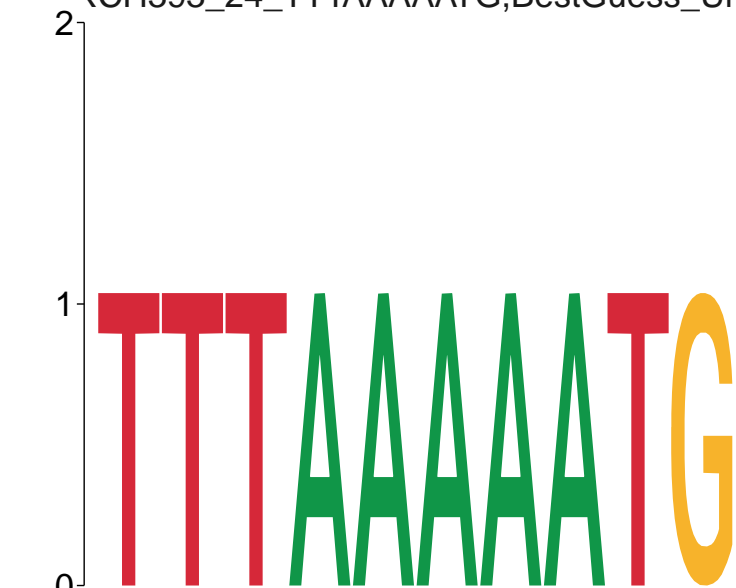


ETS:E-box

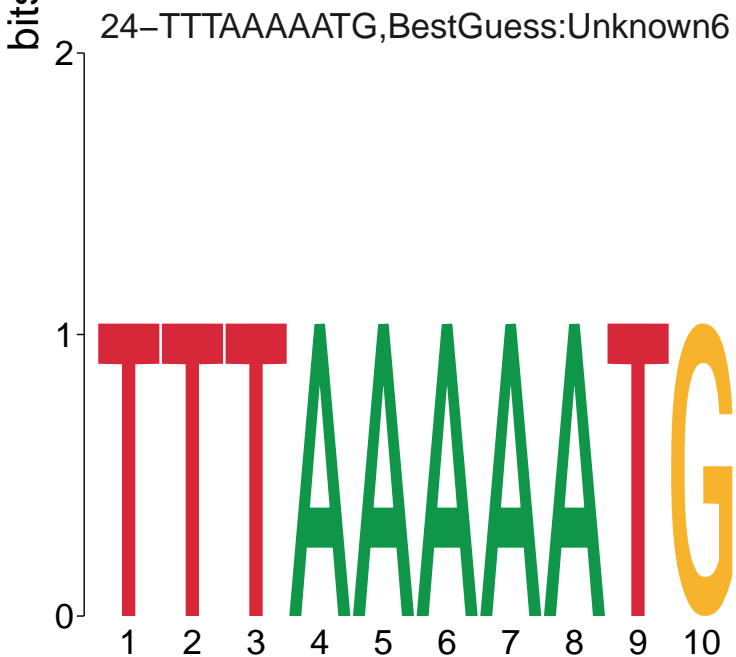




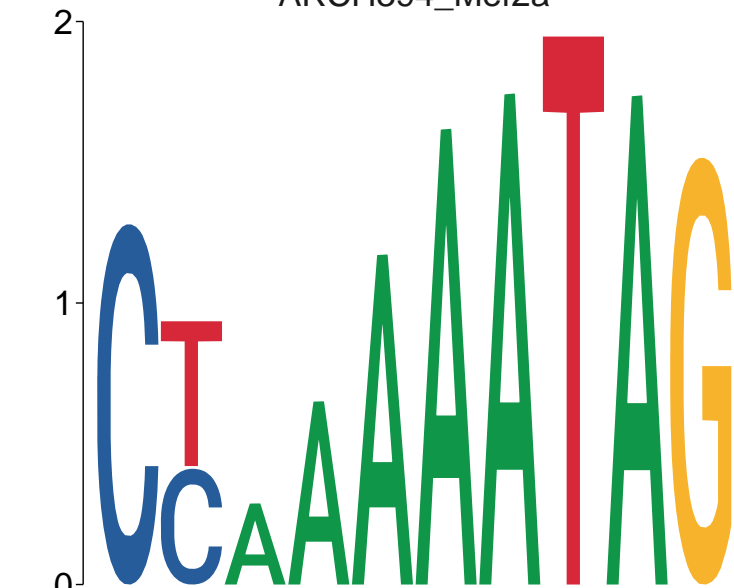
RCH393_24_TTTAAAAATG,BestGuess_Uni



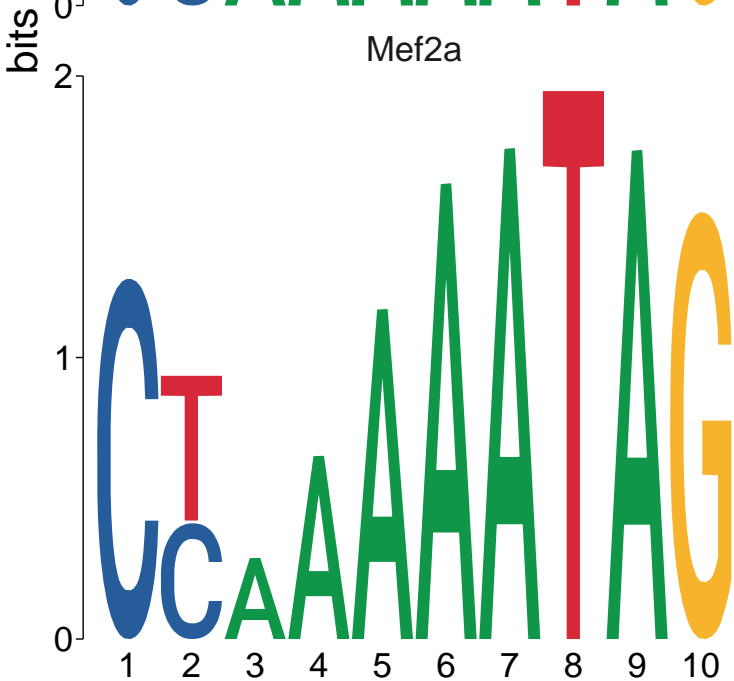
24-TTTAAAAATG,BestGuess:Unknown6



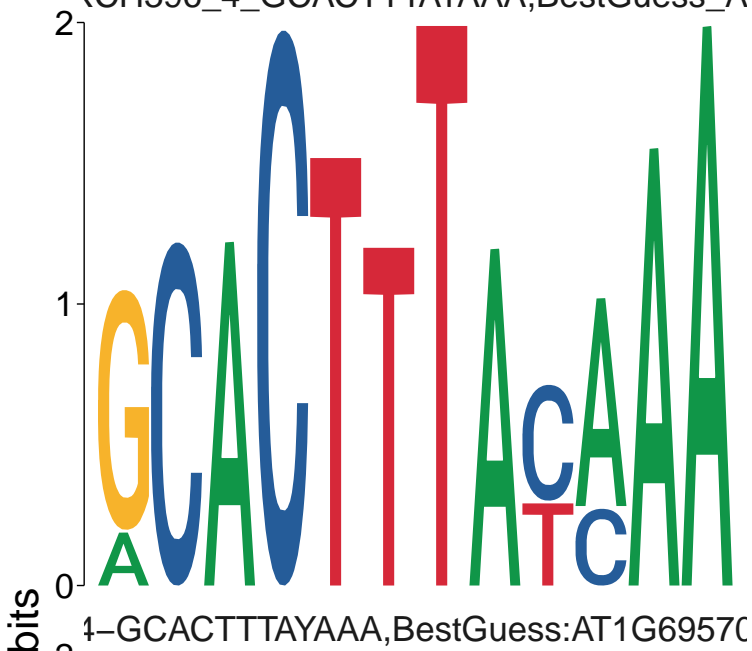
ARCH394_Mef2a



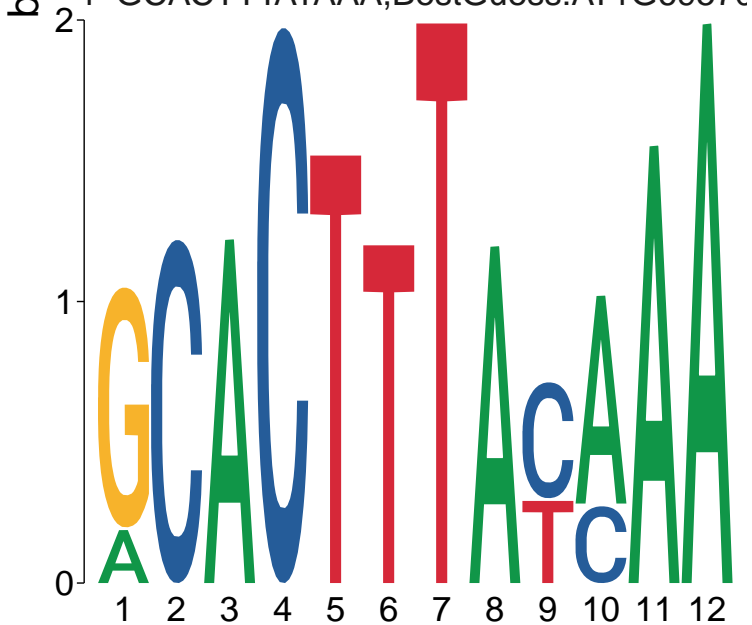
Mef2a



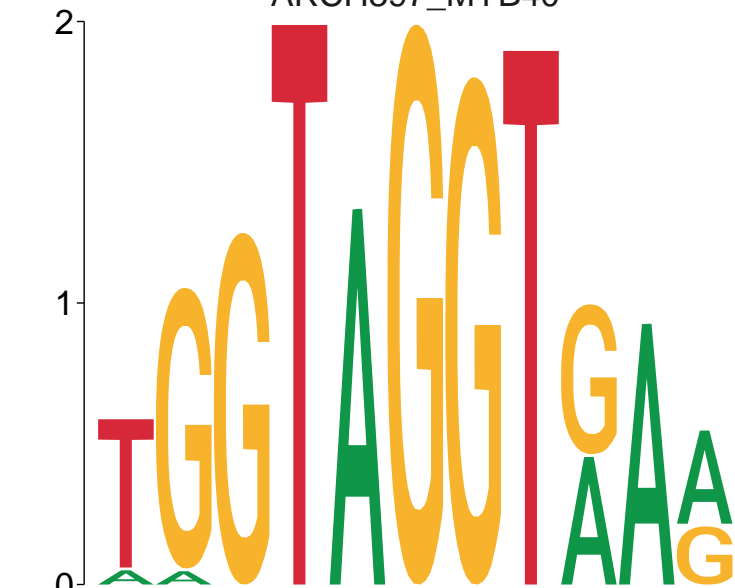
CH396_4_GCACCTTTAYAAA,BestGuess_A'



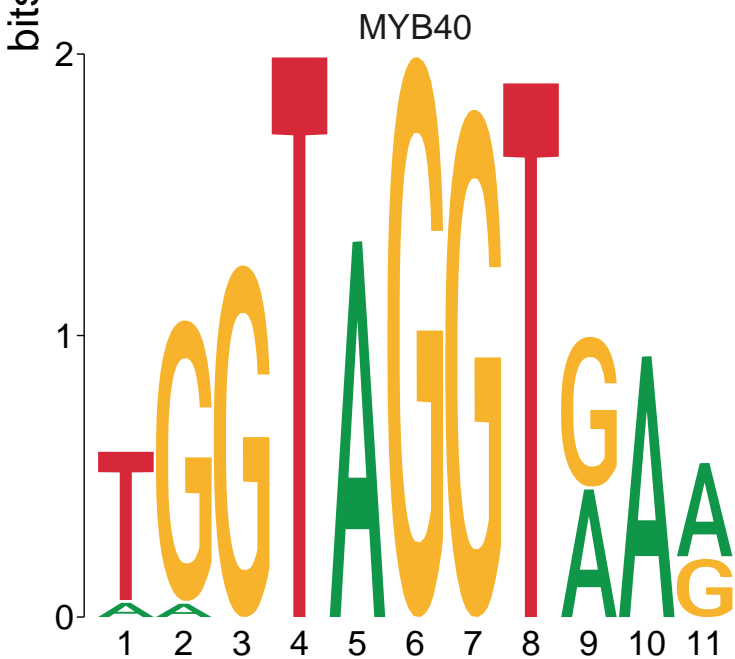
4-GCACCTTTAYAAA,BestGuess:AT1G6957C



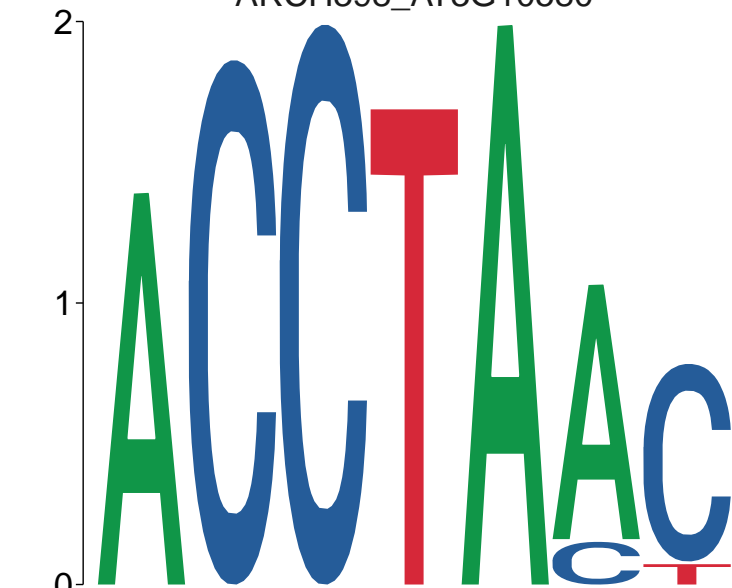
ARCH397_MYB40



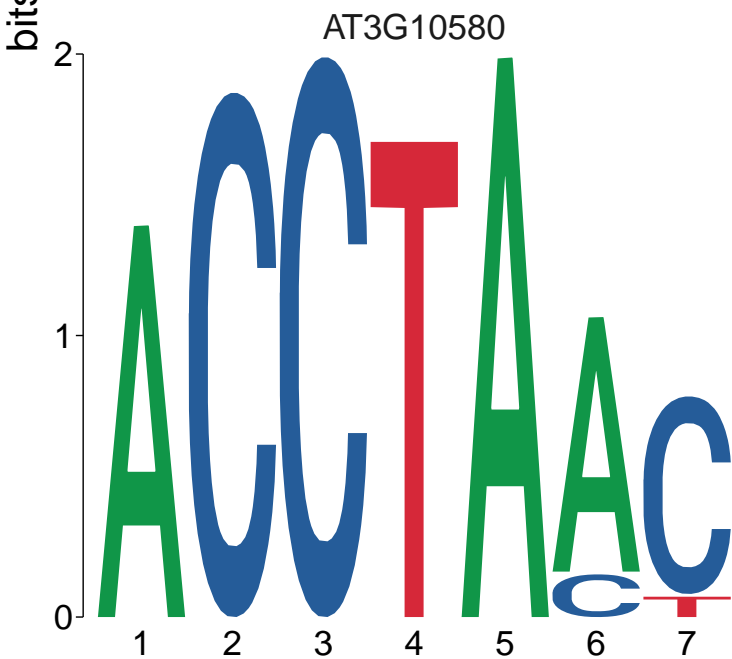
MYB40



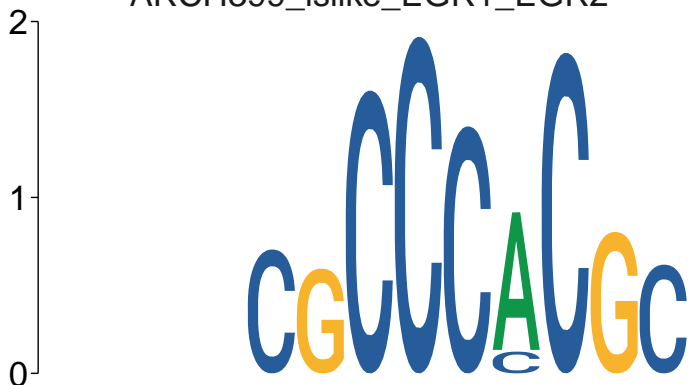
ARCH398_AT3G10580



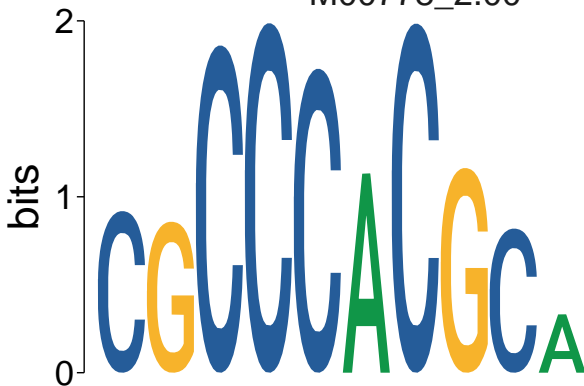
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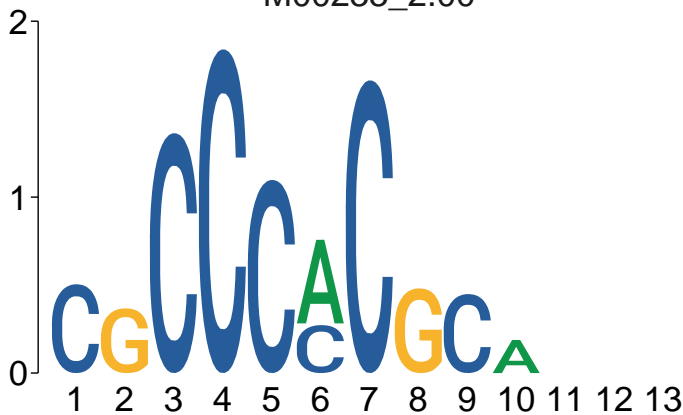
ARCH399_islike_EGR1_EGR2



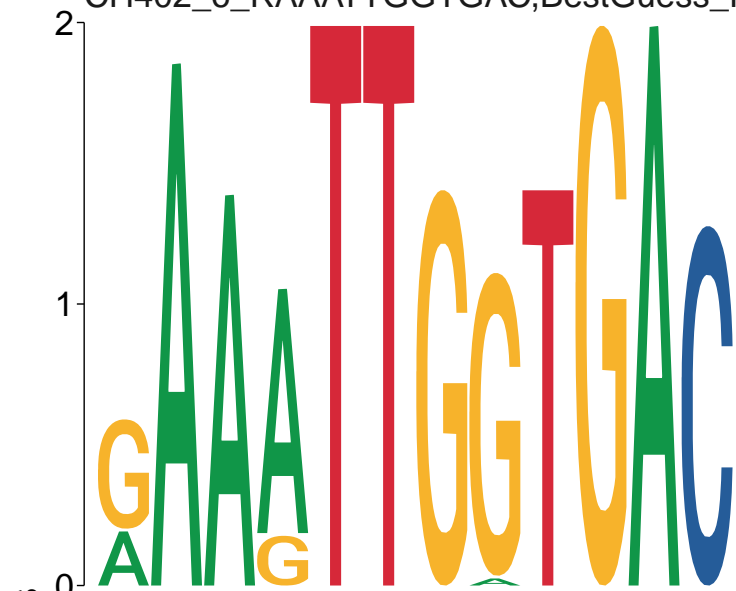
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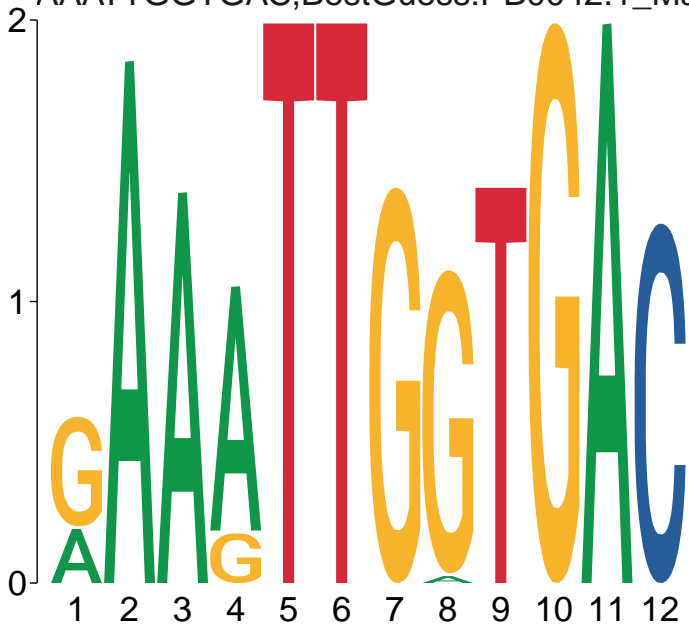
M00233_2.00



CH402_6_RAAATTGGTGAC,BestGuess_F



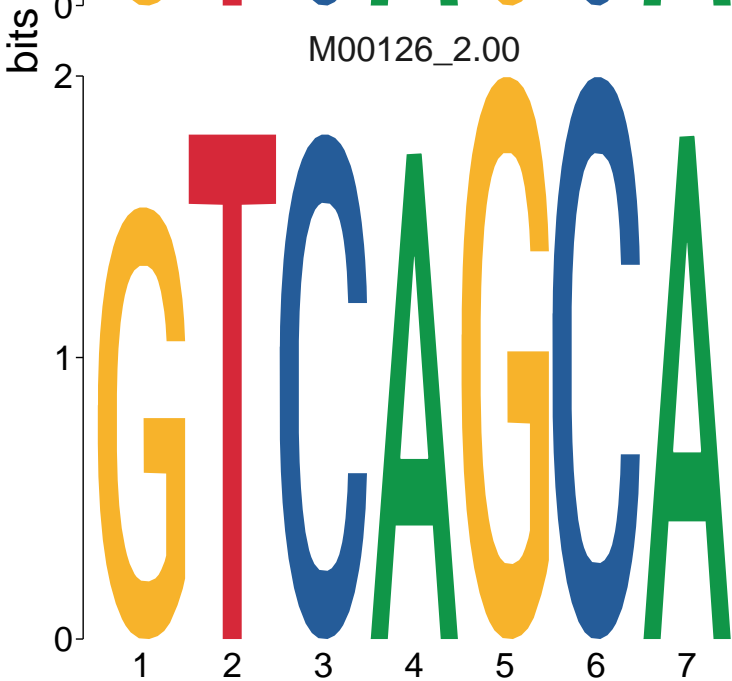
AAATTGGTGAC,BestGuess:PB0042.1_Ma



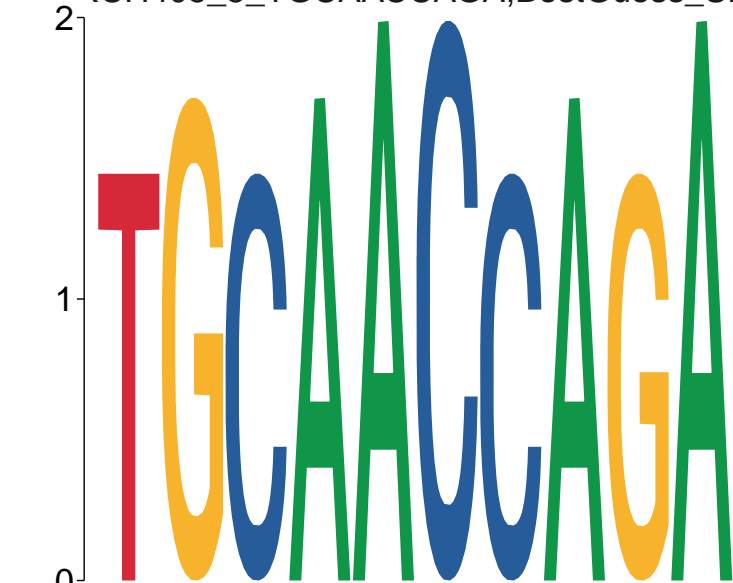
ARCH403_bZIP



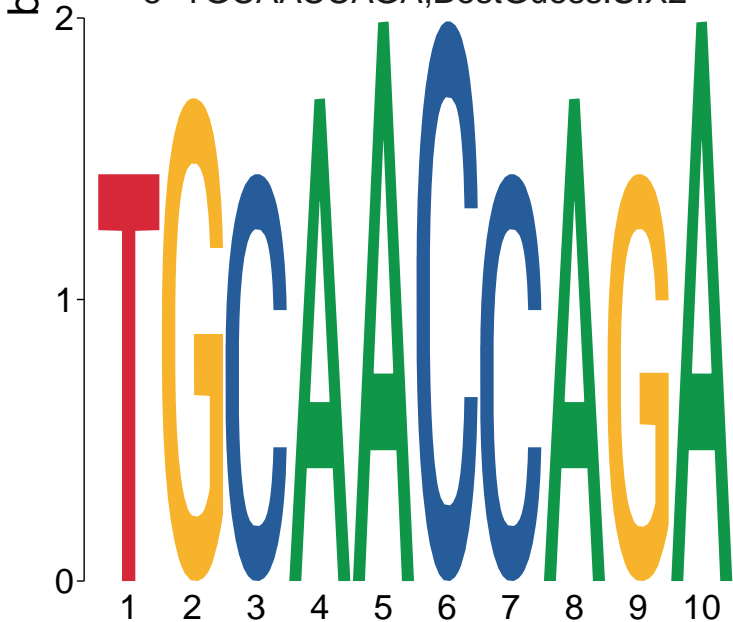
M00126_2.00



RCH405_5_TGCAACCAGA,BestGuess_SI)



5-TGCAACCAGA,BestGuess:SIX2



CH406_RUNX1_RUNX_RUNX1_RUNX_AI



RUNX-AML



RUNX1



RUNX



RUNX1.1



ARCH408_Rfx1_Rfx2_X_box_RFX_X_bo



Rfx1



X-box



RFX



X-box.1



Rfx2



Rfx1.1

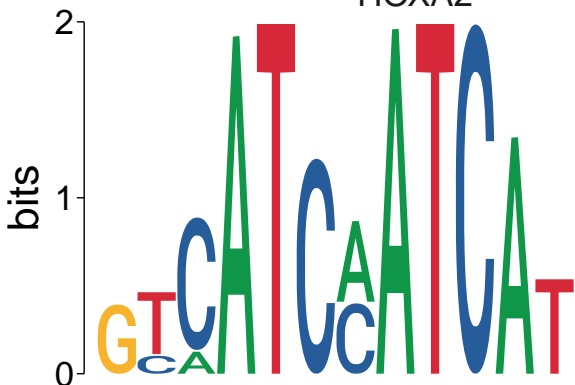


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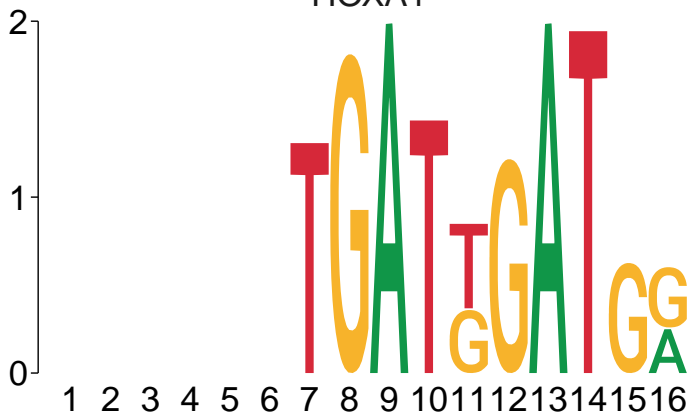
ARCH410_HOXA1_HOXA2



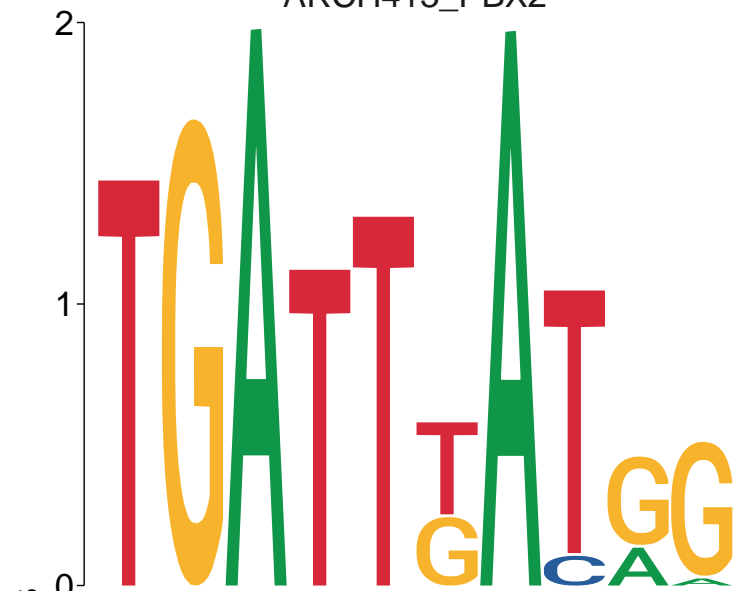
HOXA2



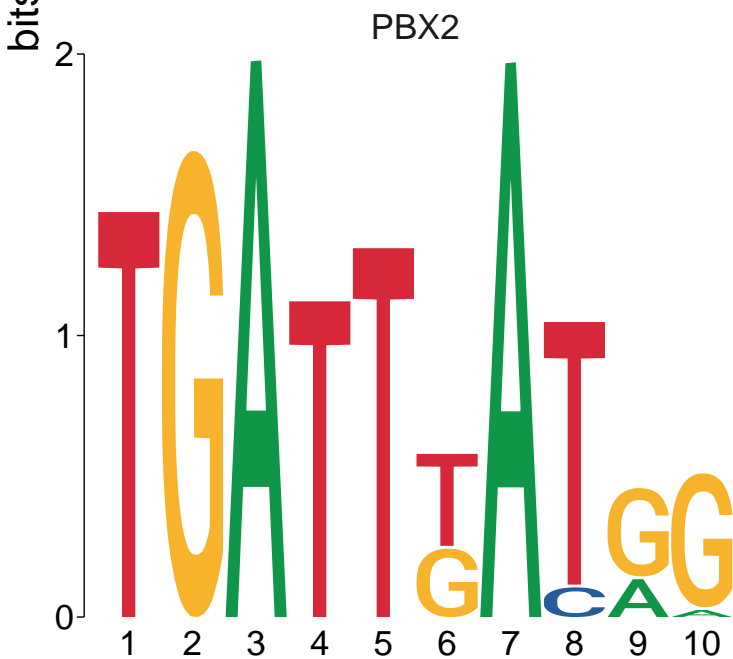
HOXA1



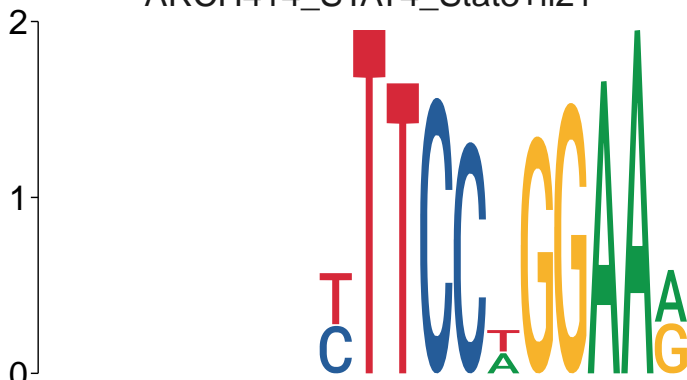
ARCH413_PBX2



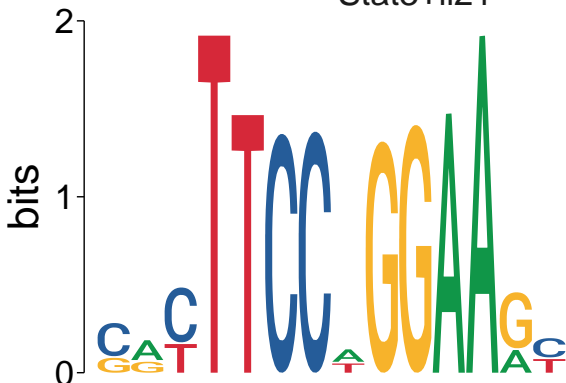
PBX2



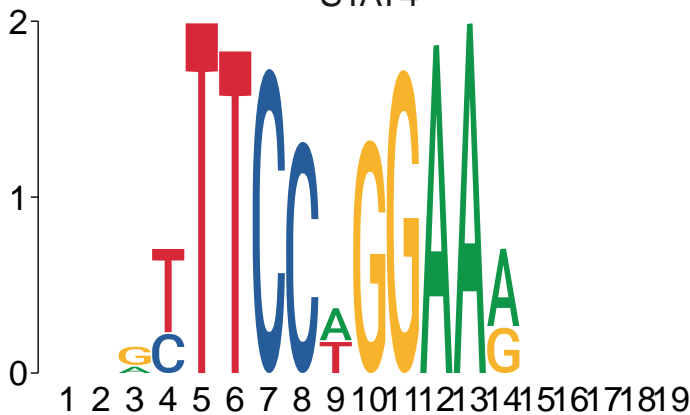
ARCH414_STAT4_Stat3+il21



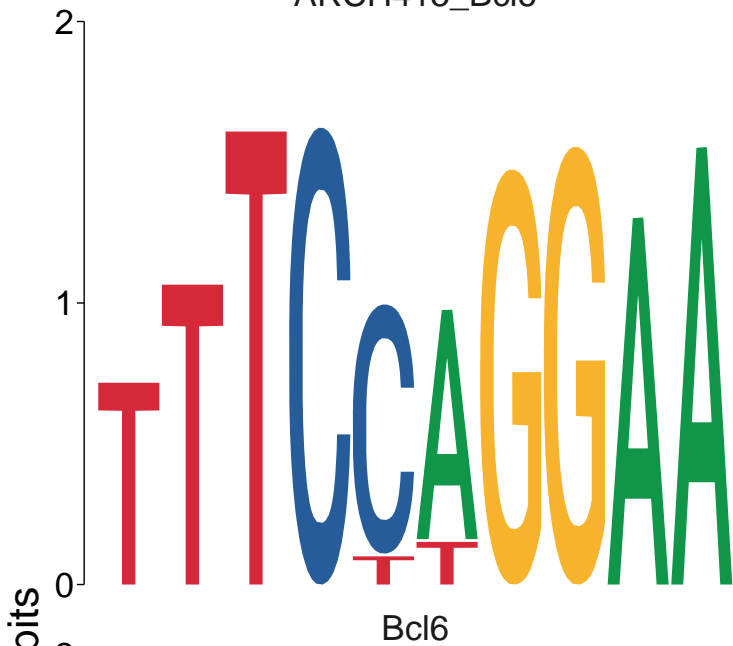
Stat3+il21



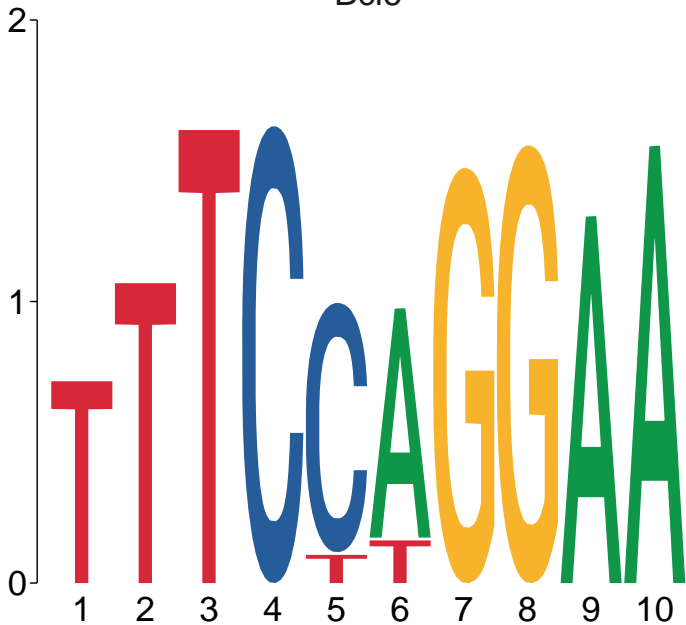
STAT4



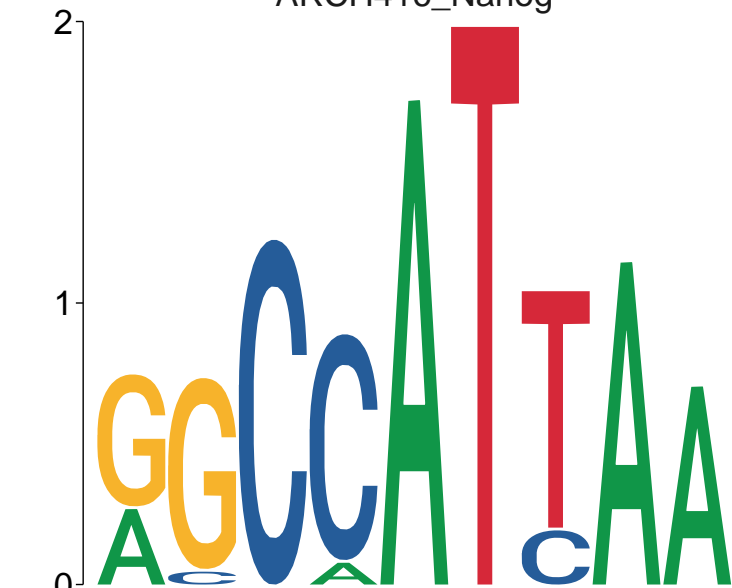
ARCH415_Bcl6



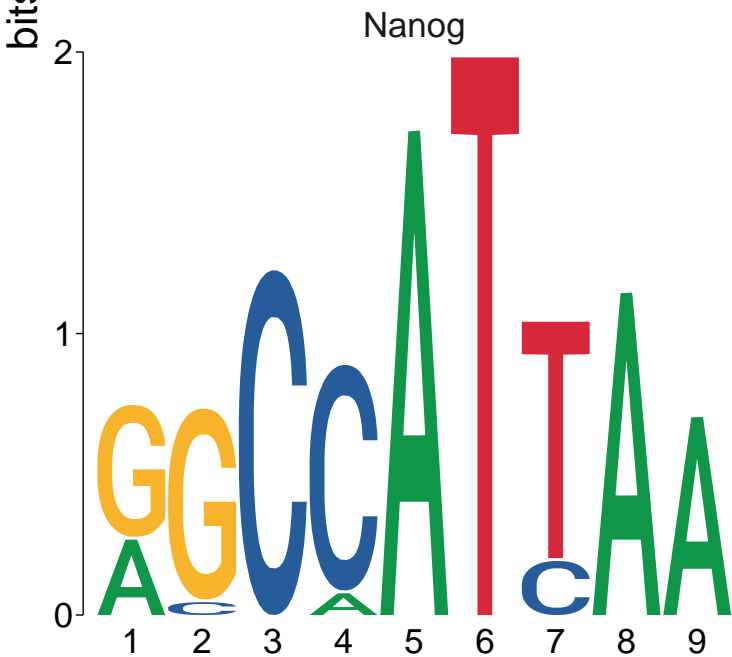
Bcl6



ARCH416_Nanog



Nanog



ARCH417_ANTP_HOXA2_ANTP_HOXA3

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M00516_2.00

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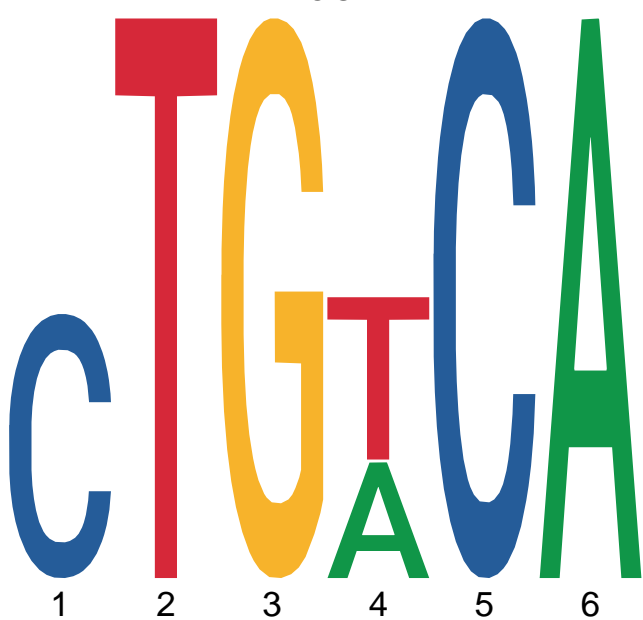
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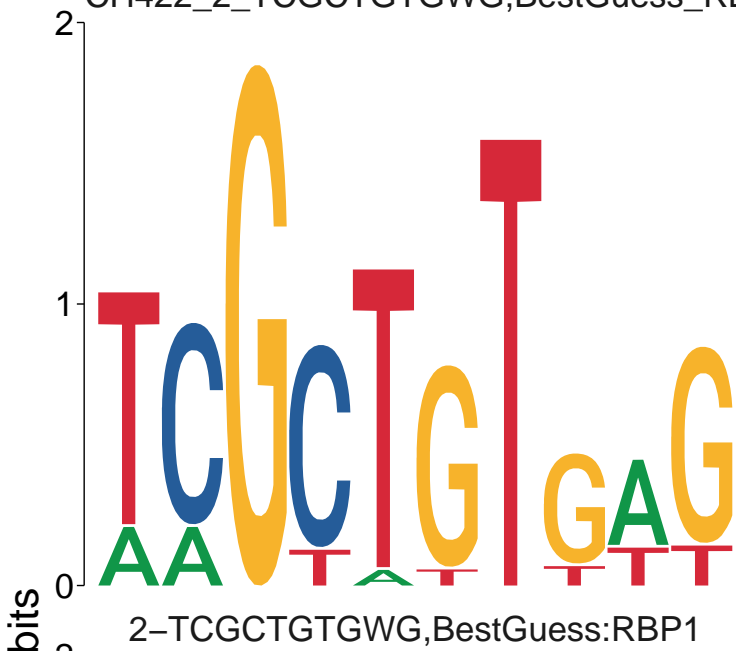
ARCH421_Meis1



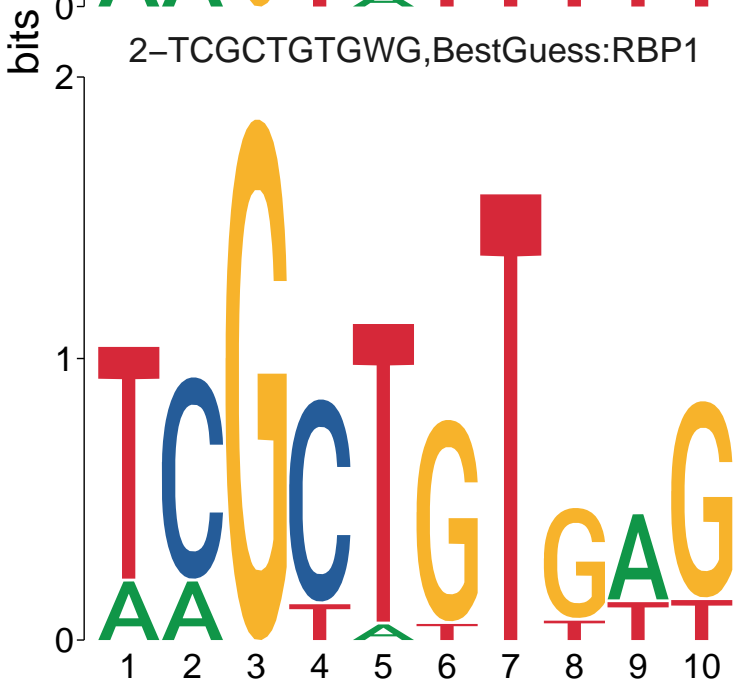
Meis1



CH422_2_TCGCTGTGWG,BestGuess_RB



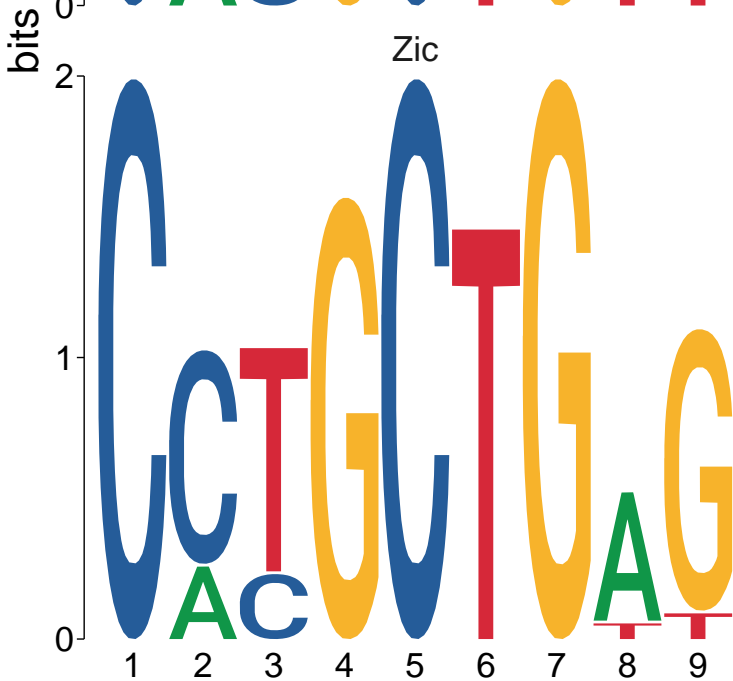
2-TCGCTGTGWG,BestGuess:RBP1



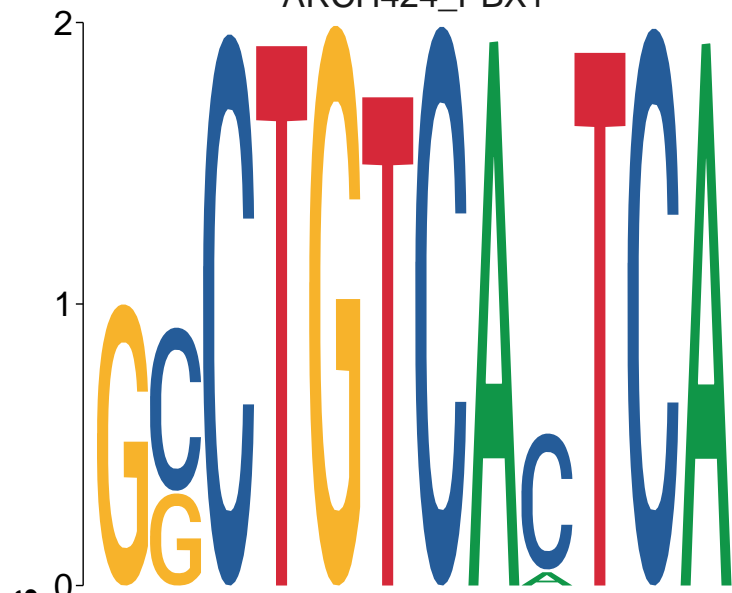
ARCH423_Zic



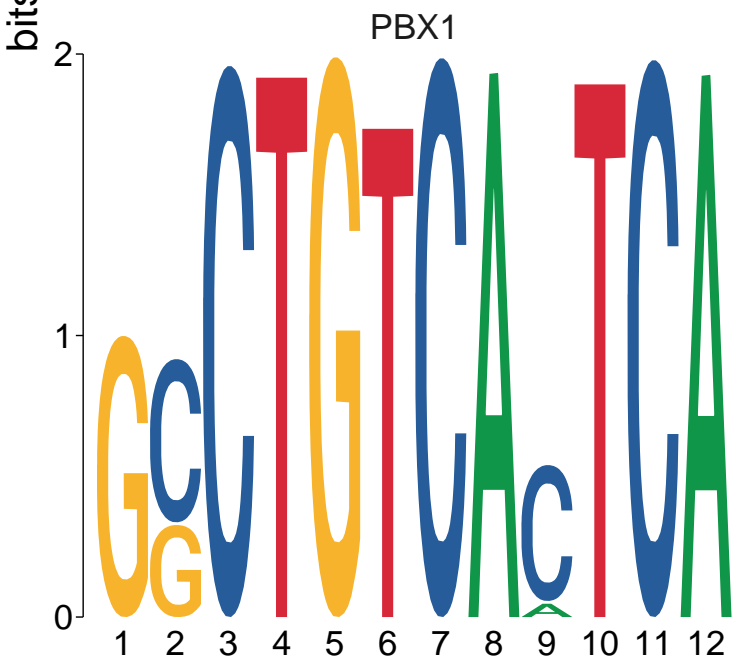
Zic



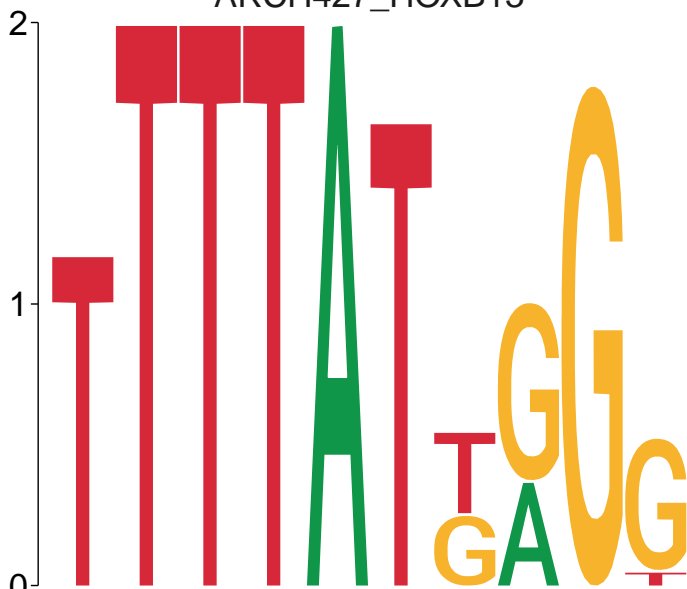
ARCH424_PBX1



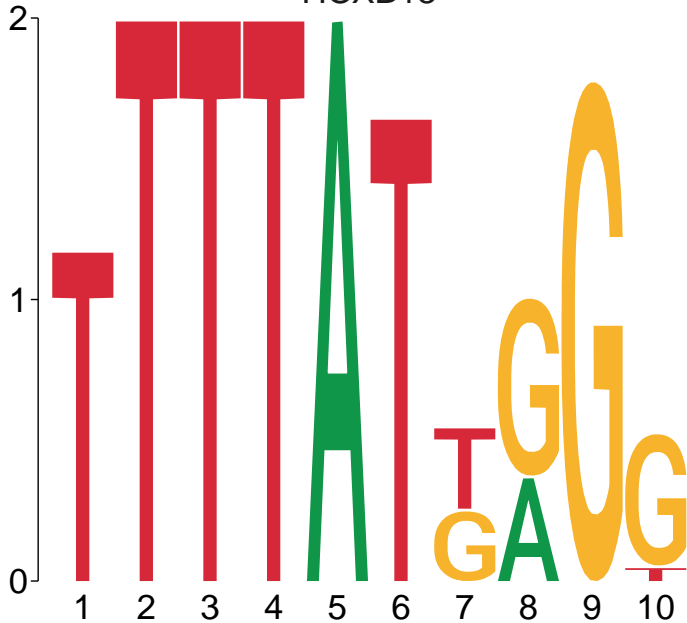
PBX1



ARCH427_HOXB13



HOXB13



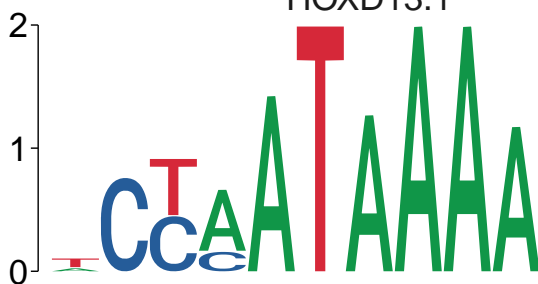
ARCH428_Hoxd13_HOXD13_HOXD13



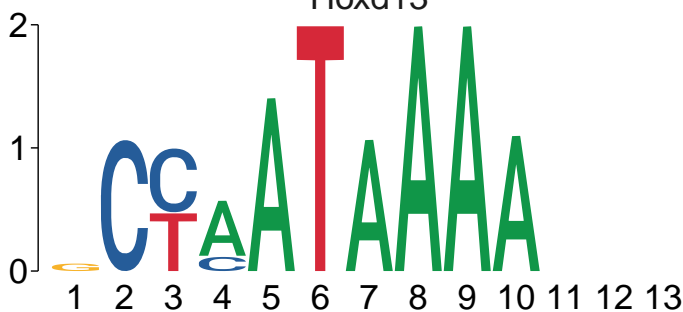
HOXD13



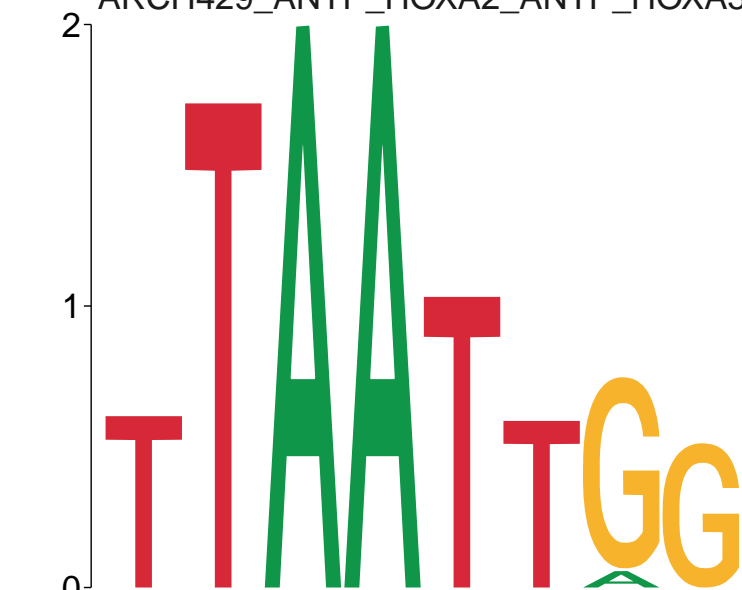
HOXD13.1



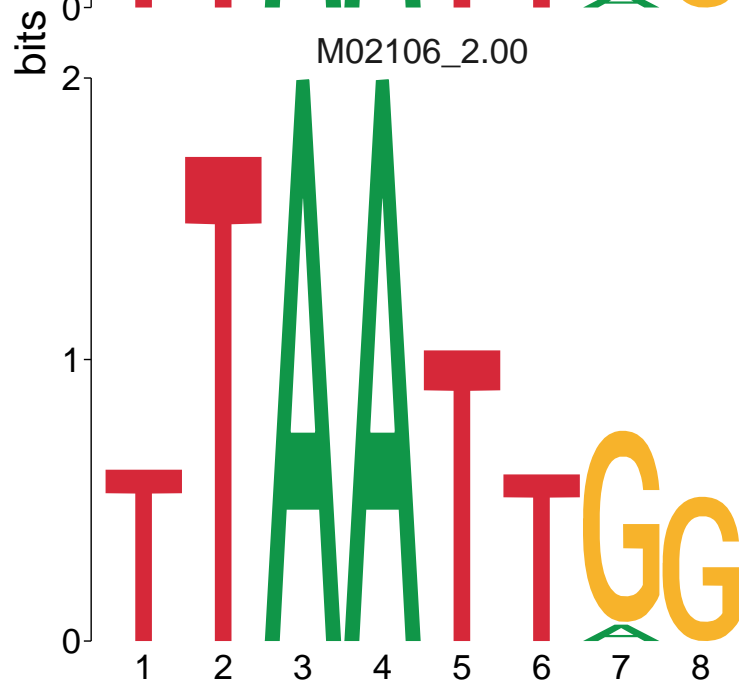
Hoxd13



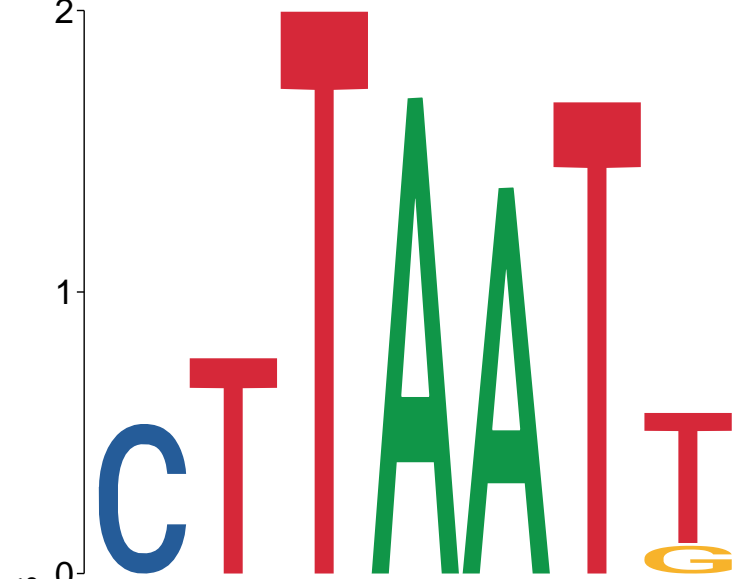
ARCH429_ANTP_HOXA2_ANTP_HOXA3



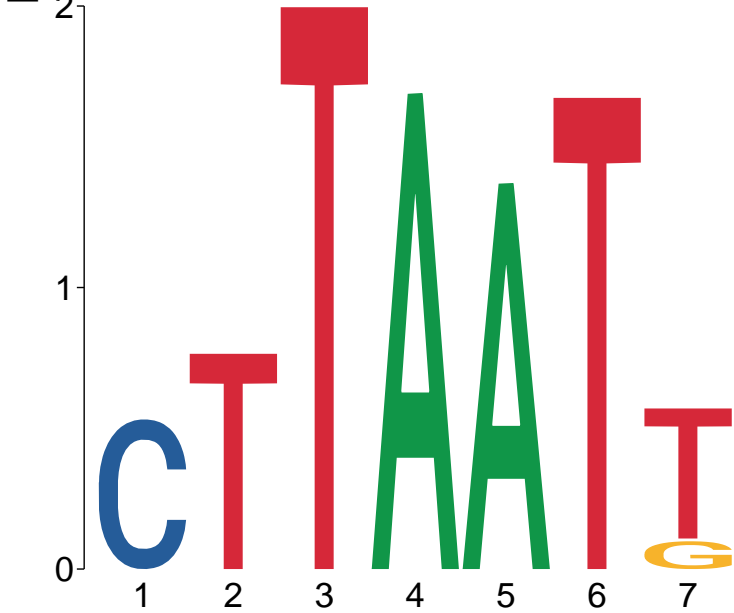
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ARCH430_Nvec_vc1.1_XM_032377135.2



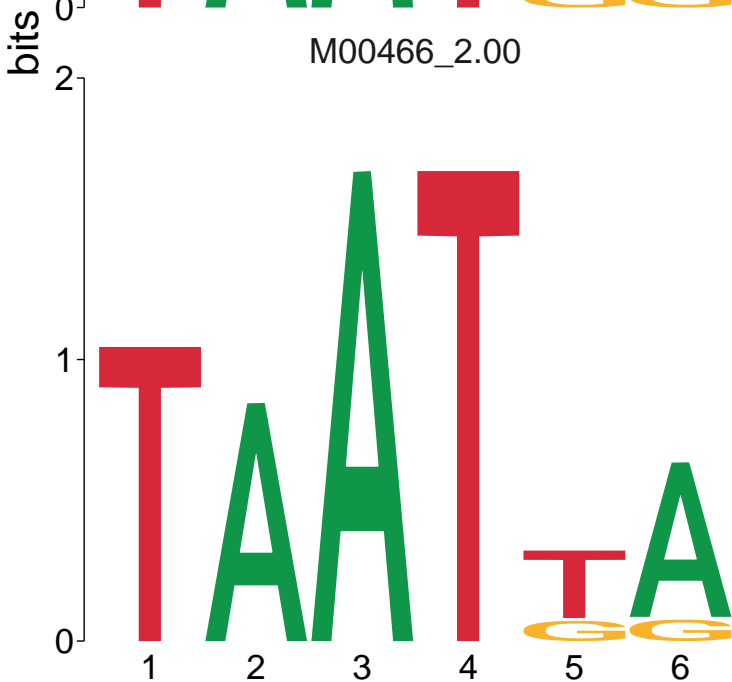
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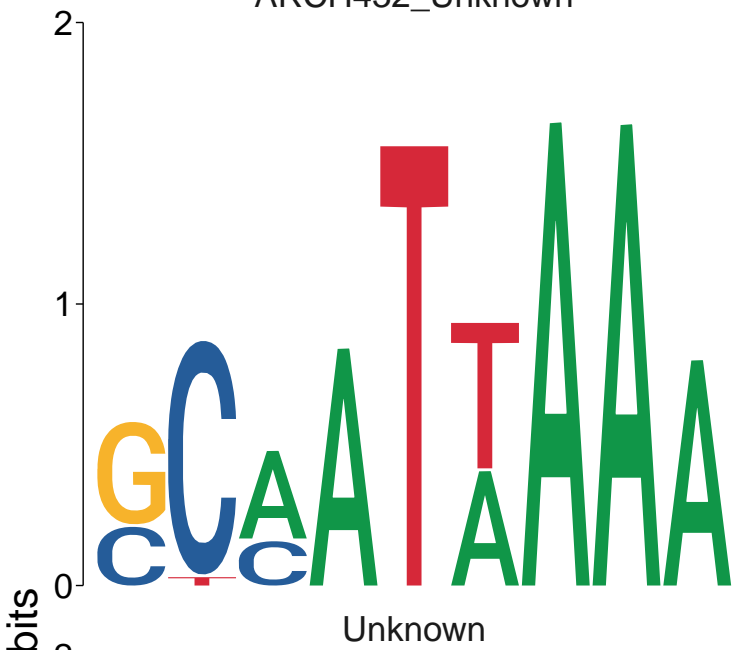
ARCH431_islike_ANTP_EN1



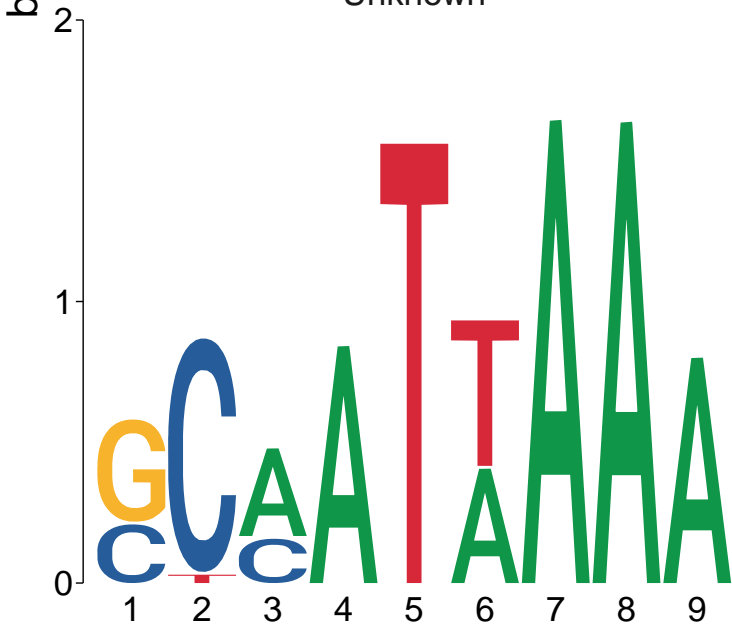
M00466_2.00



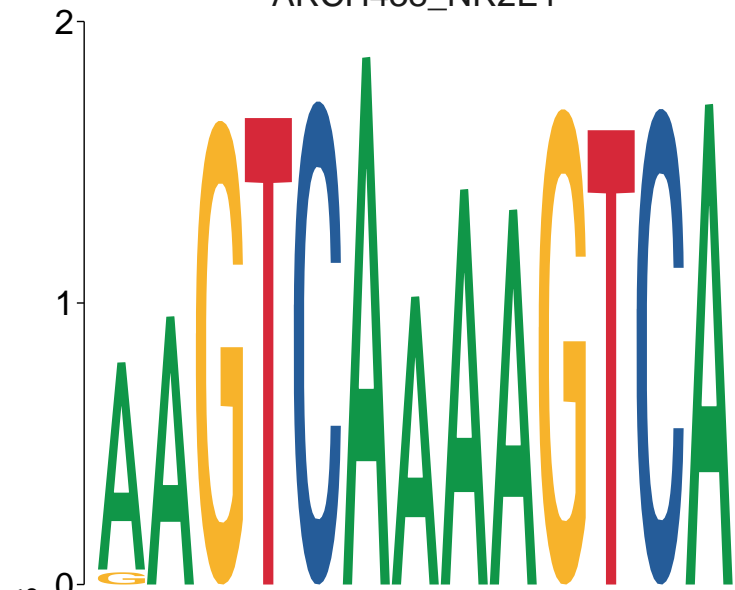
ARCH432_Unknown



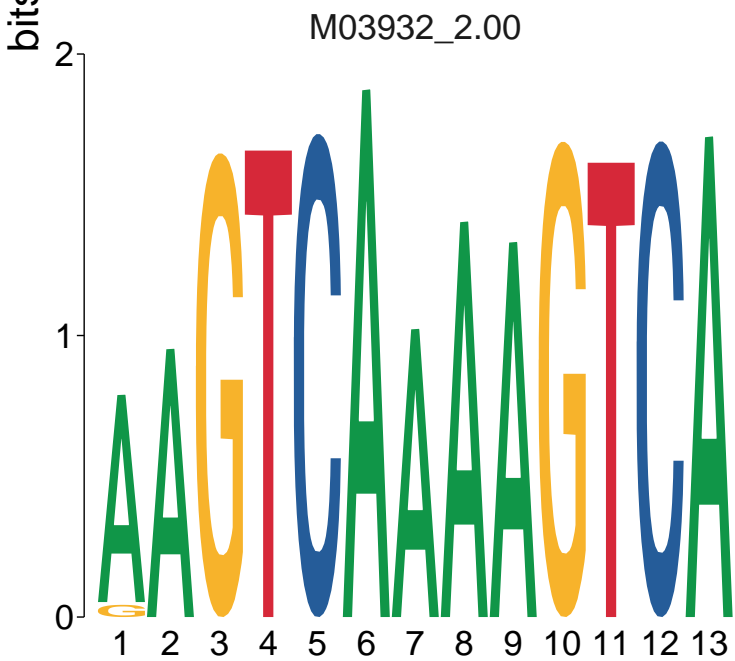
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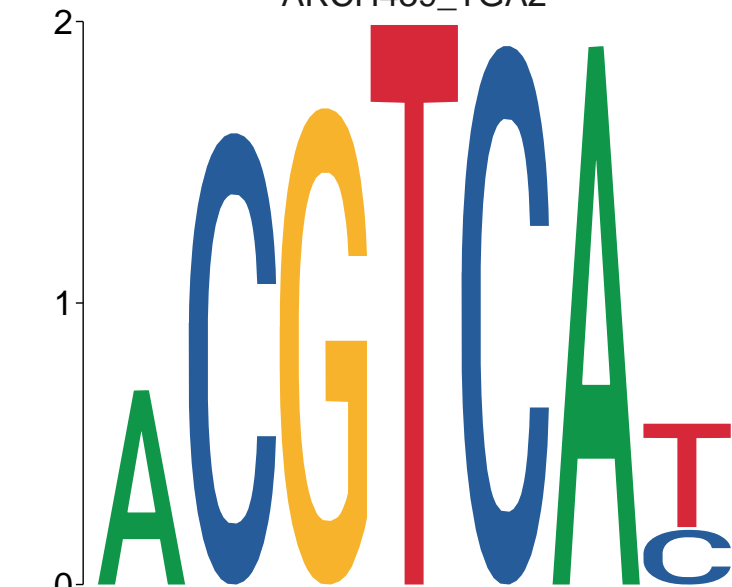
ARCH438_NR2E1



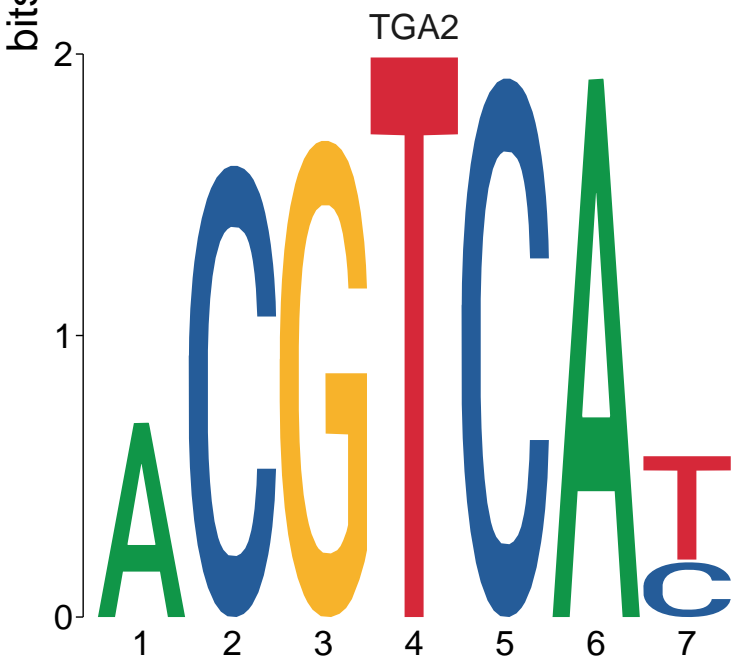
M03932_2.00



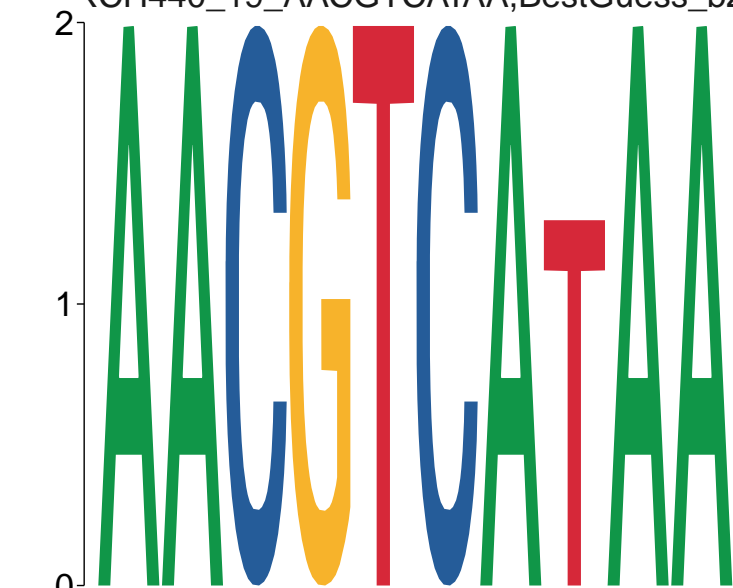
ARCH439_TGA2



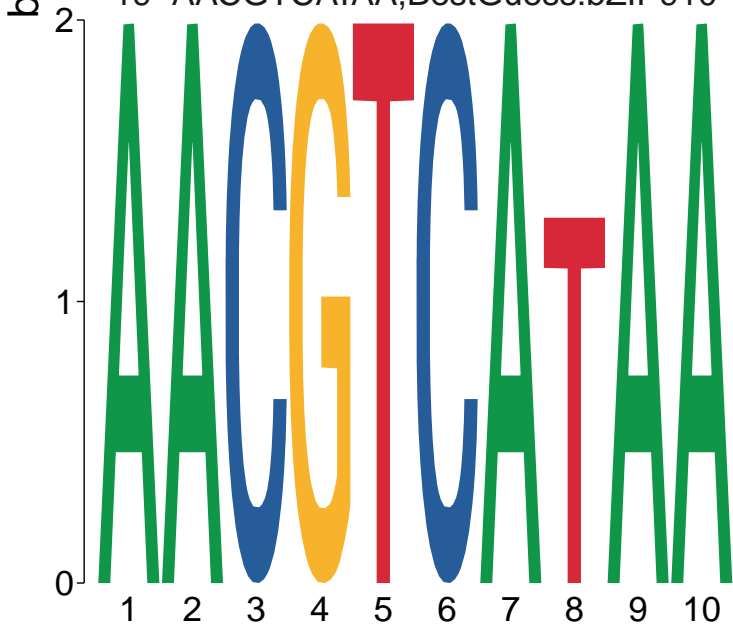
TGA2



RCH440_19_AACGTCATAA,BestGuess_bZ



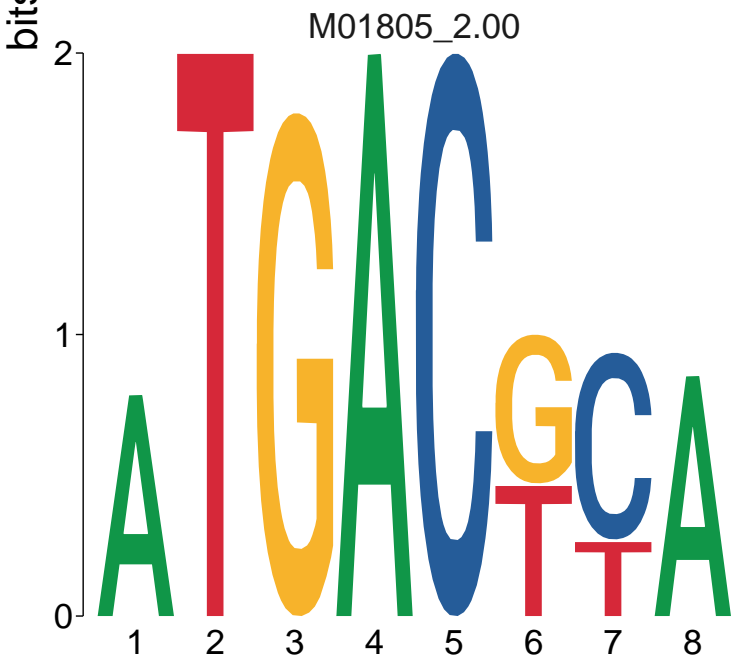
19-AACGTCATAA,BestGuess:bZIP910



ARCH441_FOS_FOSB_FOSL1



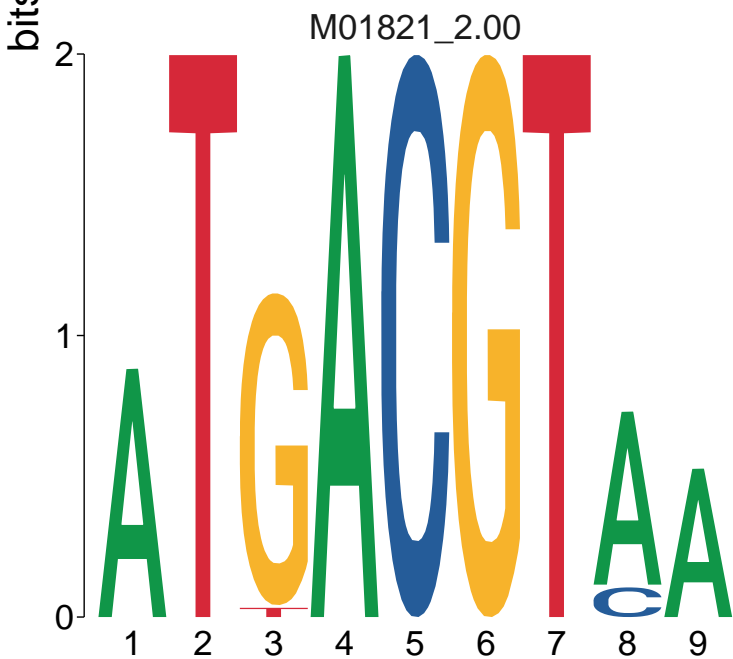
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ARCH442_ATF1_CREB1



M01821_2.00



ARCH443_ARID2

2

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0

bits



M03590_2.00

2

1

0

1

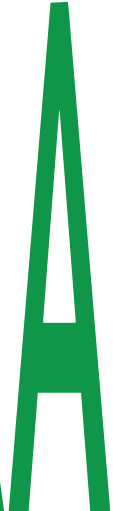
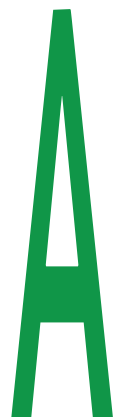
2

3

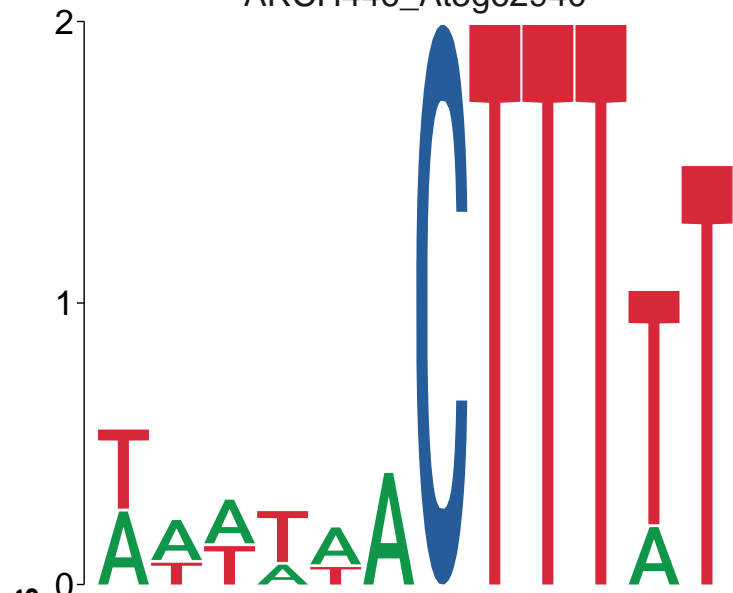
4

5

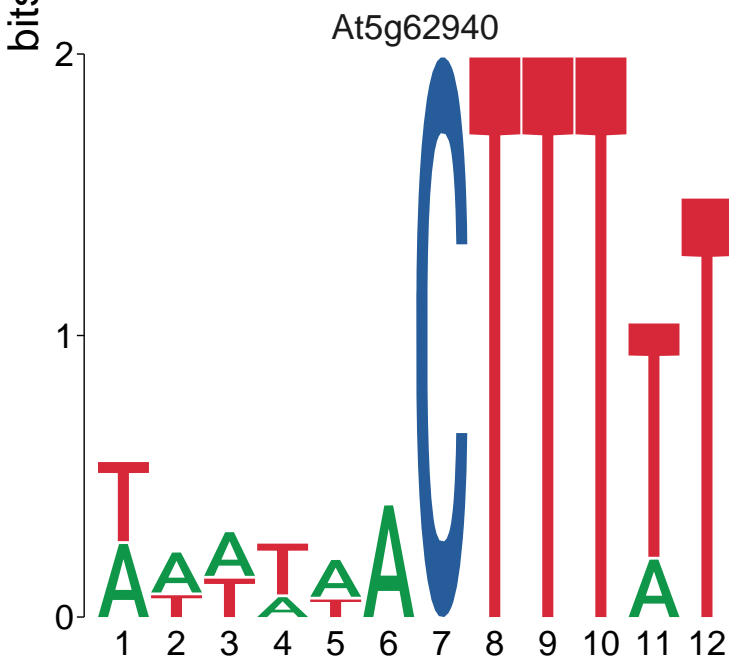
6



ARCH446_At5g62940



At5g62940



RCH447_5_TTTATATTTT,BestGuess_TSO

2

1

0

bits

5-TTTATATTTT,BestGuess:TSO1

2

1

0

1

2

3

4

5

6

7

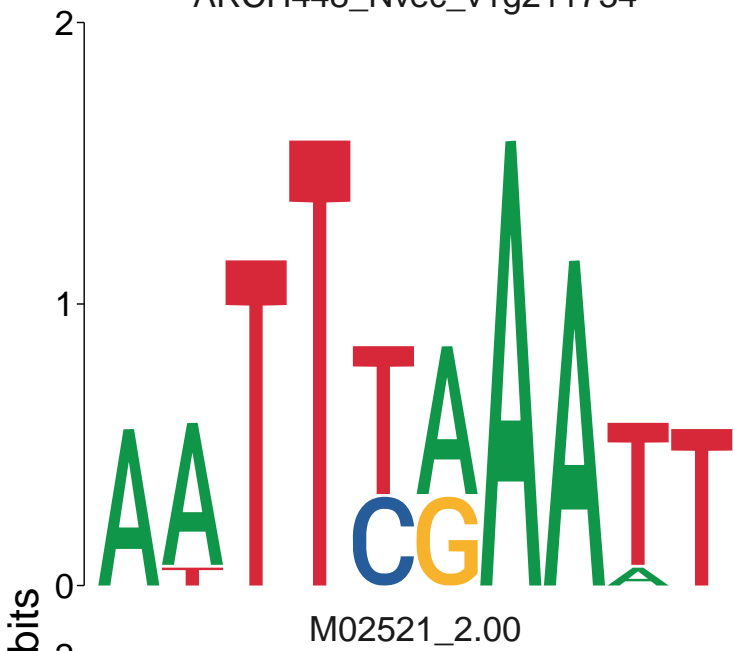
8

9

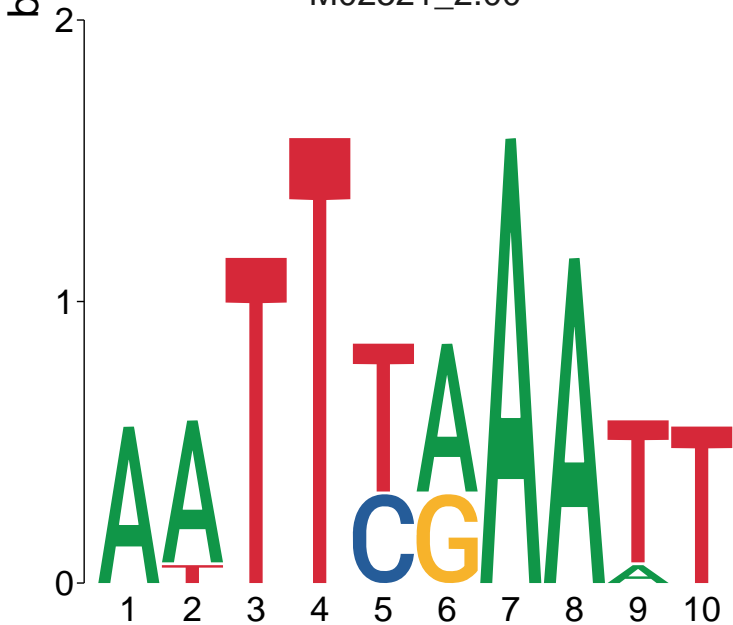
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ARCH448_Nvec_v1g211734



M02521_2.00



ARCH449_RRTF1_ERF4_RAP212_RAP26



RAP26



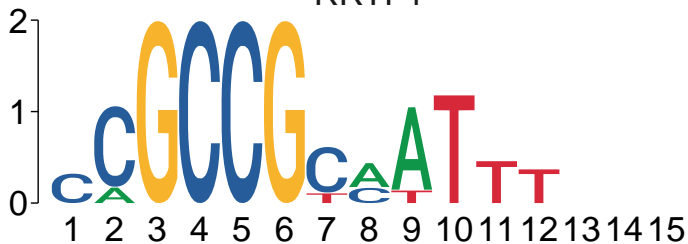
RAP212

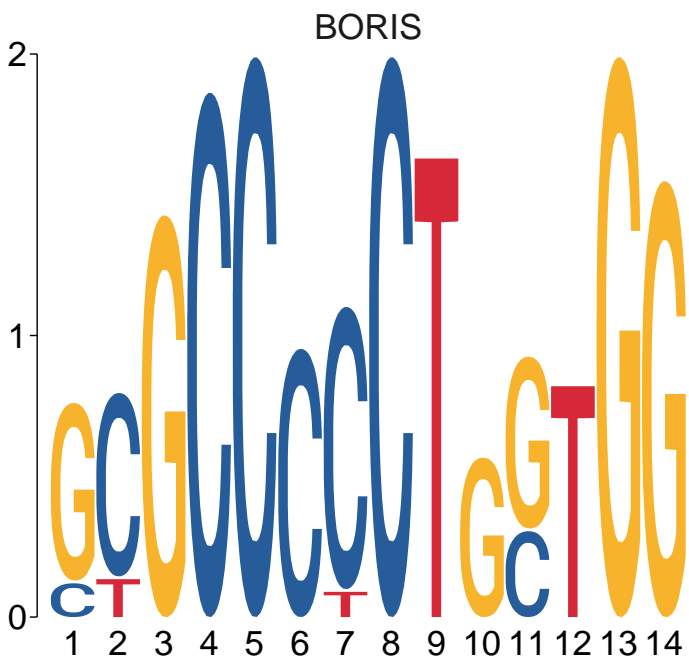
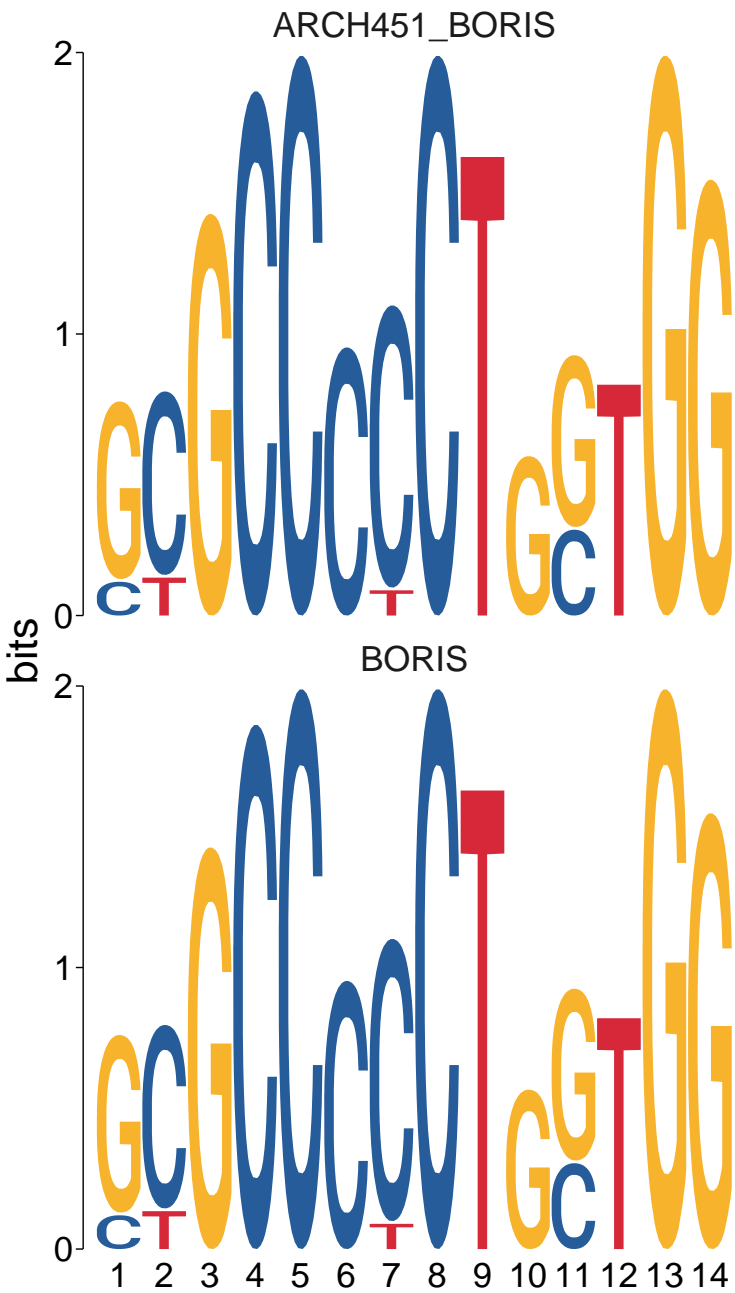


ERF4



RRTF1

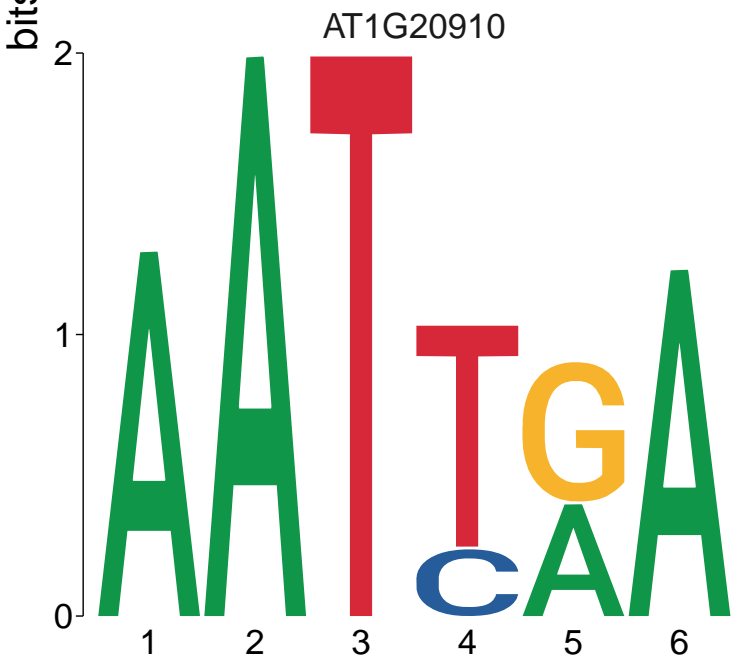




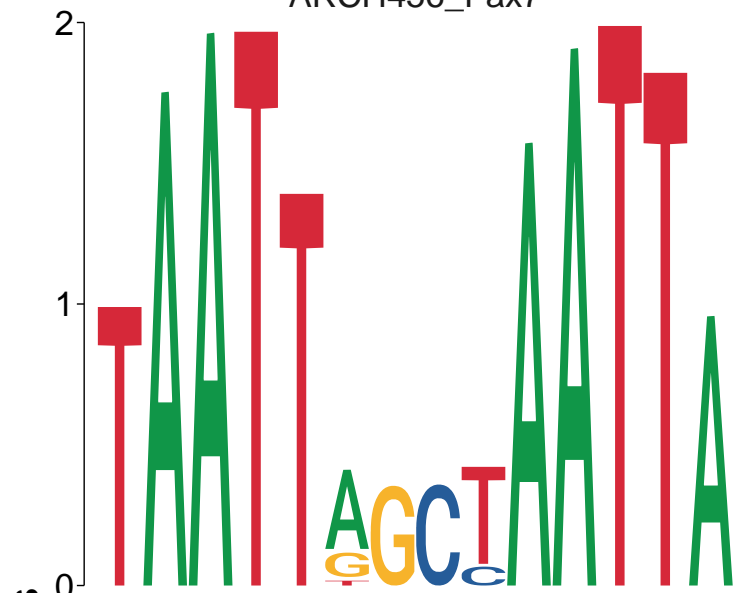
ARCH453_AT1G20910



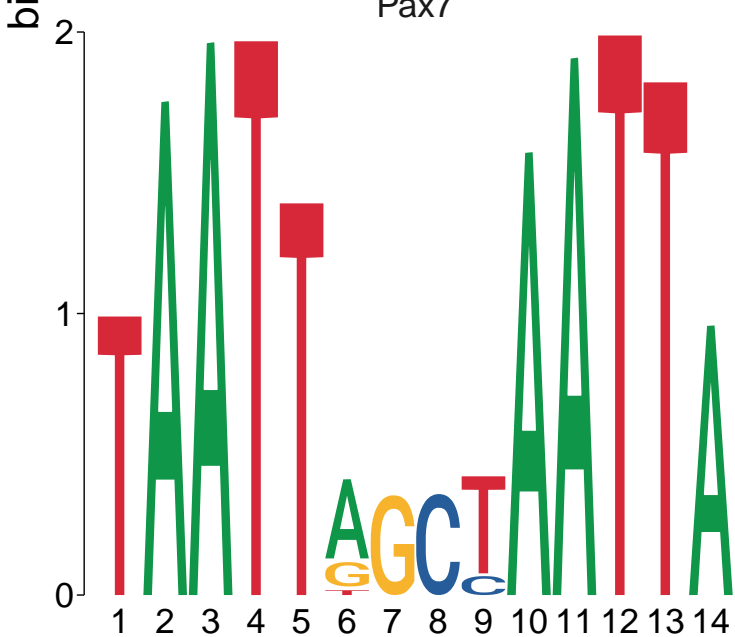
AT1G20910



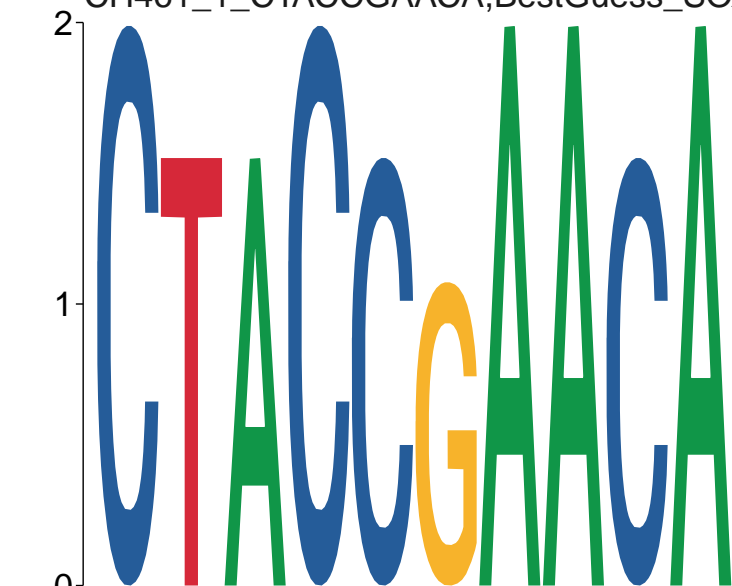
ARCH456_Pax7



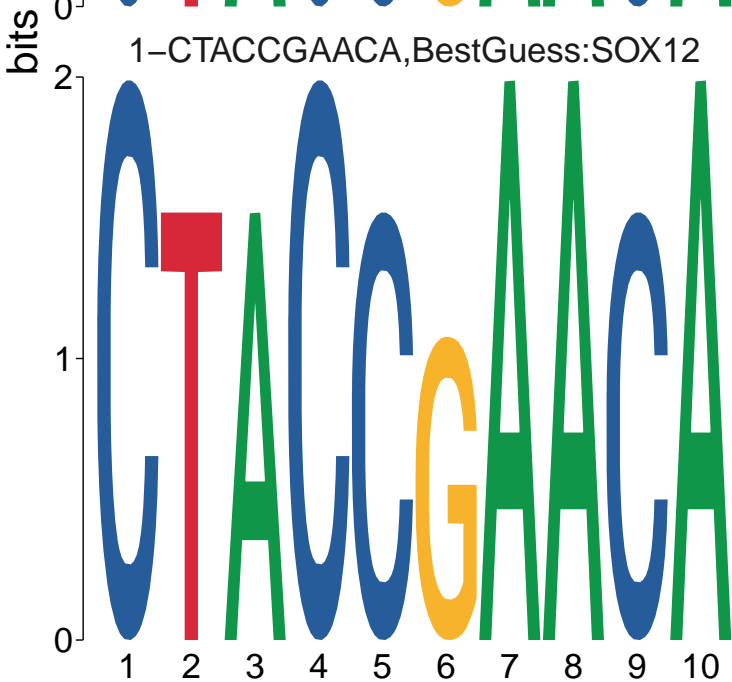
Pax7



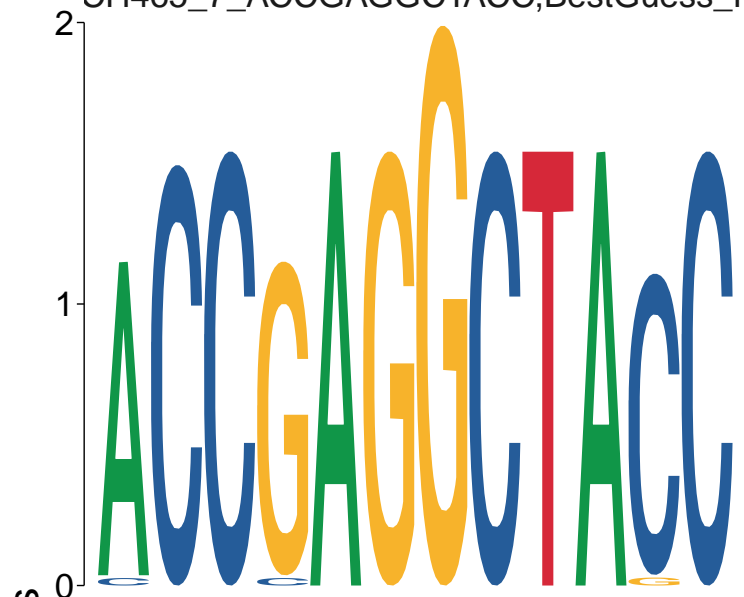
CH461_1_CTACCGAACA, BestGuess_SOX



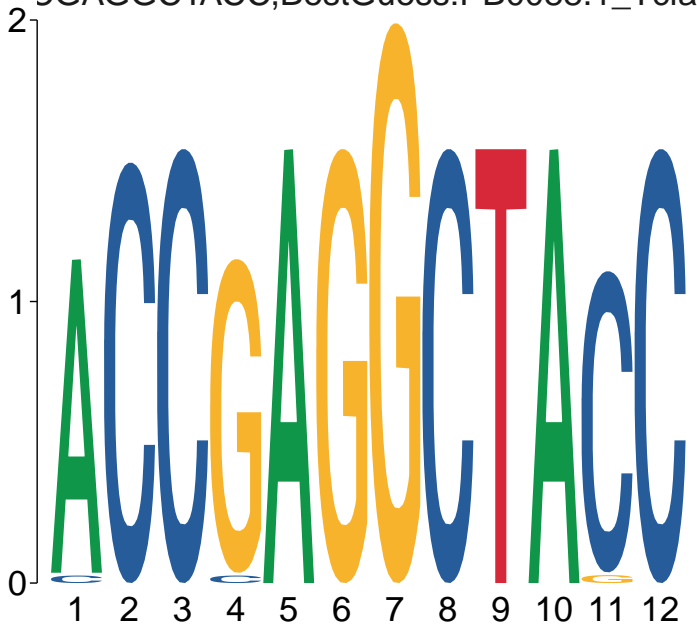
1-CTACCGAACA, BestGuess:SOX12



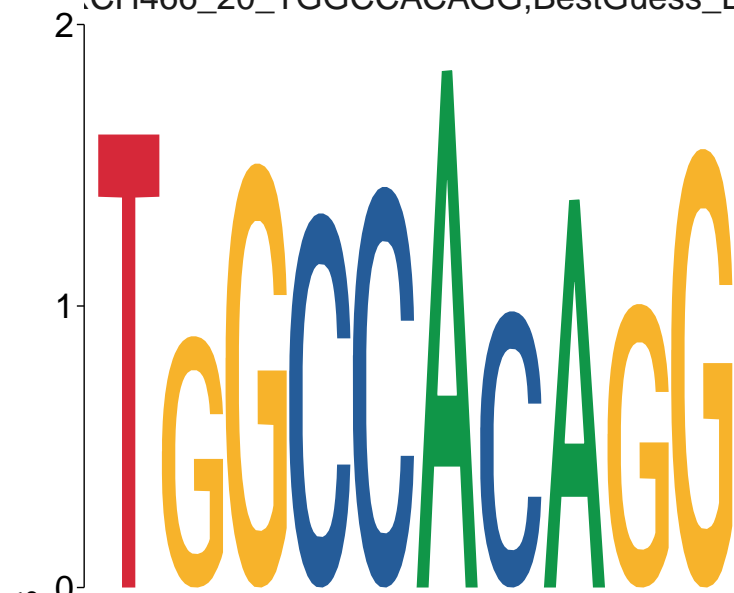
CH465_7_ACCGAGGCTACC,BestGuess_F



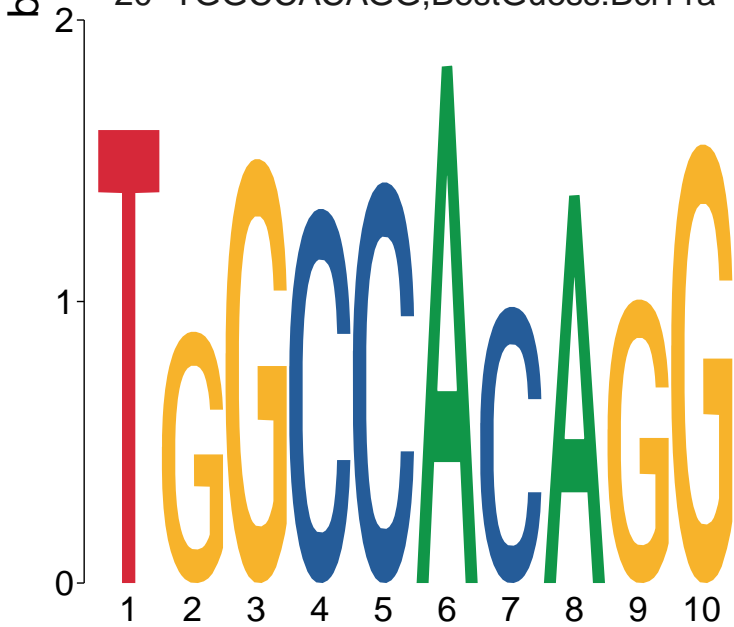
CGAGGCTACC,BestGuess:PB0088.1_Tcfa



CH466_20_TGGCCACAGG,BestGuess_B



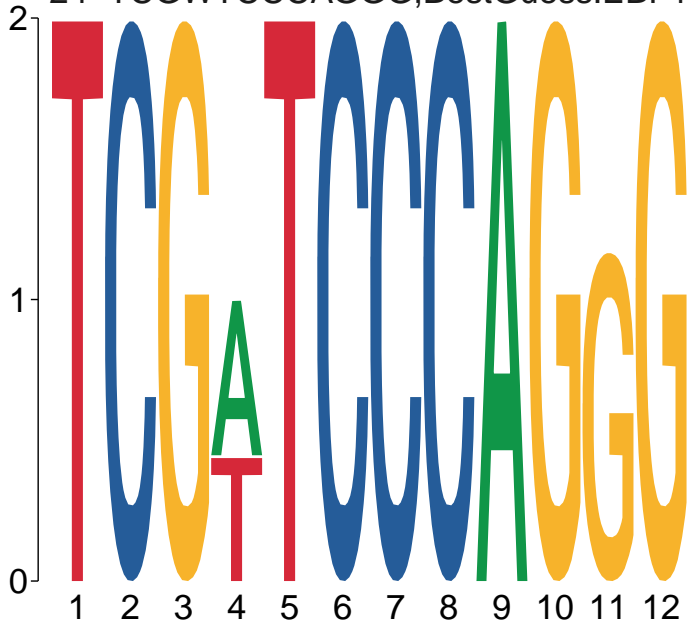
20-TGGCCACAGG,BestGuess:Bcl11a



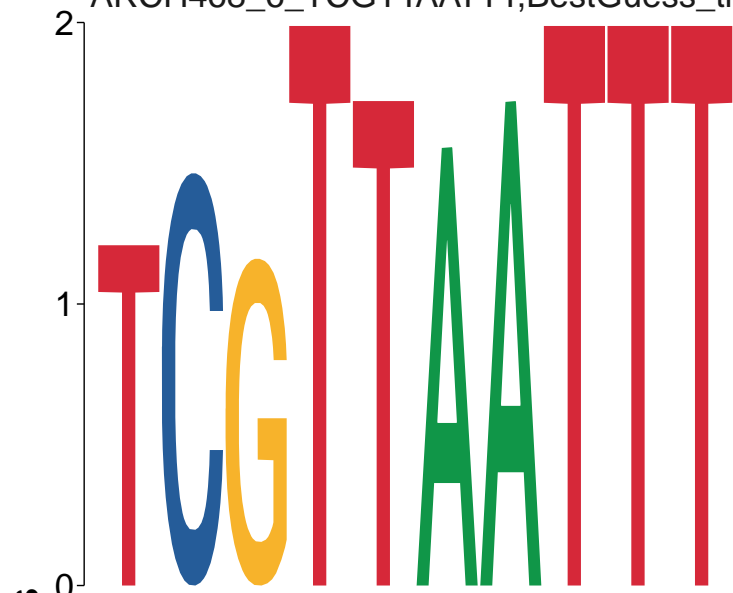
CH467_24_TCGWTCCCAGGG,BestGuess_



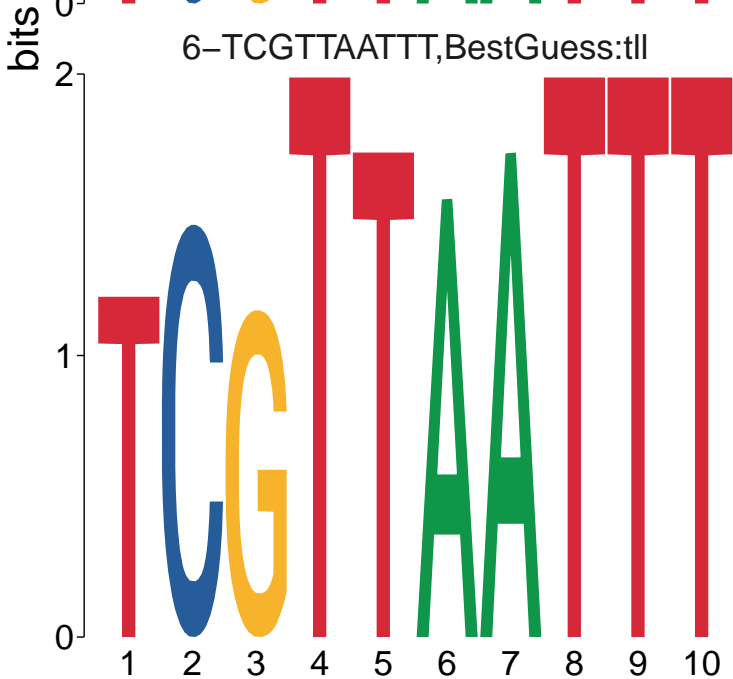
24-TCGWTCCCAGGG,BestGuess:EBF1



ARCH468_6_TCGTTAATTT,BestGuess_tll



6-TCGTTAATTT,BestGuess:tll



ARCH469_ANTP_HOXA2_ANTP_HOXA3

2

1

0

bits



M03166_2.00

2

1

0

1 2 3 4 5 6



ARCH470_ANTP_HOXA2_ANTP_HOXA3

2

1

0

bits

M03221_2.00

2

1

0

1

2

3

4

5

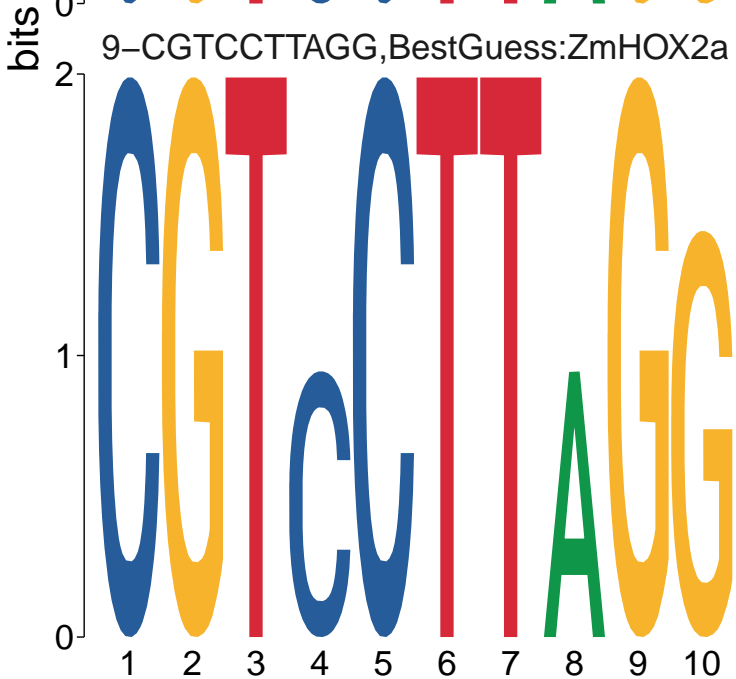
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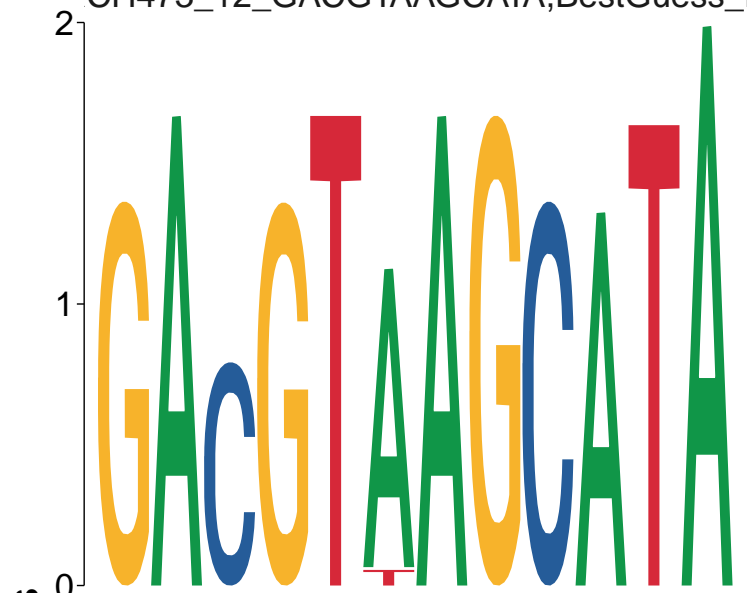
CH471_9_CGTCCTTAGG,BestGuess_ZmH



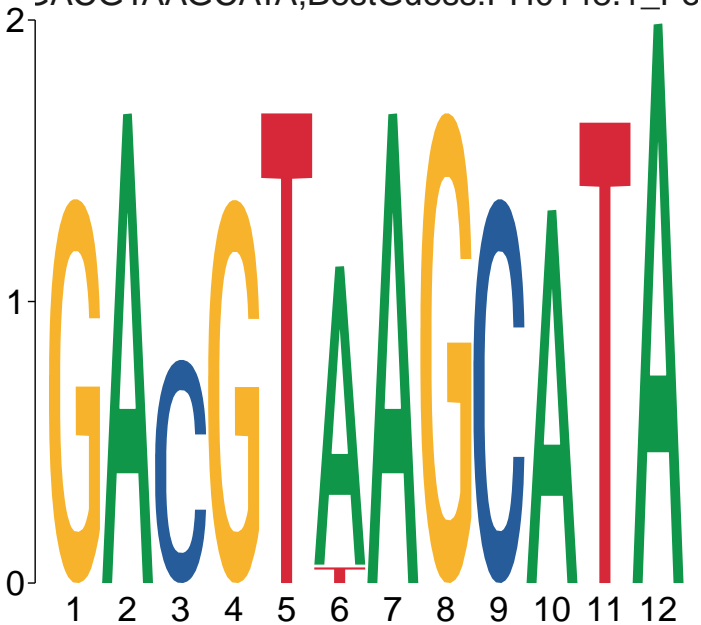
9-CGTCCTTAGG,BestGuess:ZmHOX2a



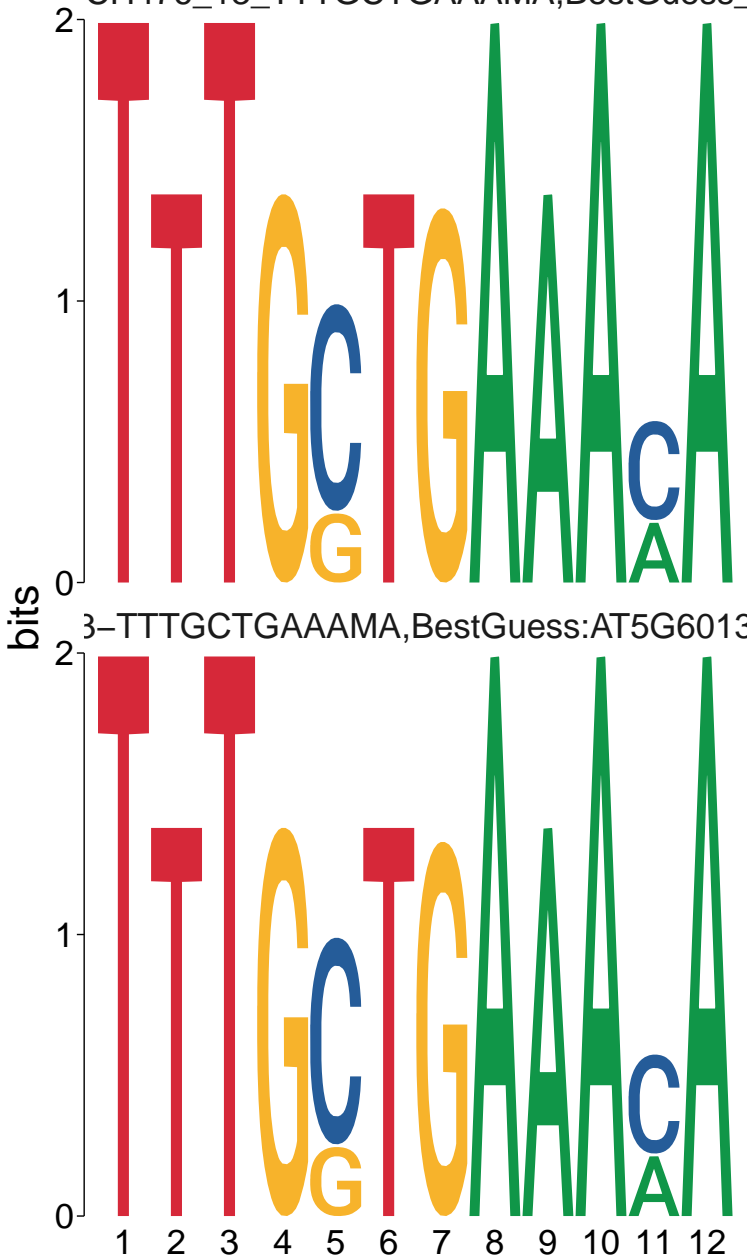
CH473_12_GACGTAAGCATA,BestGuess_



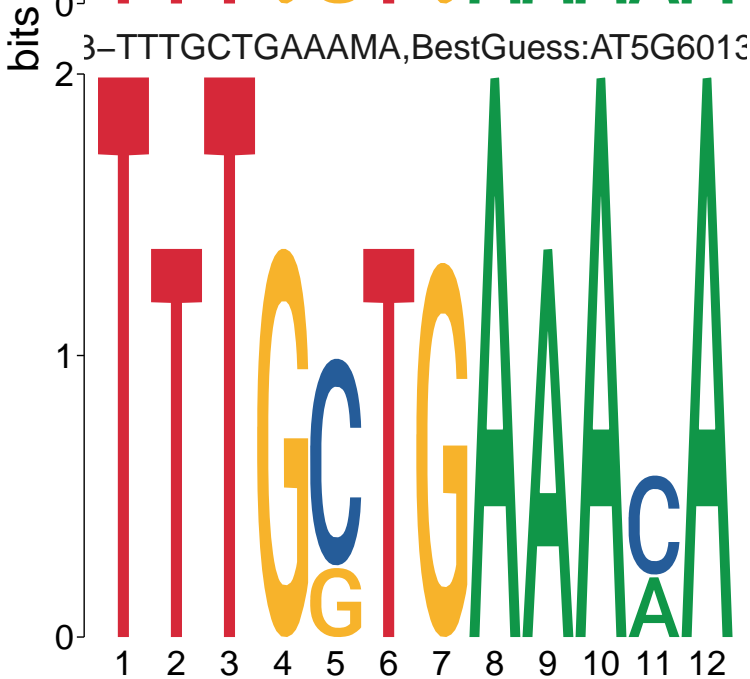
3ACGTAAGCATA,BestGuess:PH0148.1_Po



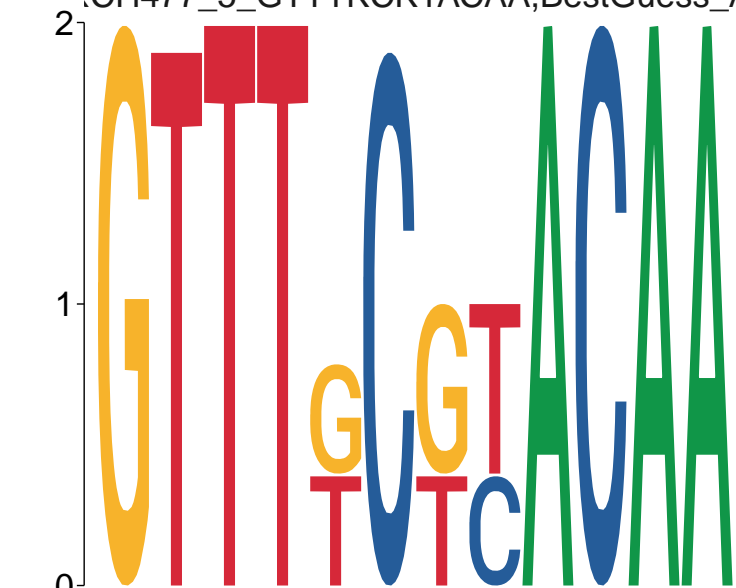
CH476_13_TTTGCTGAAAMA,BestGuess_



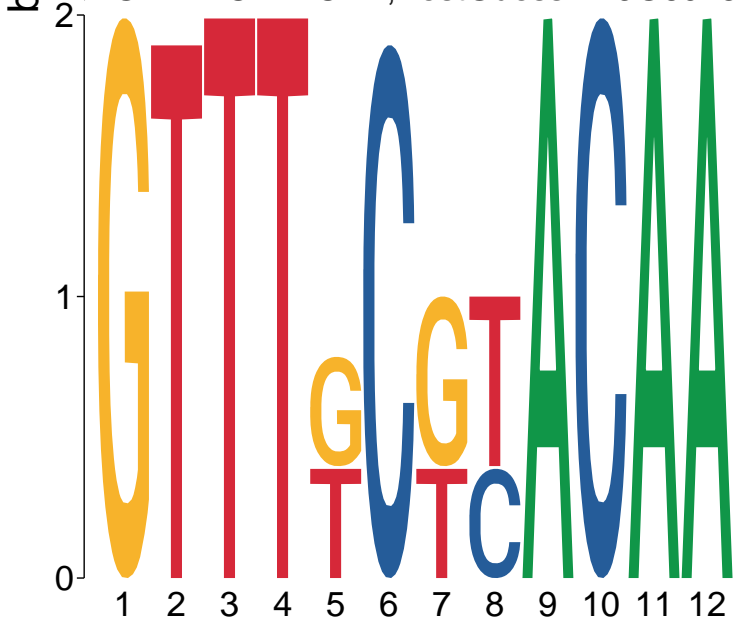
3-TTTGCTGAAAMA,BestGuess:AT5G6013



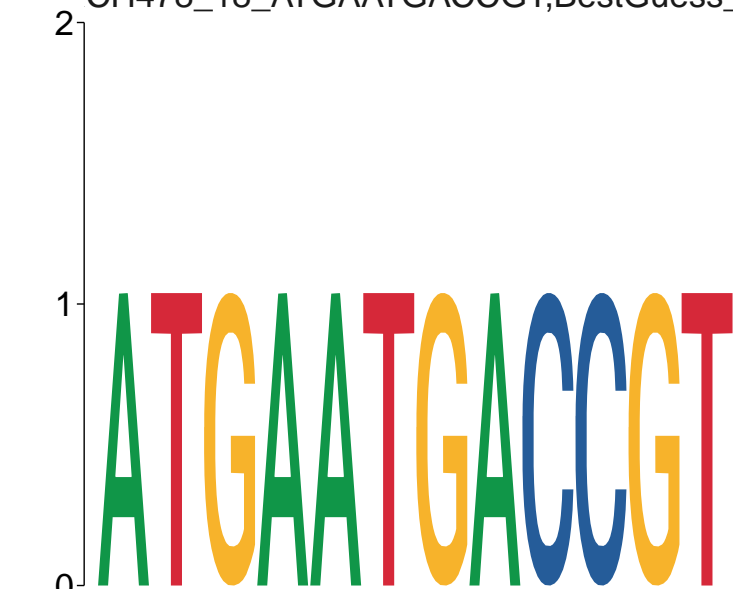
CH477_5_GTTTKCKYACAA,BestGuess_A



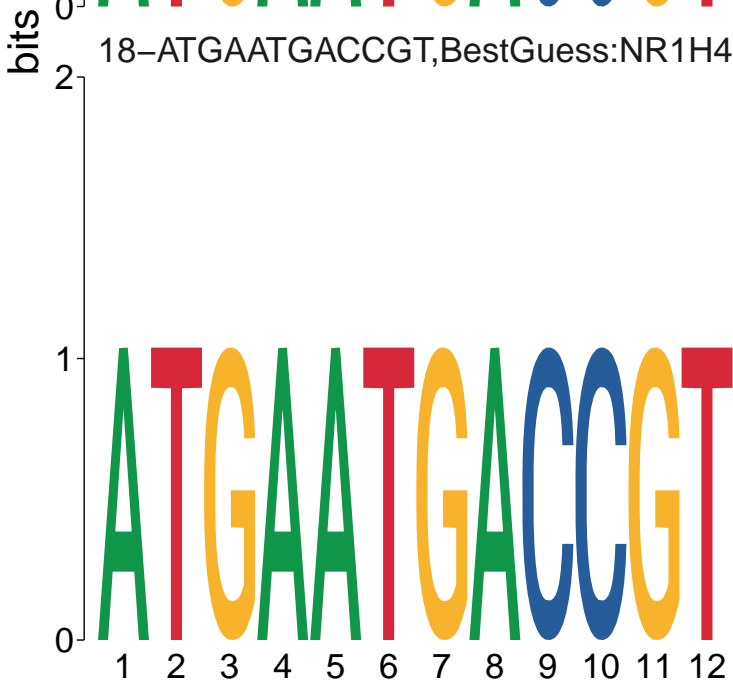
5-GTTTKCKYACAA,BestGuess:AT5G60130



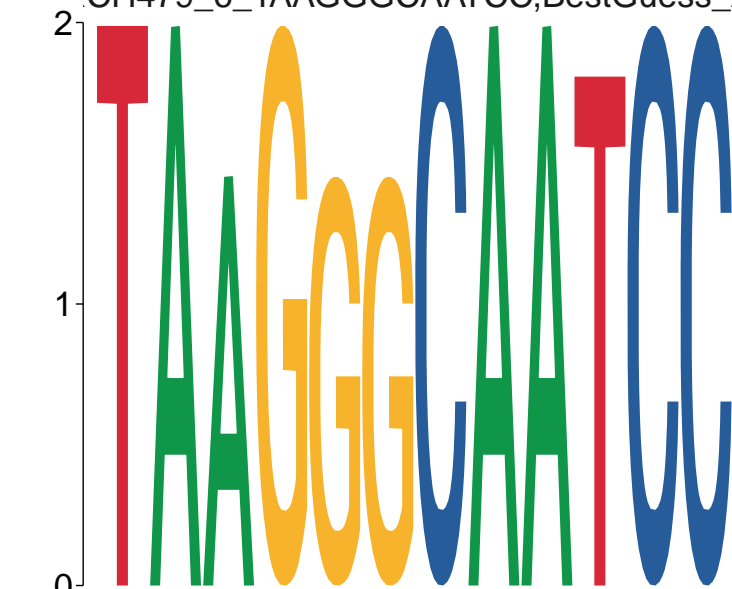
CH478_18_ATGAATGACCGT,BestGuess_I



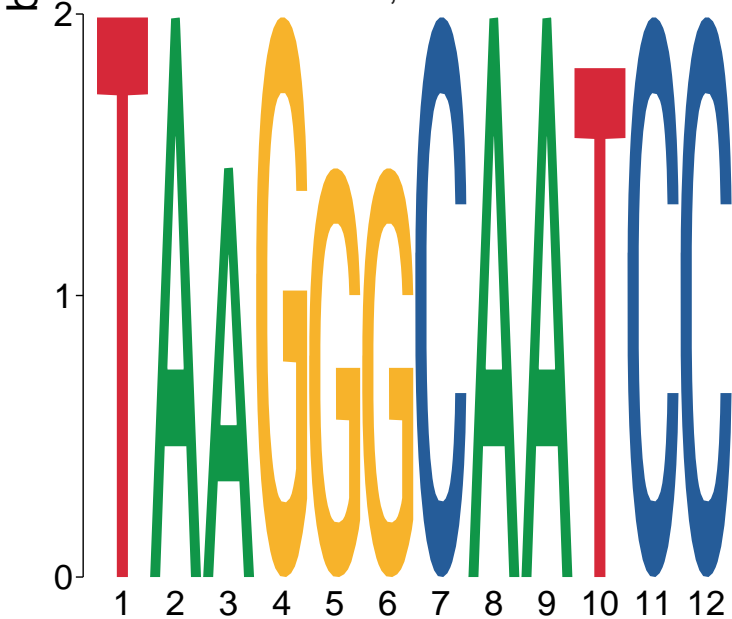
18-ATGAATGACCGT,BestGuess:NR1H4



CH479_6_TAAGGGCAATCC,BestGuess_A



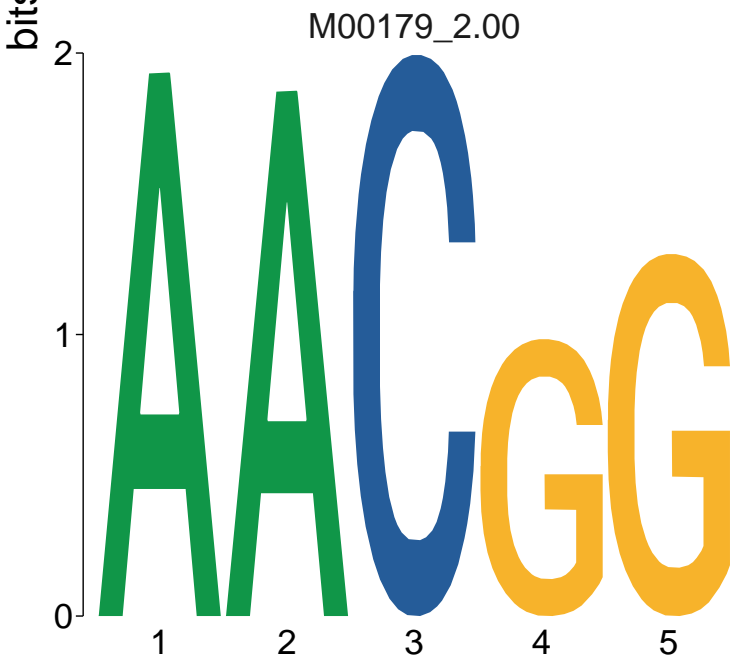
TAAGGGCAATCC,BestGuess:AT2G3355



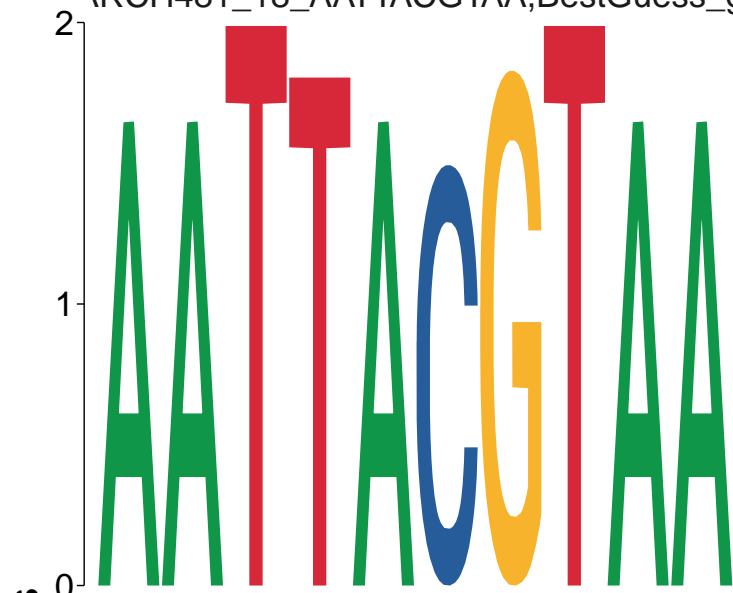
ARCH480_MYB_MYBL1



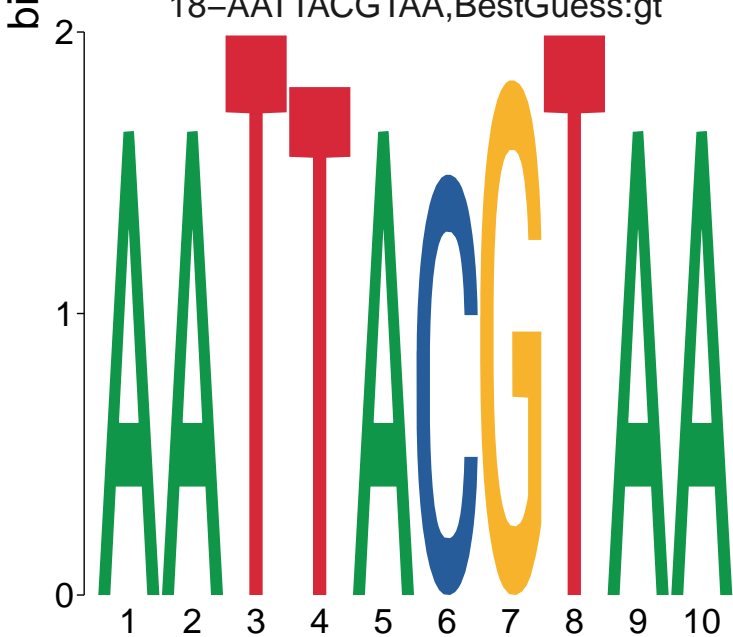
M00179_2.00



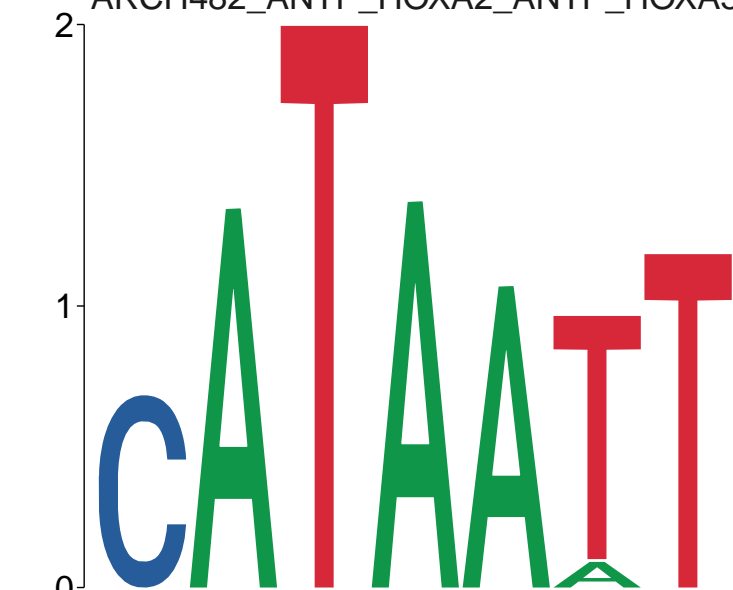
ARCH481_18_AATTACGTAA,BestGuess_g



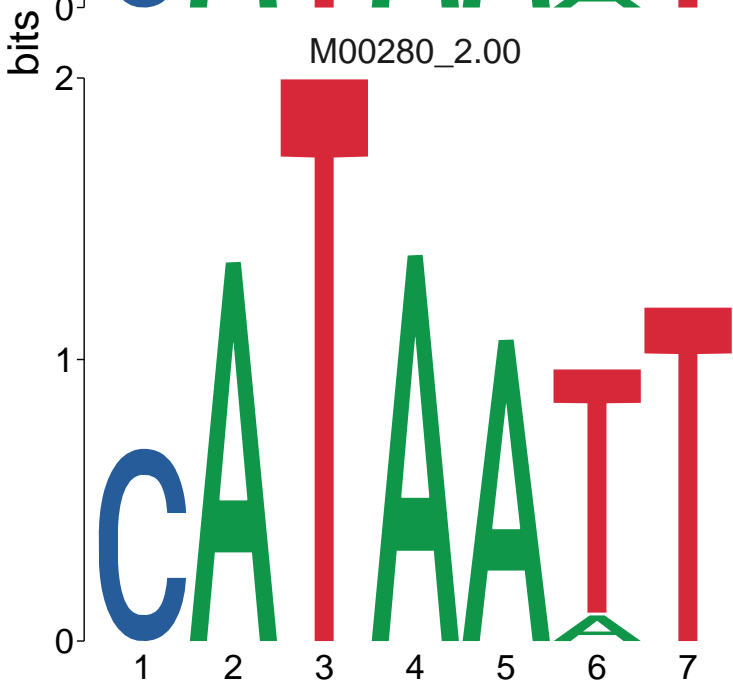
18-AATTACGTAA,BestGuess:gt



ARCH482_ANTP_HOXA2_ANTP_HOXA3



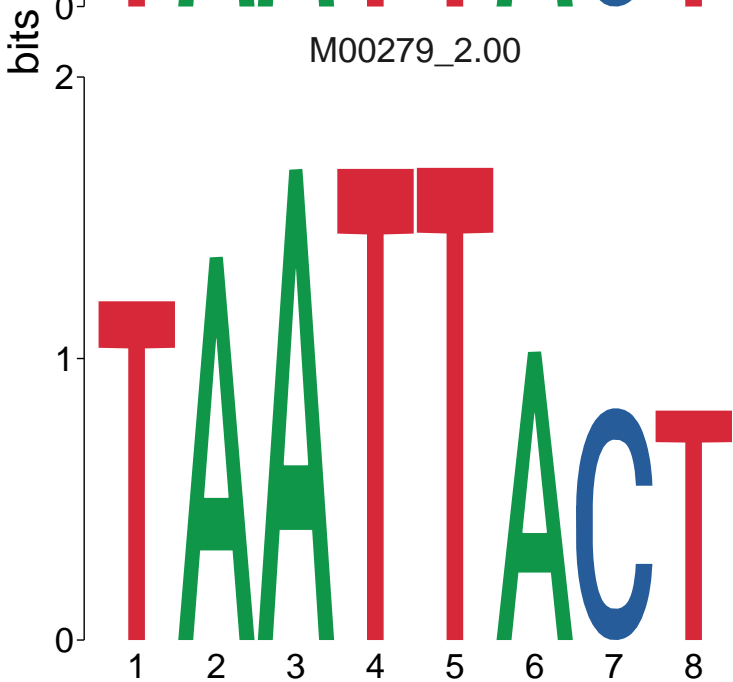
M00280_2.00



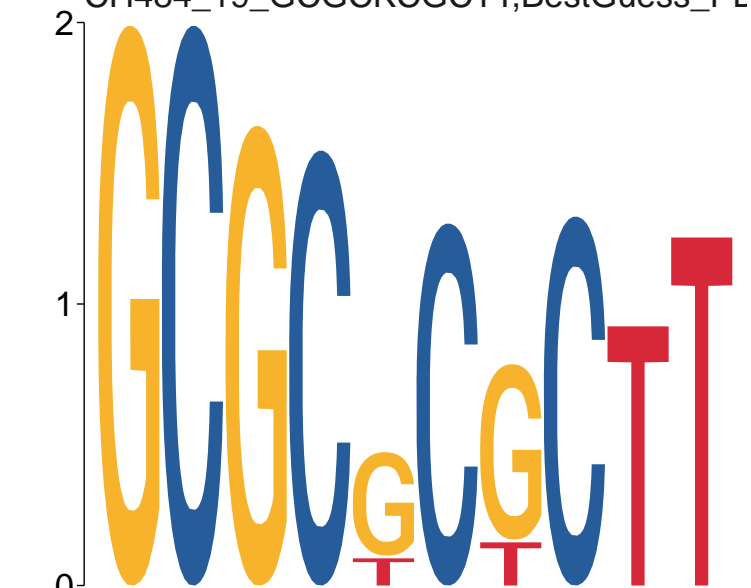
ARCH483_ANTP_MNX1



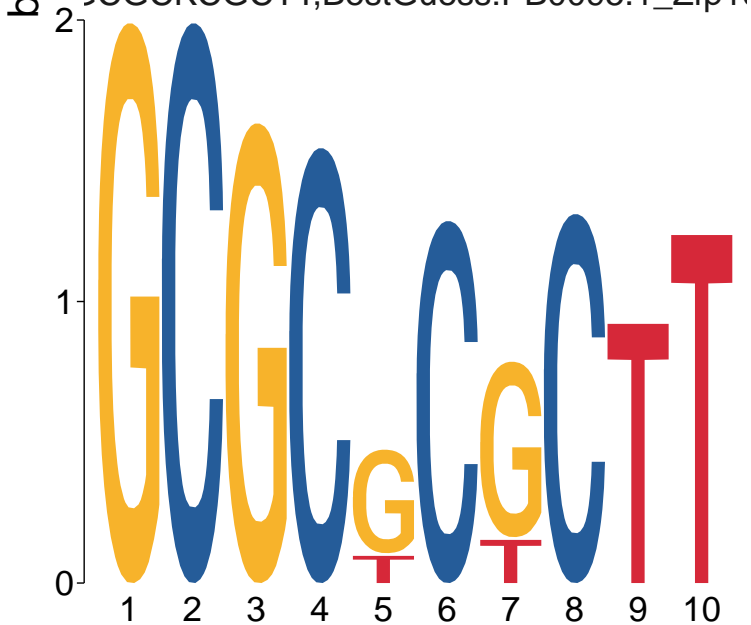
M00279_2.00



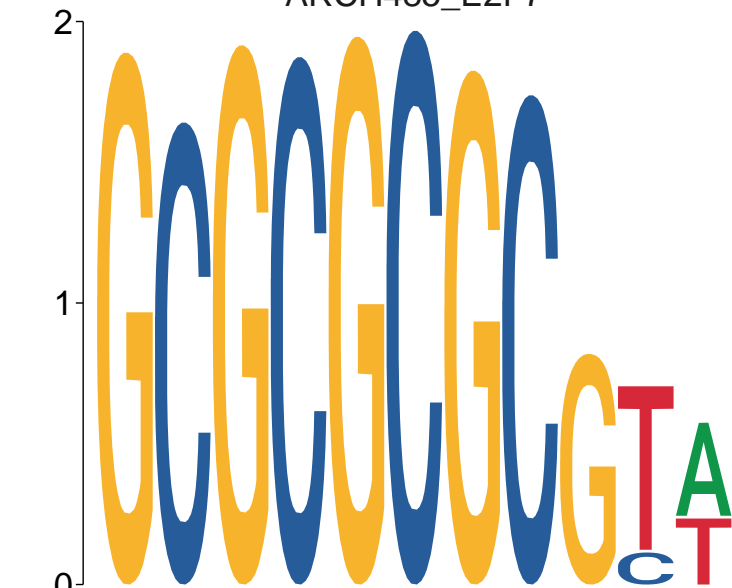
CH484_19_GCGCKCGCTT,BestGuess_PE



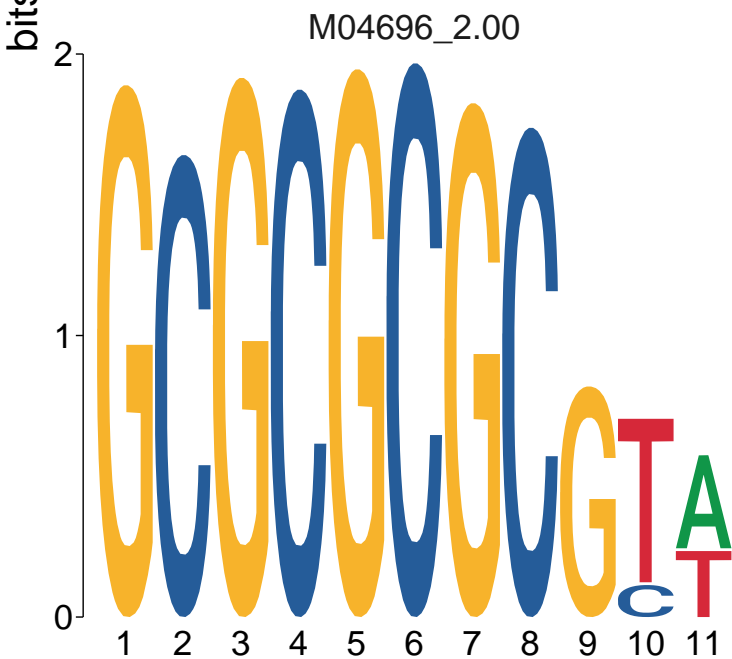
GCGCKCGCTT,BestGuess:PB0095.1_Zfp10



ARCH485_E2F7



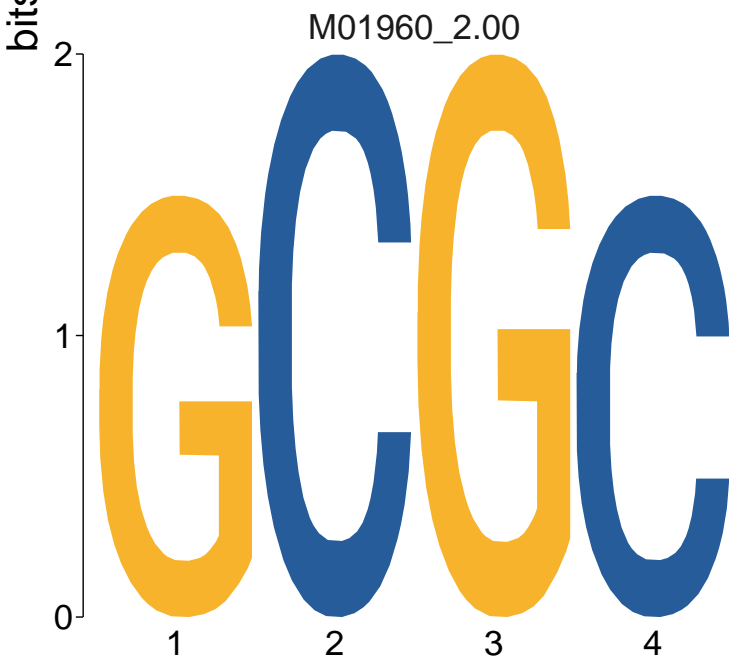
M04696_2.00



ARCH486_Nvec_v1g200125



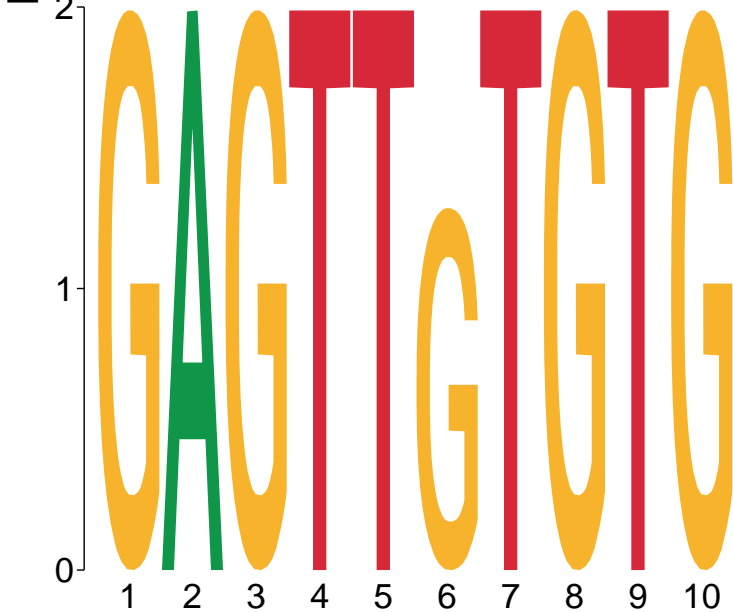
M01960_2.00



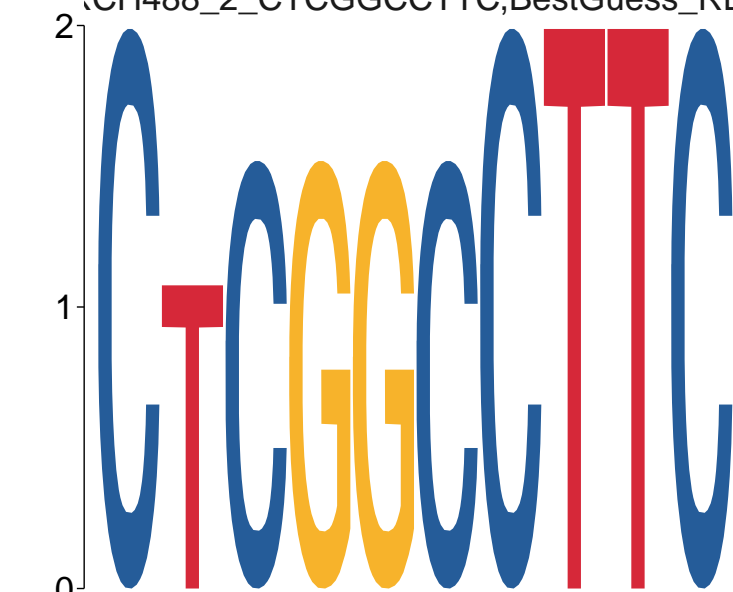
CH487_21_GAGTTGTGTG,BestGuess_PE



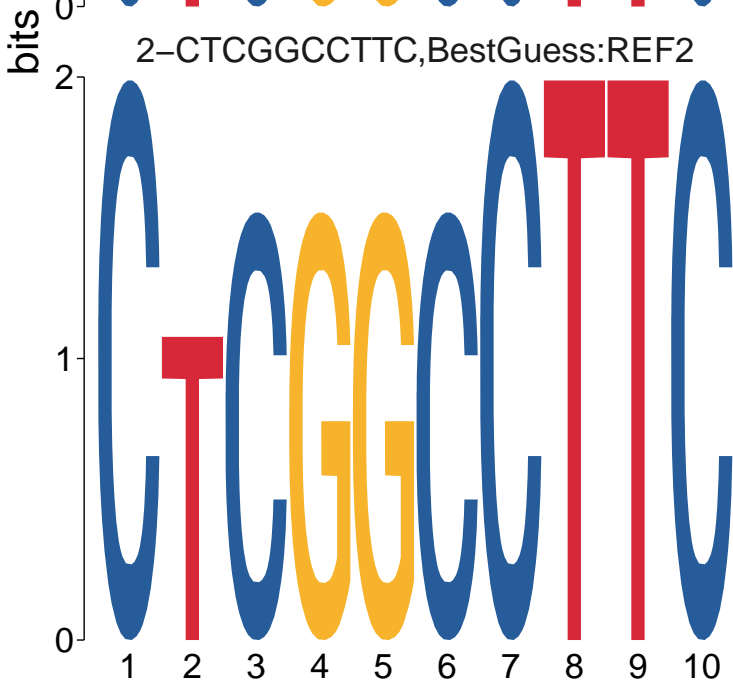
GAGTTGTGTG,BestGuess:PB0208.1_Zsca



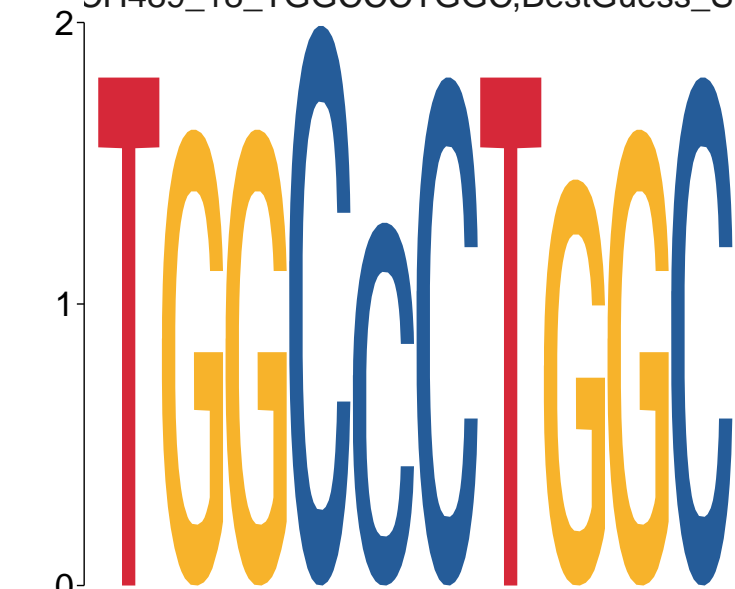
CH488_2_CTCGGCCTTC,BestGuess_RE



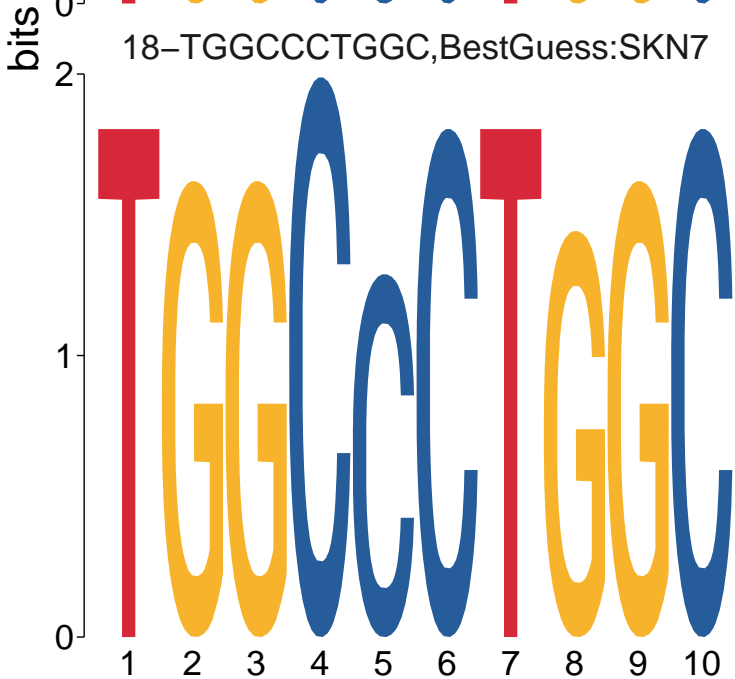
2-CTCGGCCTTC,BestGuess:REF2



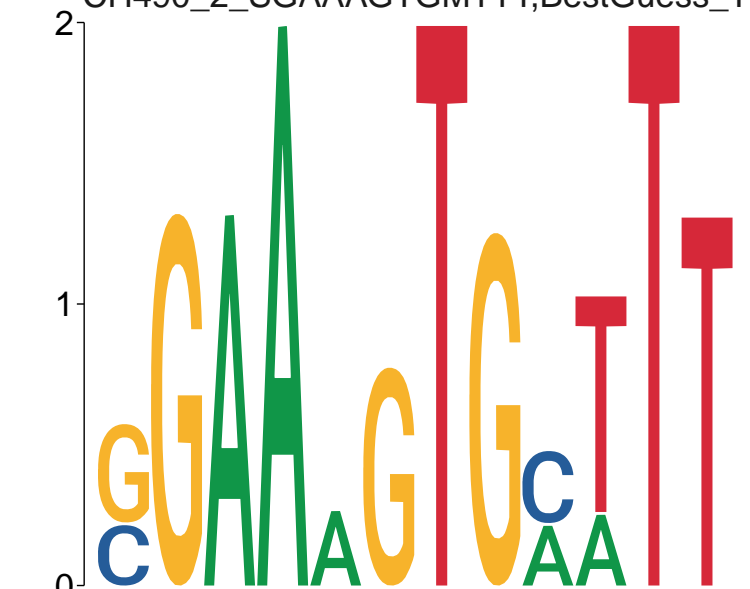
CH489_18_TGGCCCTGGC,BestGuess_Sk



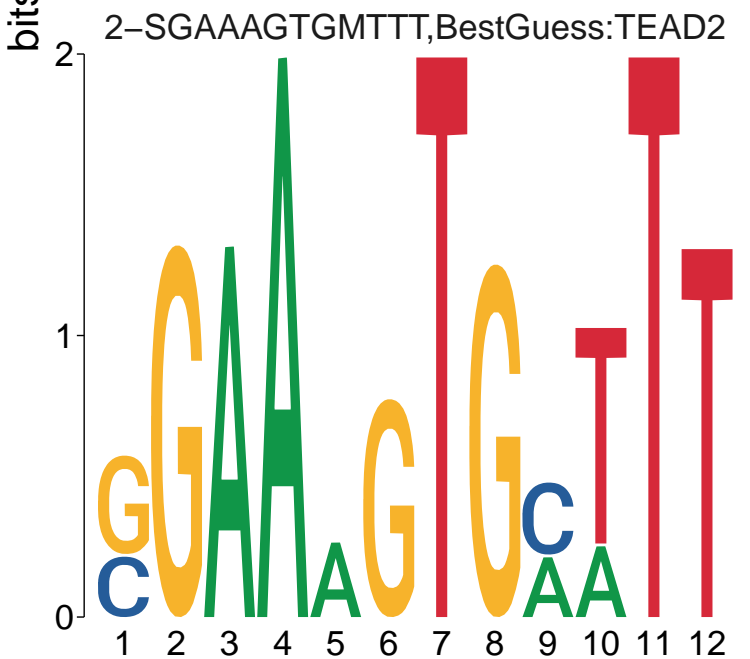
18-TGGCCCTGGC,BestGuess:SKN7



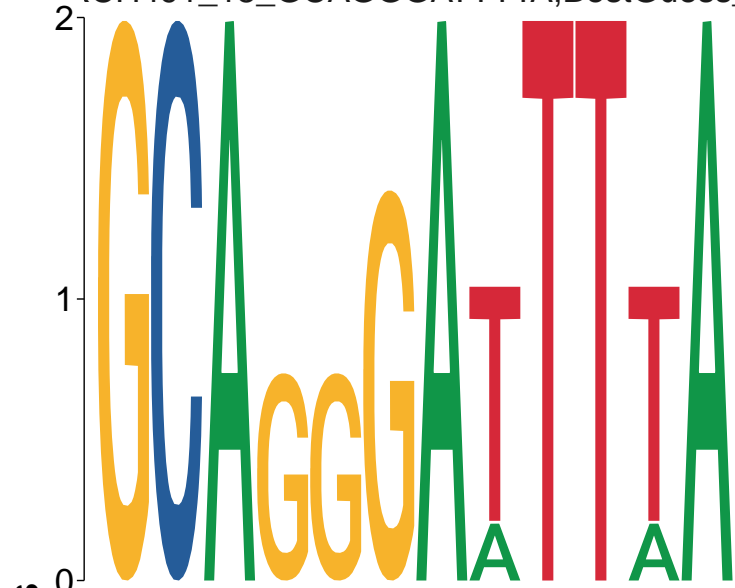
CH490_2_SGAAAGTGMTTT,BestGuess_T



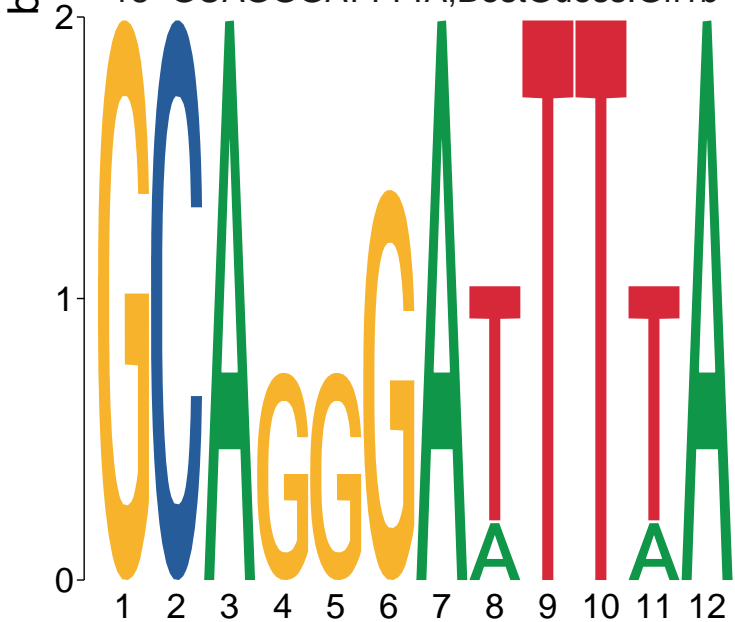
2-SGAAAGTGMTTT,BestGuess:TEAD2



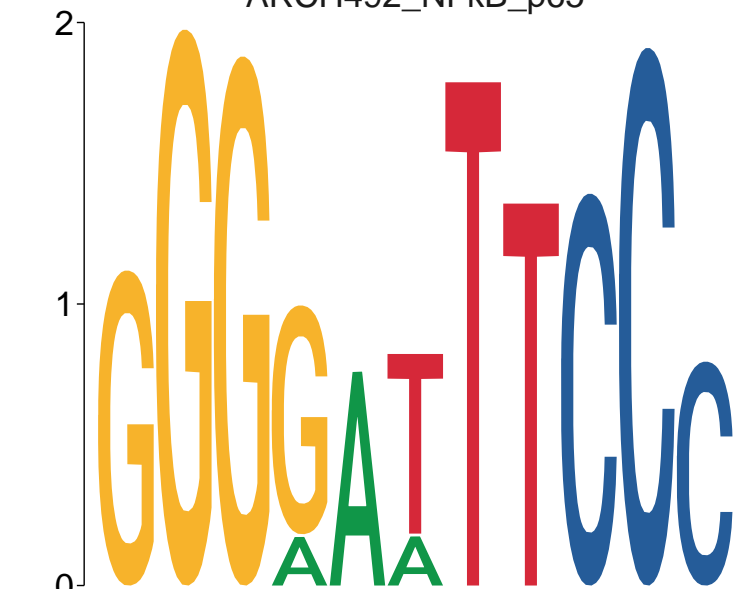
CH491_18_GCAGGGATTTTA,BestGuess_



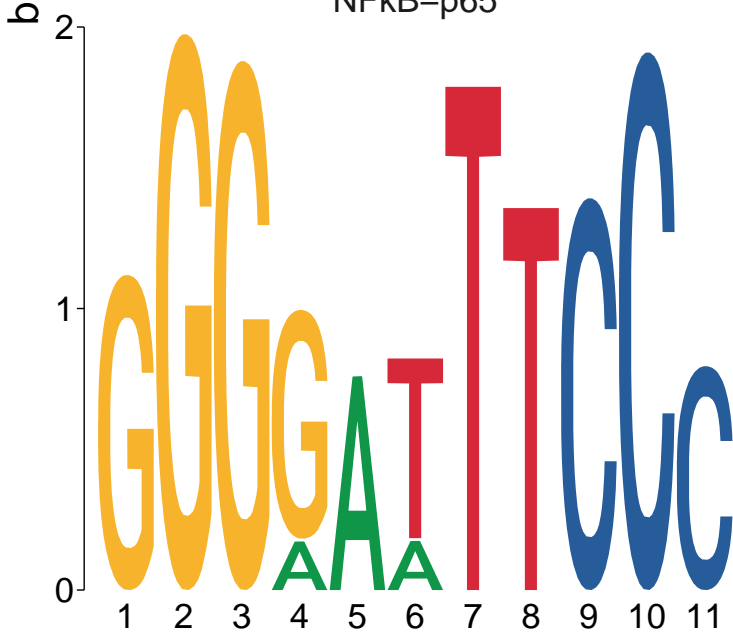
18-GCAGGGATTTTA,BestGuess:Gfi1b



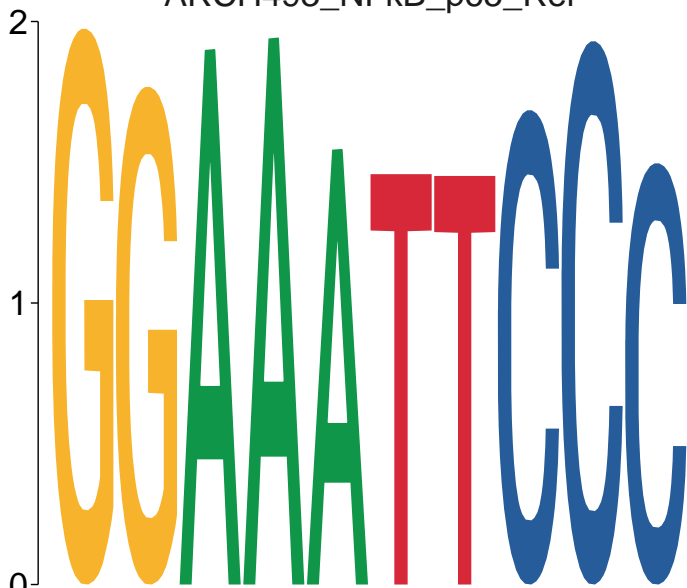
ARCH492_NFkB_p65



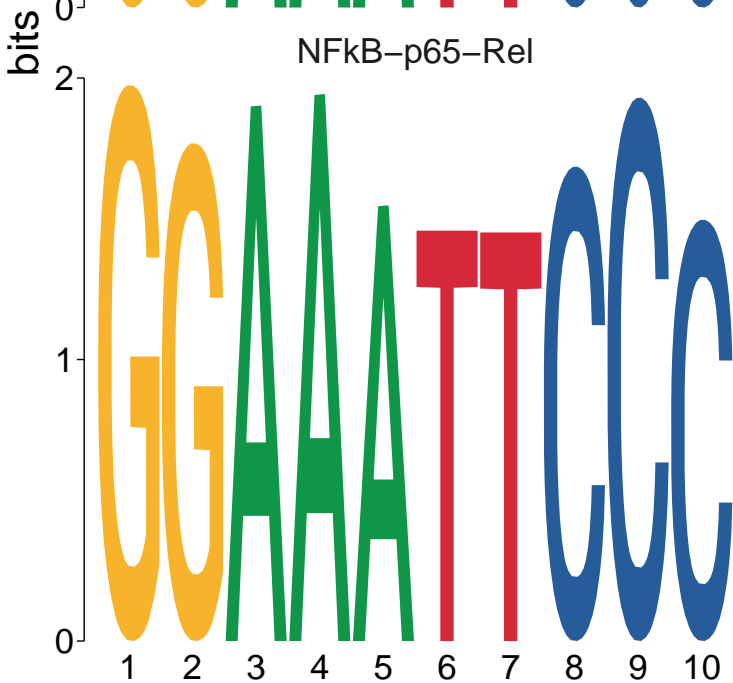
NFkB-p65



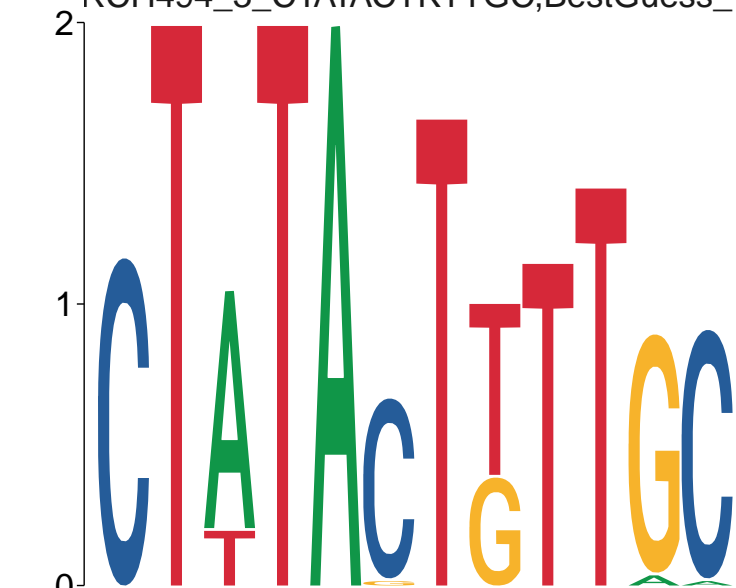
ARCH493_NFkB_p65_Rel



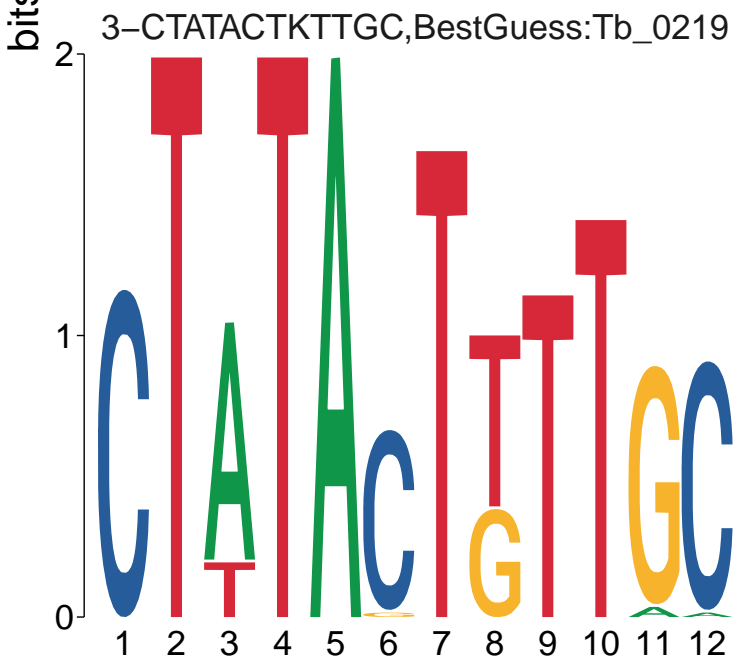
NFkB-p65-Rel



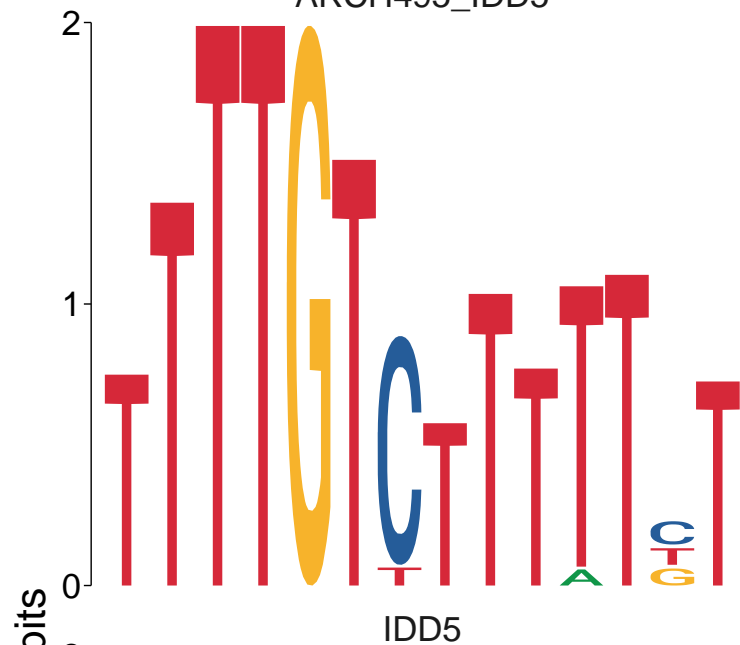
RCH494_3_CTATACTKTTGC,BestGuess_T



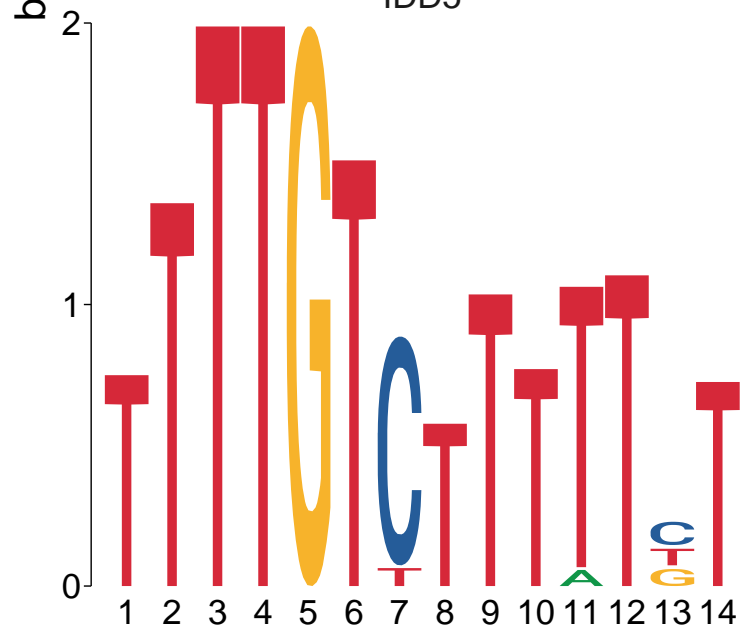
3-CTATACTKTTGC,BestGuess:Tb_0219



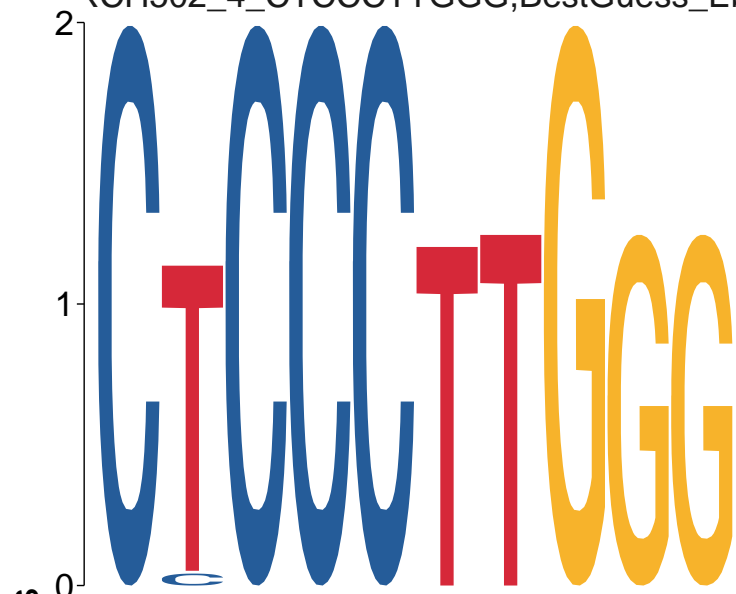
ARCH495_IDD5



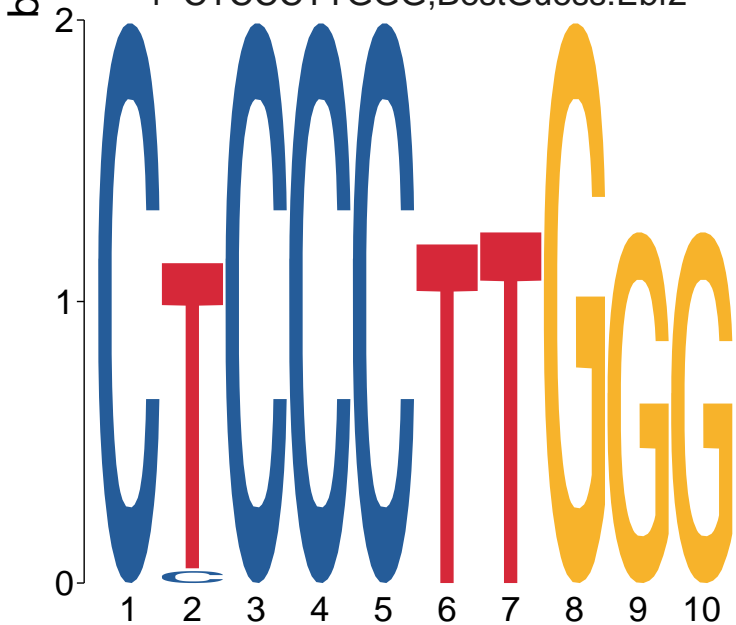
IDD5



RCH502_4_CTCCCTTGGG,BestGuess_Eb



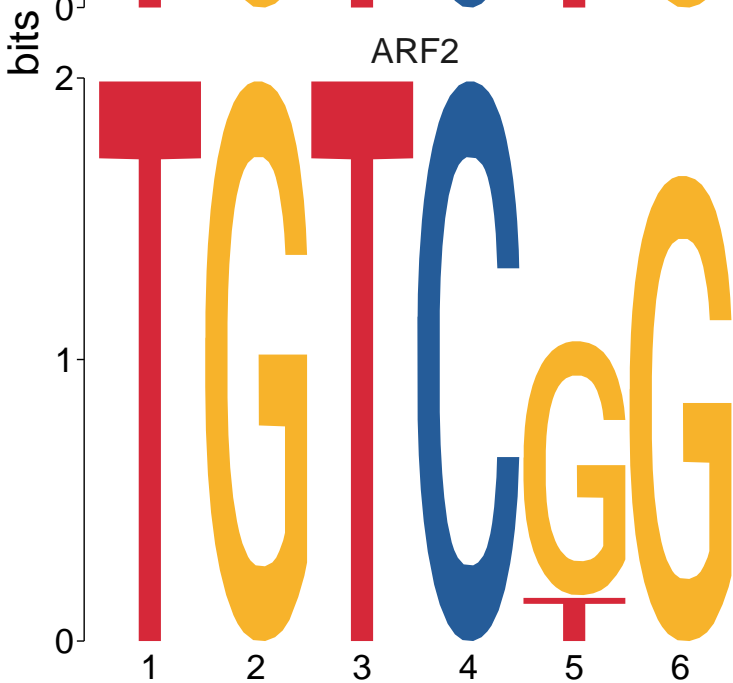
4-CTCCCTTGGG,BestGuess:Ebf2



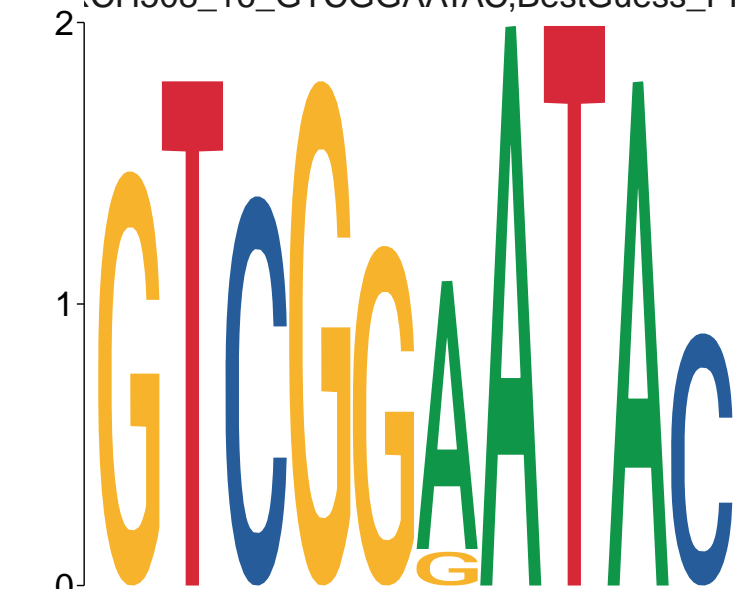
ARCH507_ARF2



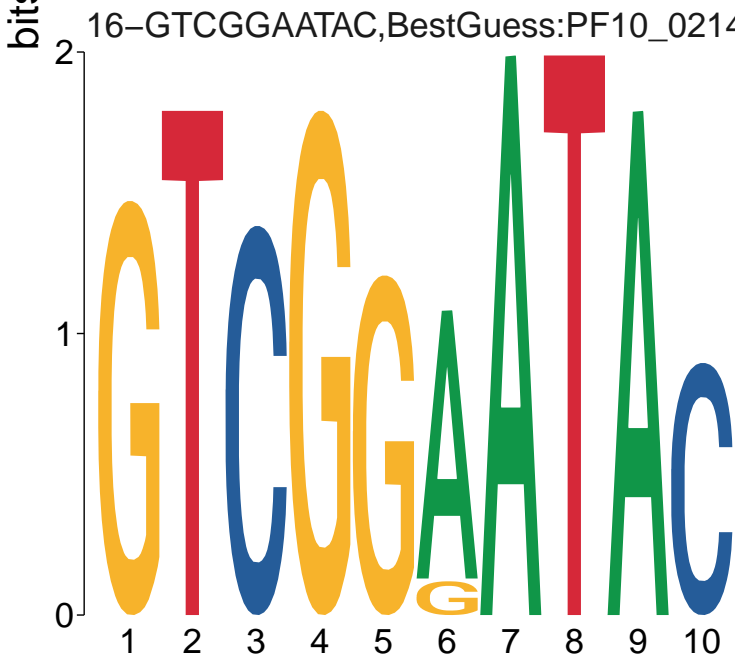
ARF2



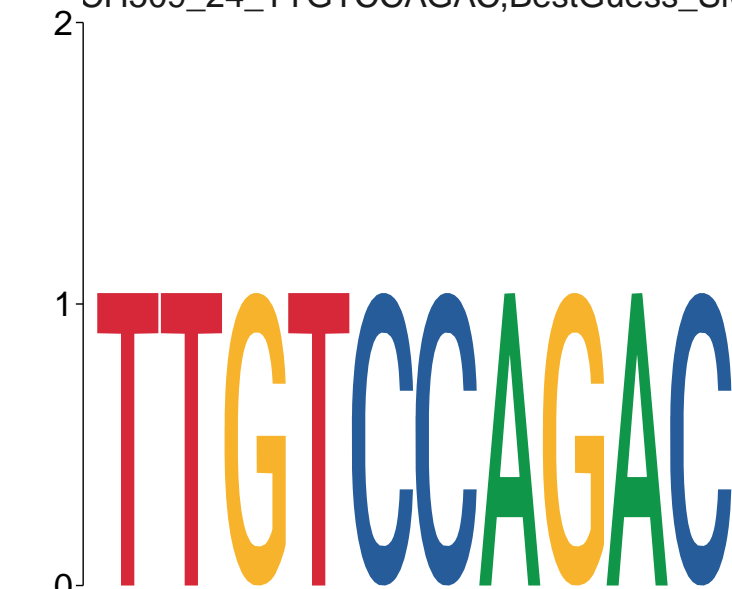
CH508_16_GTCGGAATAC,BestGuess_PF



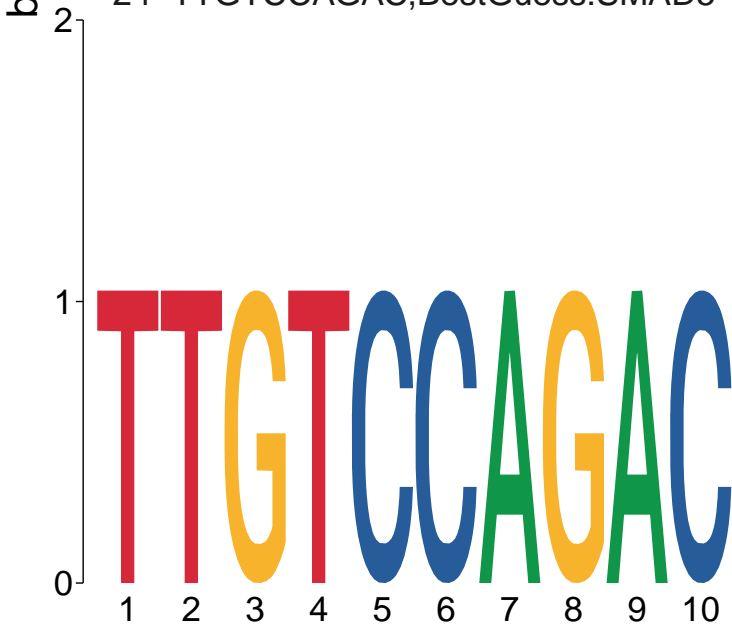
16-GTCGGAATAC,BestGuess:PF10_0214



CH509_24_TTGTCCAGAC,BestGuess_SM



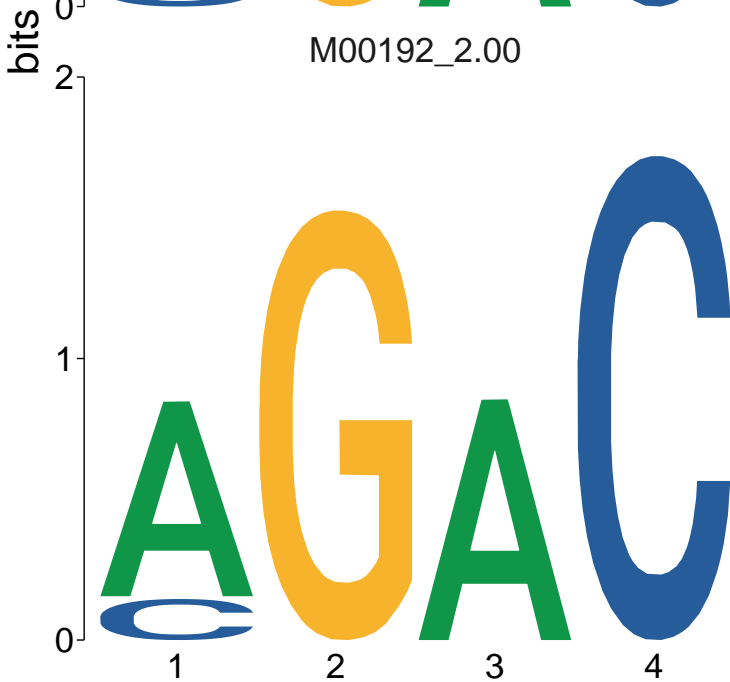
24-TTGTCCAGAC,BestGuess:SMAD5



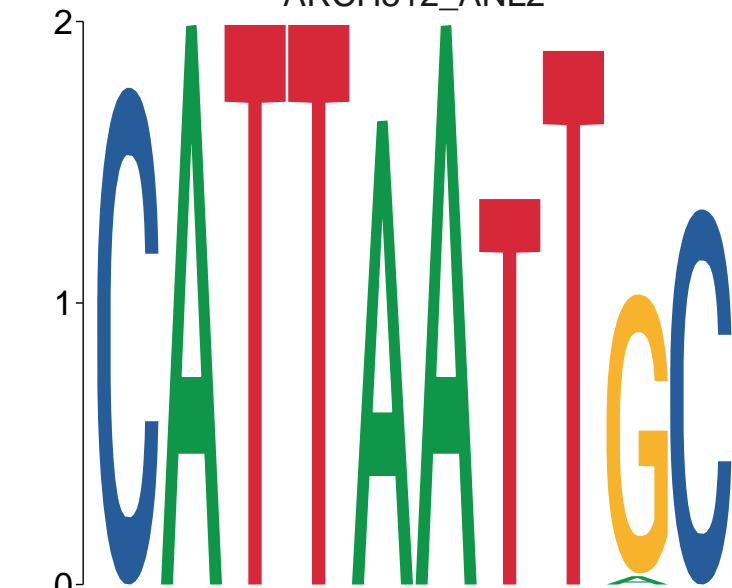
ARCH510_SMAD2



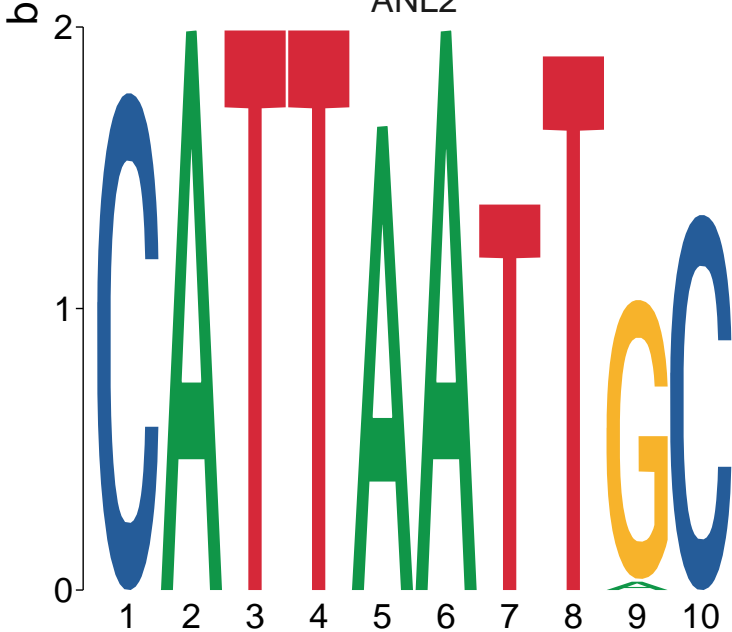
M00192_2.00



ARCH512_ANL2



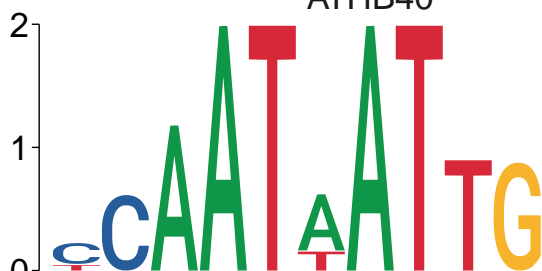
ANL2



ARCH513_HAT5_HAT5_ATHB40



ATHB40



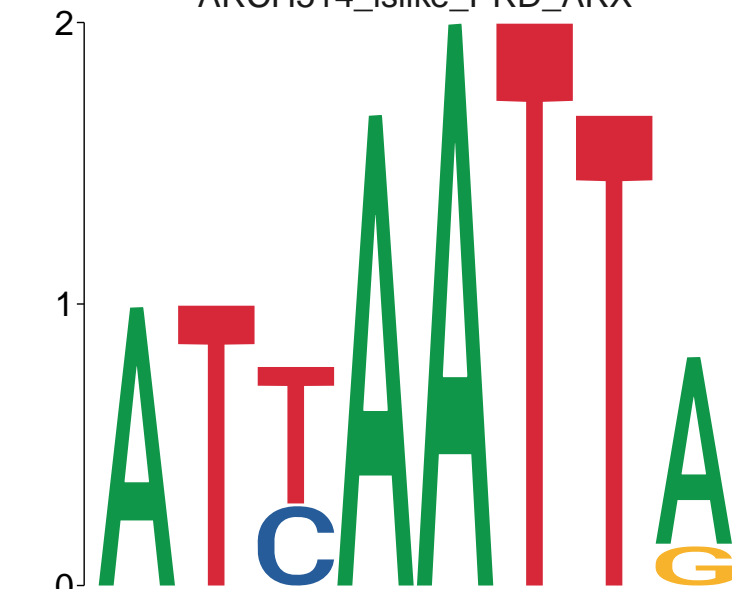
HAT5



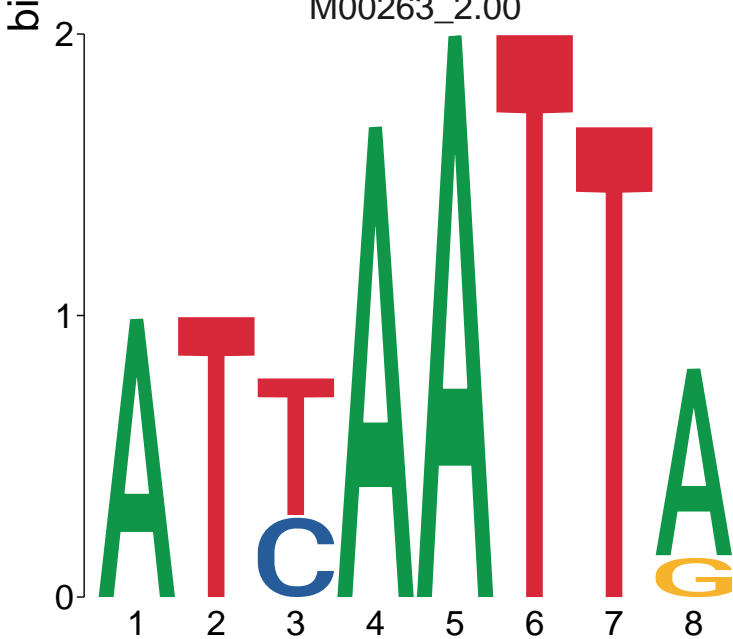
HAT5.1



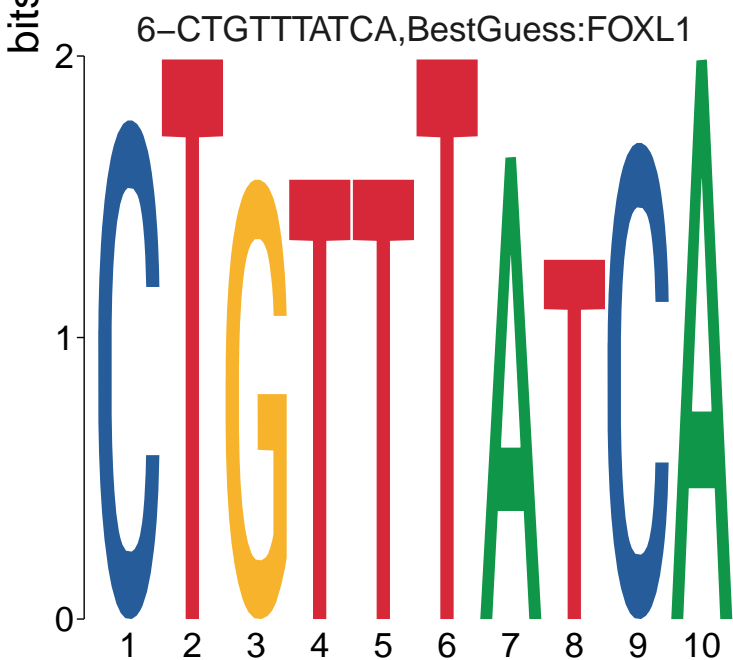
ARCH514_islike_PRD_ARX



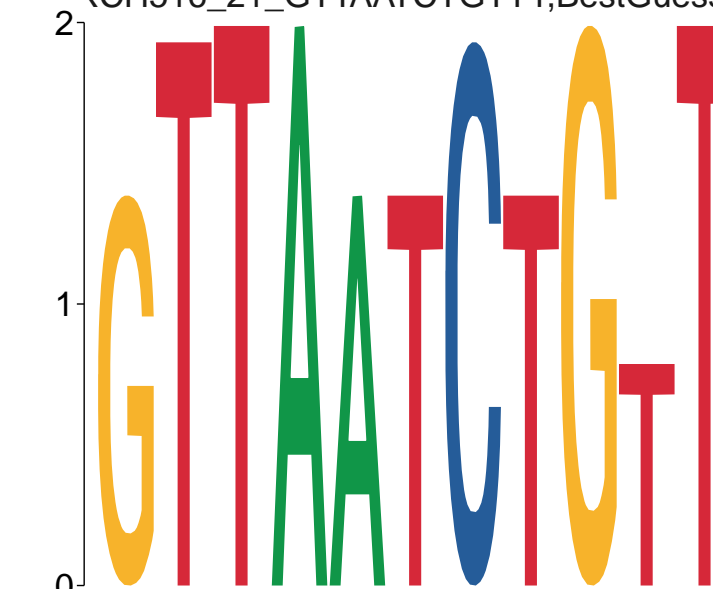
M00263_2.00



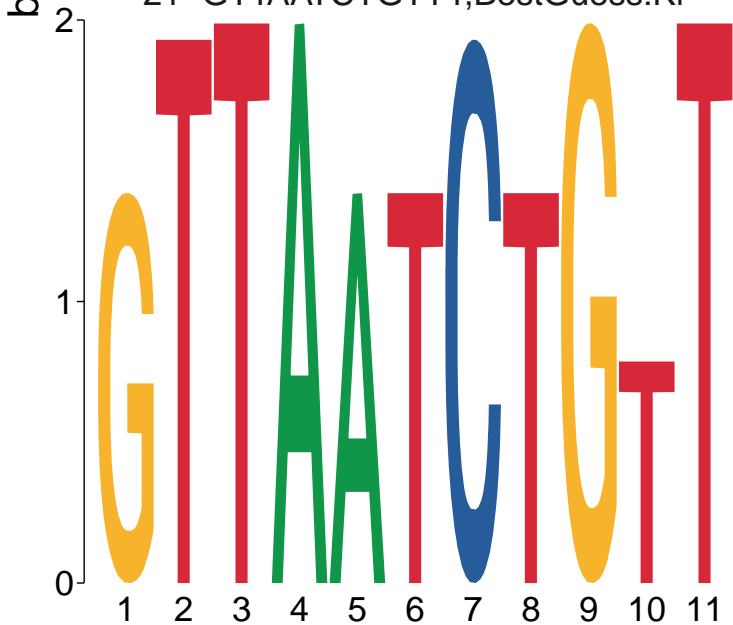
CH515_6_CTGTTTATCA,BestGuess_FOX



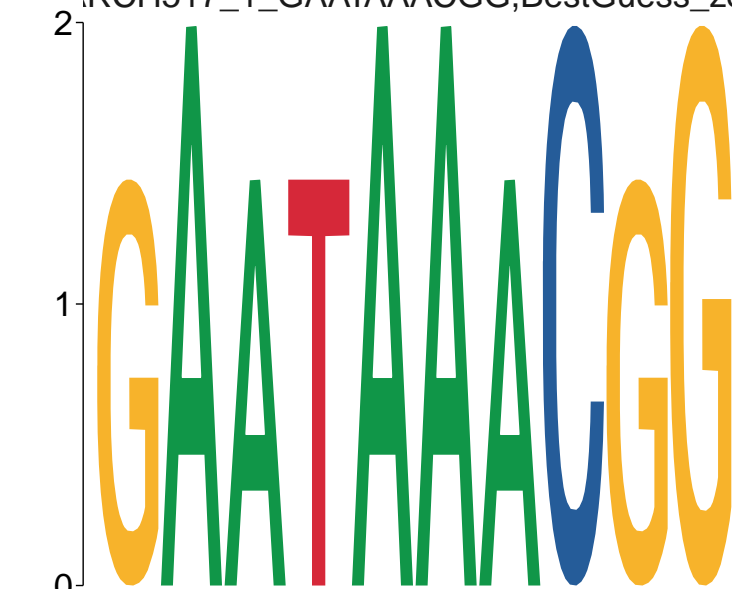
RCH516_21_GTTAATCTGTTY,BestGuess_



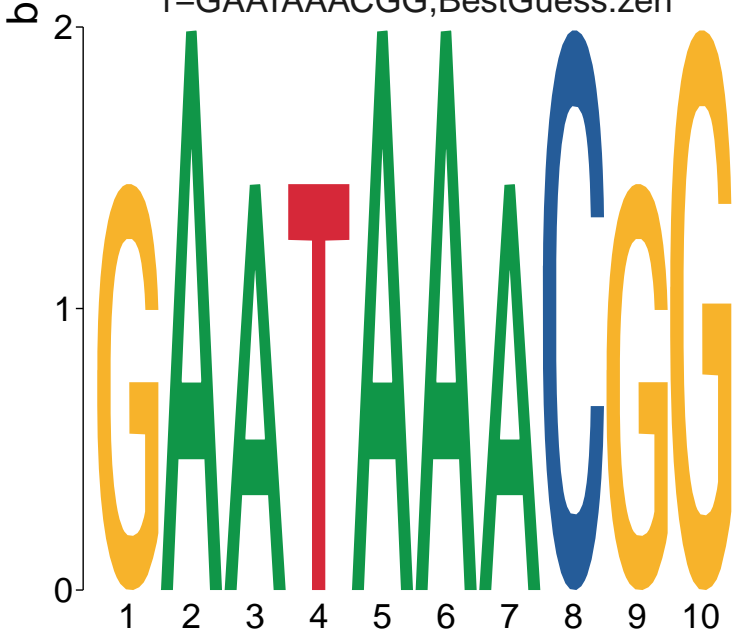
21-GTTAATCTGTTY,BestGuess:Kr



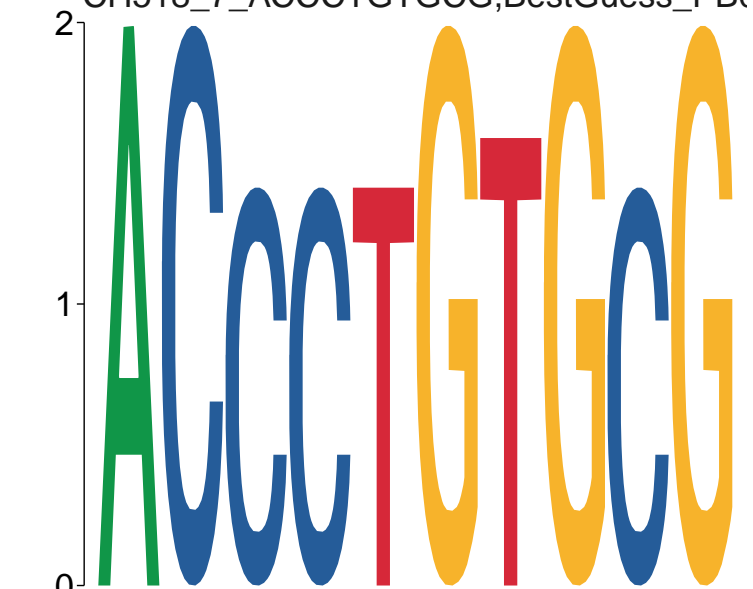
RCH517_1_GAATAAACGG,BestGuess_ze



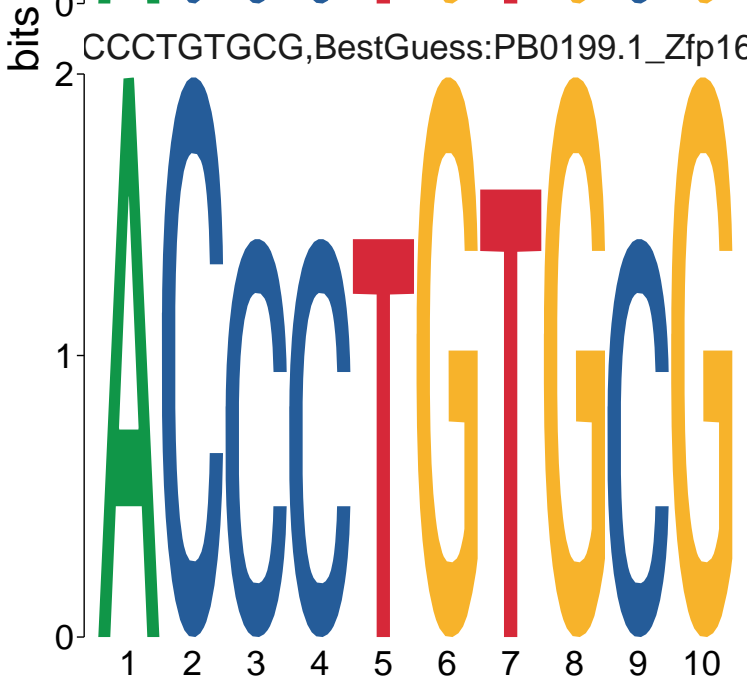
1-GAATAAACGG,BestGuess:zen



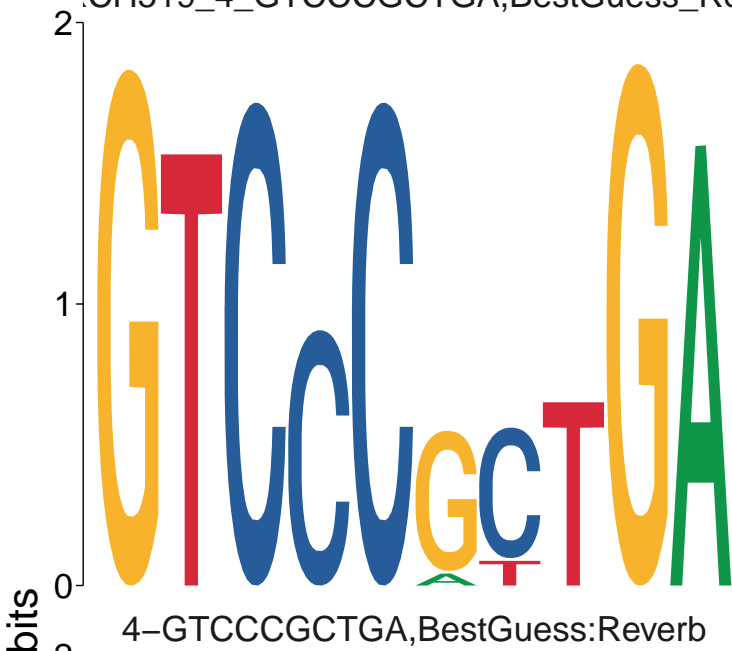
CH518_7_ACCCTGTGCG,BestGuess_PB0



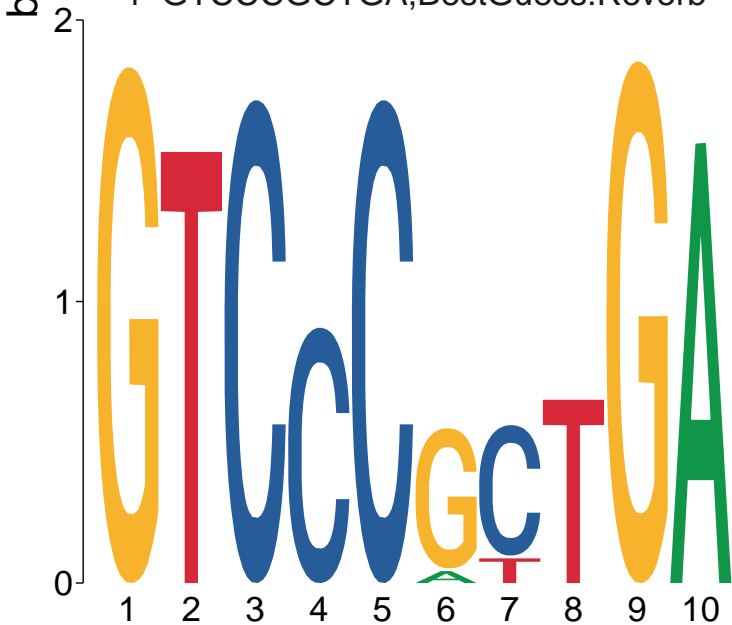
CCCTGTGCG,BestGuess:PB0199.1_Zfp16



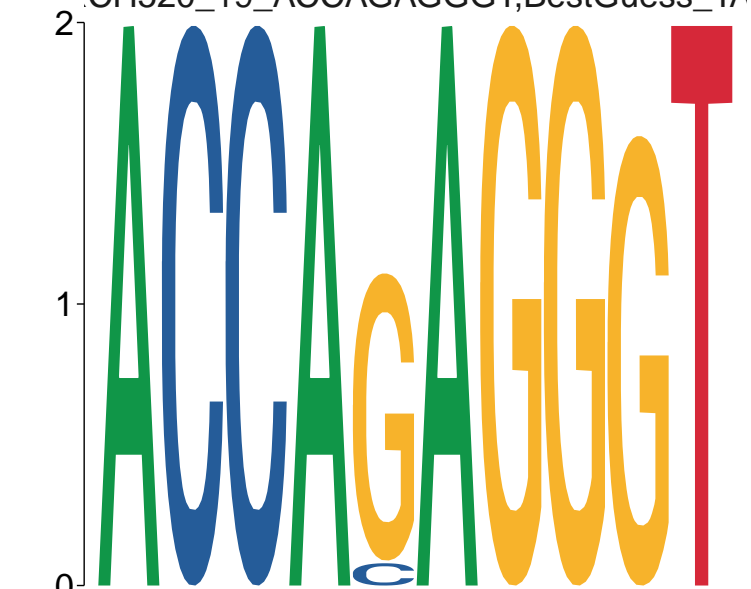
CH519_4_GTCCCGCTGA,BestGuess_Re'



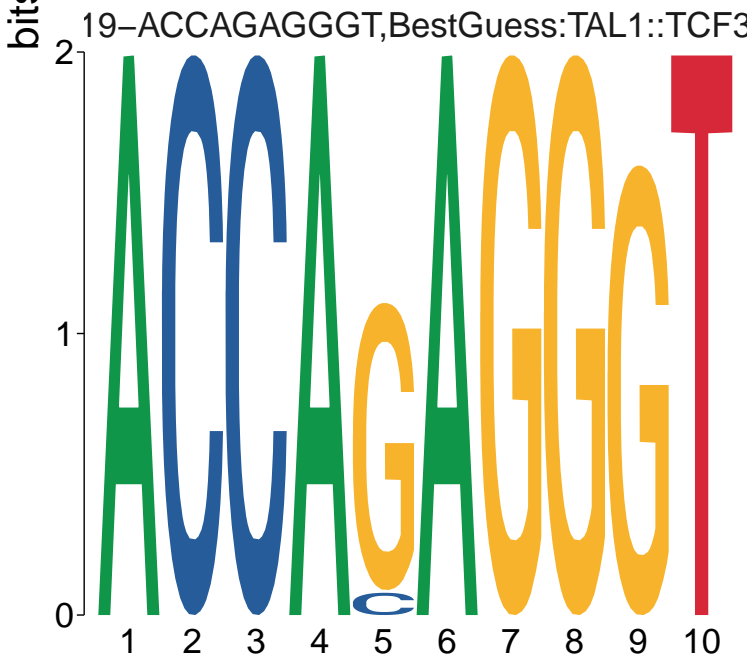
4-GTCCCGCTGA,BestGuess:Reverb



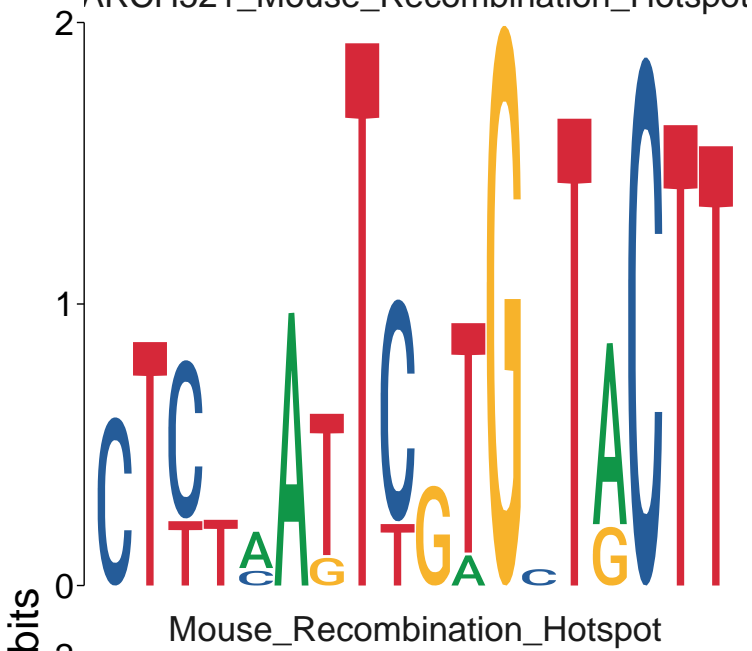
CH520_19_ACCAGAGGGGT,BestGuess_TA



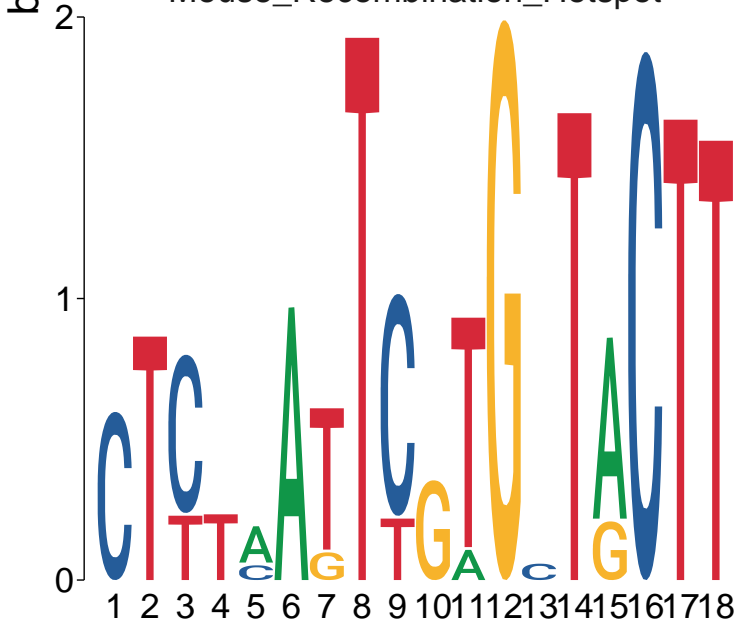
19-ACCAGAGGGGT,BestGuess:TAL1::TCF3

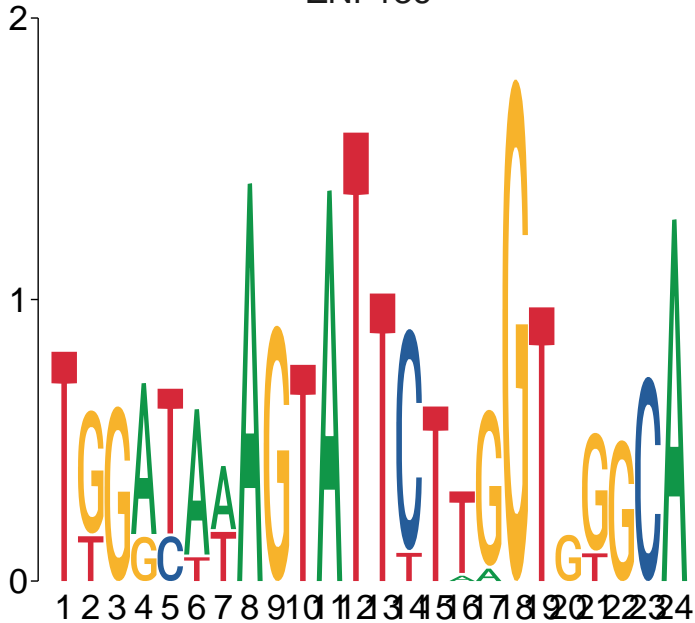
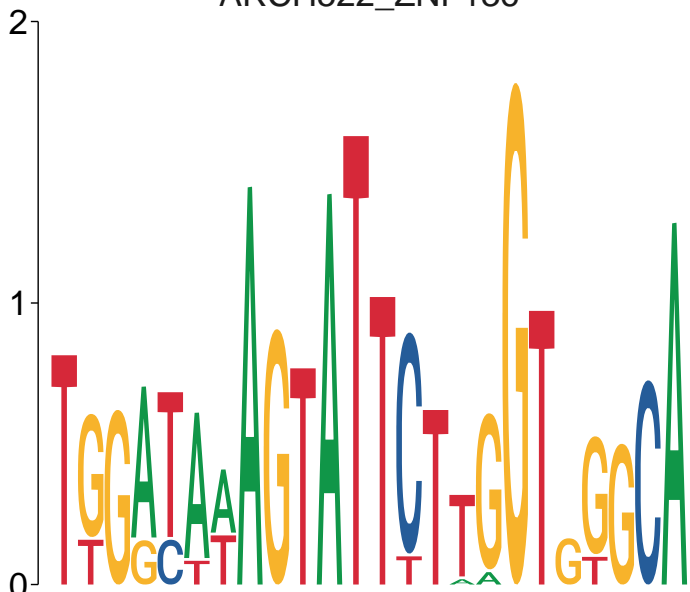


ARCH521_Mouse_Recombination_Hotspot

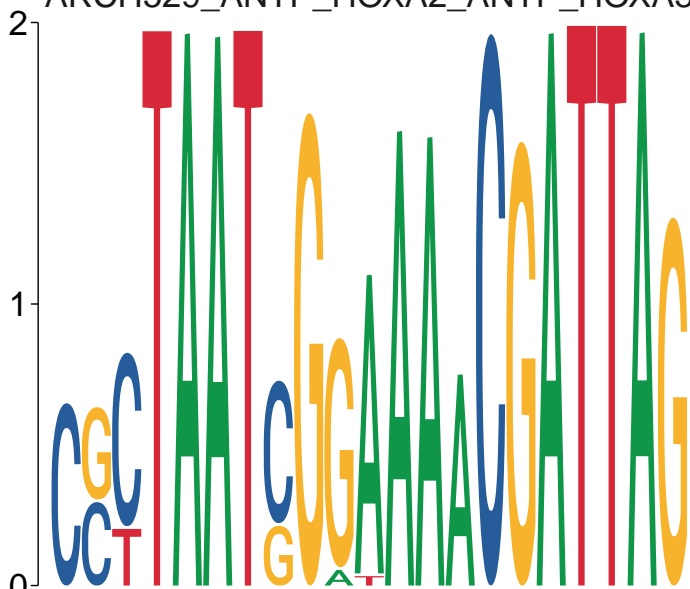


Mouse_Recombination_Hotspot

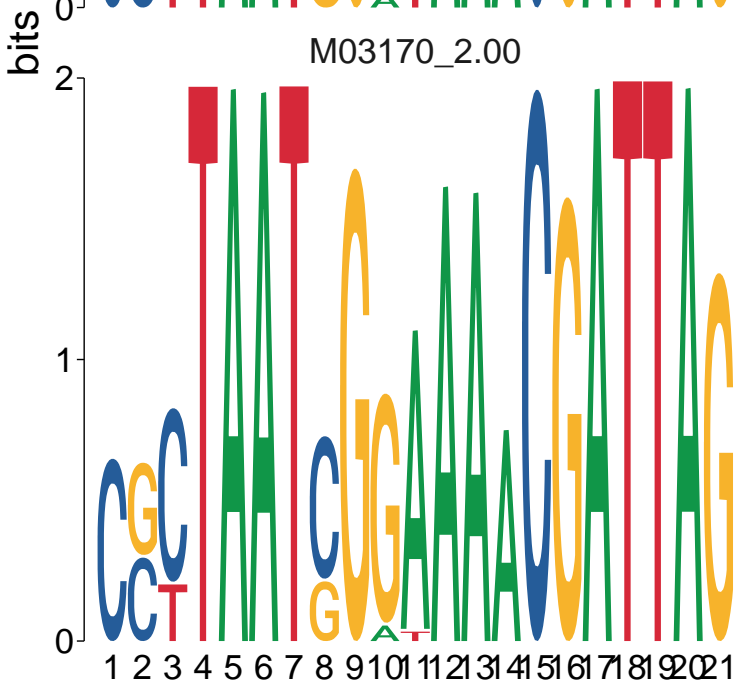




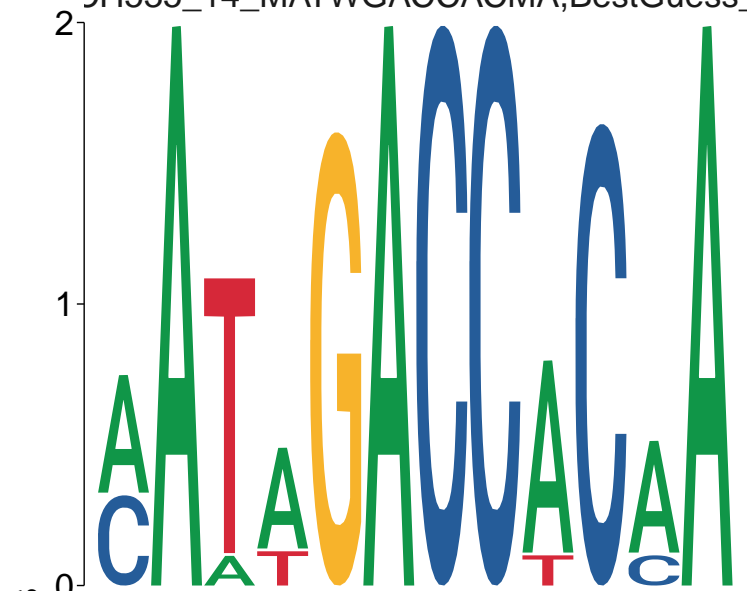
ARCH529_ANTP_HOXA2_ANTP_HOXA3



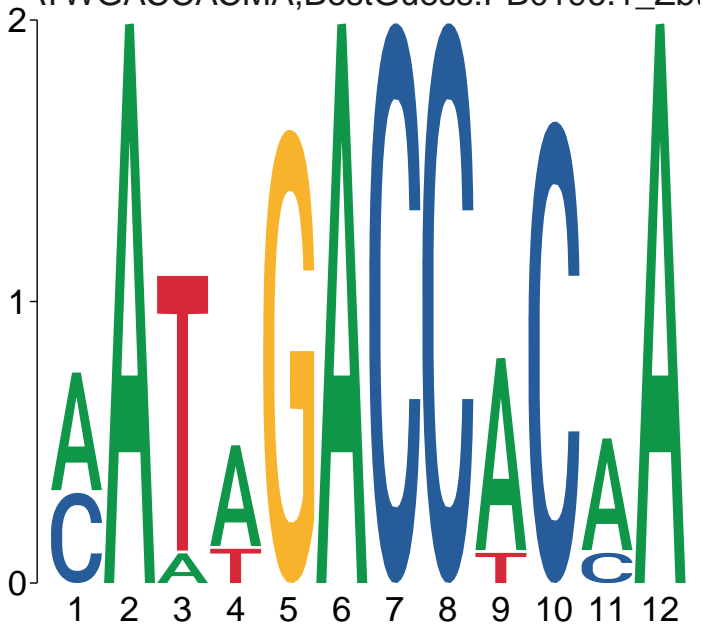
M03170_2.00



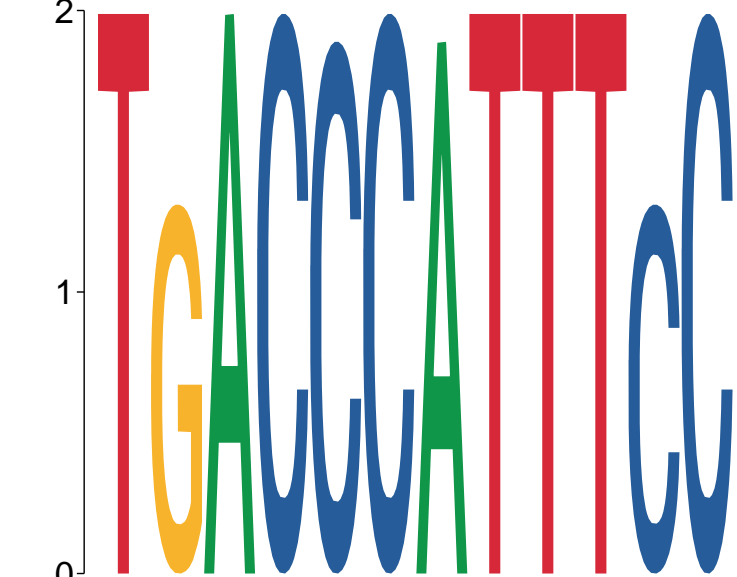
CH535_14_MATWGACCACMA,BestGuess_



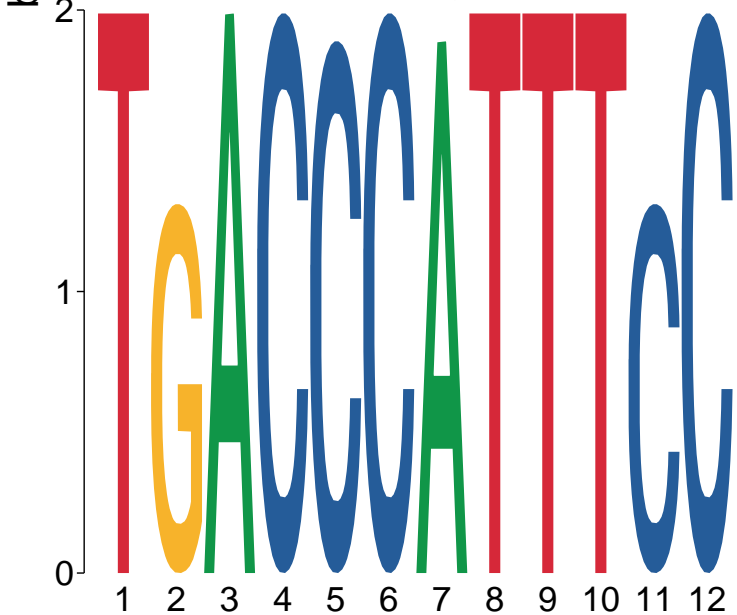
ATWGACCACMA,BestGuess:PB0196.1_Zb1



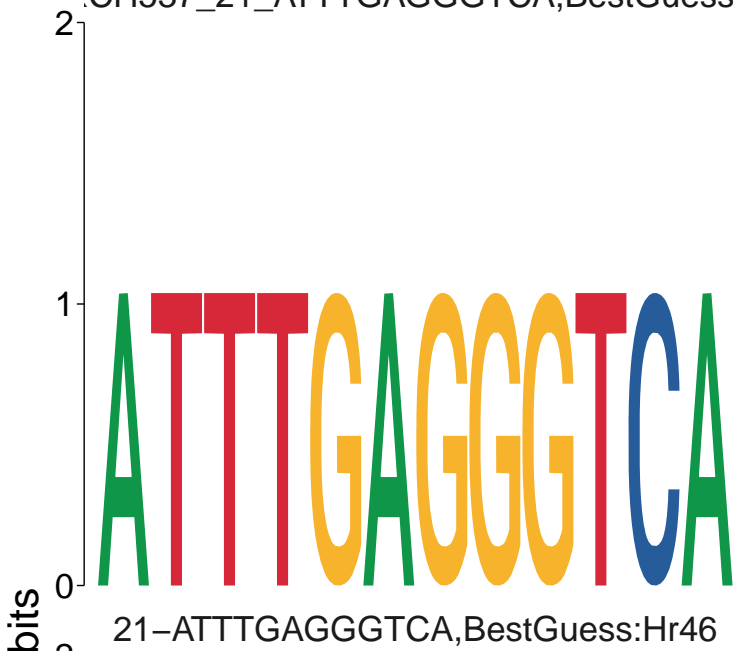
CH536_17_TGACCCATTTC, BestGuess_



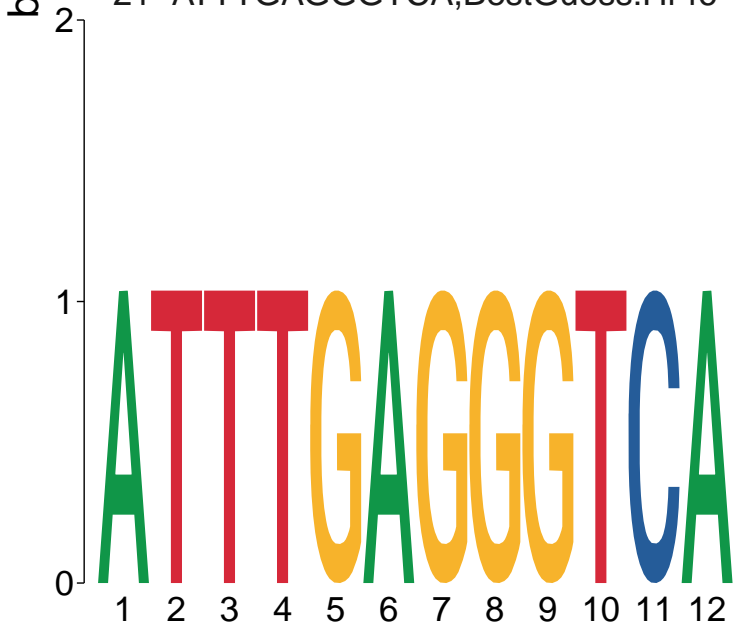
17-TGACCCATTTC, BestGuess:Hr46



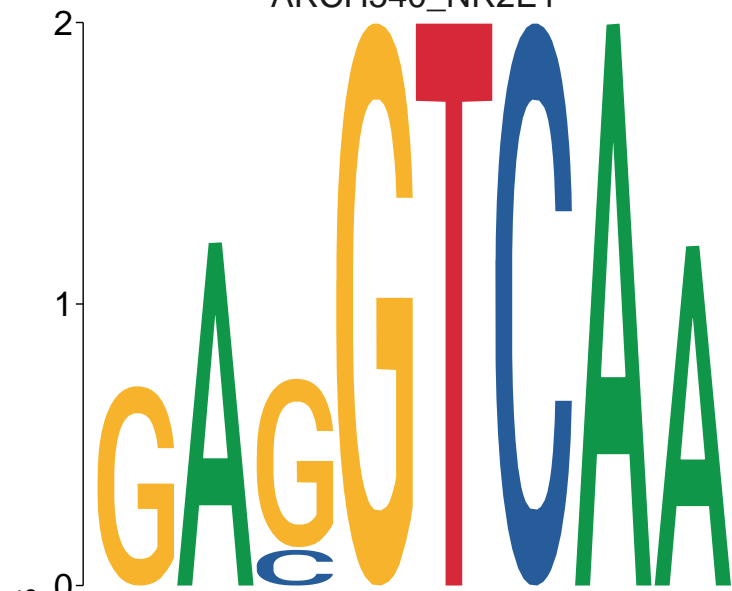
CH537_21_ATTTGAGGGTCA,BestGuess_



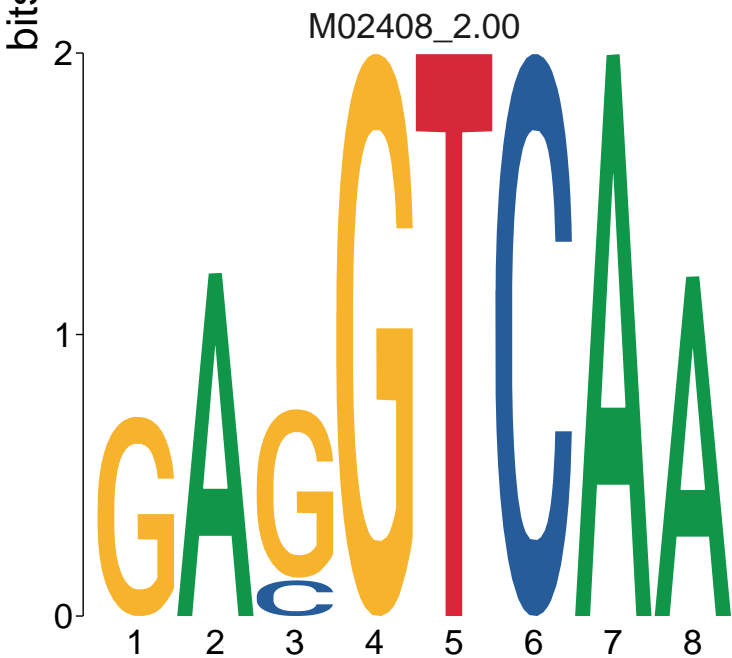
21-ATTTGAGGGTCA,BestGuess:Hr46



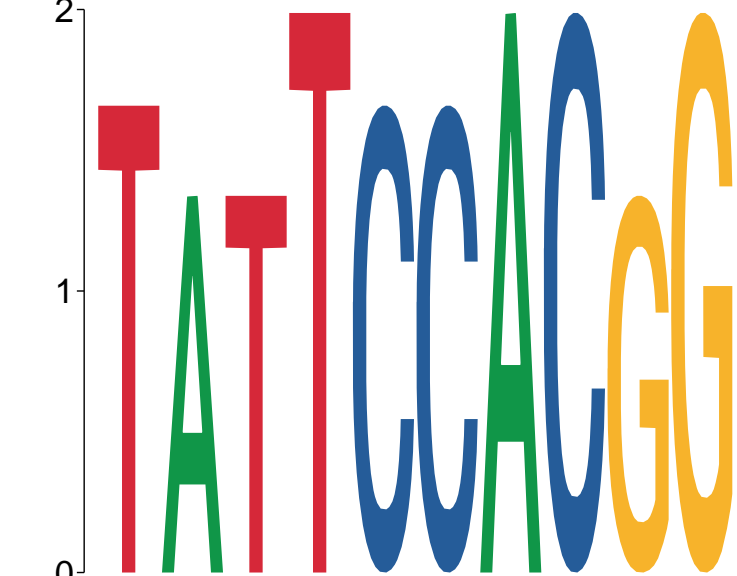
ARCH540_NR2E1



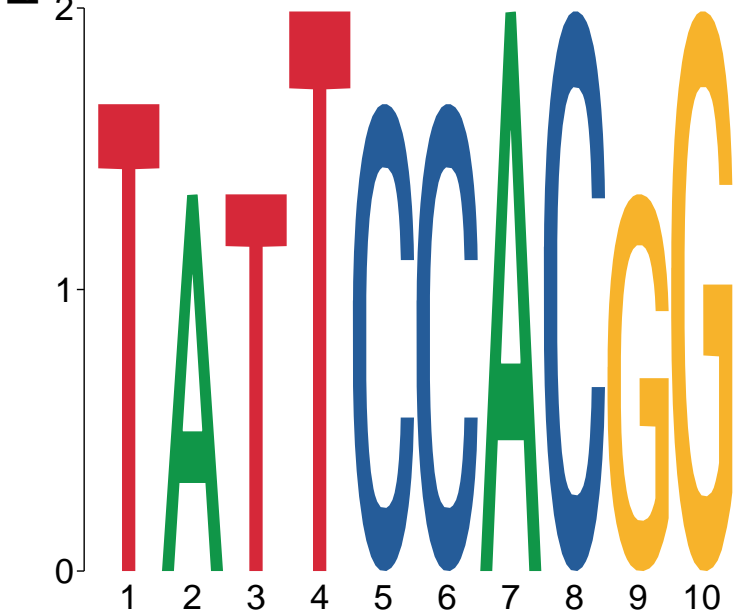
M02408_2.00



RCH541_5_TATTCCACGG,BestGuess_nit_



5-TATTCCACGG,BestGuess:nit-4



ARCH542_TEAD3_TEAD2_TEAD4



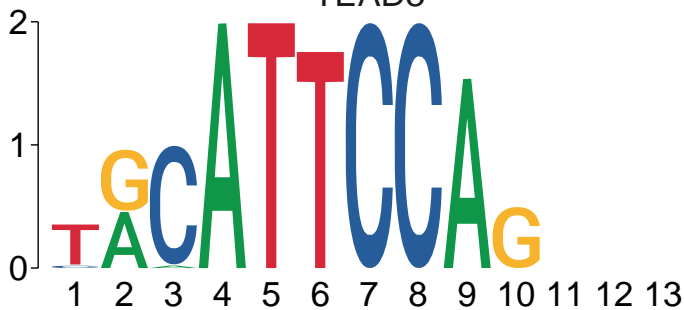
TEAD4



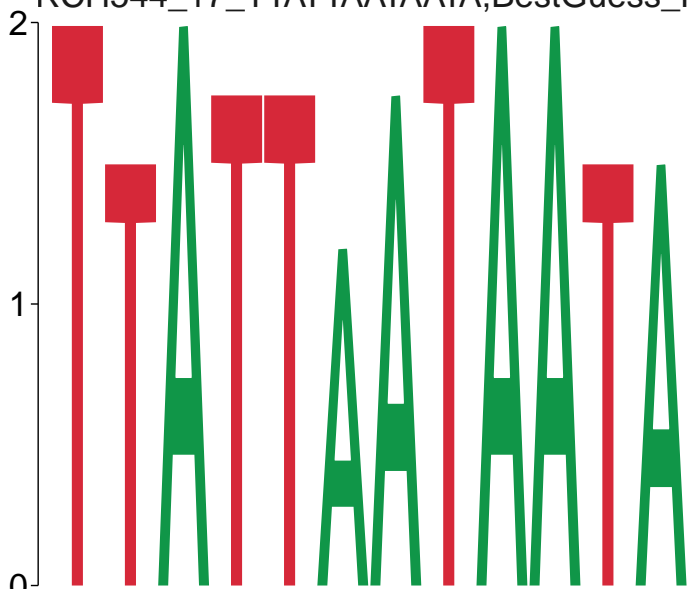
TEAD2



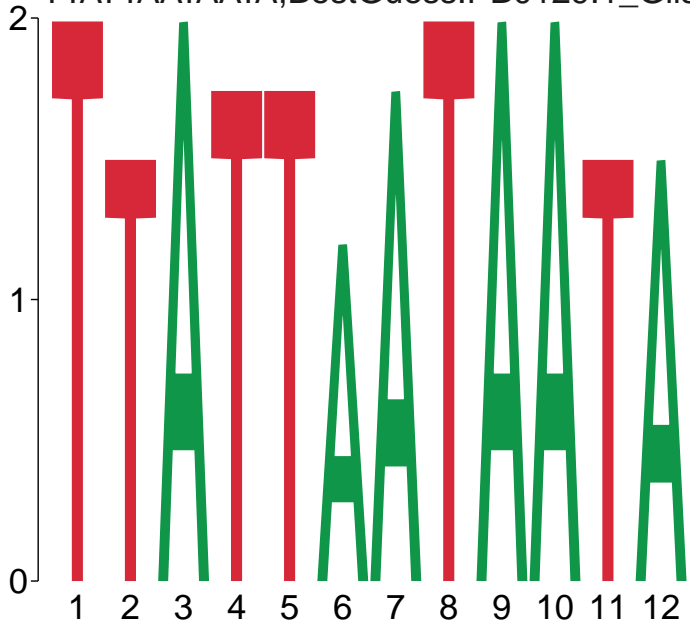
TEAD3



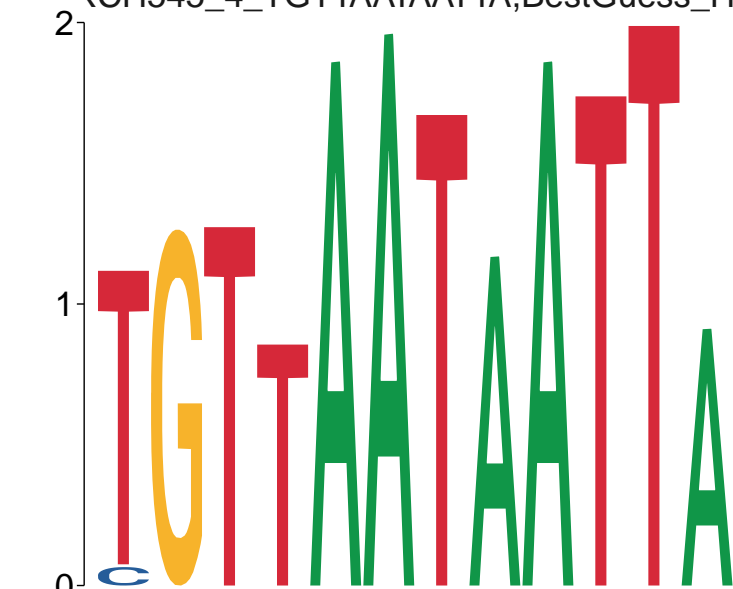
RCH544_17_TTATTAATAATA, BestGuess_P



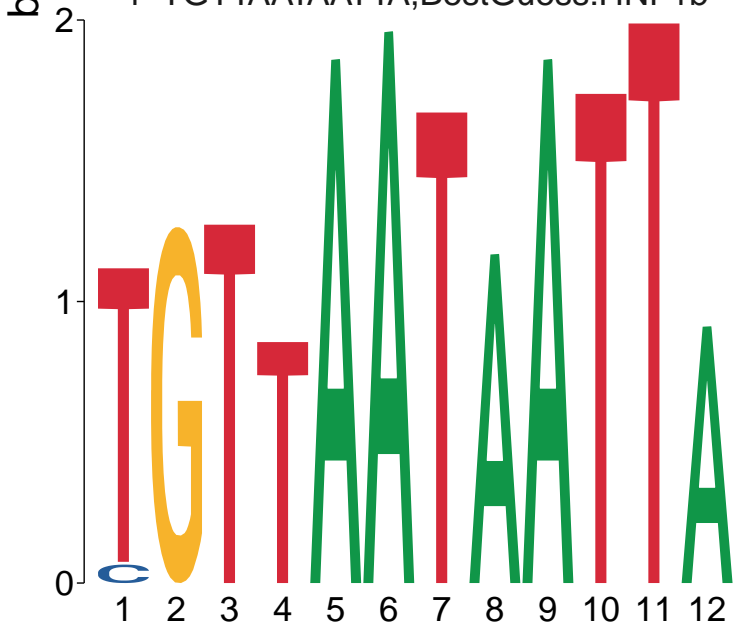
TTATTAATAATA, BestGuess:PB0129.1_Glis



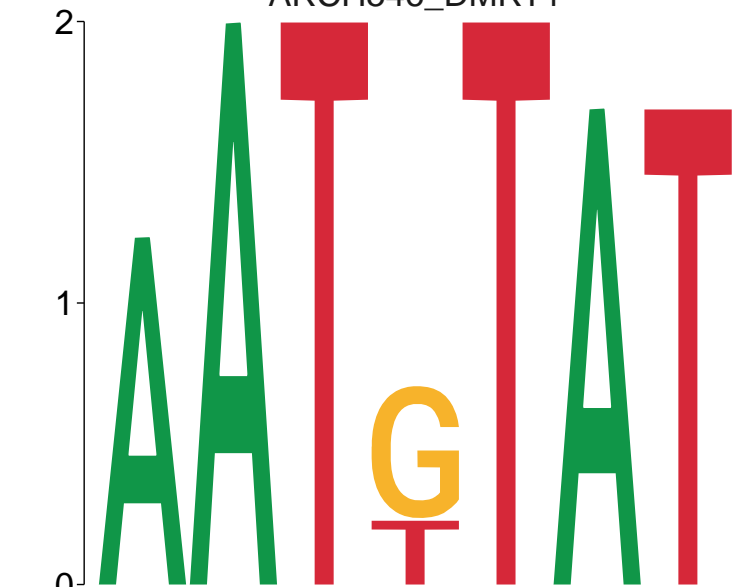
RCH545_4_TGTTAATAATTA,BestGuess_HNF1b



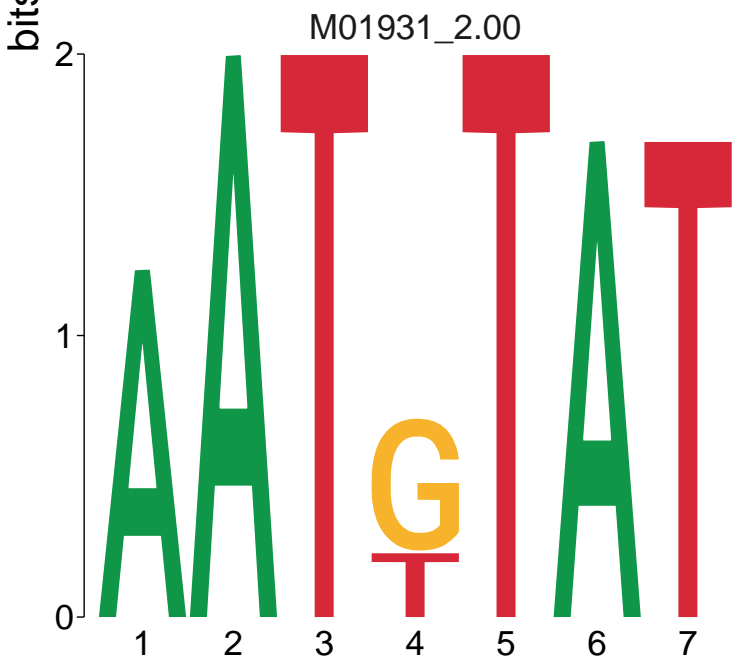
4-TGTTAATAATTA,BestGuess:HNF1b



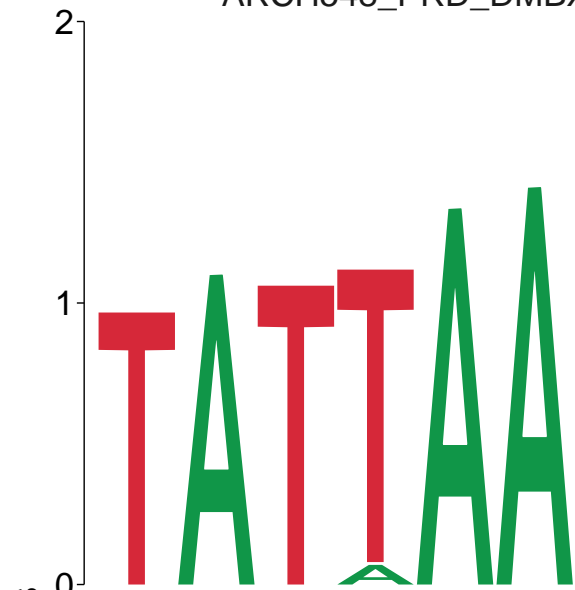
ARCH546_DMRT1



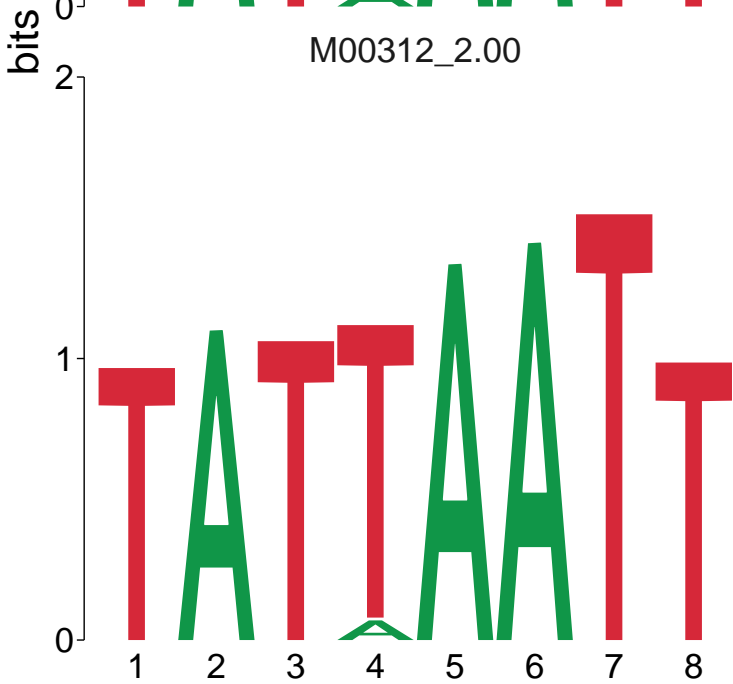
M01931_2.00



ARCH548_PRD_DMBX1



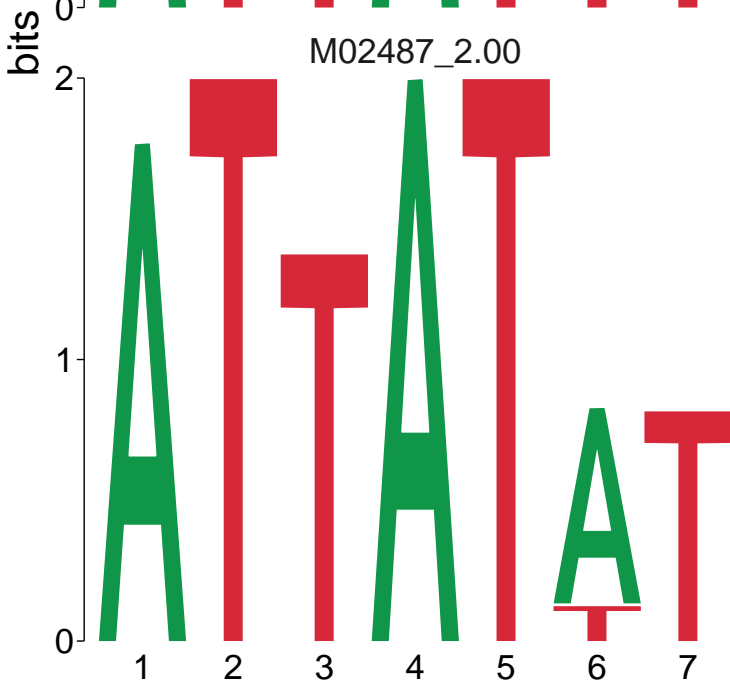
M00312_2.00



ARCH549_islike_HMG20A



M02487_2.00



ARCH550_Nvec_vc1.1_XM_032368623.2

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1

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M00889_2.00

2

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1

2

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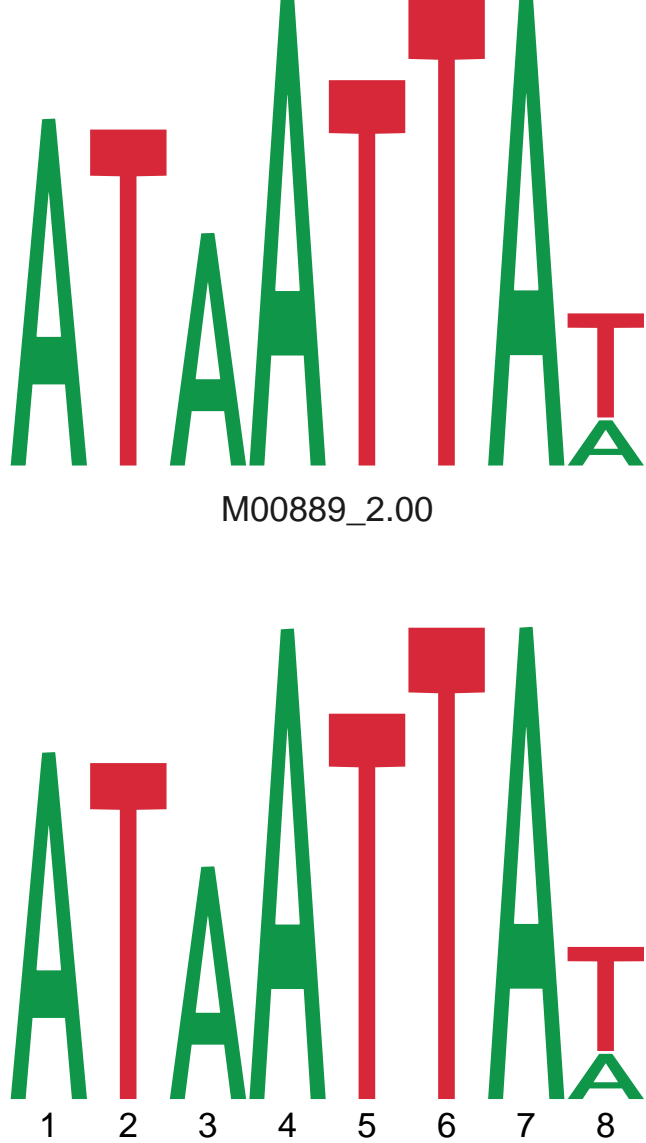
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6

7

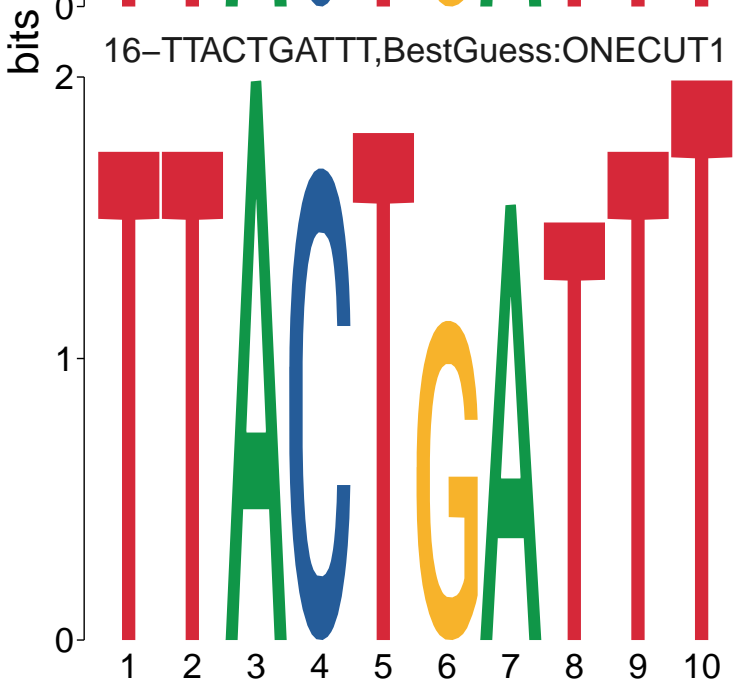
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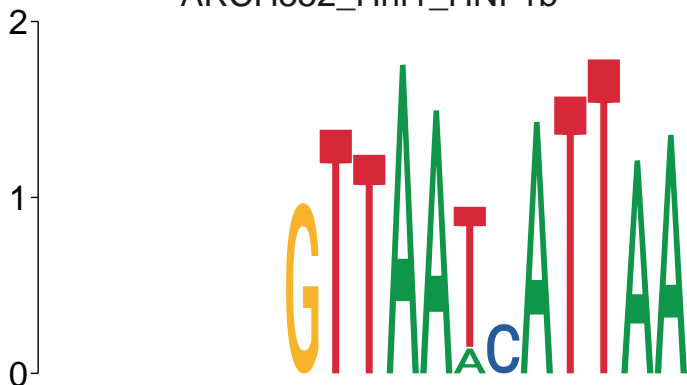
CH551_16_TTACTGATTT,BestGuess_ONI



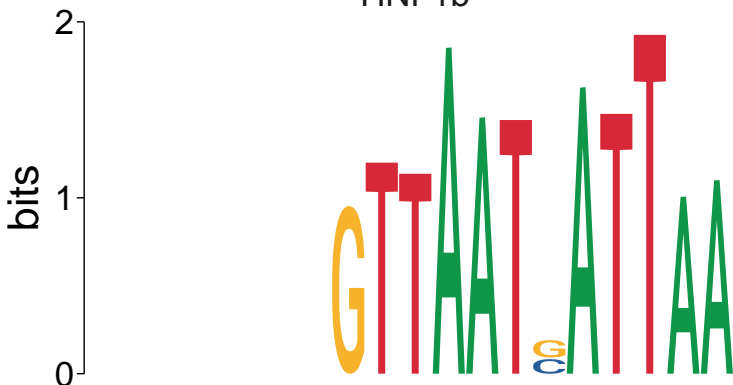
16-TTACTGATTT,BestGuess:ONECUT1



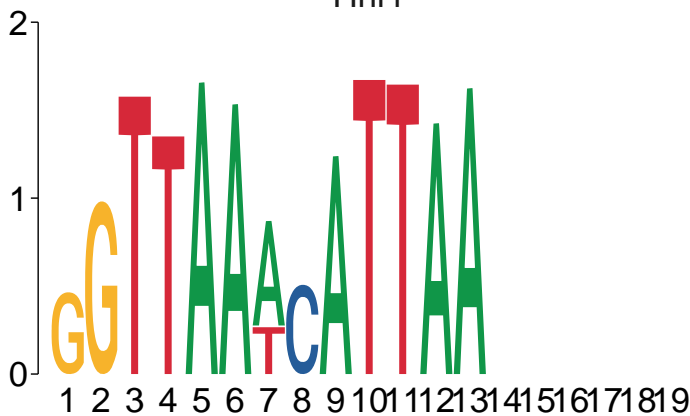
ARCH552_Hnf1_HNF1b



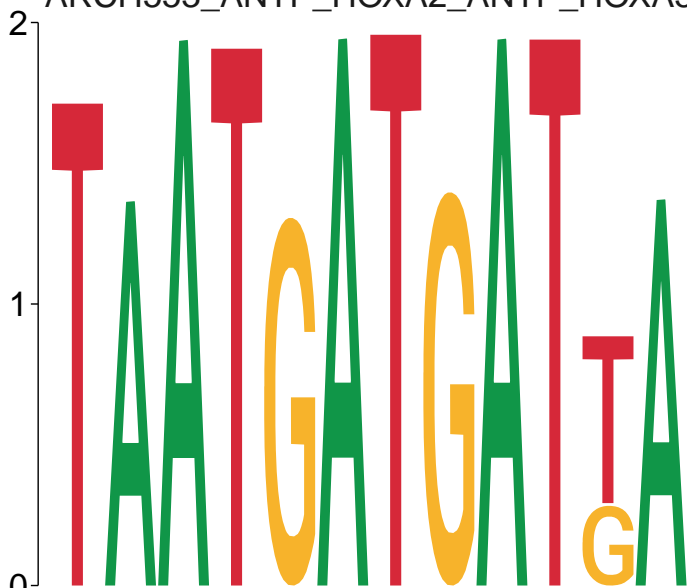
HNF1b



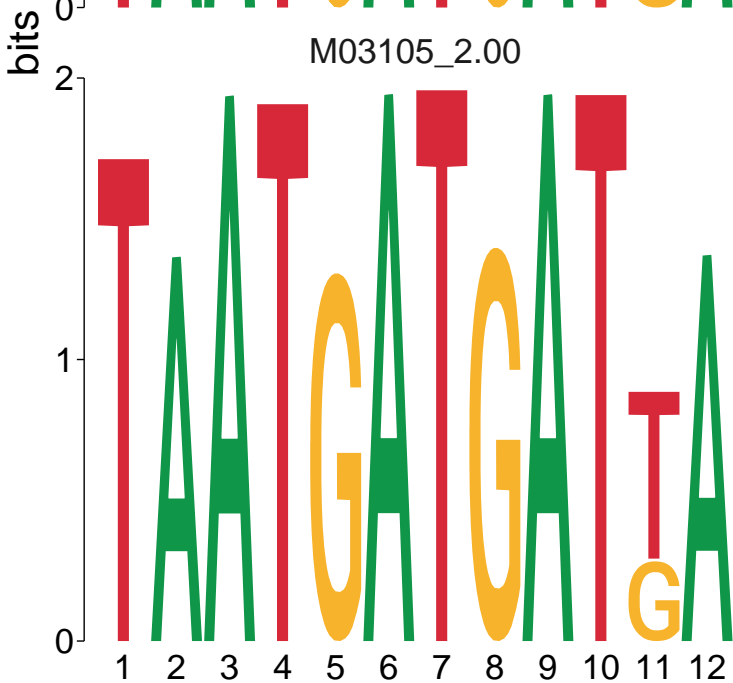
Hnf1



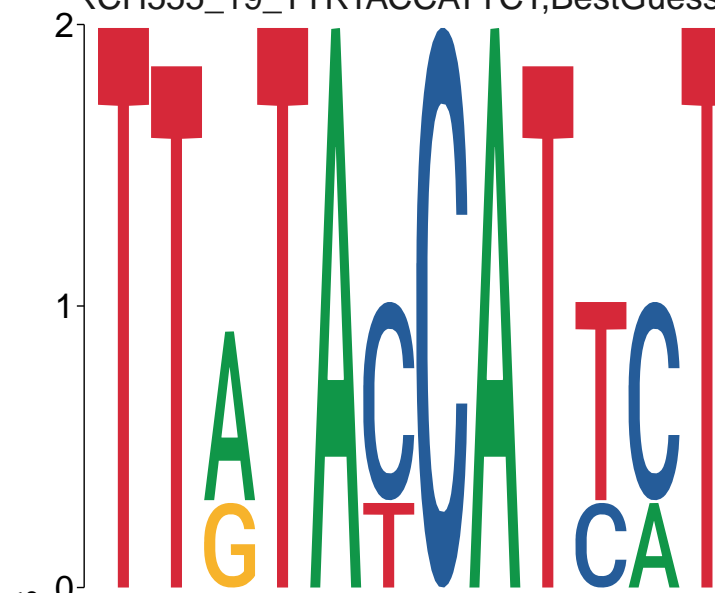
ARCH553_ANTP_HOXA2_ANTP_HOXA3



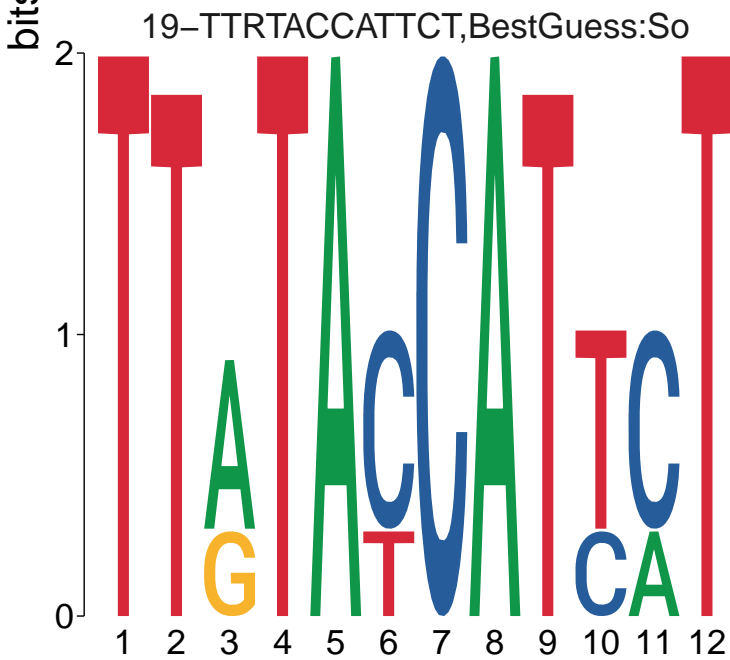
M03105_2.00



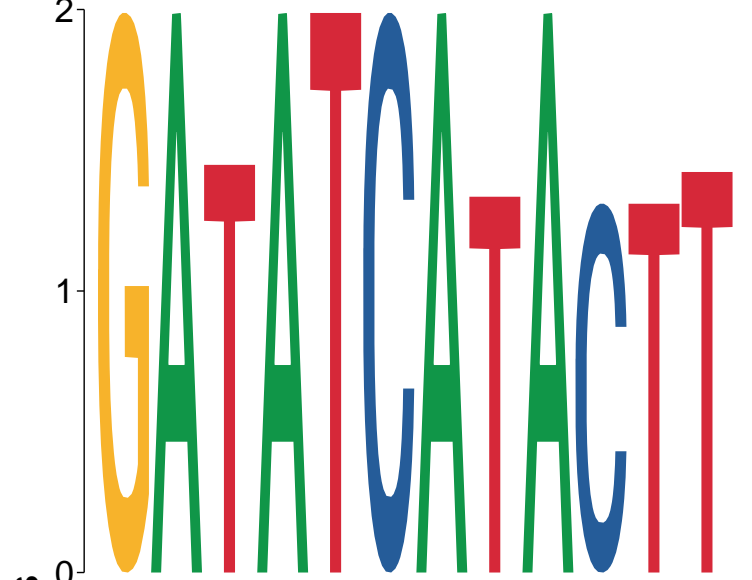
RCH555_19_TTRTACCATTCT,BestGuess_5



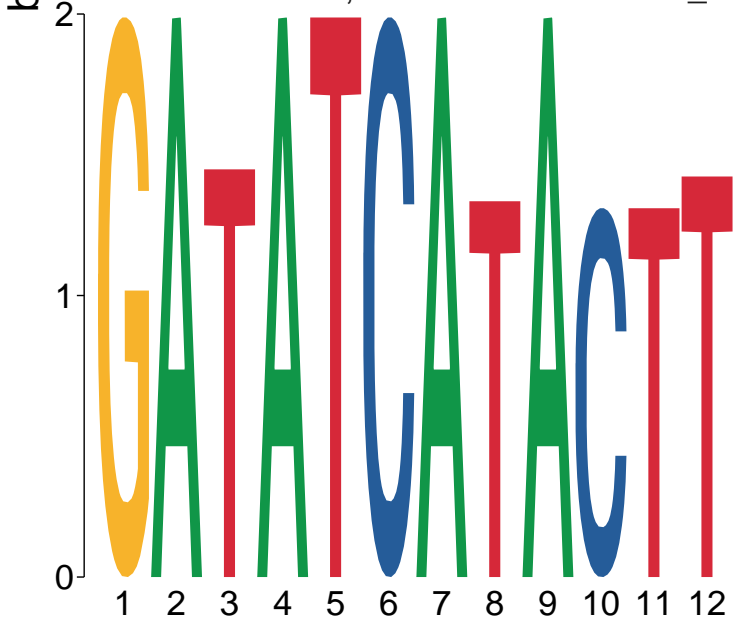
19-TTRTACCATTCT,BestGuess:So



RCH556_4_GATATCATACTT,BestGuess_Pf



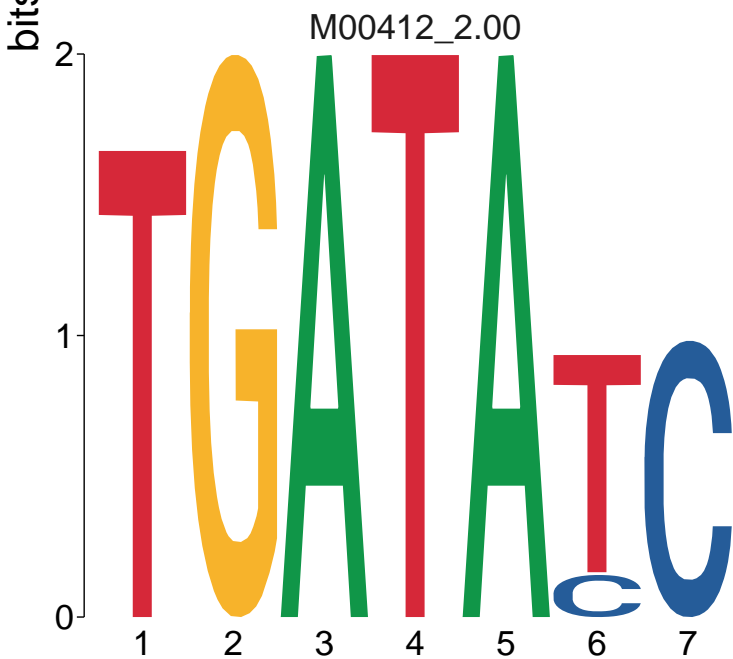
-GATATCATACTT,BestGuess:PH0161.1_Si



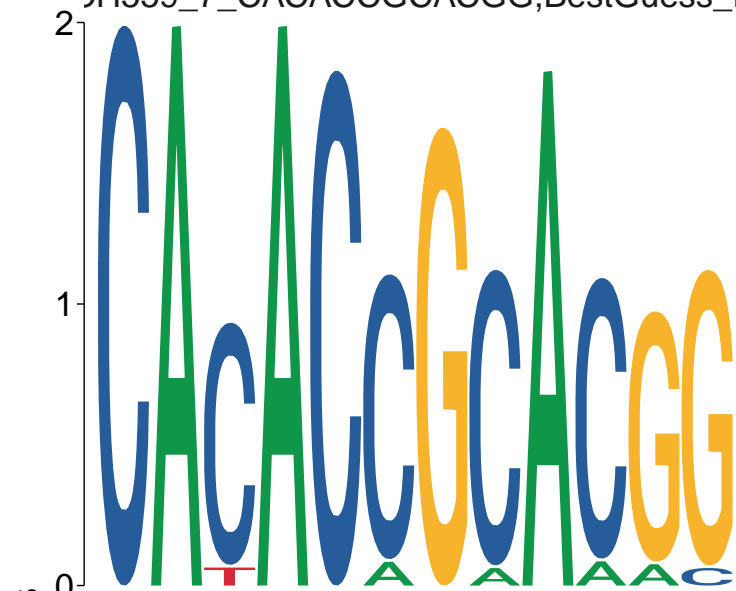
ARCH558_SINE_SIX3



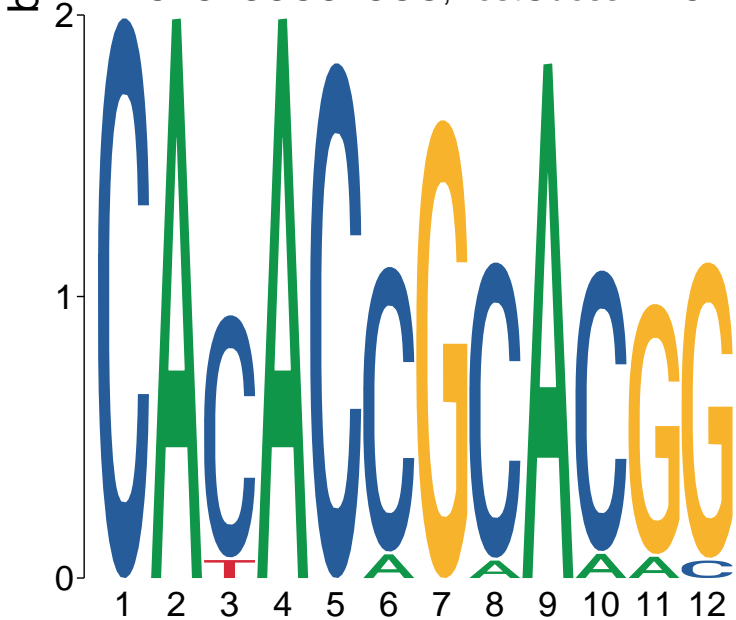
M00412_2.00



CH559_7_CACACCGCACGG,BestGuess_F



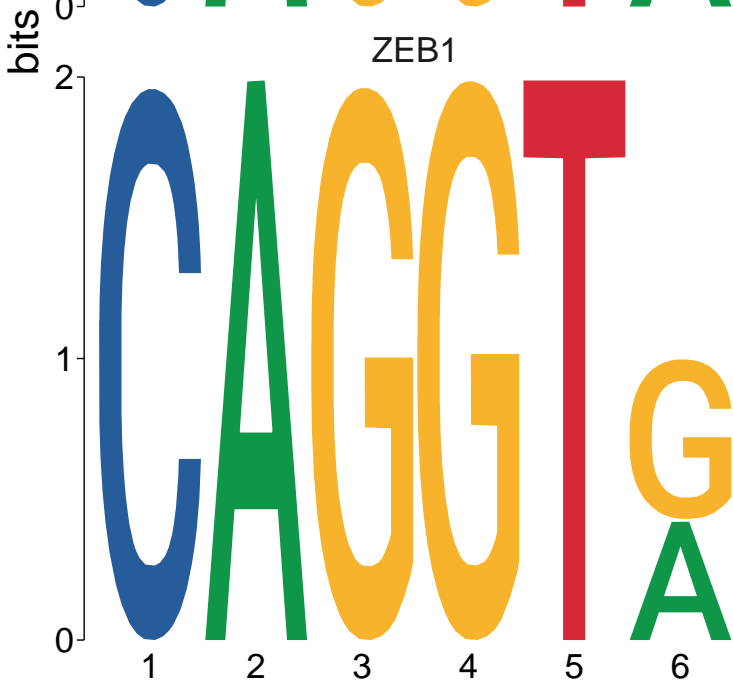
7-CACACCGCACGG,BestGuess:PHO2



ARCH562_ZEB1



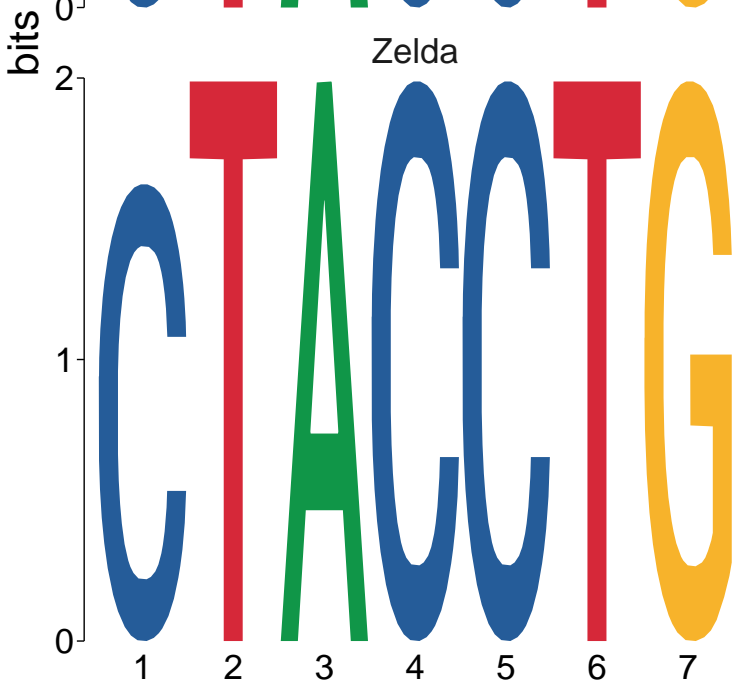
ZEB1



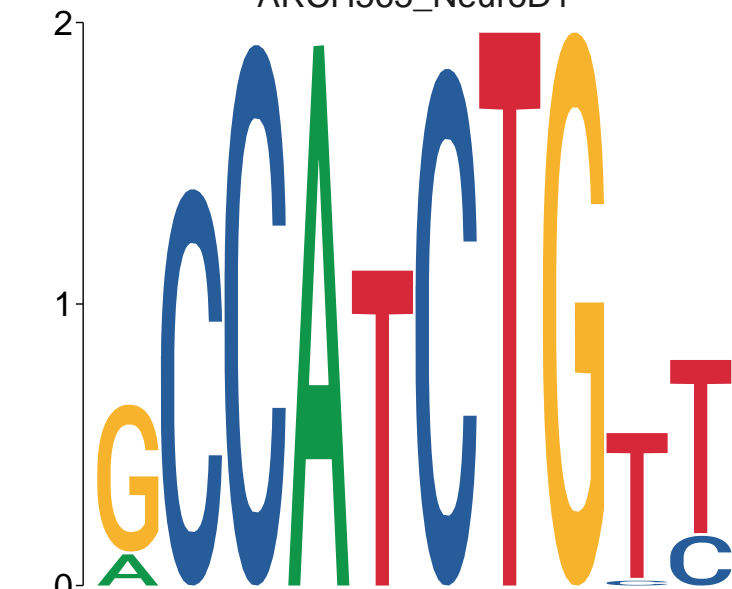
ARCH563_Zelda



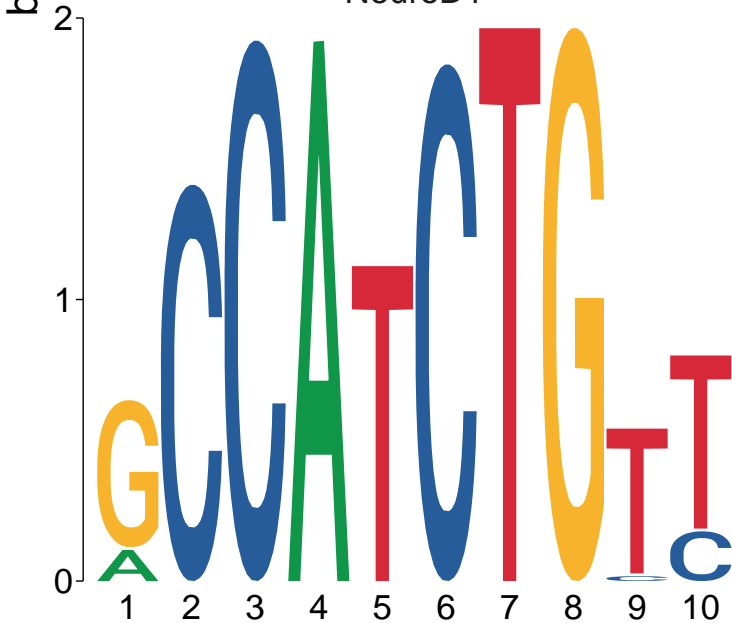
Zelda

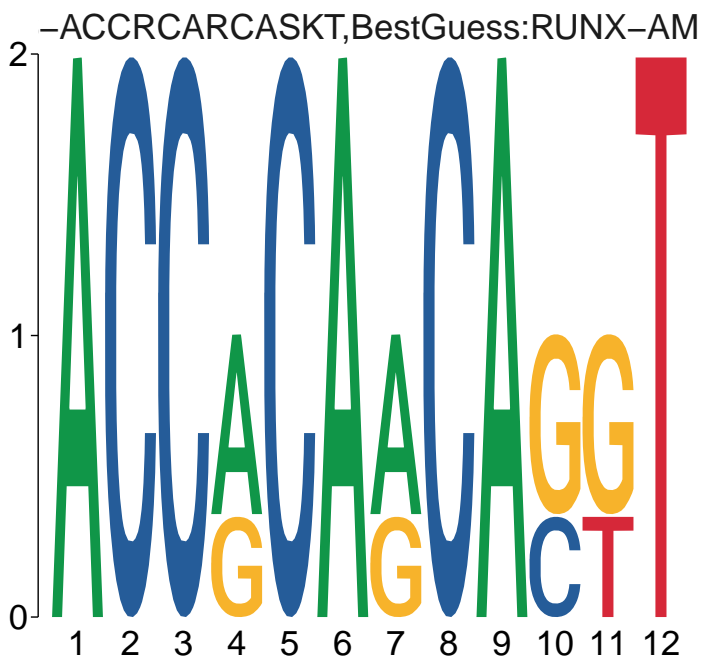
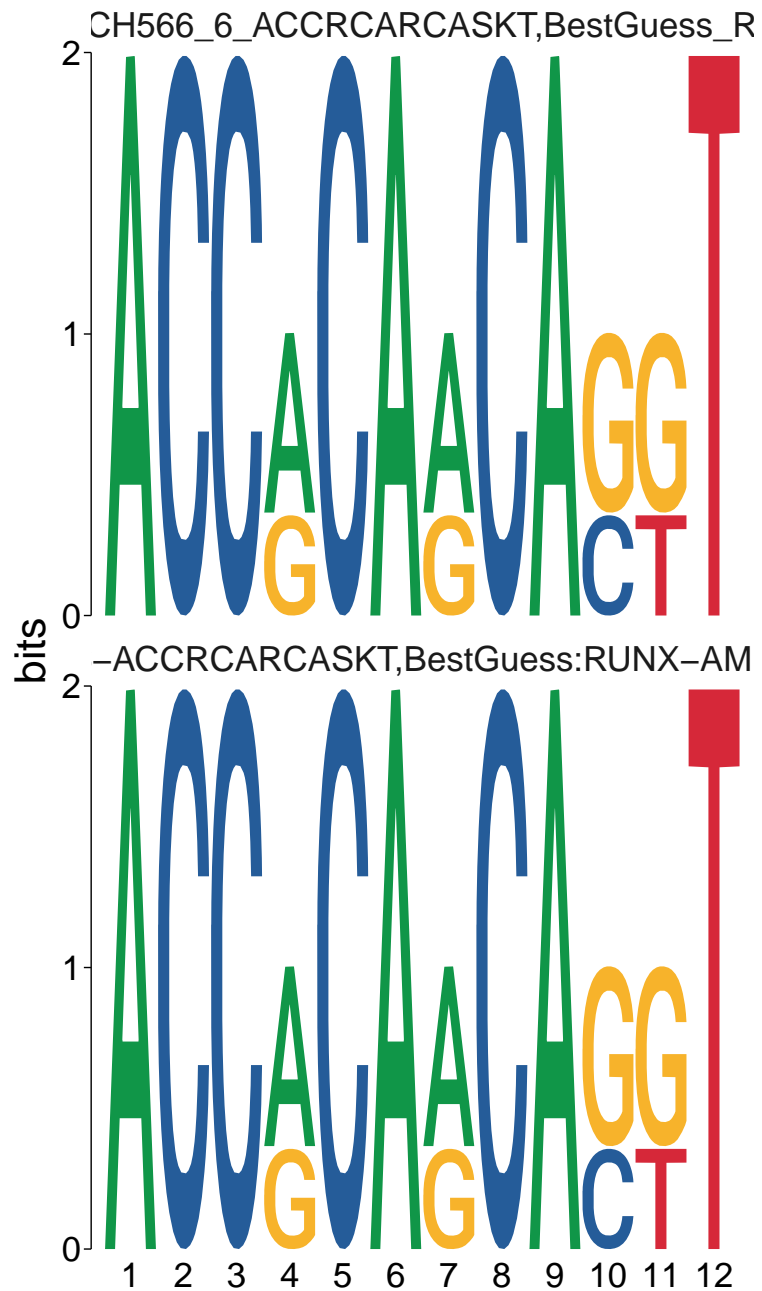


ARCH565_NeuroD1

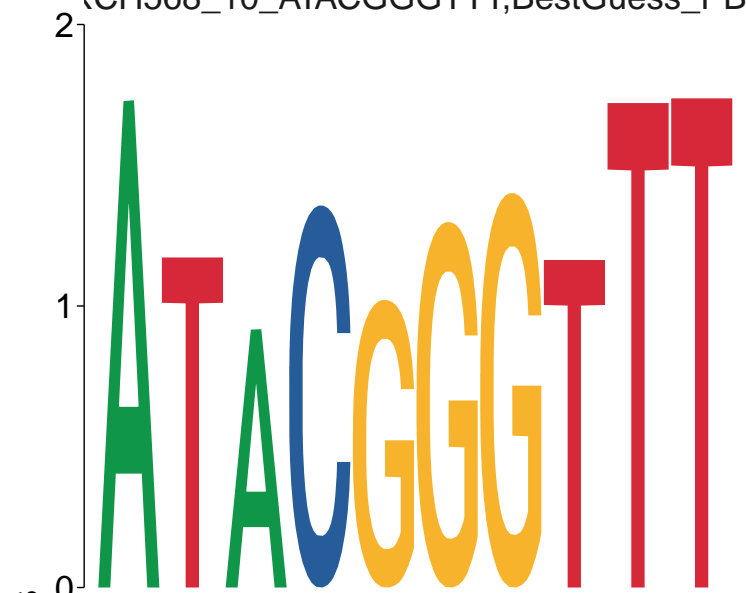


NeuroD1

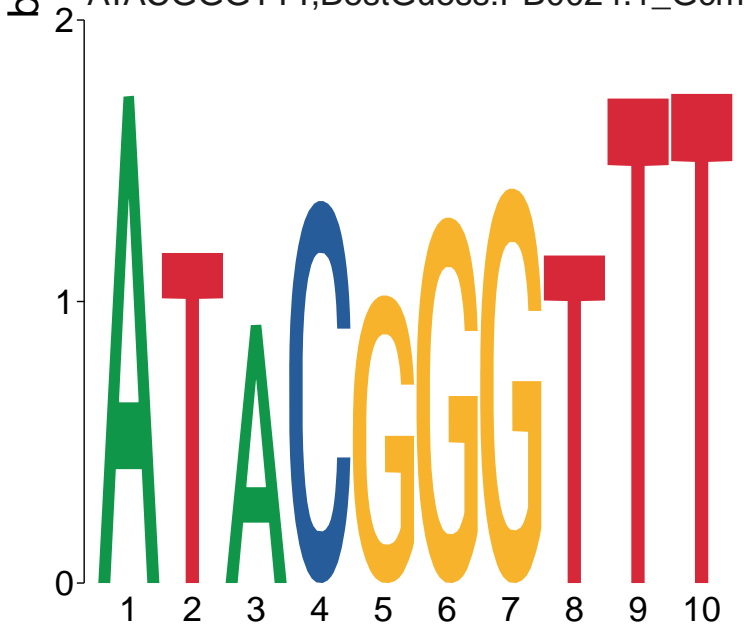




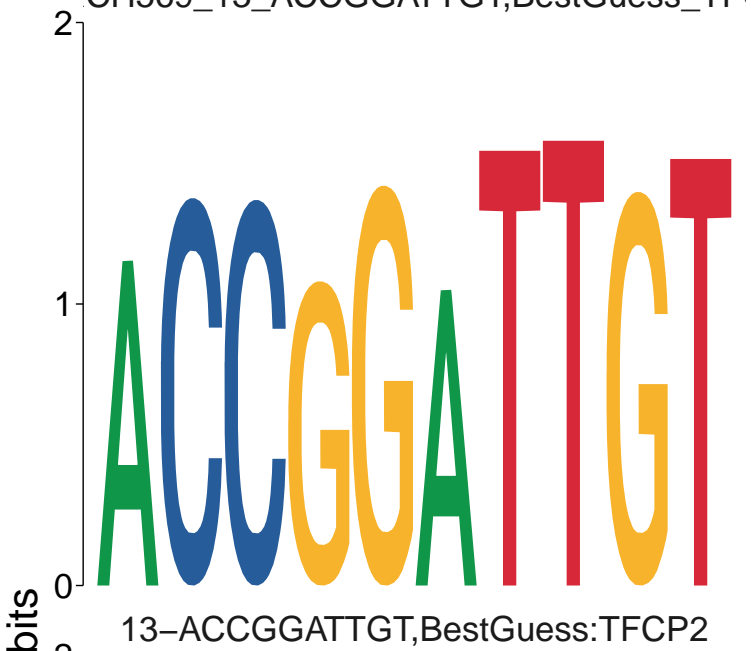
CH568_10_ATACGGGTTT,BestGuess_PB



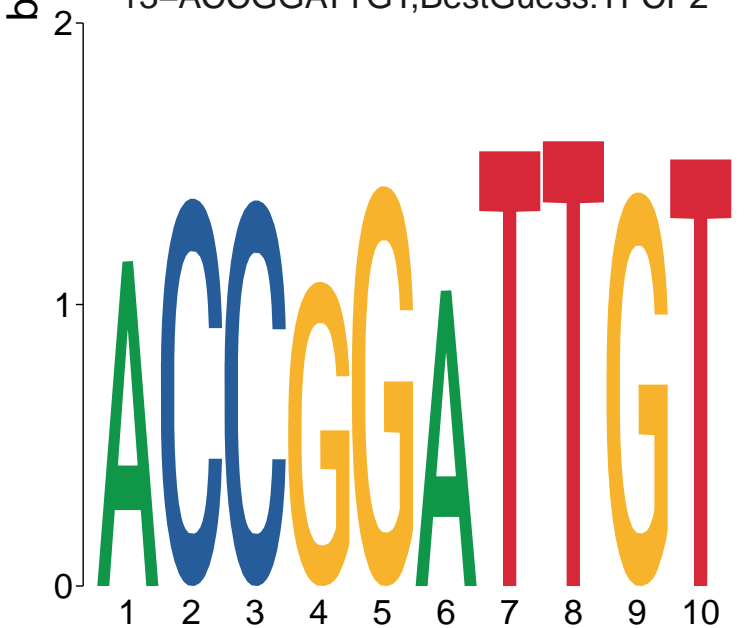
ATACGGGTTT,BestGuess:PB0024.1_Gcm



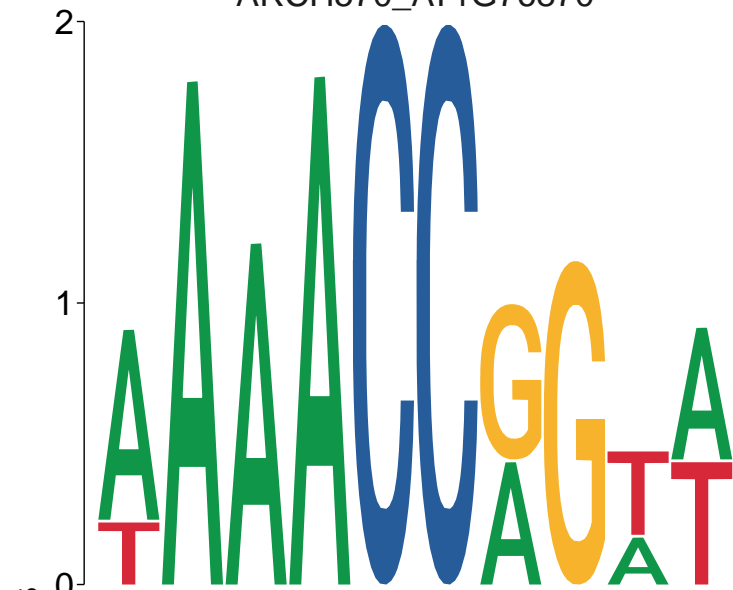
CH569_13_ACCGGATTGT,BestGuess_TF



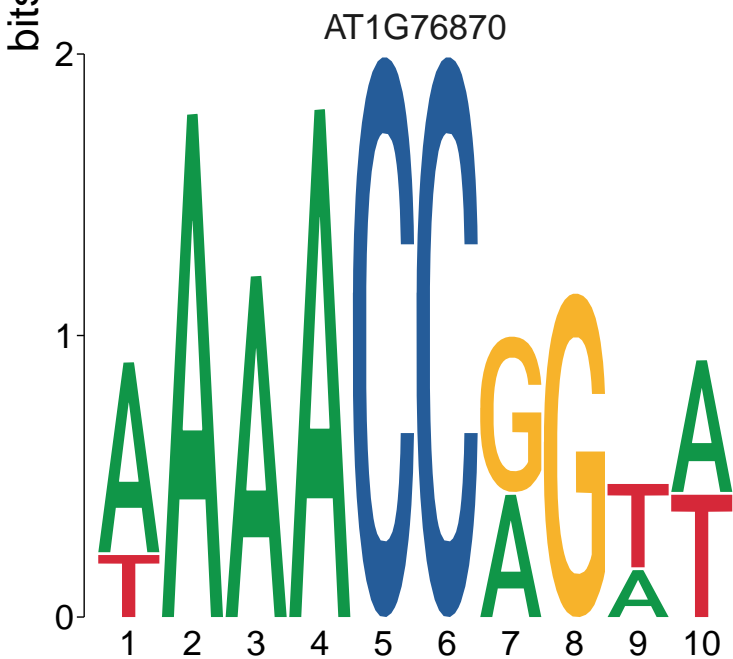
13-ACCGGATTGT,BestGuess:TFCP2



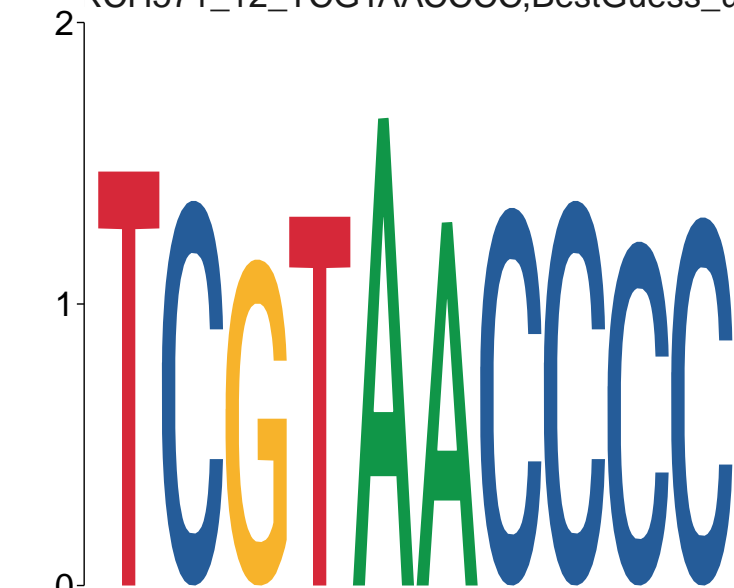
ARCH570_AT1G76870



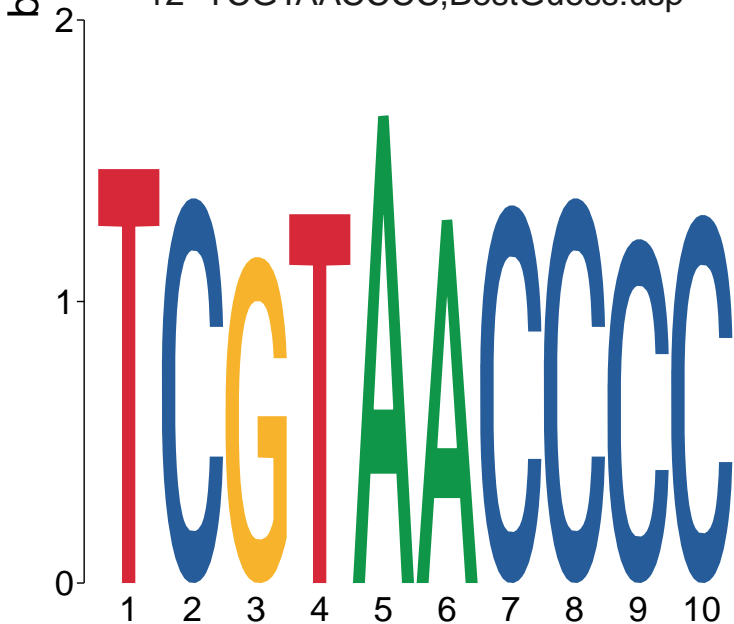
AT1G76870



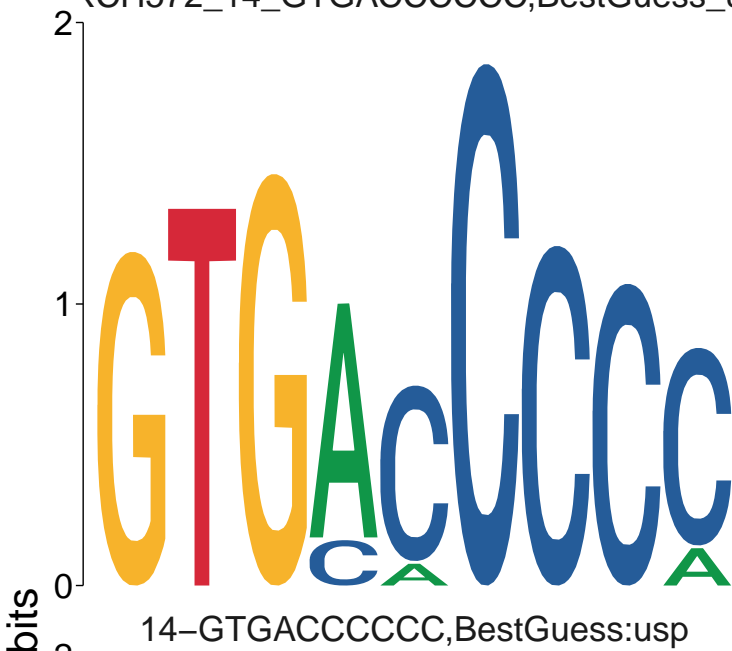
RCH571_12_TCGTAACCCC,BestGuess_us



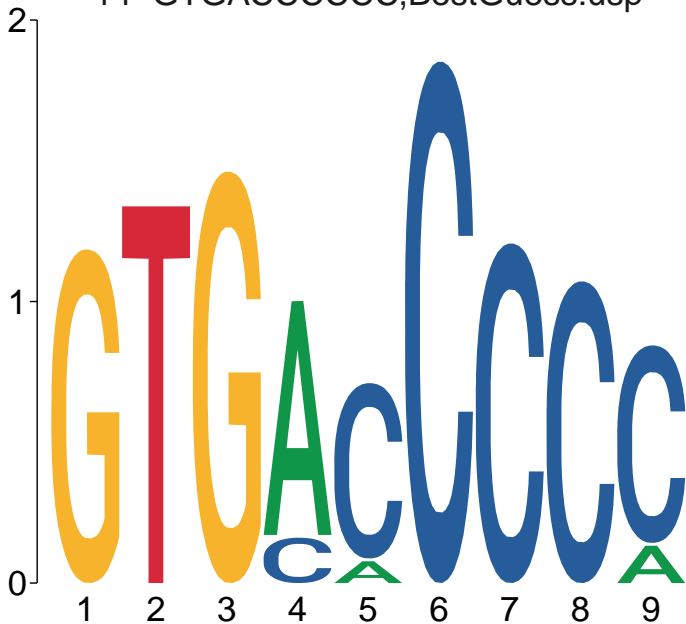
12-TCGTAACCCC,BestGuess:usp



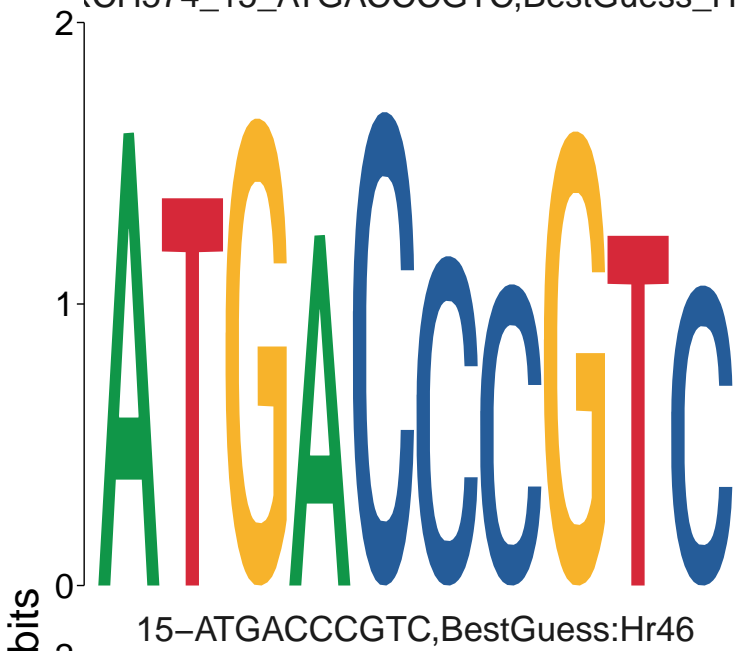
RCH572_14_GTGACCCCCC, BestGuess_u



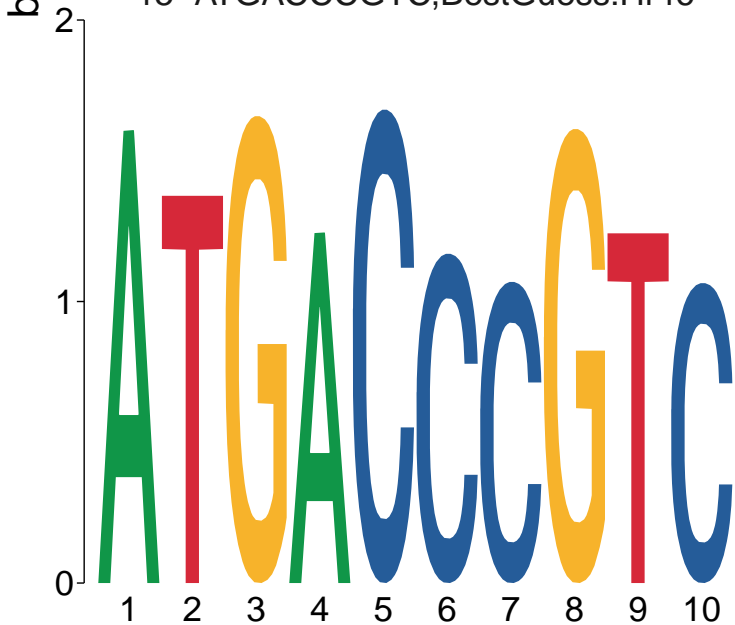
14-GTGACCCCCC, BestGuess:usp



CH574_15_ATGACCCGTC,BestGuess_Hr



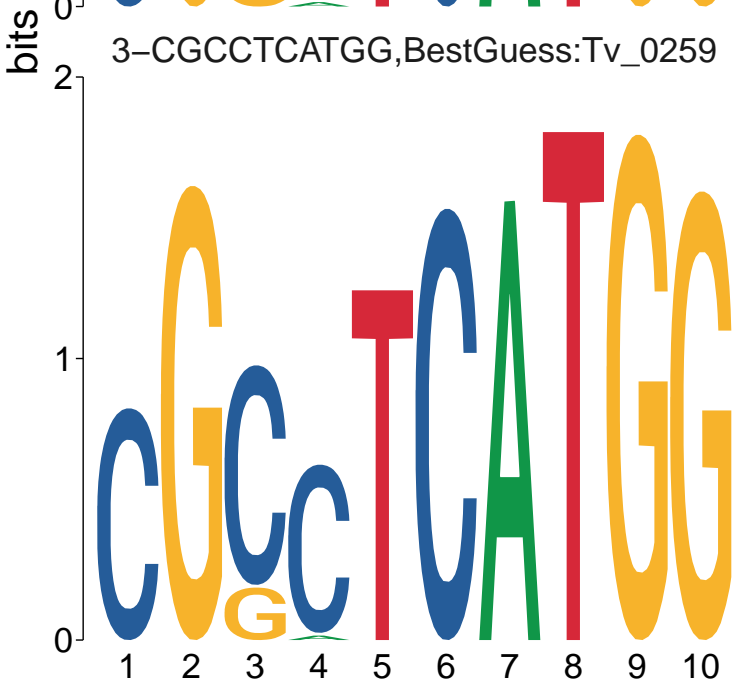
15-ATGACCCGTC,BestGuess:Hr46



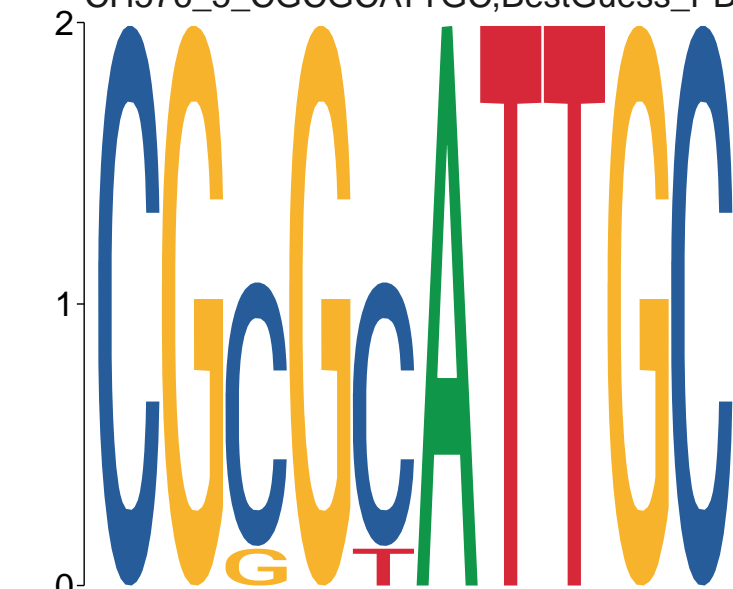
CH575_3_CGCCTCATGG,BestGuess_Tv_



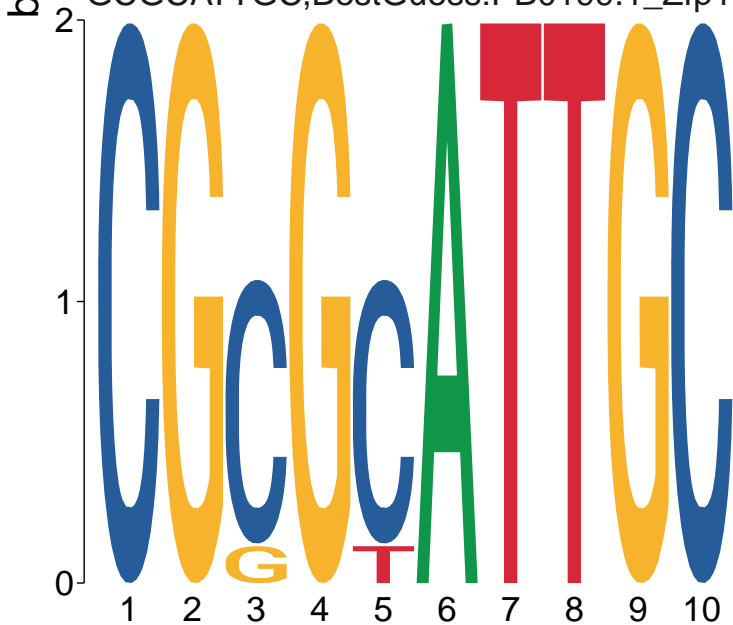
3-CGCCTCATGG,BestGuess:Tv_0259



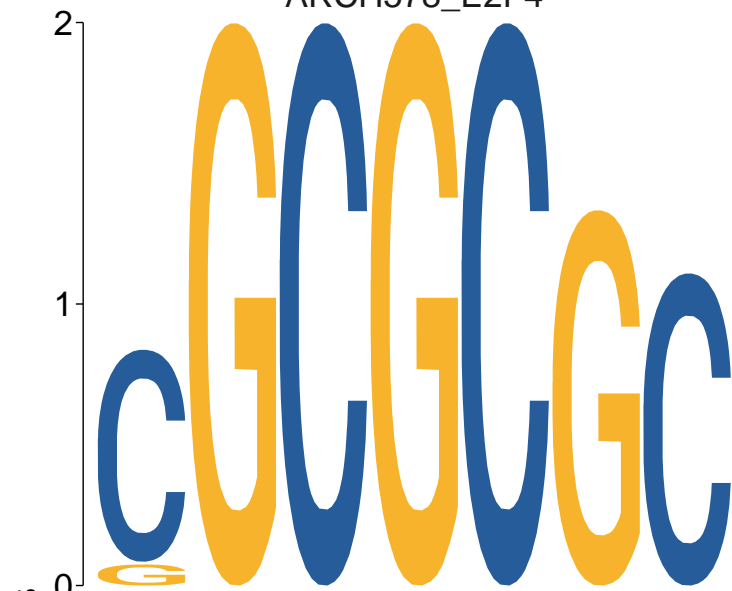
CH576_5_CGCGCATTGC,BestGuess_PBC



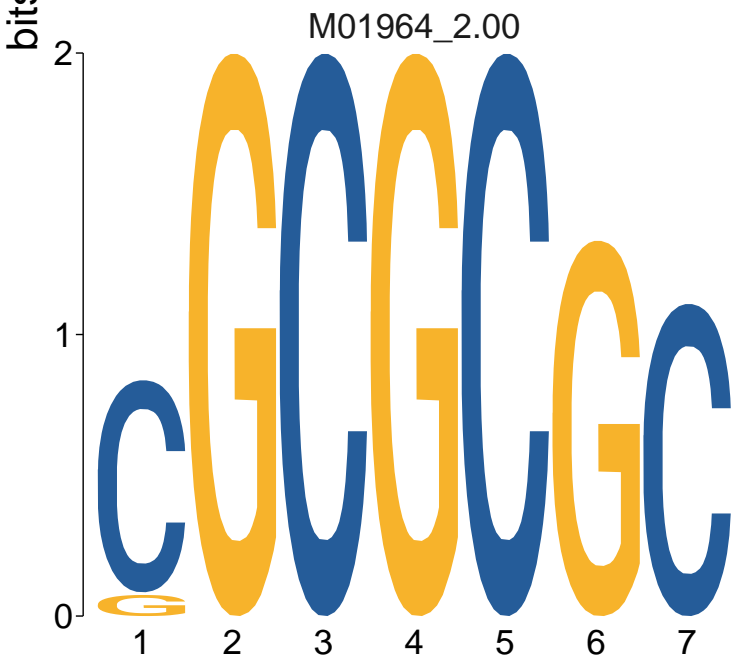
GCGCATTGC,BestGuess:PB0199.1_Zfp16



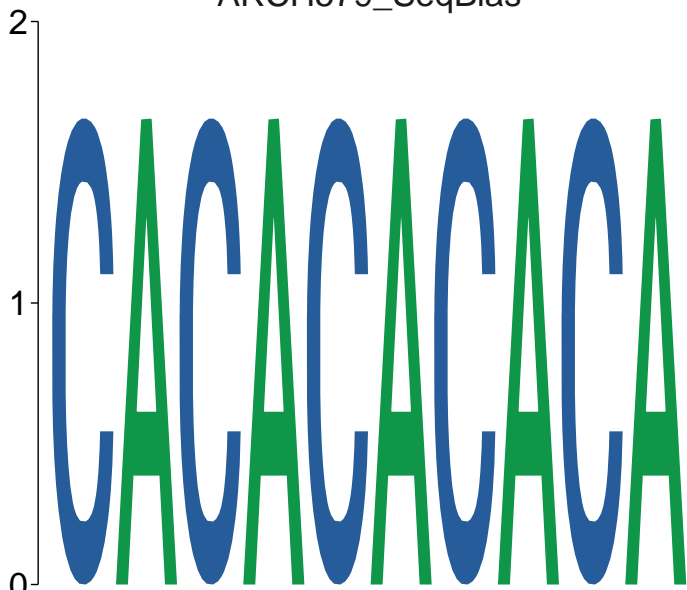
ARCH578_E2F4



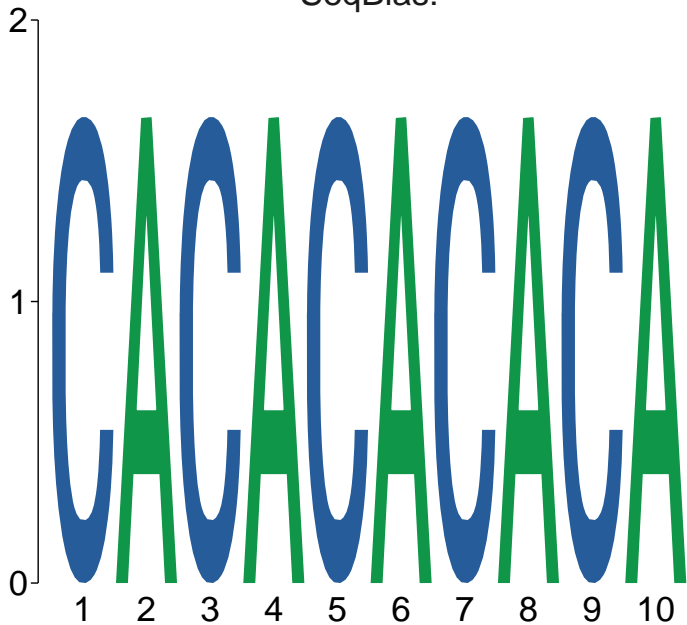
M01964_2.00



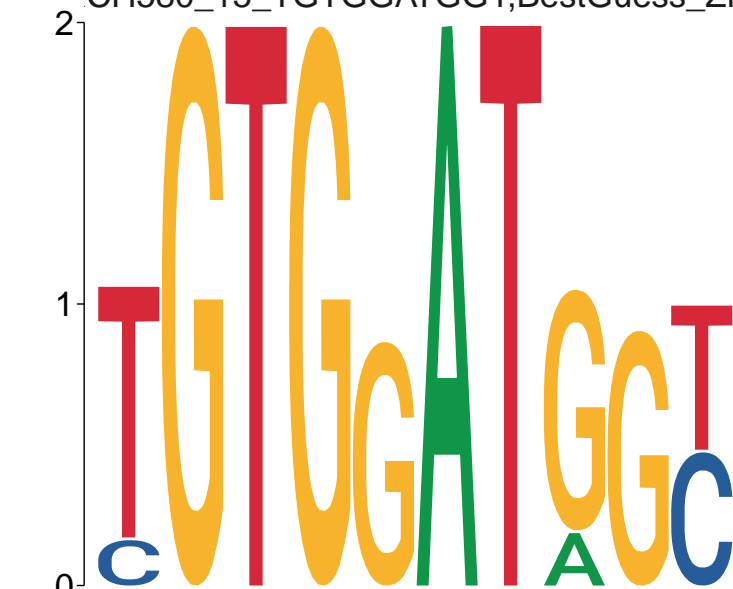
ARCH579_SeqBias



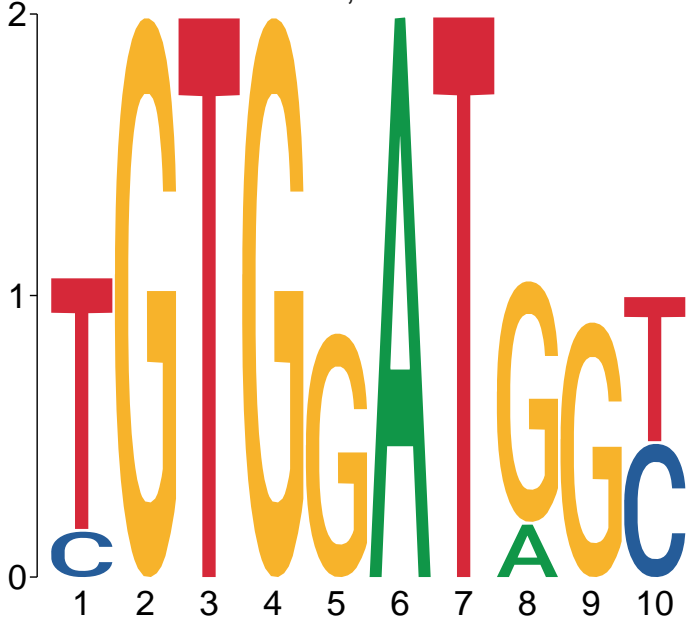
SeqBias:



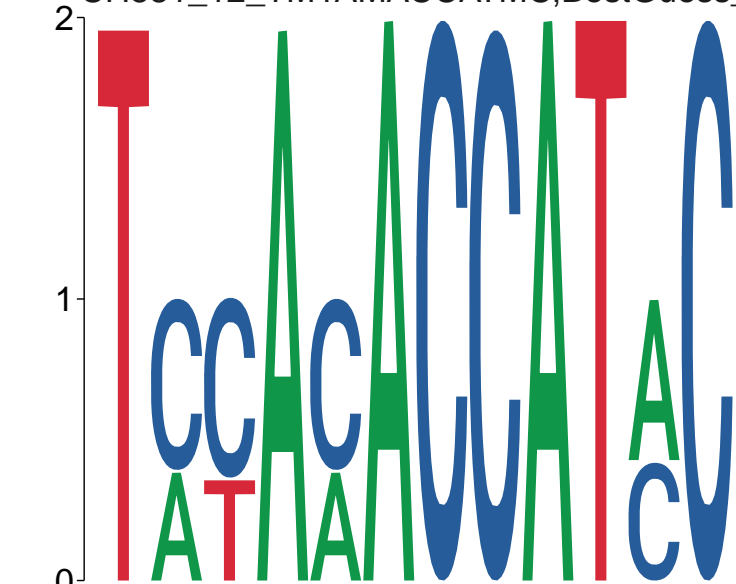
CH580_15_TGTGGATGGY,BestGuess_ZN



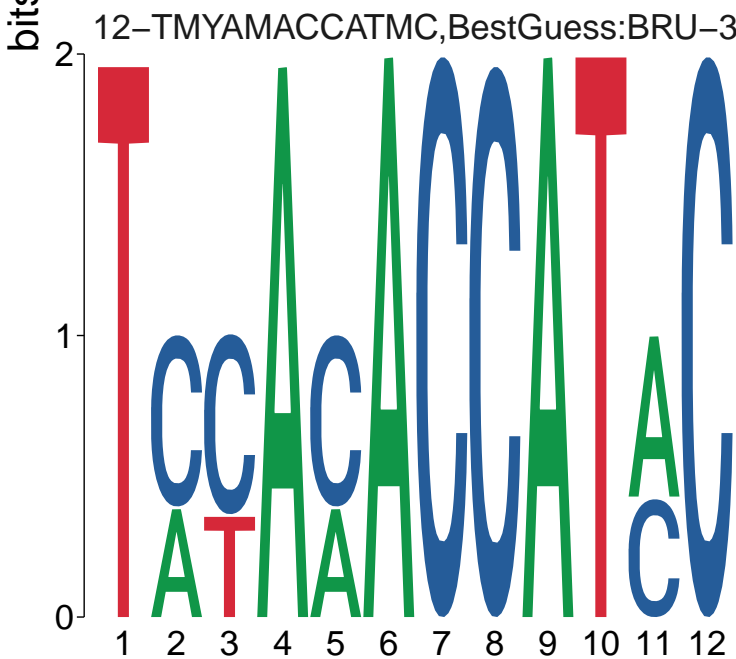
15-TGTGGATGGY,BestGuess:ZNF354C



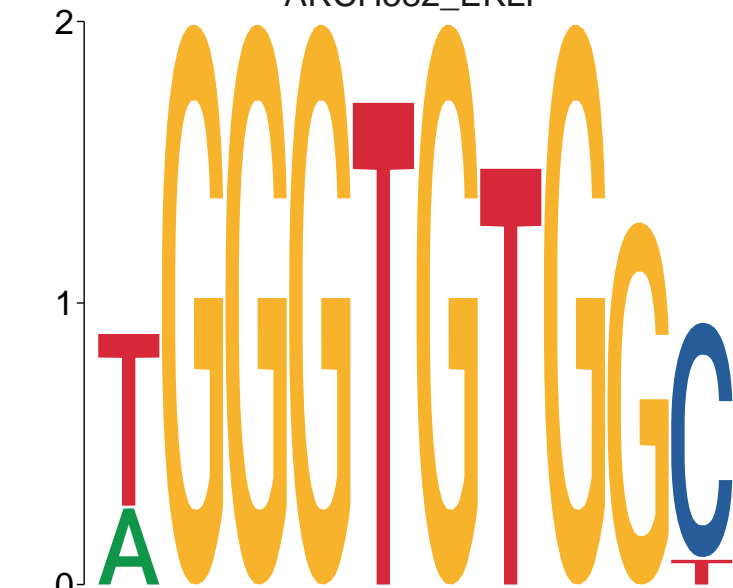
CH581_12_TMYAMACCATMC,BestGuess_



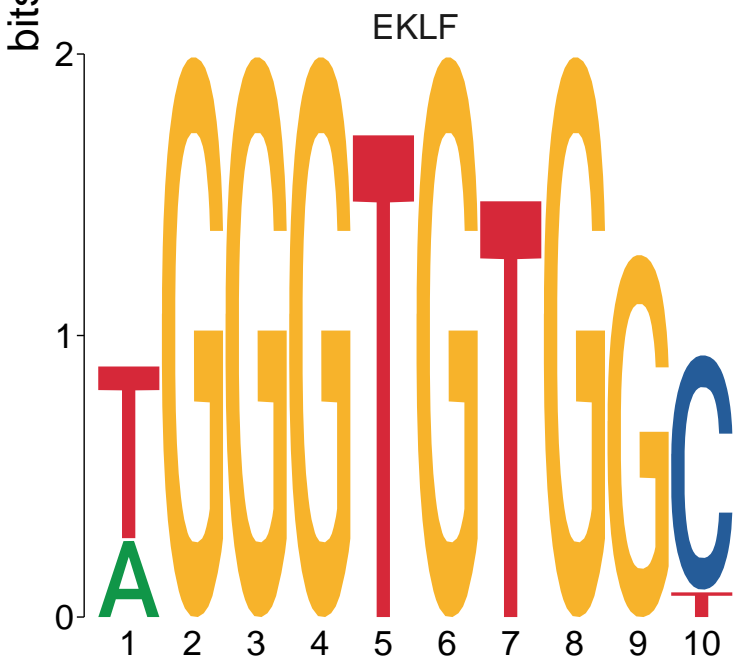
12-TMYAMACCATMC,BestGuess:BRU-3



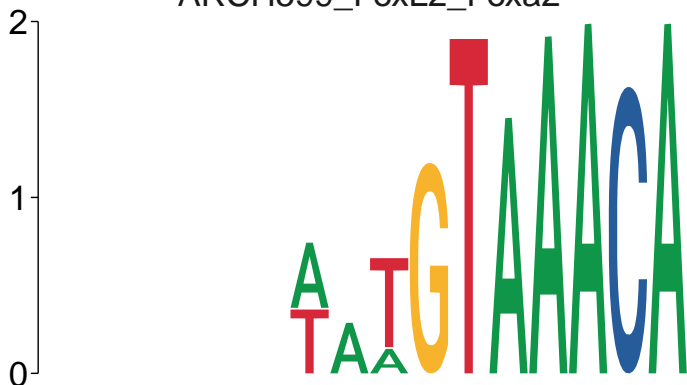
ARCH582_EKLF



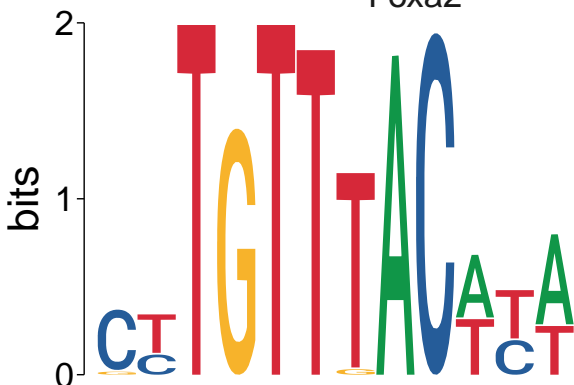
EKLF



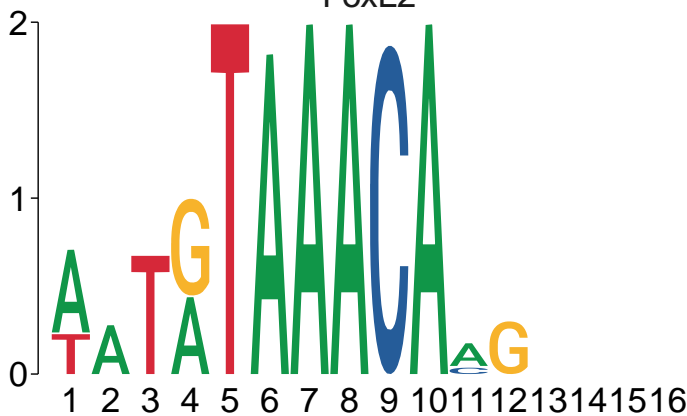
ARCH599_FoxL2_Foxa2



Foxa2



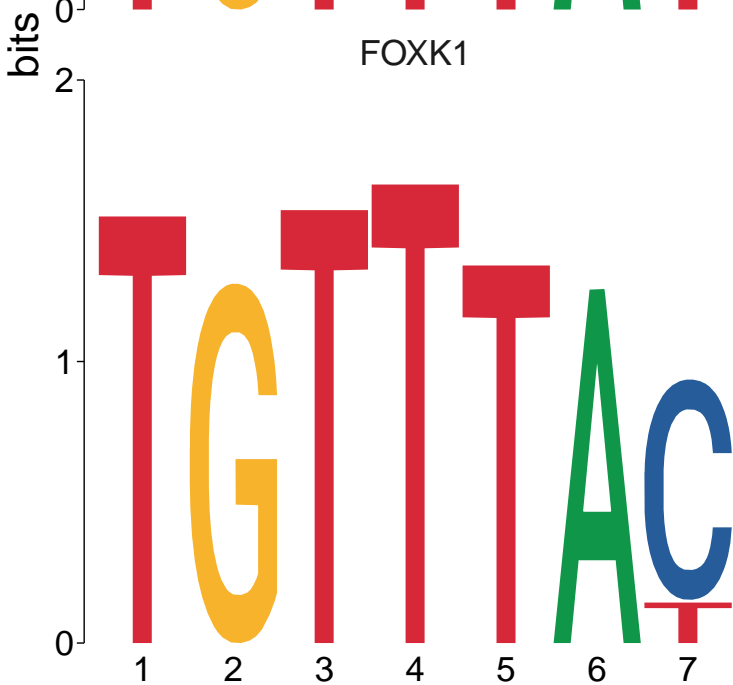
FoxL2



ARCH600_FOXK1



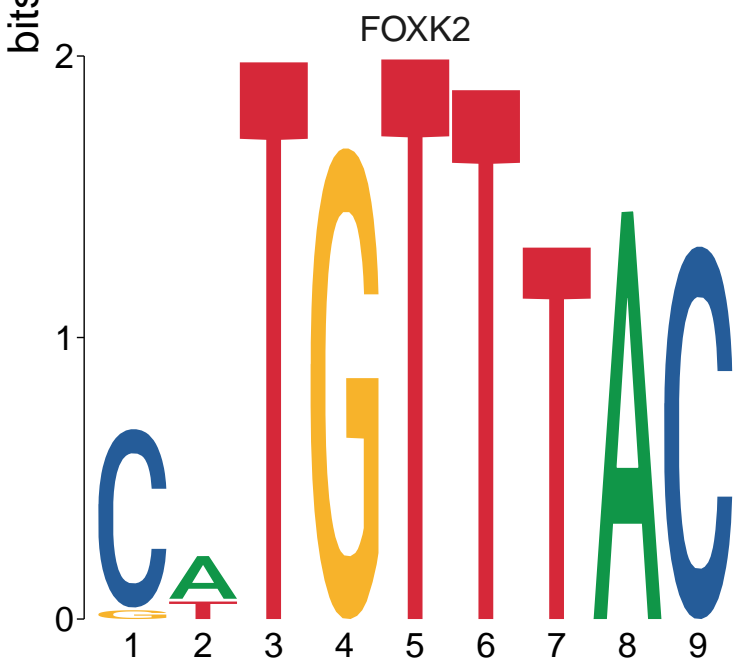
FOXK1



ARCH601_FOXK2



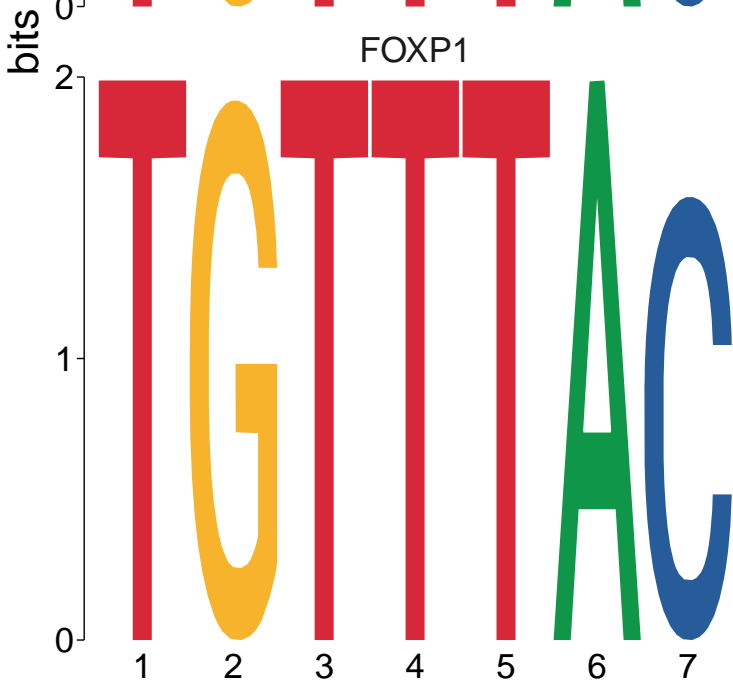
FOXK2



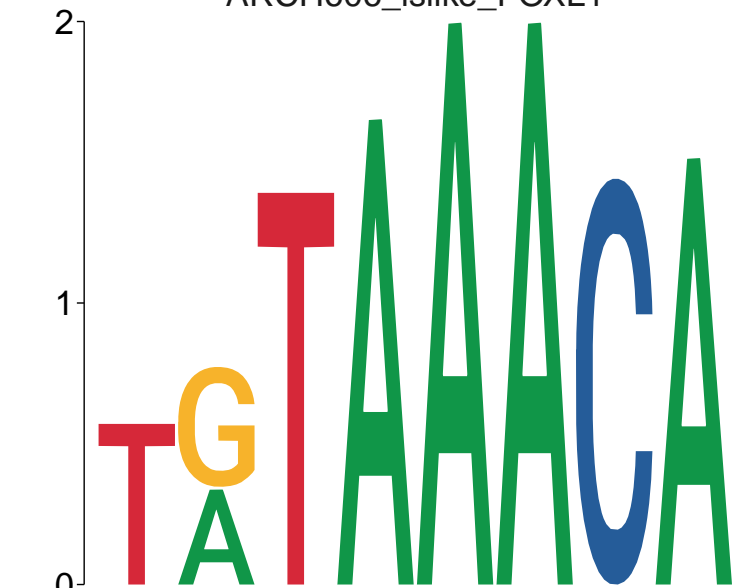
ARCH602_FOXP1



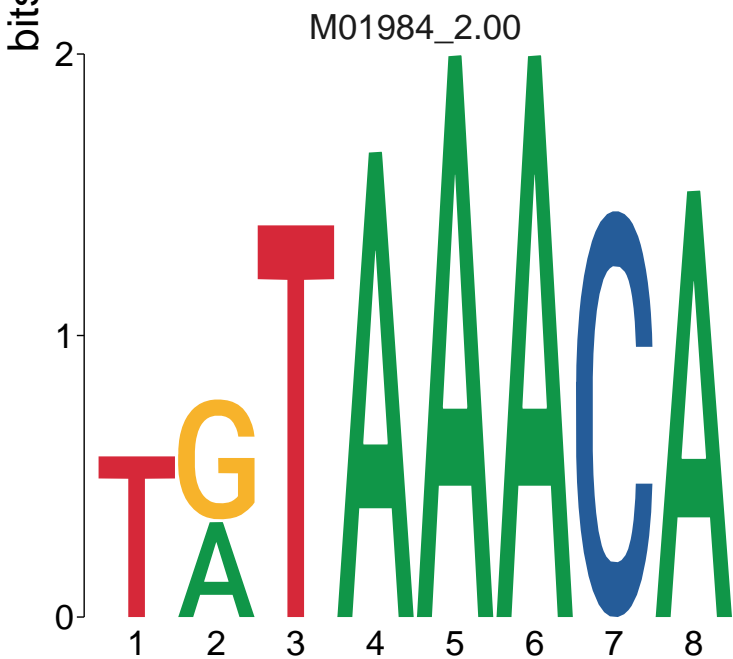
FOXP1



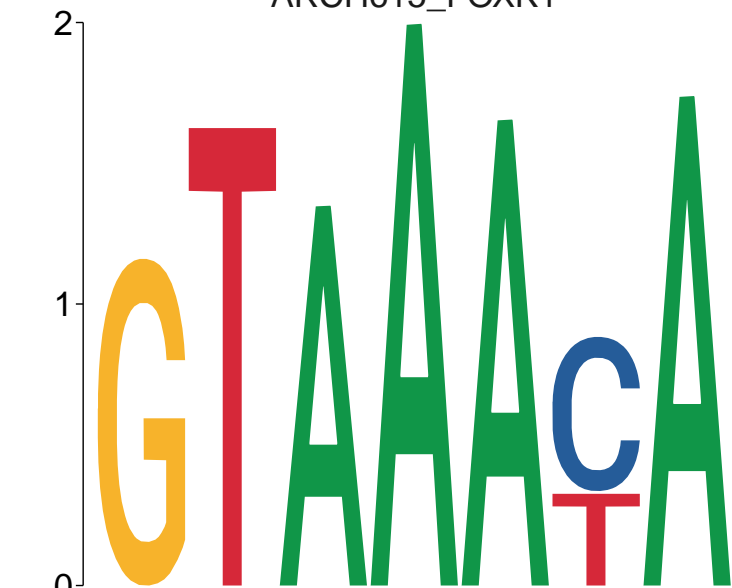
ARCH606_islike_FOXL1



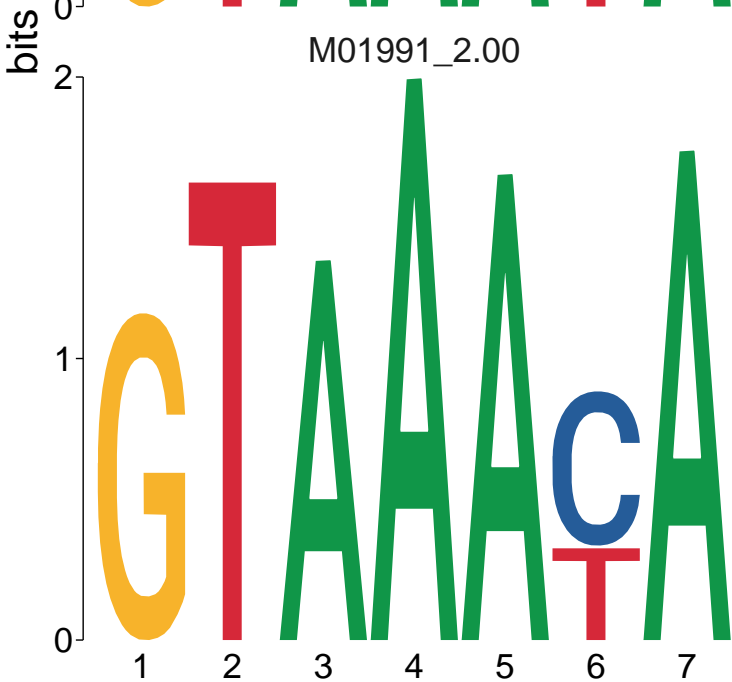
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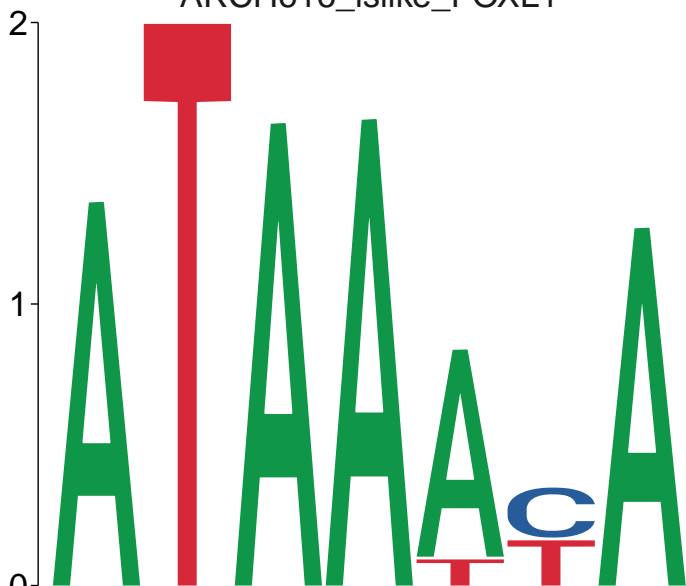
ARCH615_FOXK1



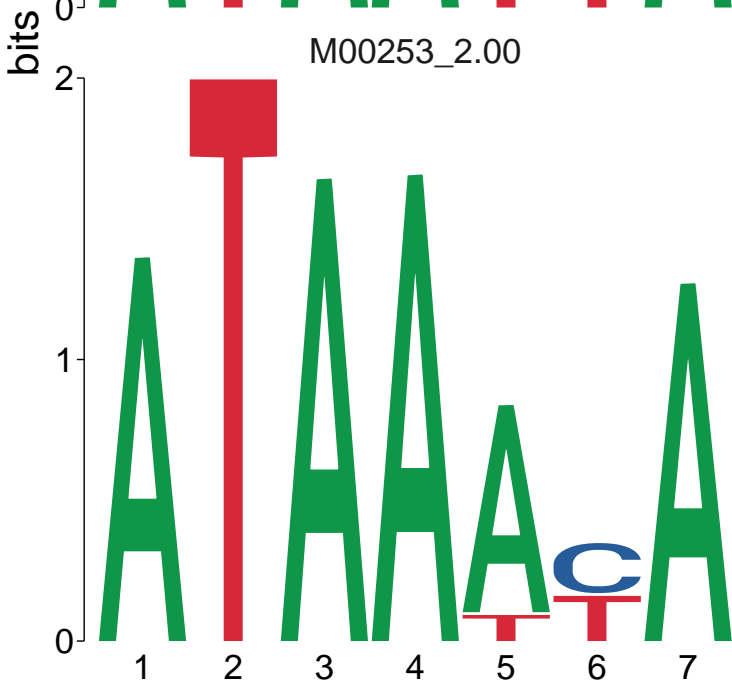
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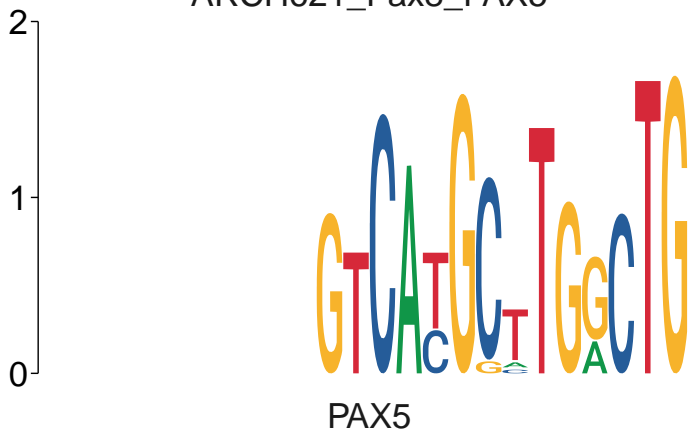
ARCH616_islike_FOXL1



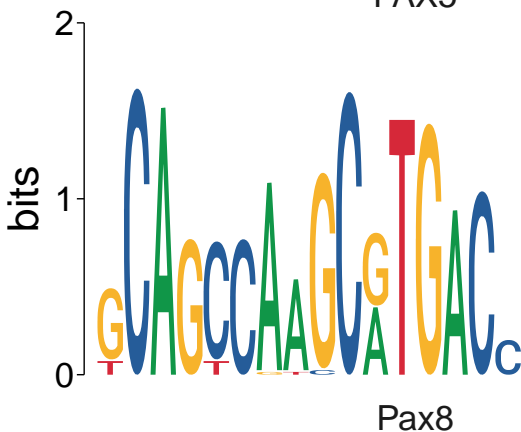
M00253_2.00



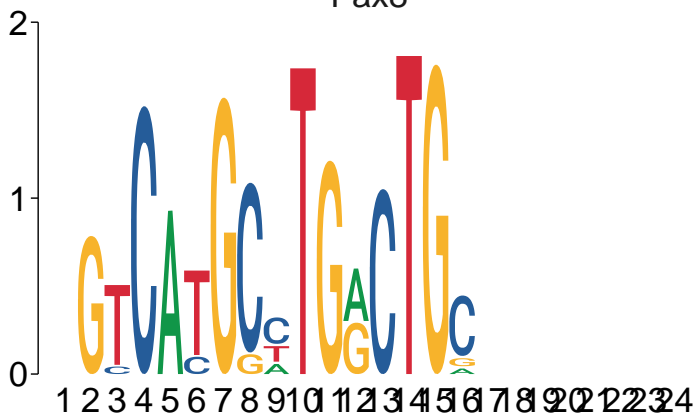
ARCH621_Pax8_PAX5



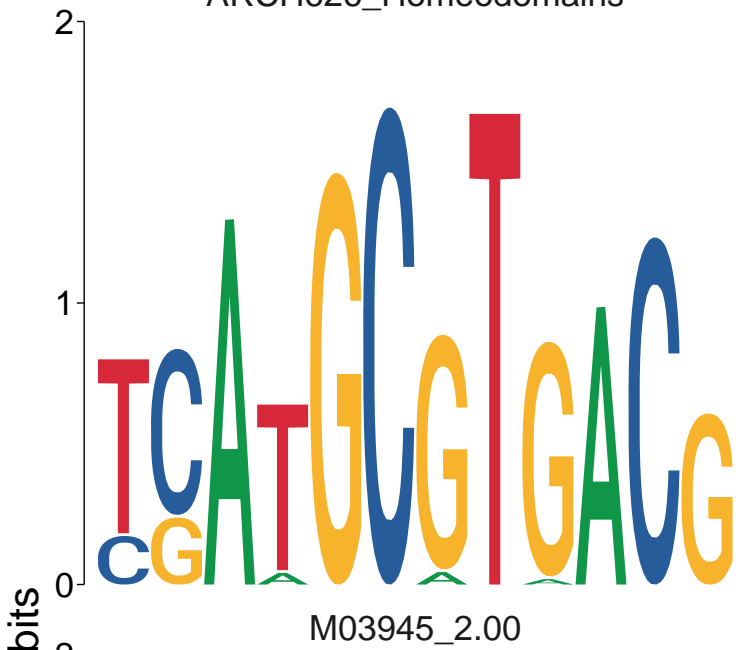
PAX5



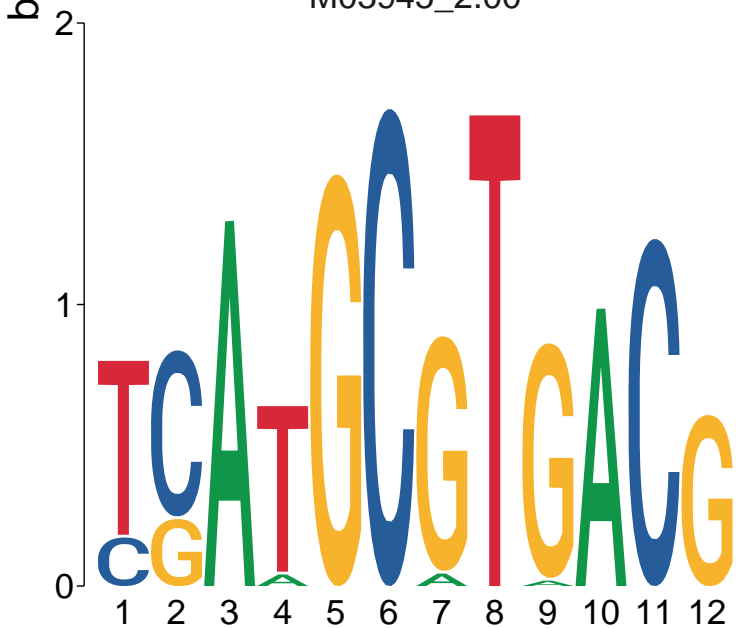
Pax8



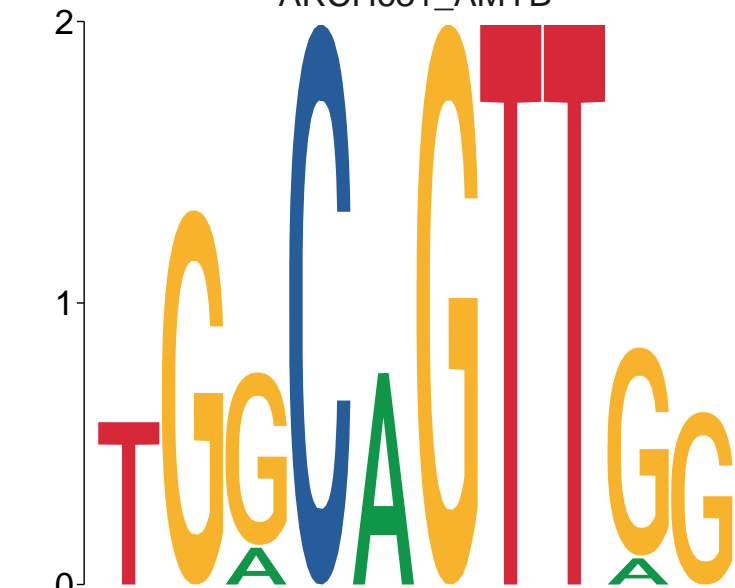
ARCH626_Homeodomains



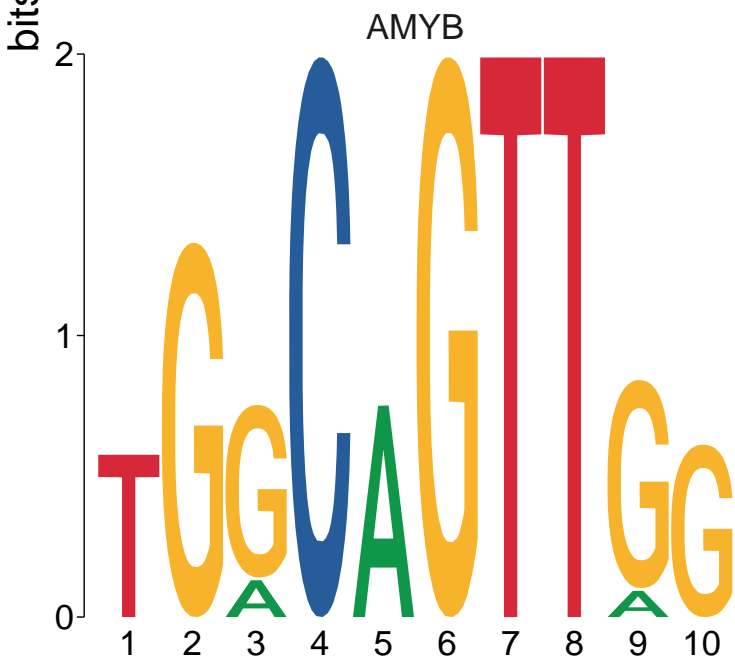
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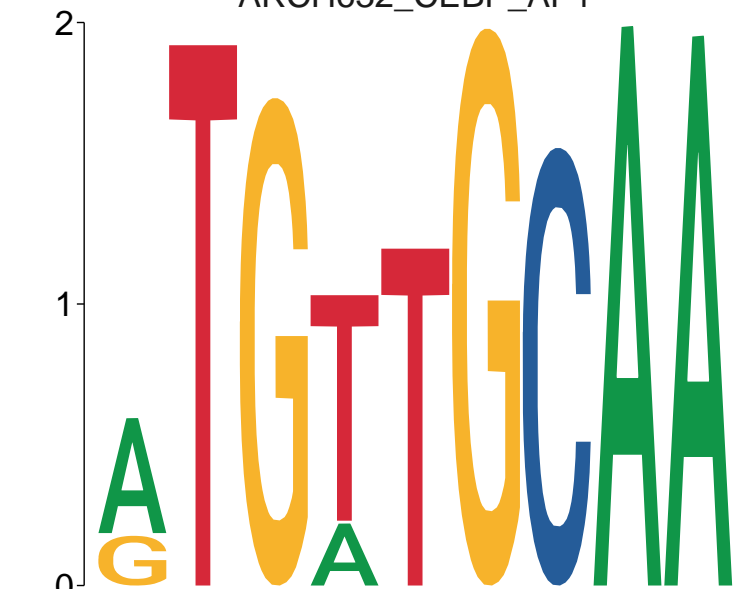
ARCH631_AMYB



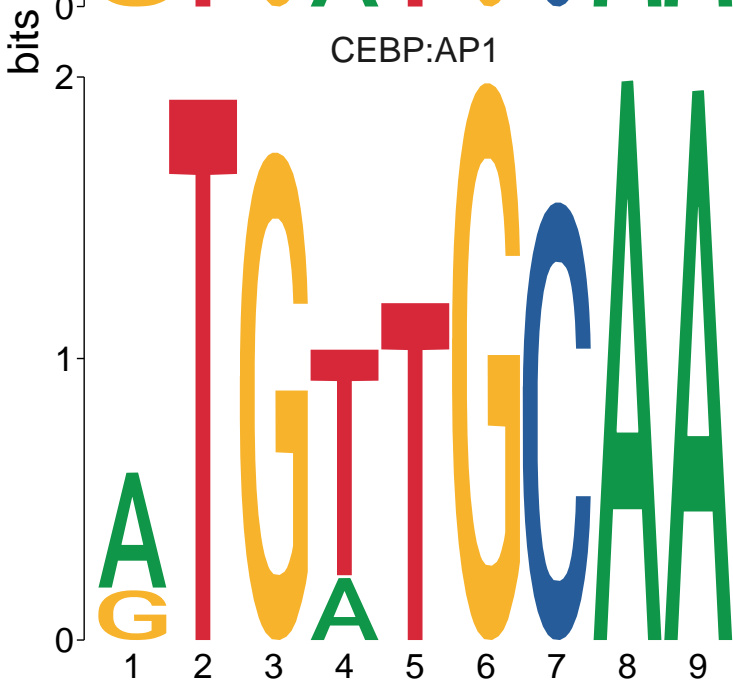
AMYB



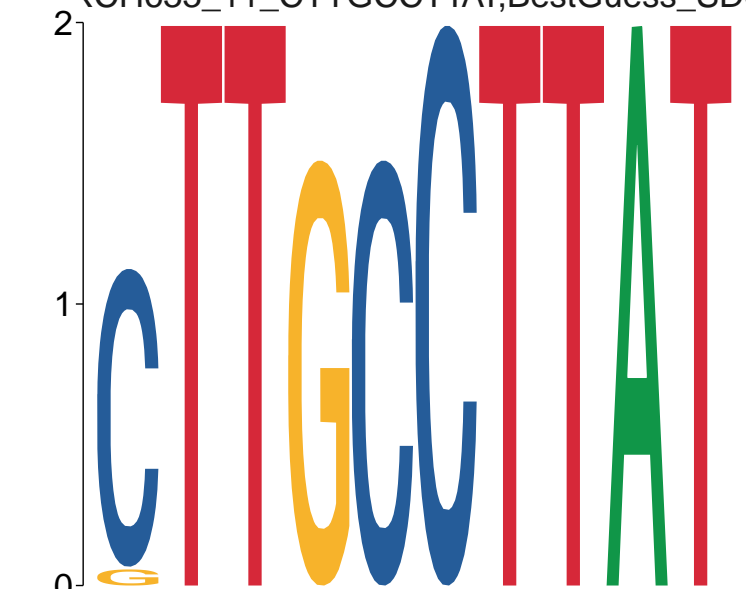
ARCH632_CEBP_AP1



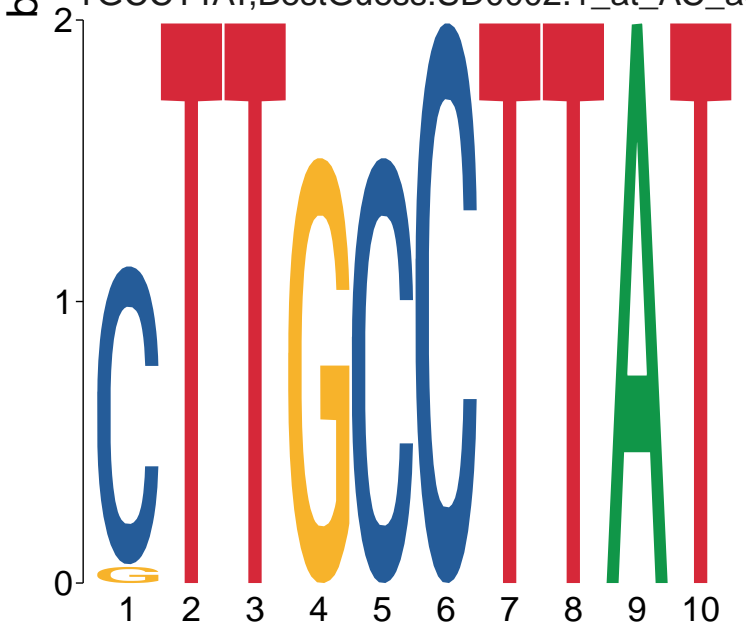
CEBP:AP1



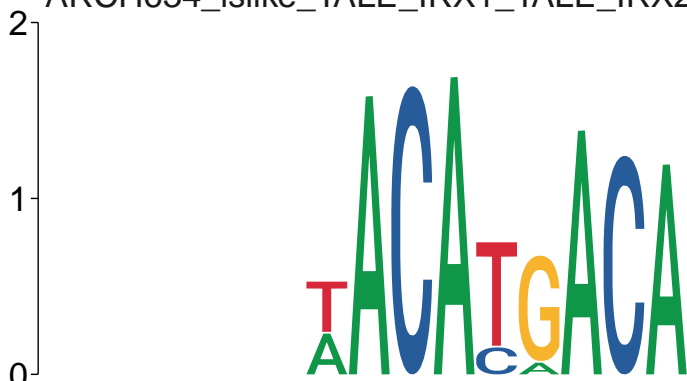
ACH633_11_CTTGCCTTAT,BestGuess_SD



TGCCTTAT,BestGuess:SD0002.1_at_AC_a



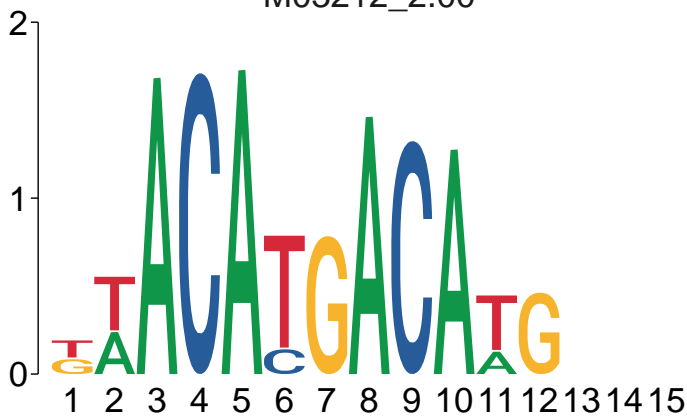
ARCH634_islike_TALE_IRX1_TALE_IRX2



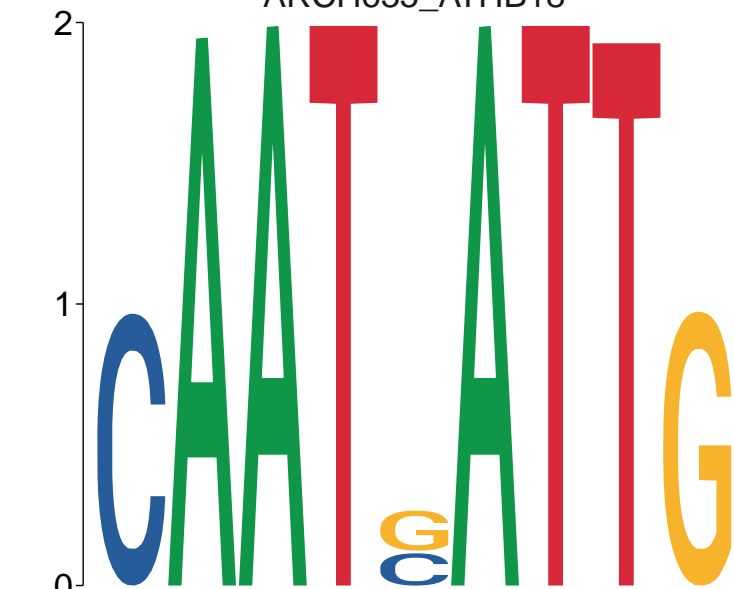
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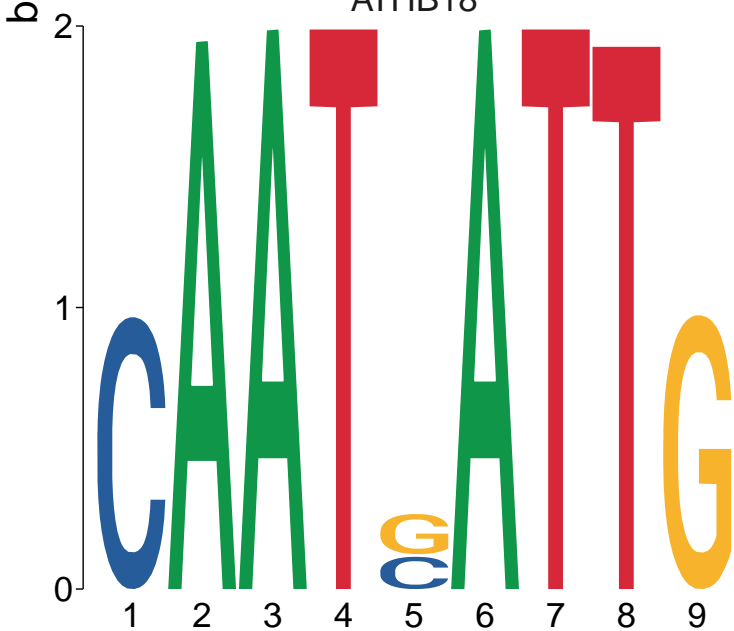
M03212_2.00



ARCH635_ATHB18



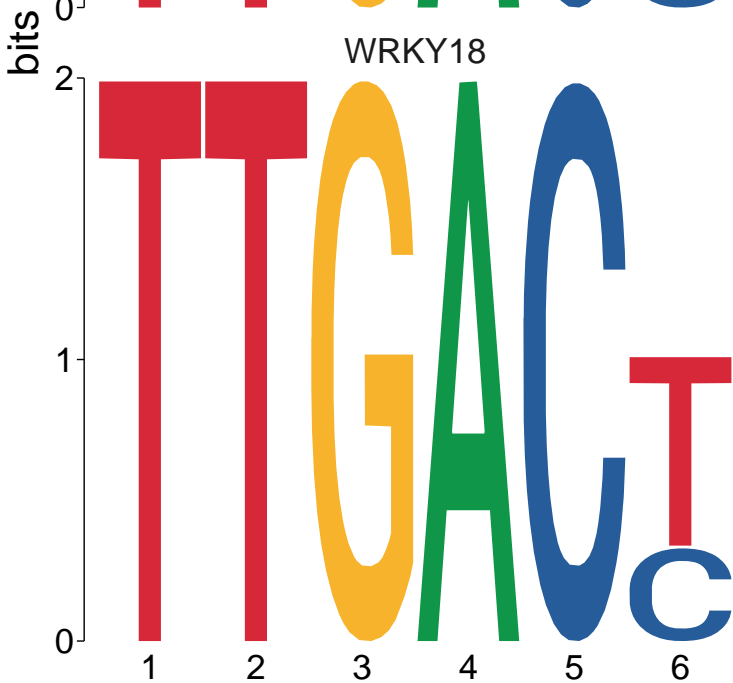
ATHB18



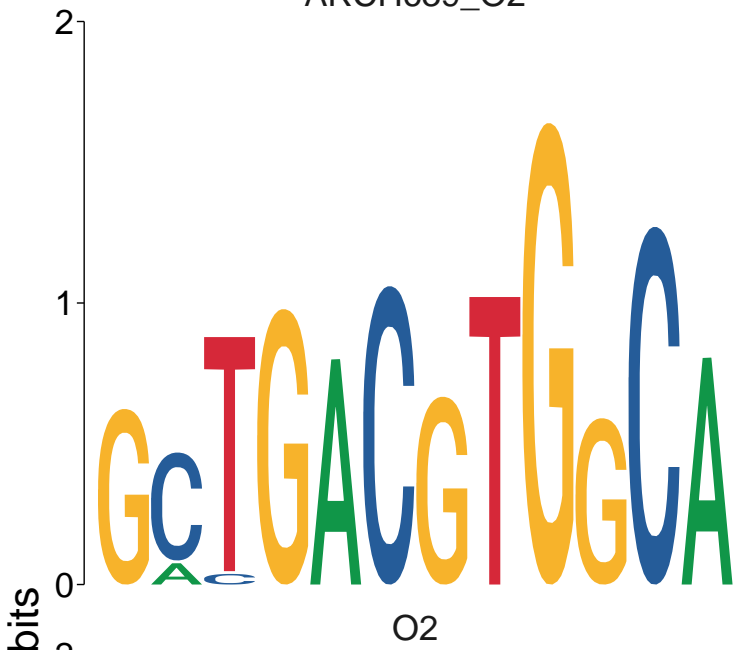
ARCH636_WRKY18



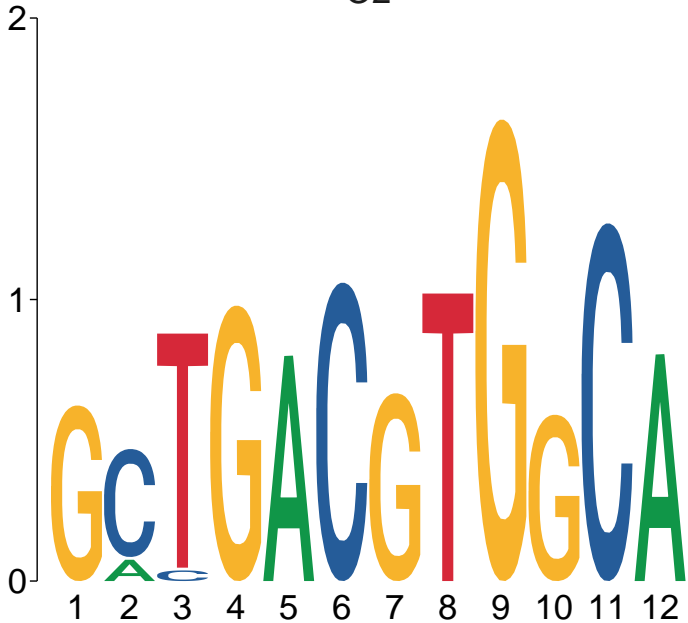
WRKY18



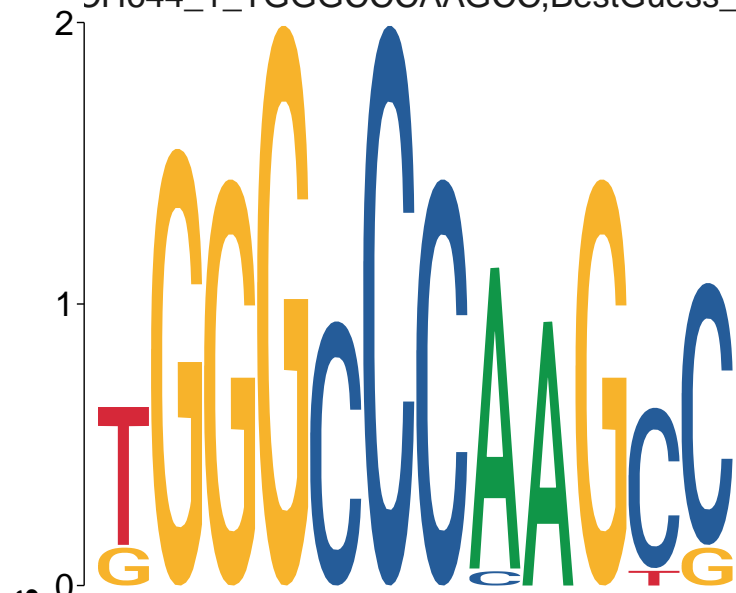
ARCH639_O2



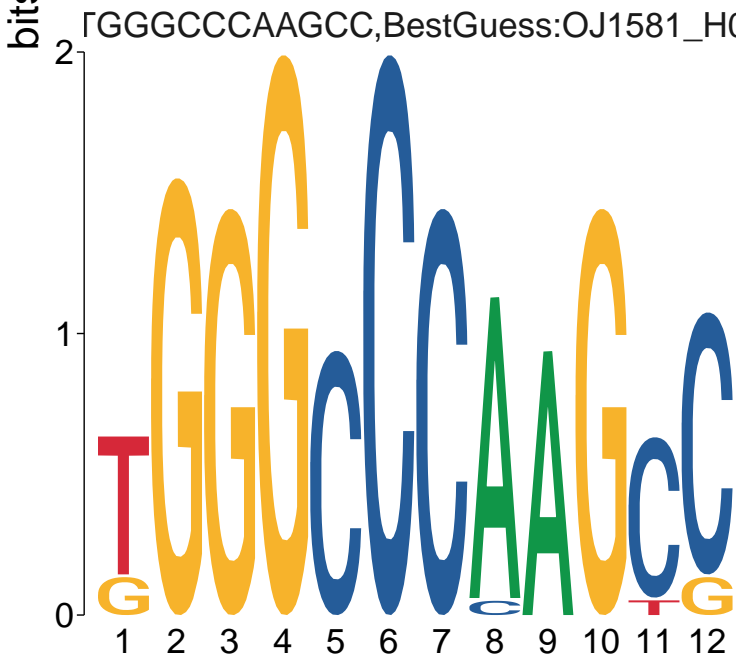
O2



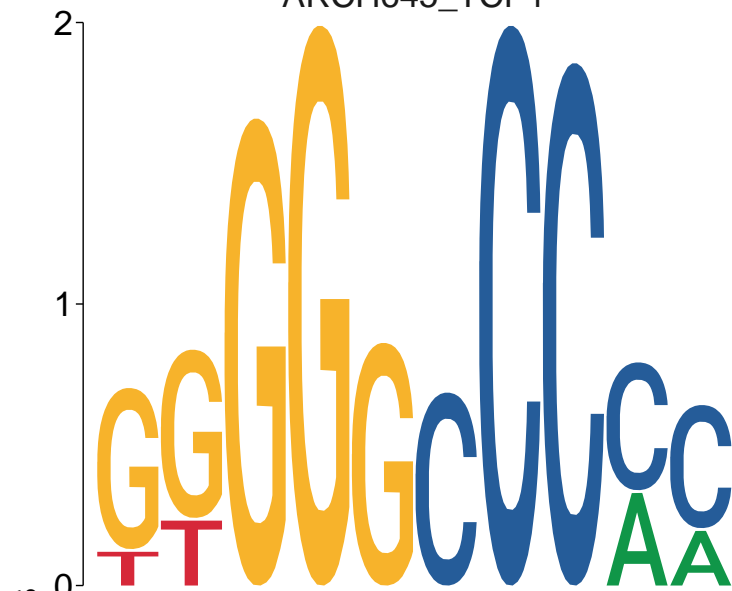
CH644_1_TGGGCCCAAGCC,BestGuess_(



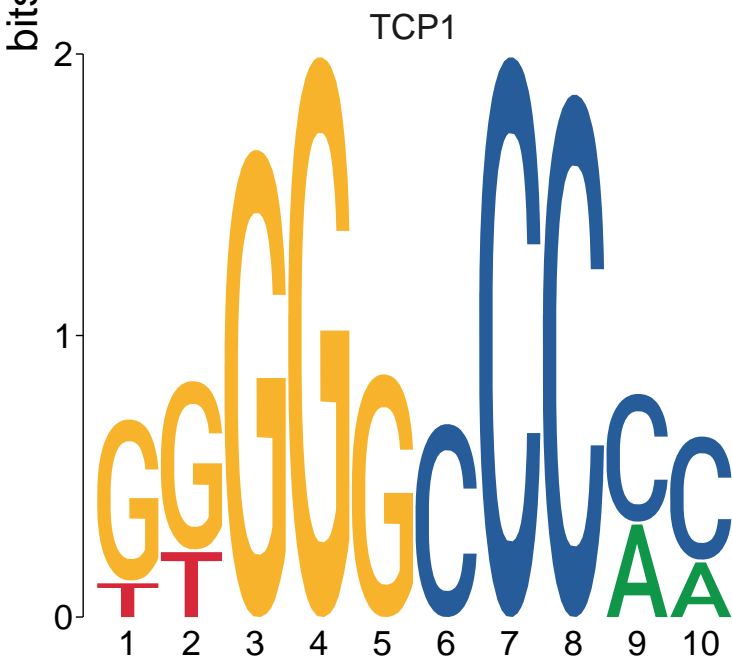
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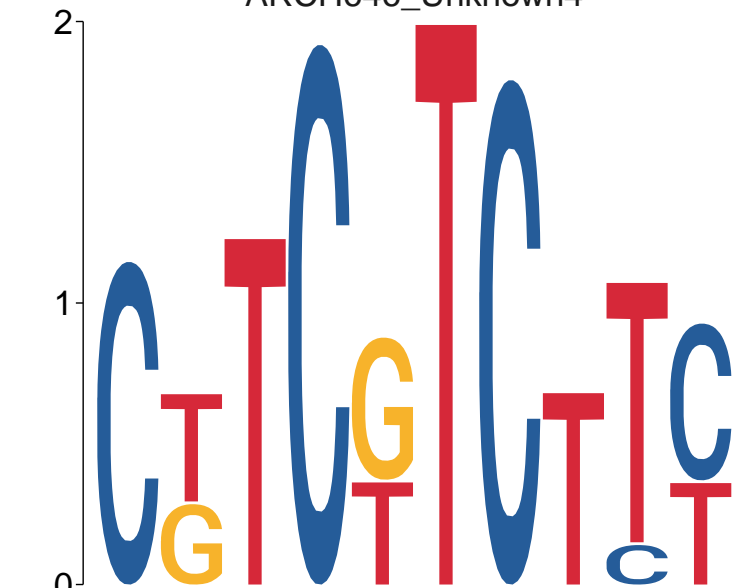
ARCH645_TCP1



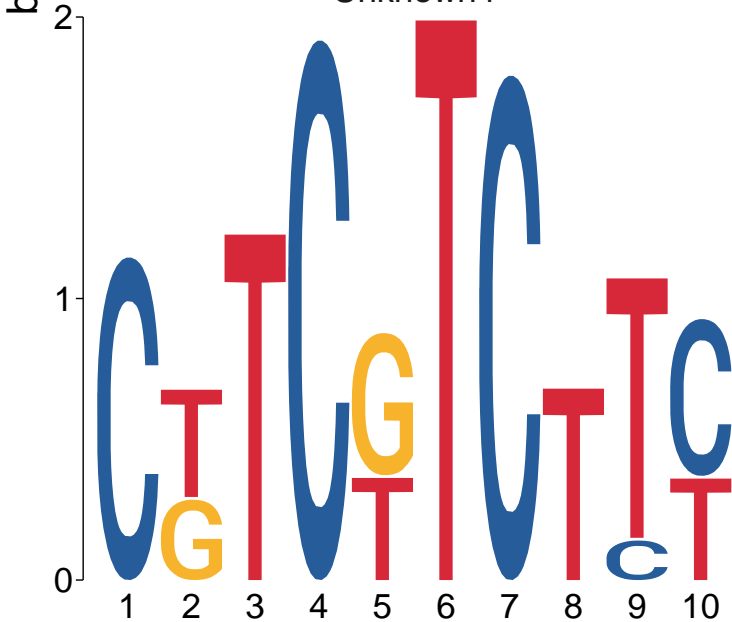
TCP1



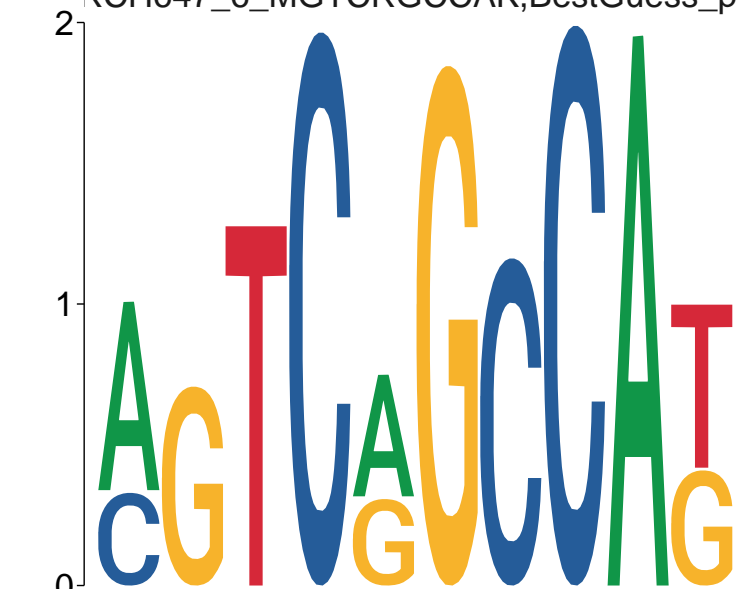
ARCH646_Unknown4



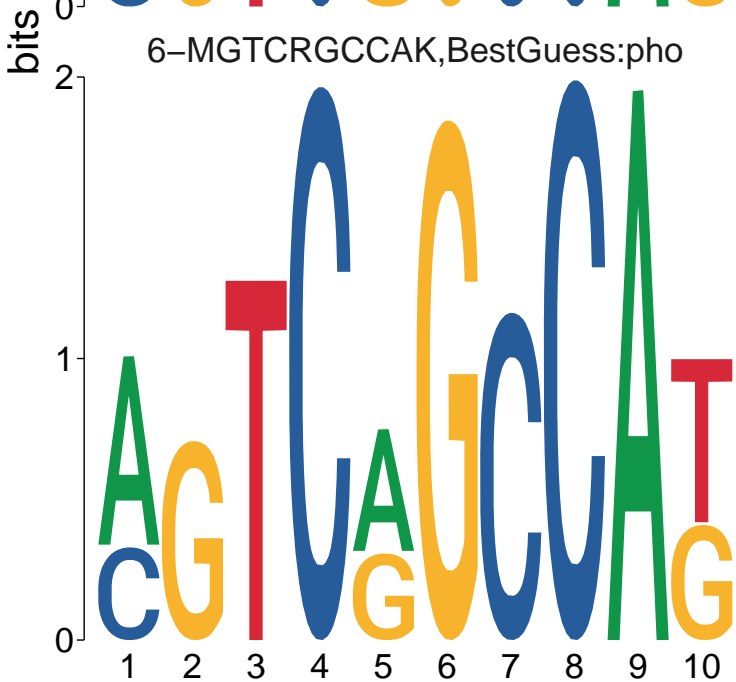
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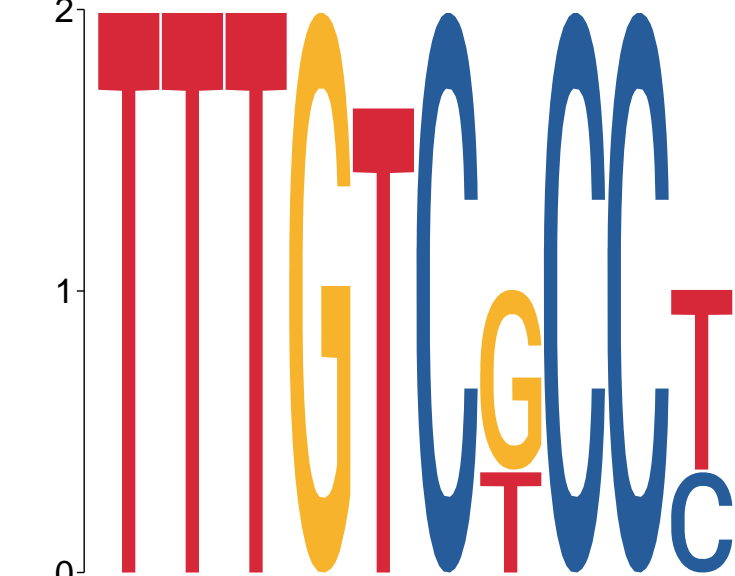
RCH647_6_MGTCRGCCAK,BestGuess_pr



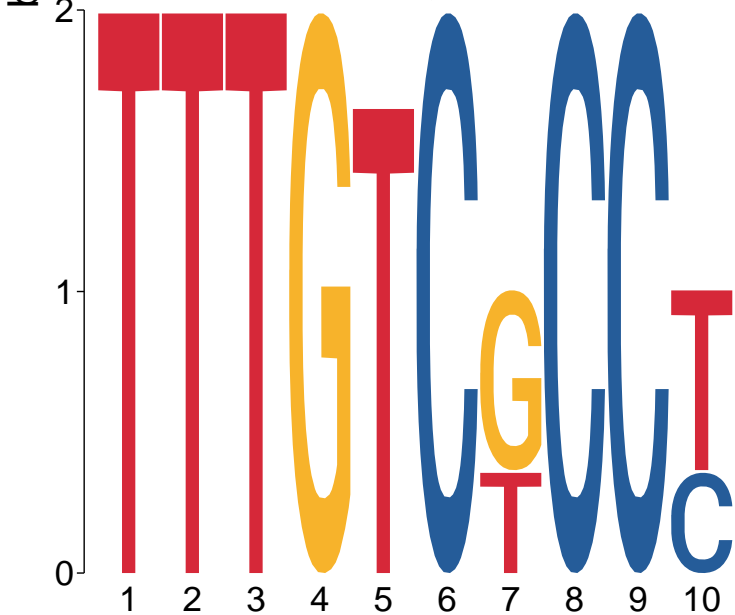
6-MGTCRGCCAK,BestGuess:pho



RCH648_7_TTTGTCKCCY,BestGuess_ID



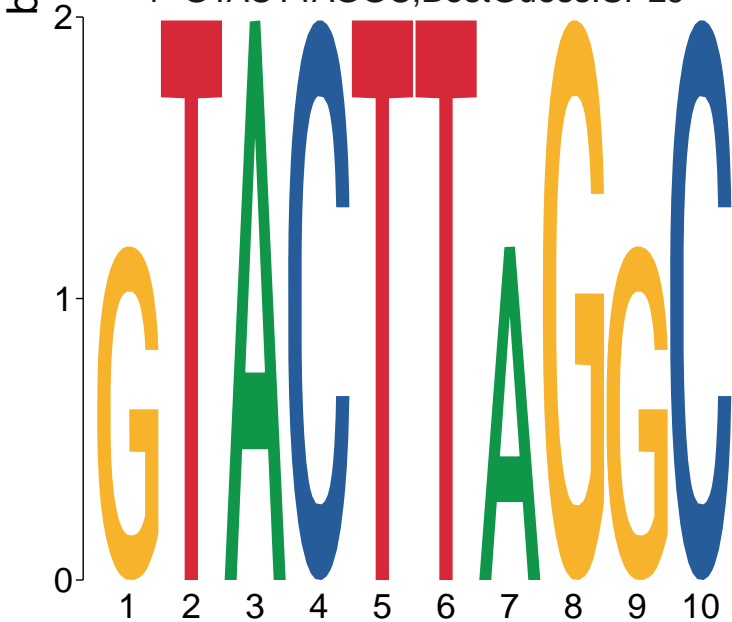
7-TTTGTCKCCY,BestGuess:ID1



RCH649_1_GTACTTAGGC,BestGuess_SPL



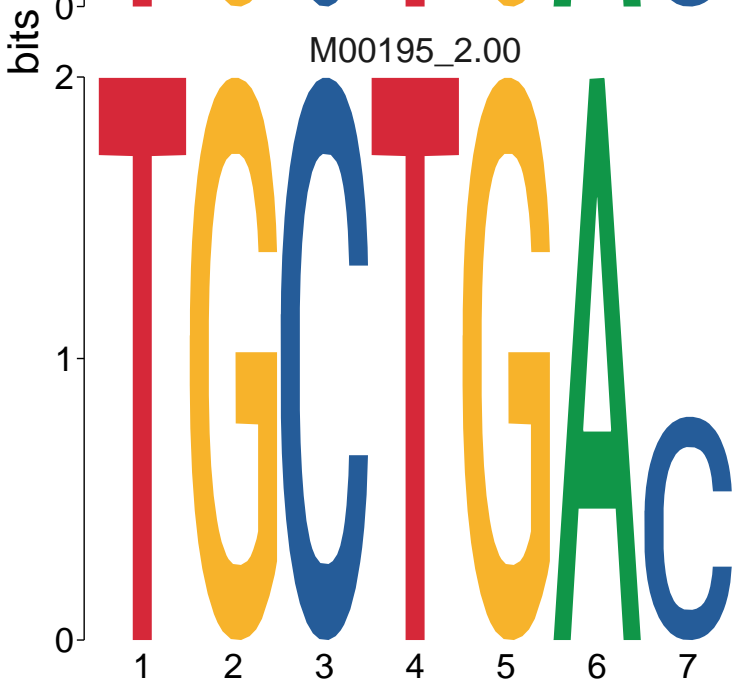
1-GTACTTAGGC,BestGuess:SPL9



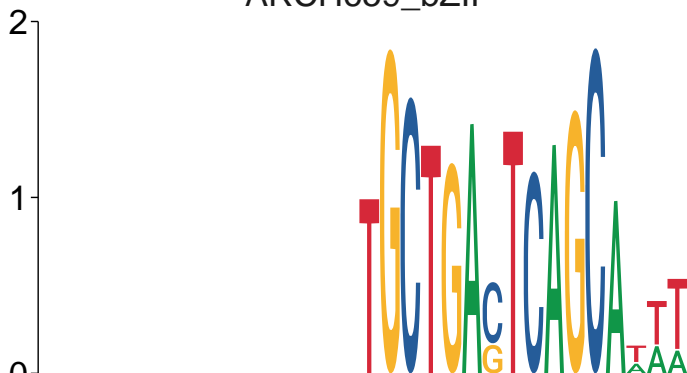
ARCH658_HBP1



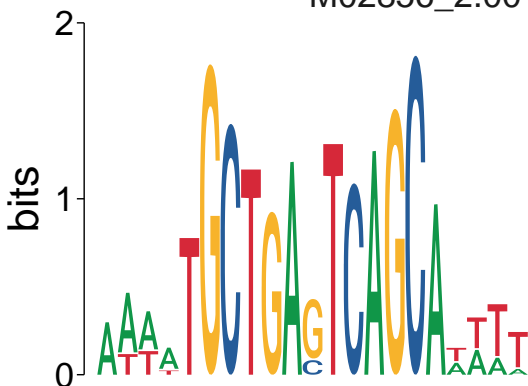
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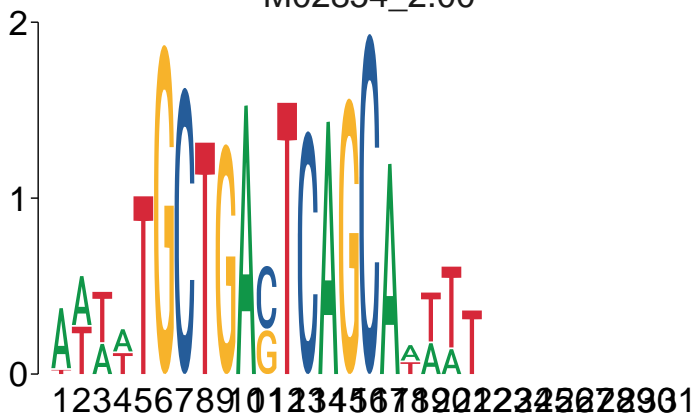
ARCH659_bZIP



M02856_2.00



M02854_2.00



ARCH660_bZIP



M04369 2.00



M04366_2.00



M04354 2.00



M04352 2.00



M04349 2.00



M04346_2.00



M02858_2.00



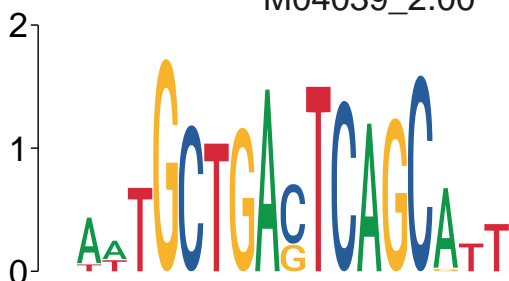
ARCH661_bZIP



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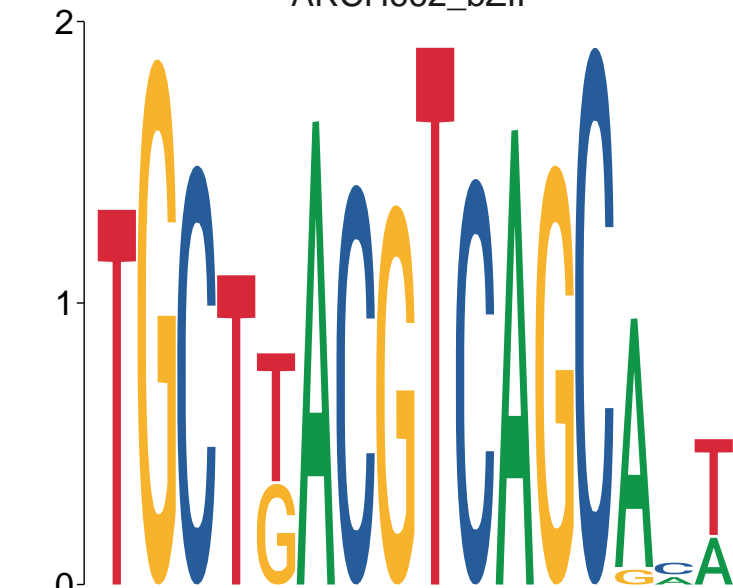
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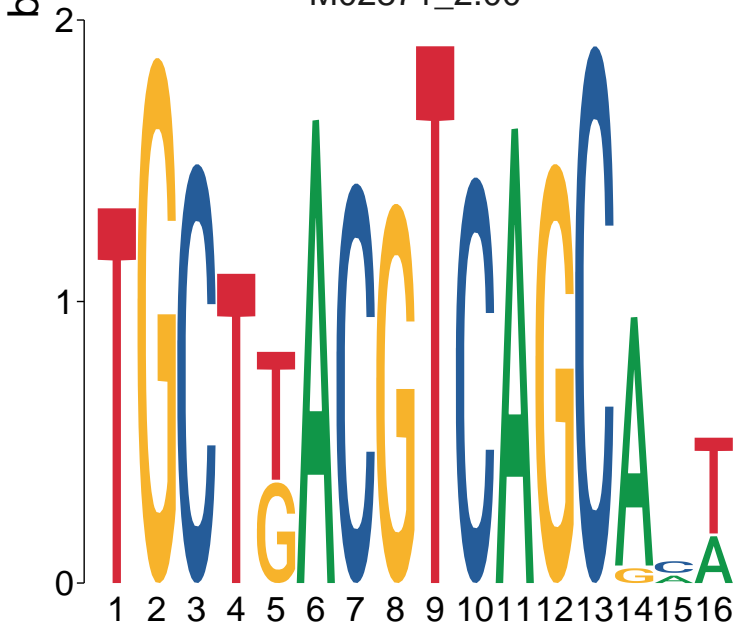
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ARCH662_bZIP



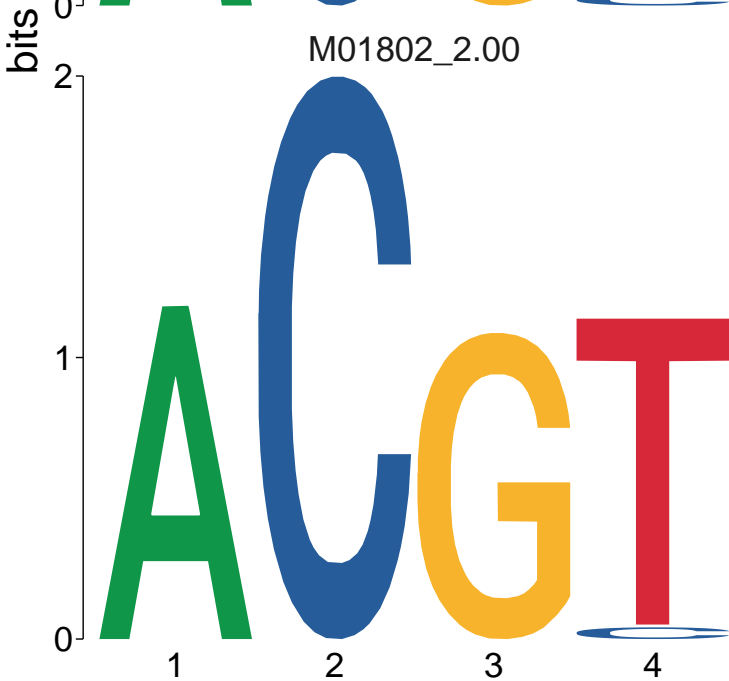
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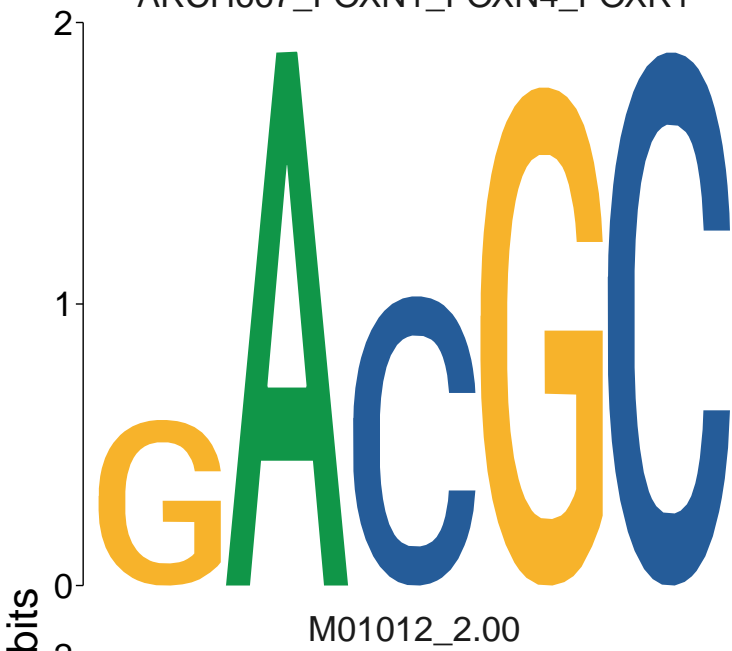
ARCH664_XBP1



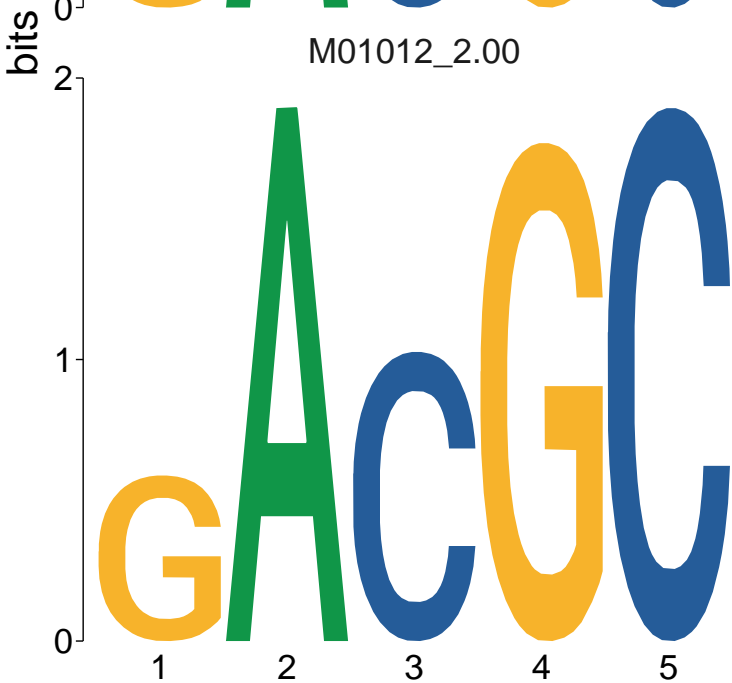
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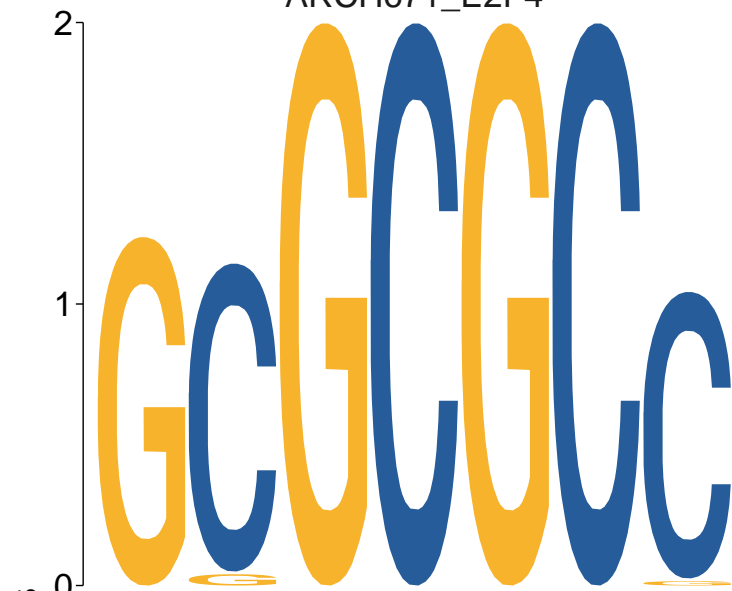
ARCH667_FOXN1_FOXN4_FOXR1



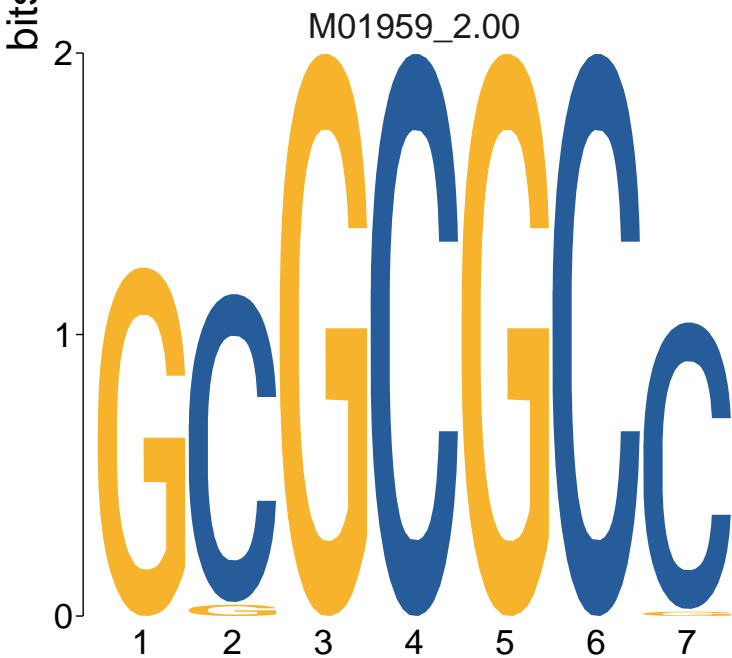
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ARCH671_E2F4



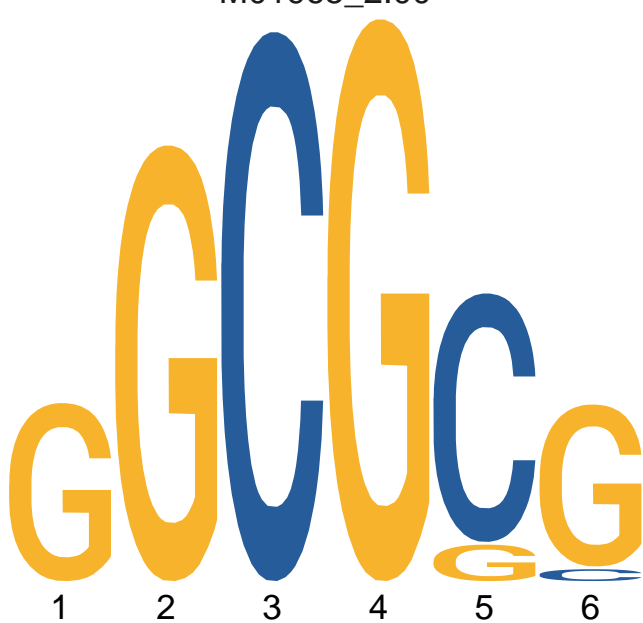
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ARCH672_E2F4



M01965_2.00



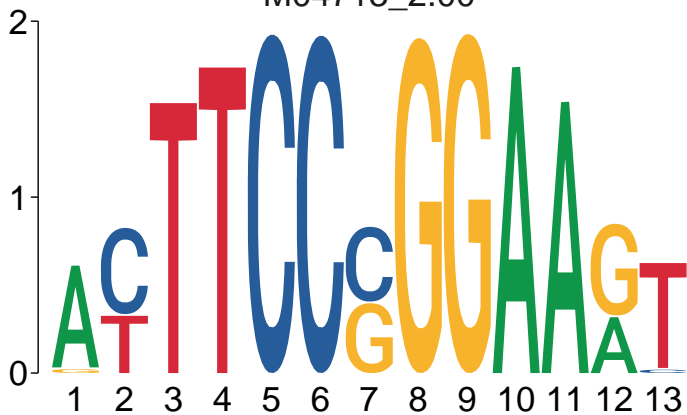
ARCH685_ETV6



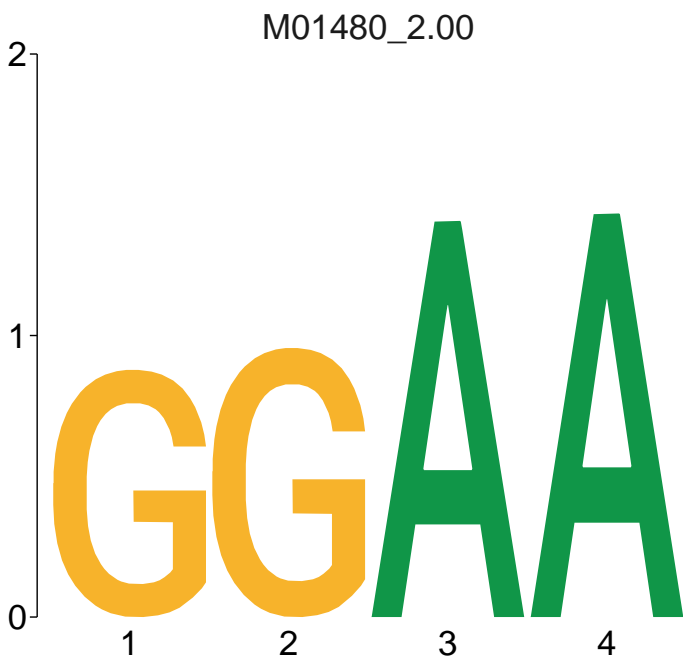
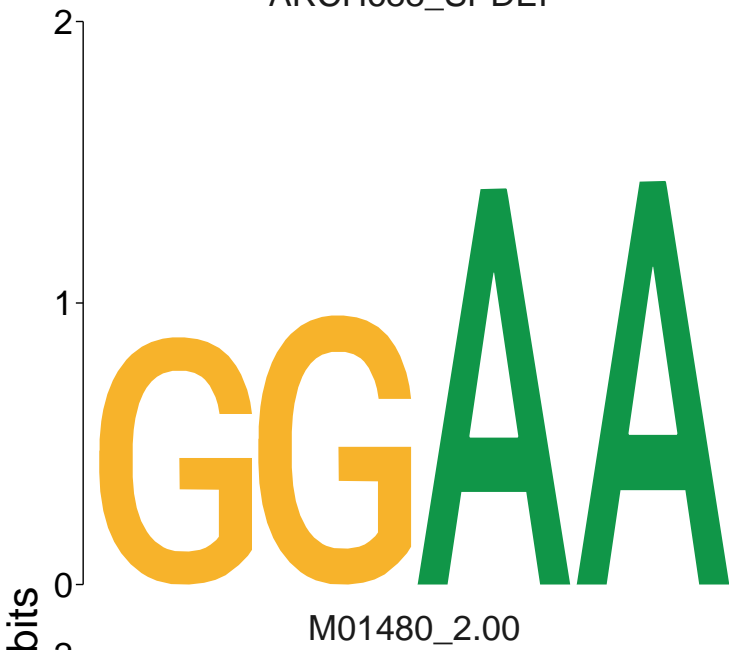
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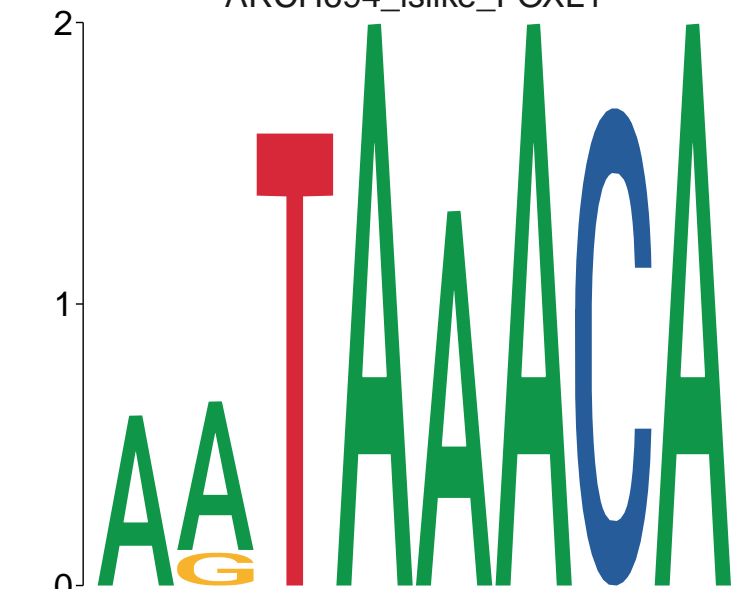
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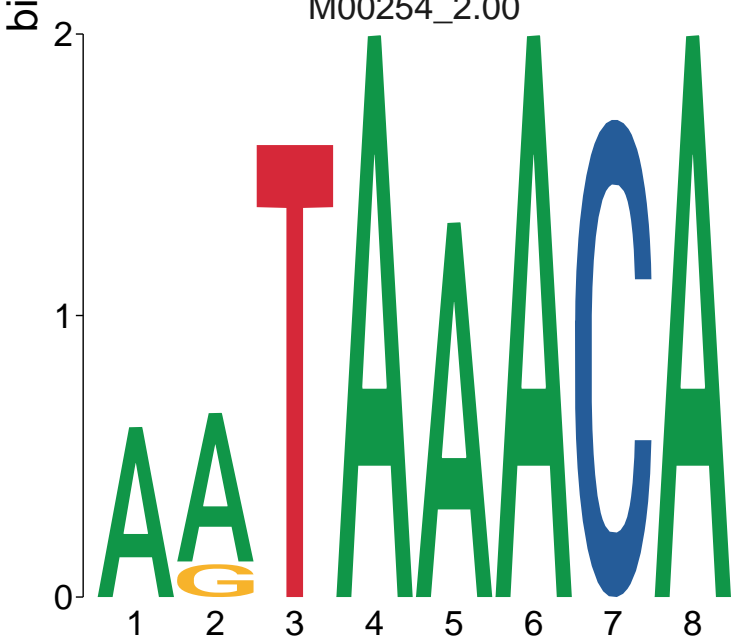
ARCH688_SPDEF



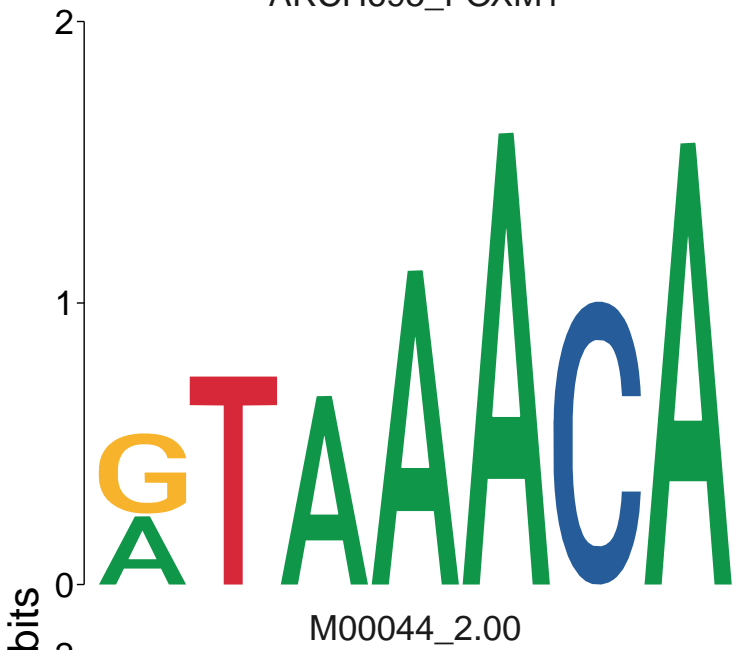
ARCH694_islike_FOXL1



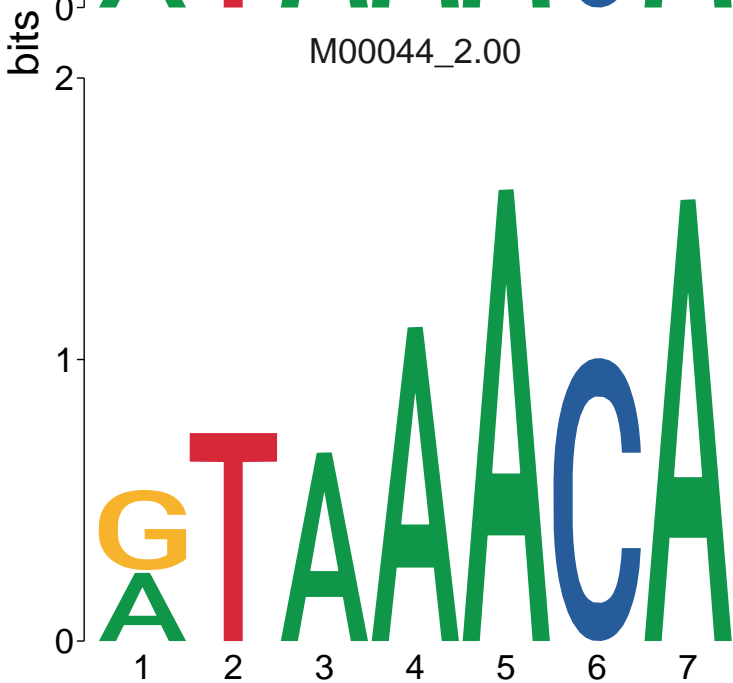
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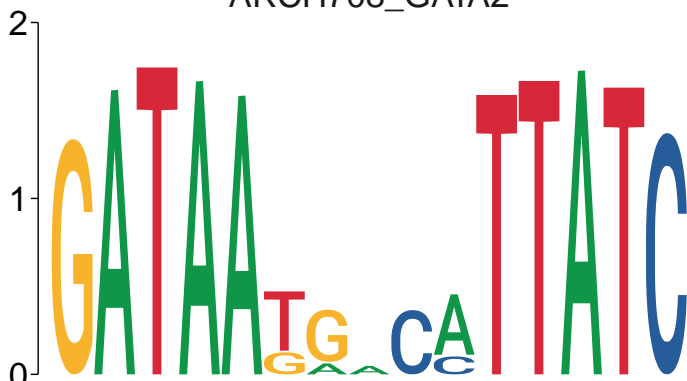
ARCH695_FOXM1



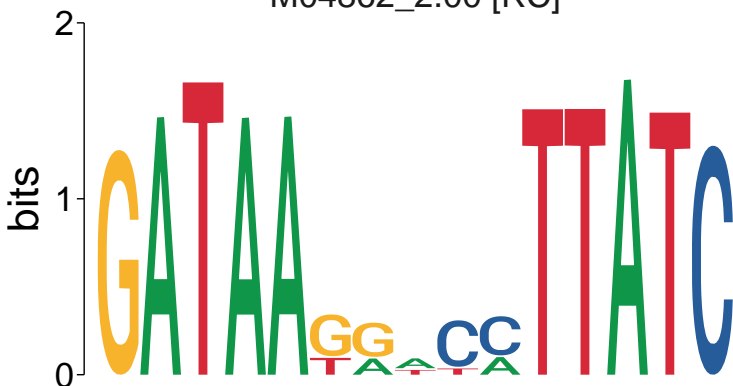
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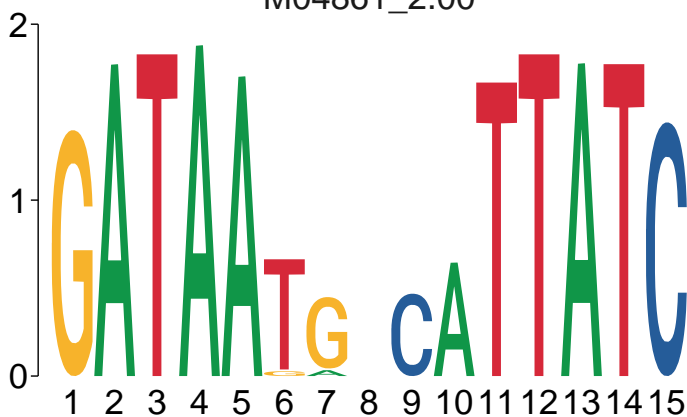
ARCH708_GATA2



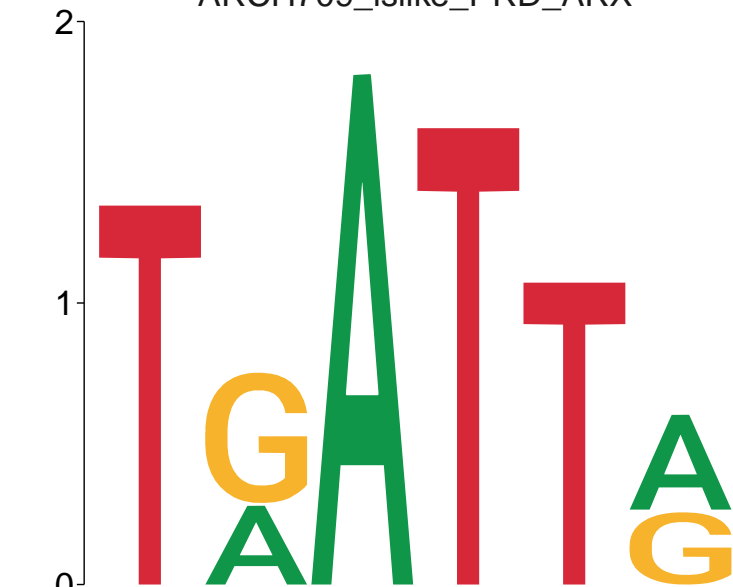
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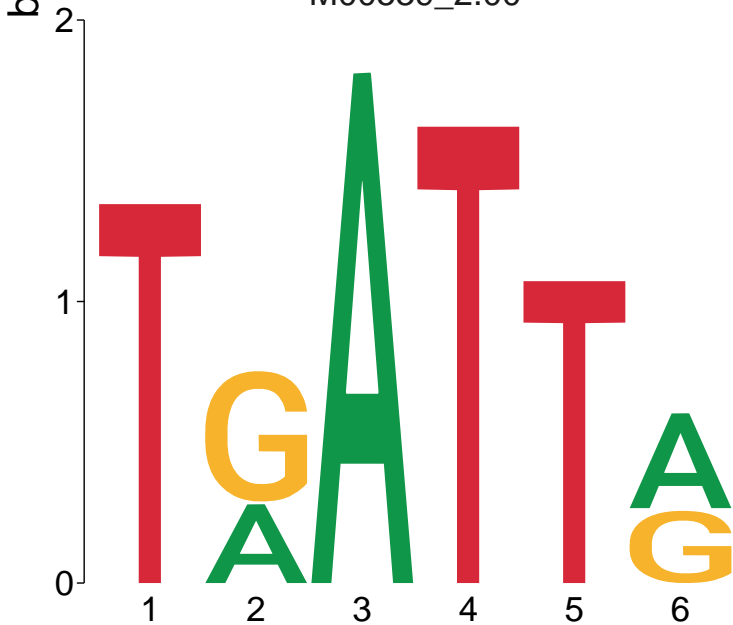
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ARCH709_islike_PRD_ARX



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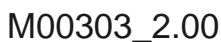




1

0

bits



2

1

0

1

2

3

4

5

6

7

8

ARCH714_ANTP_HOXA2_ANTP_HOXA3

2

1

0

bits

1

2

3

4

5

6

M02112_2.00

2

1

0

1

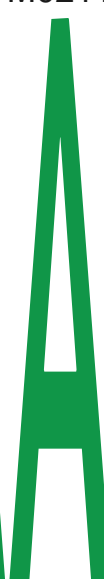
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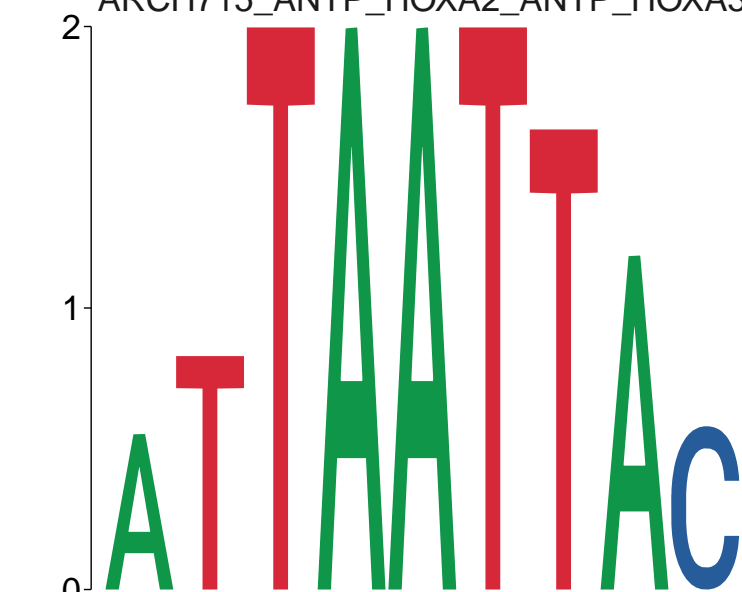
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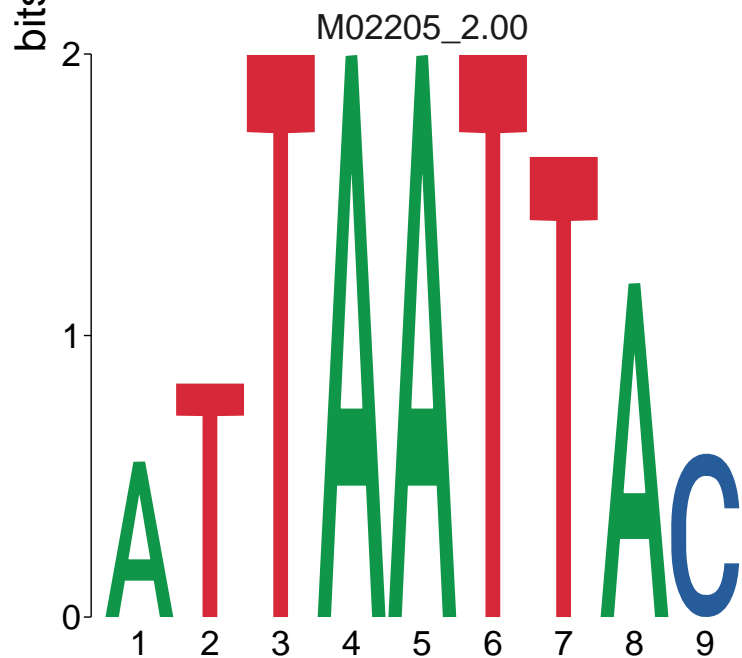
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ARCH715_ANTP_HOXA2_ANTP_HOXA3



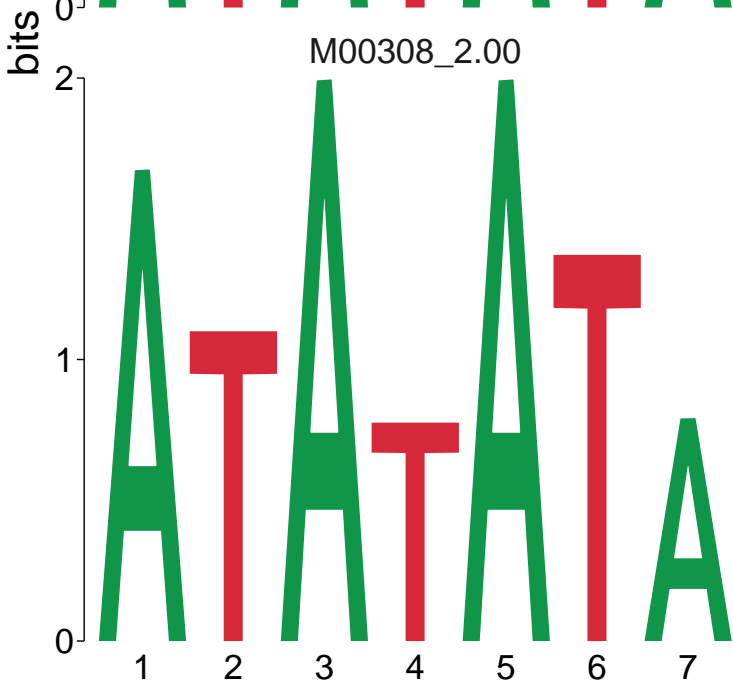
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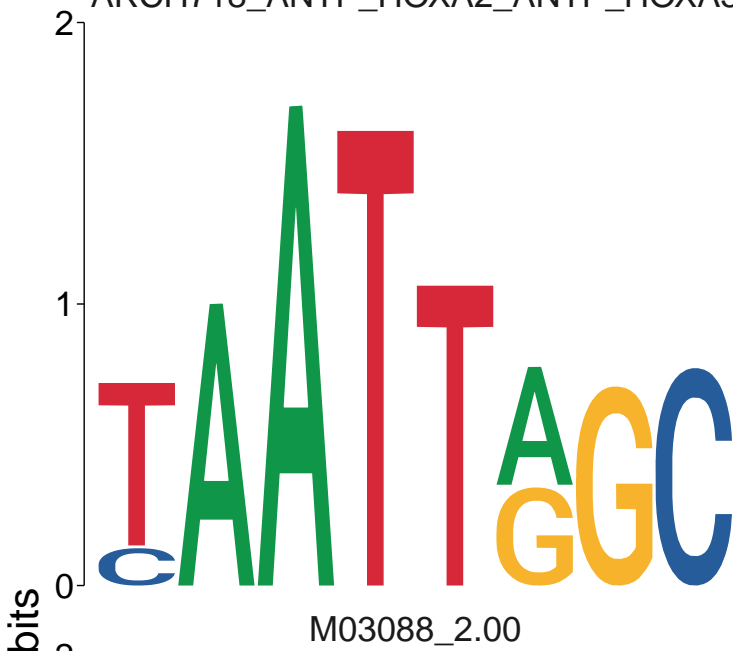
ARCH717_ANTP_MNX1



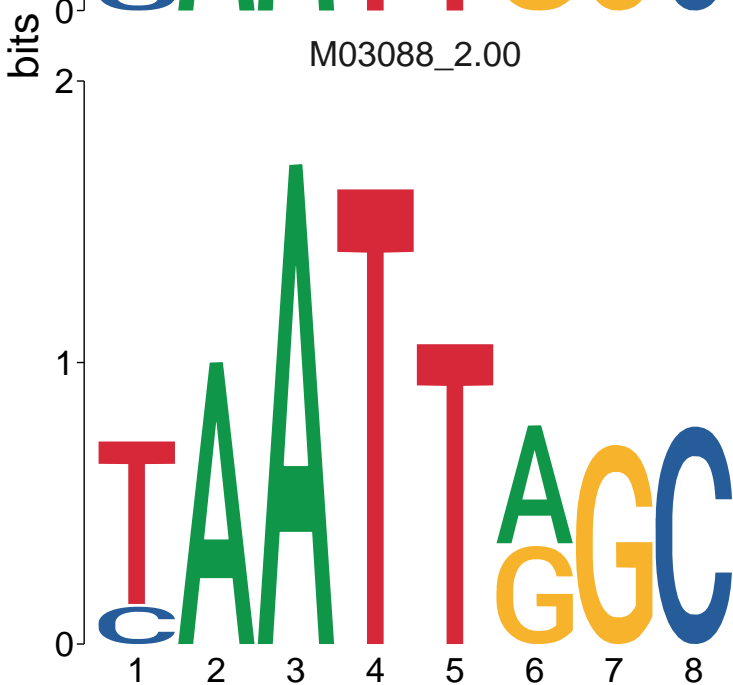
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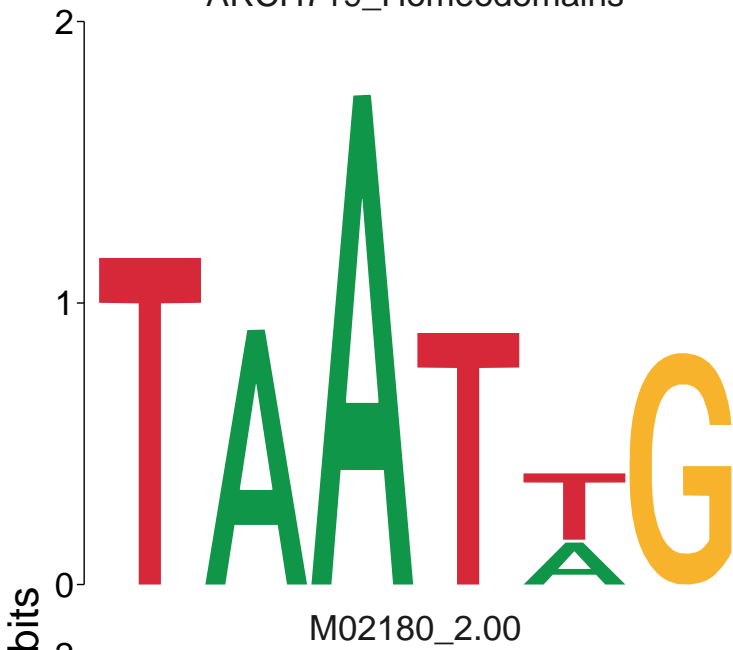
ARCH718_ANTP_HOXA2_ANTP_HOXA3



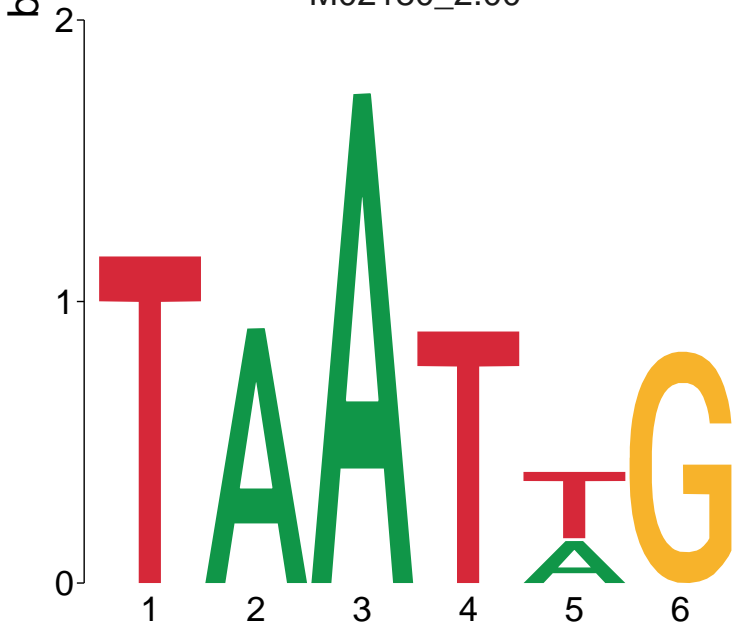
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ARCH719_Homeodomains



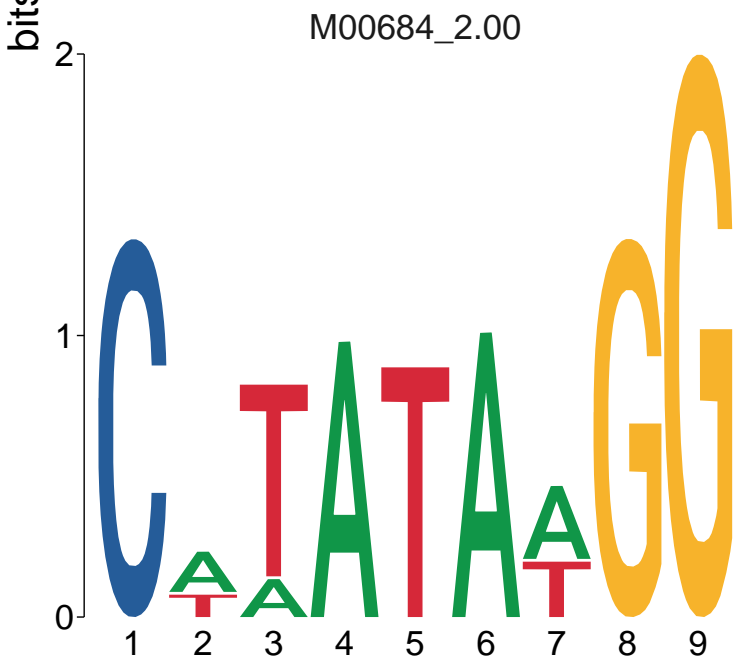
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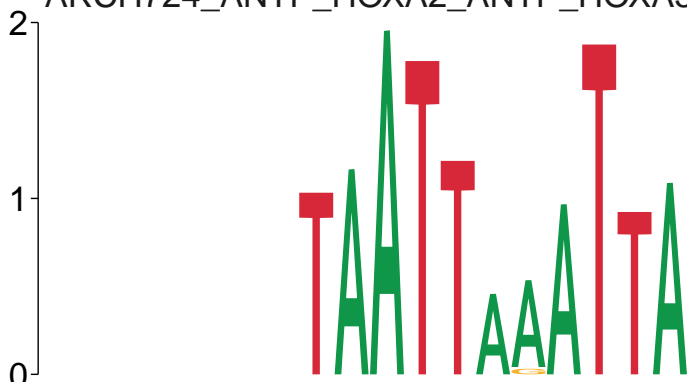
ARCH721_SRF



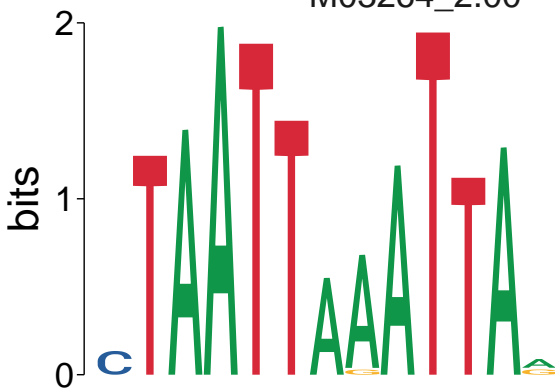
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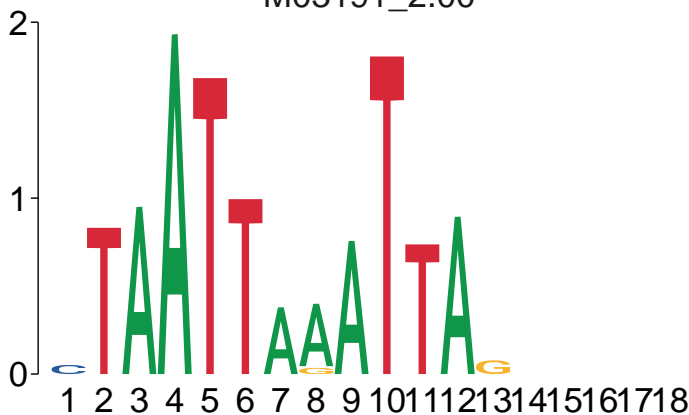
ARCH724_ANTP_HOXA2_ANTP_HOXA3



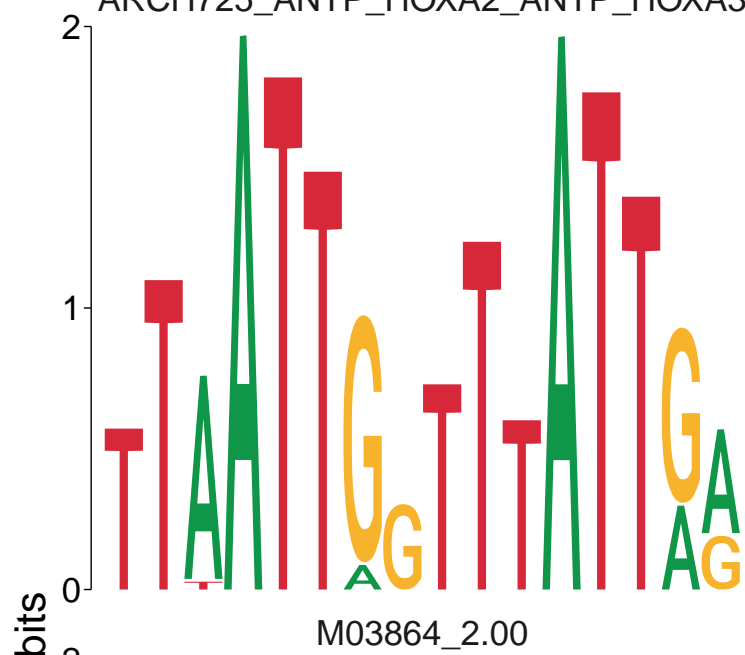
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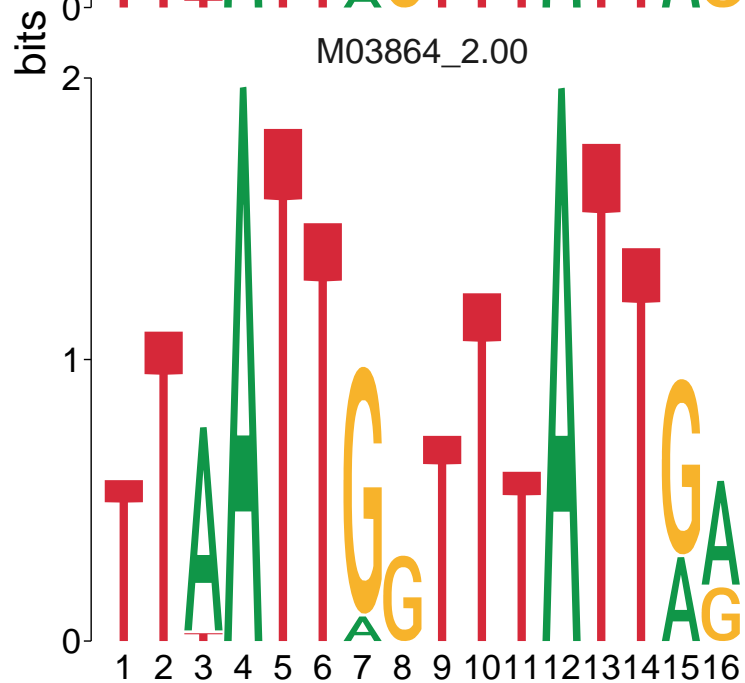
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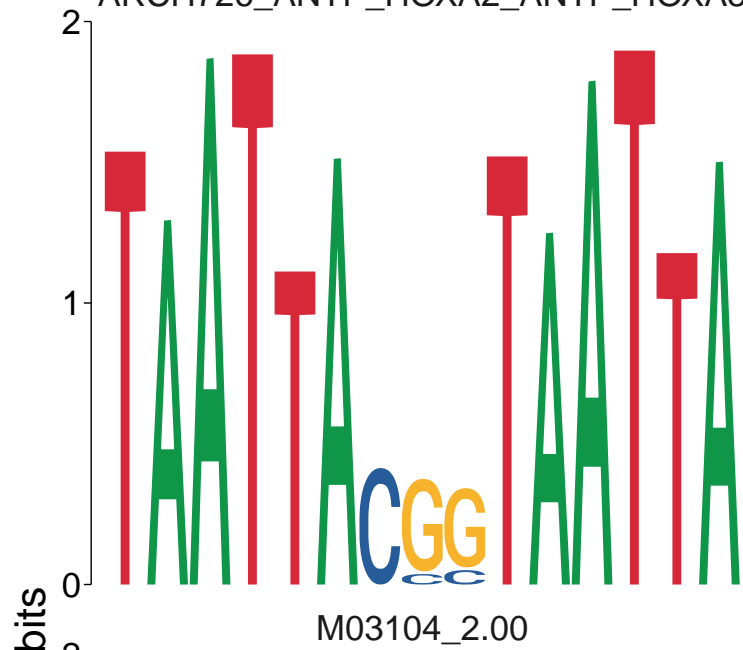
ARCH725_ANTP_HOXA2_ANTP_HOXA3



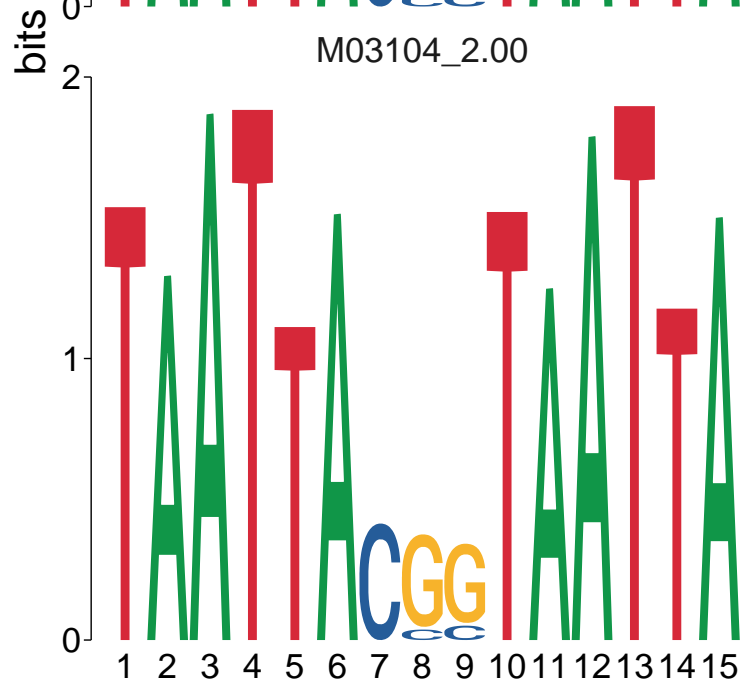
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ARCH726_ANTP_HOXA2_ANTP_HOXA3



M03104_2.00



ARCH727_ANTP_HOXA2_ANTP_HOXA3



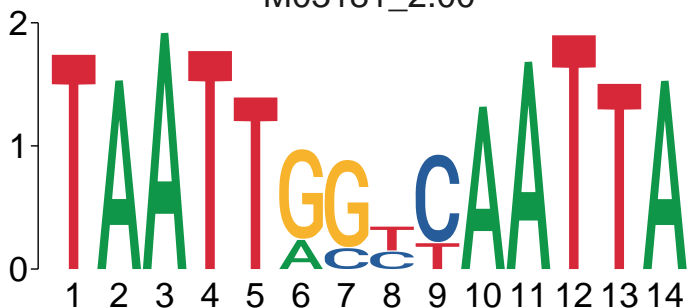
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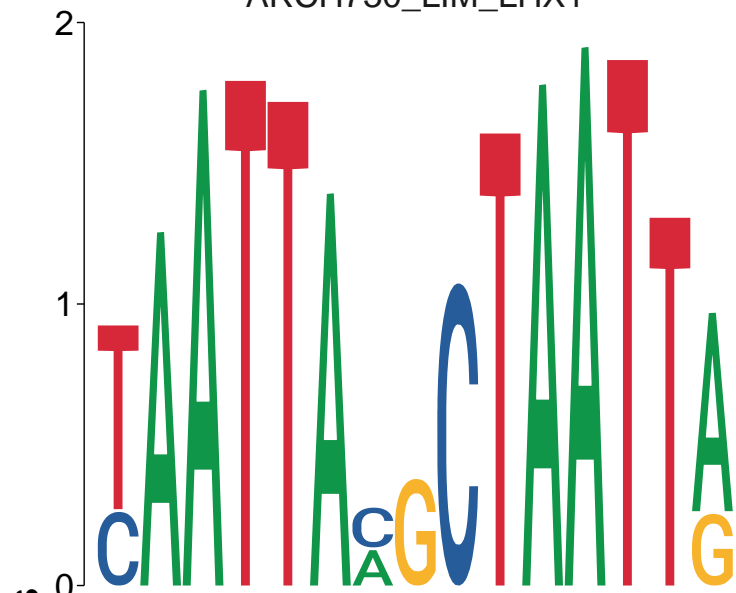
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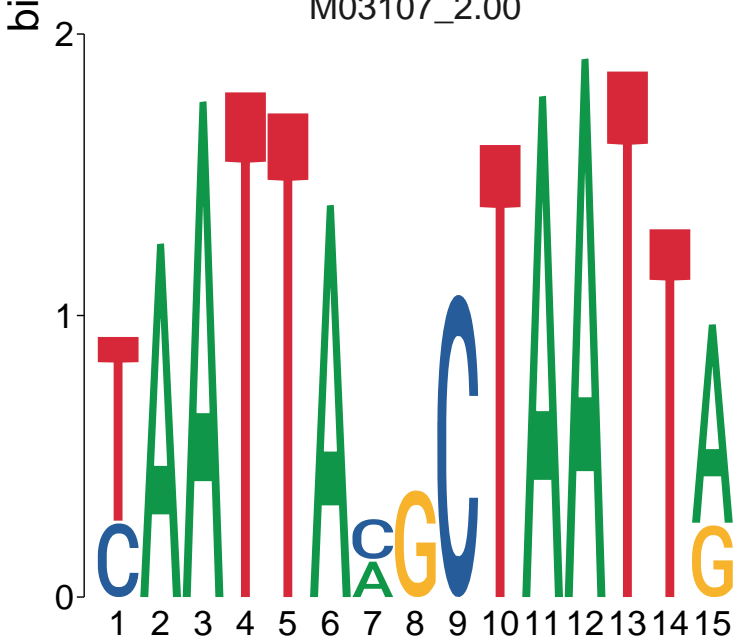
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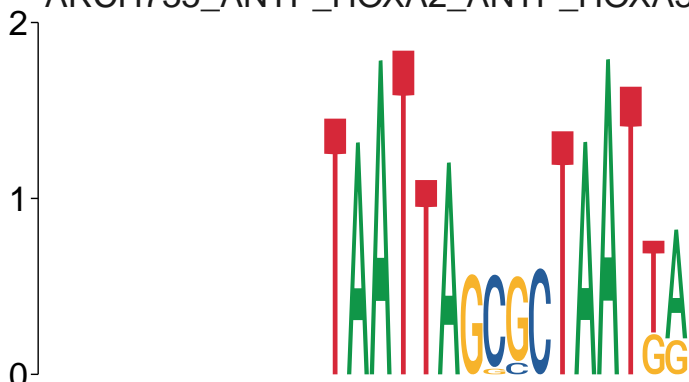
ARCH730_LIM_LHX1



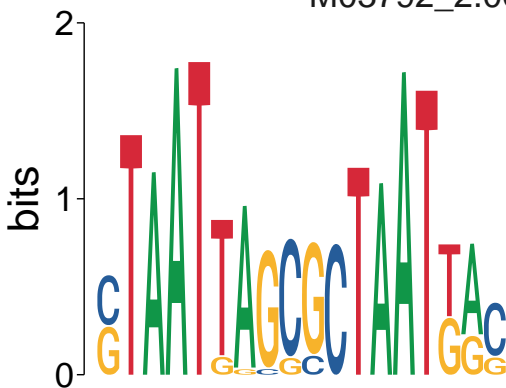
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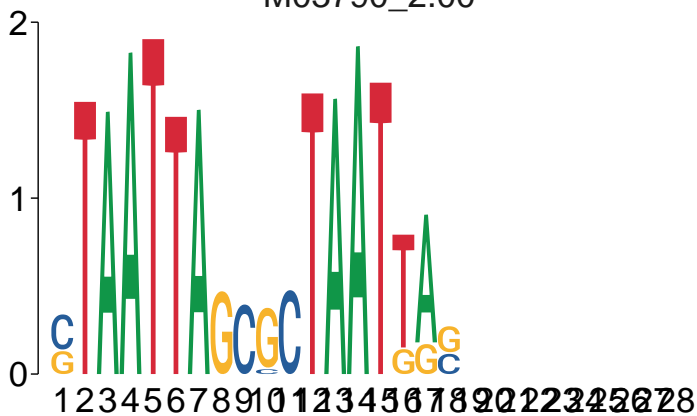
ARCH735_ANTP_HOXA2_ANTP_HOXA3



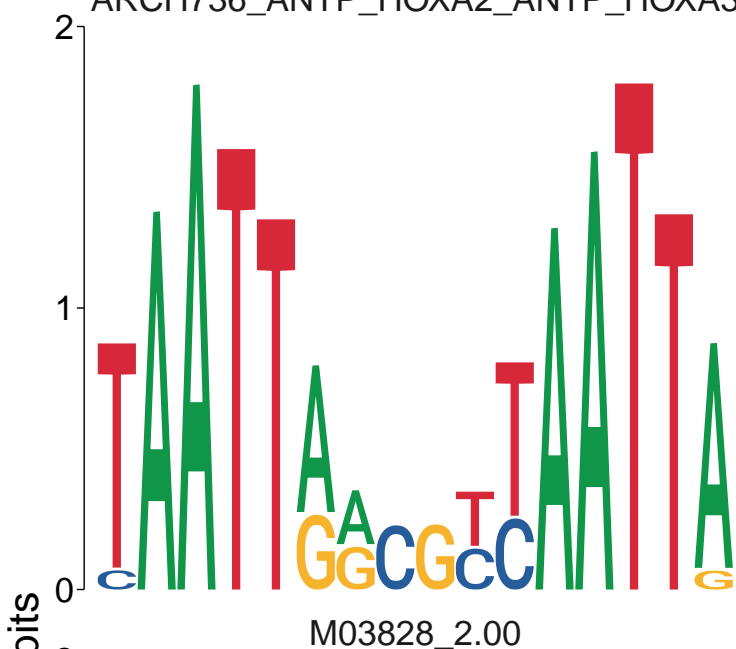
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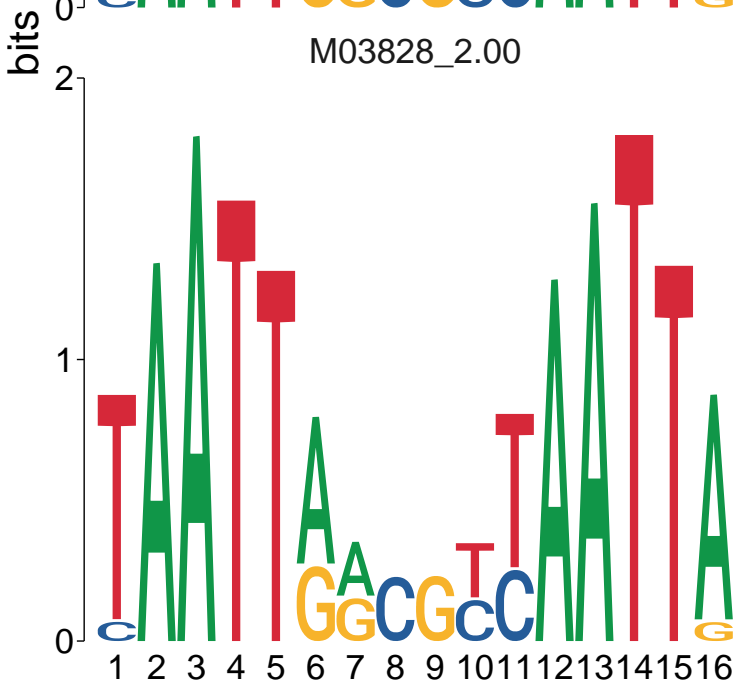
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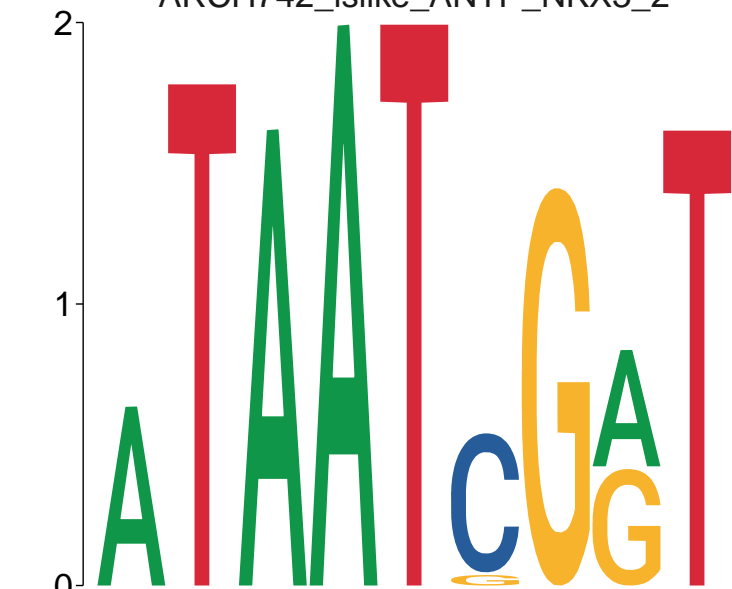
ARCH736_ANTP_HOXA2_ANTP_HOXA3



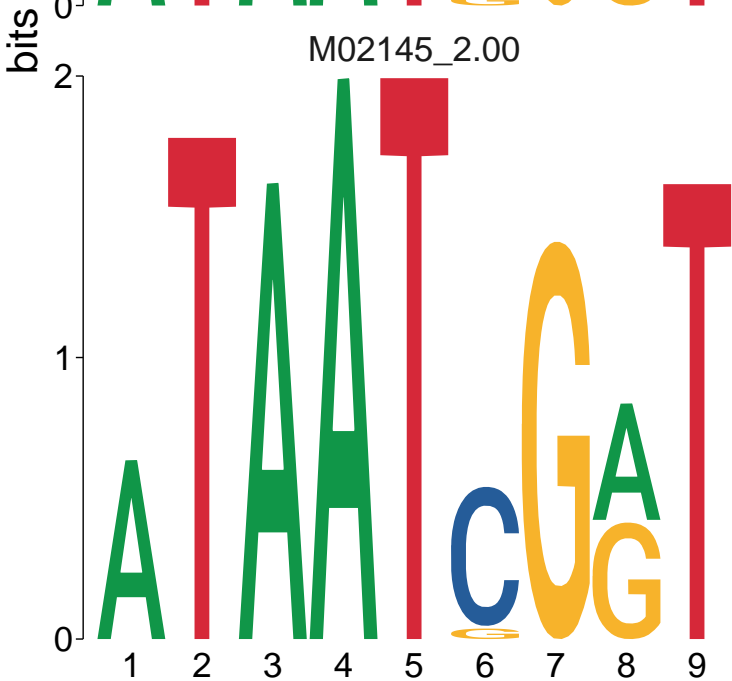
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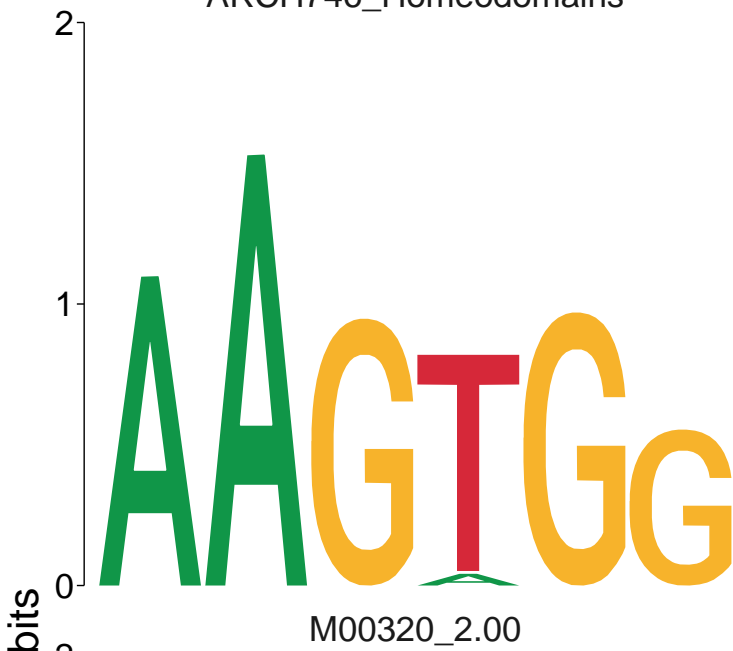
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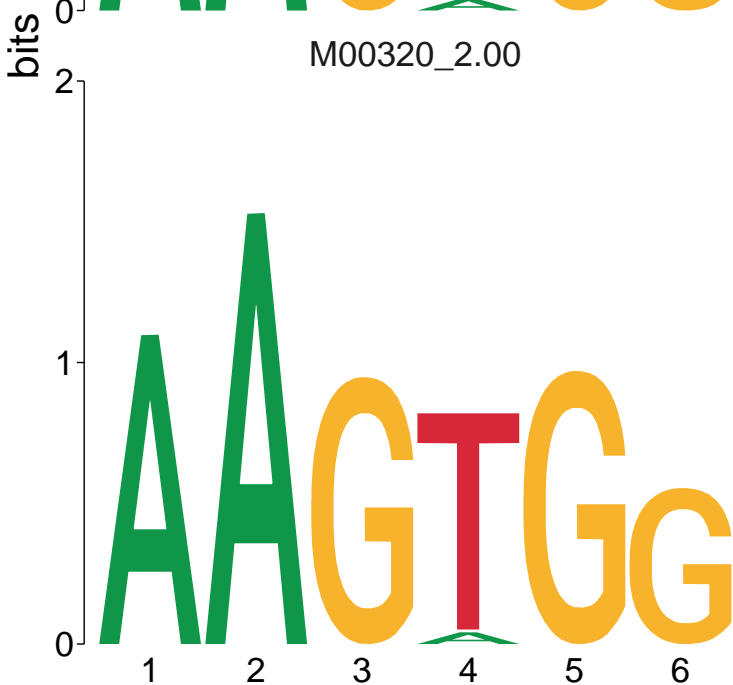
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ARCH746_Homeodomains



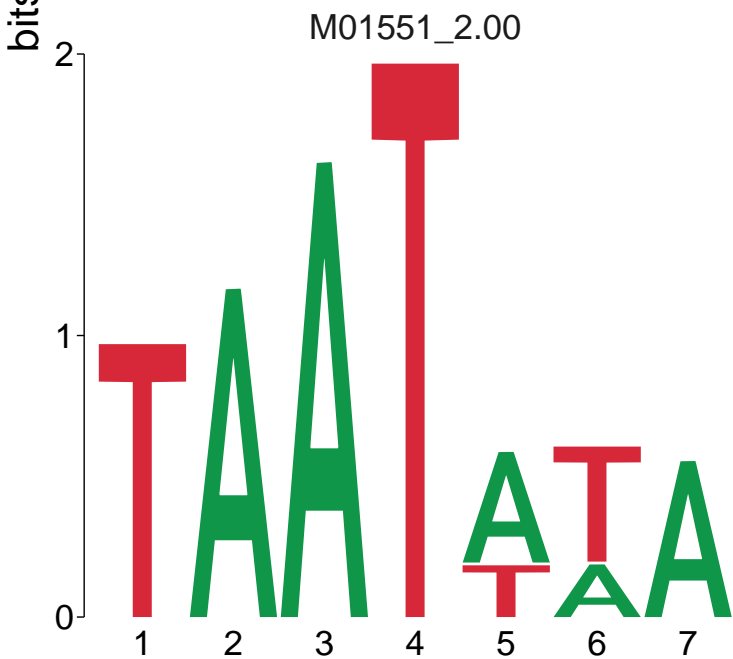
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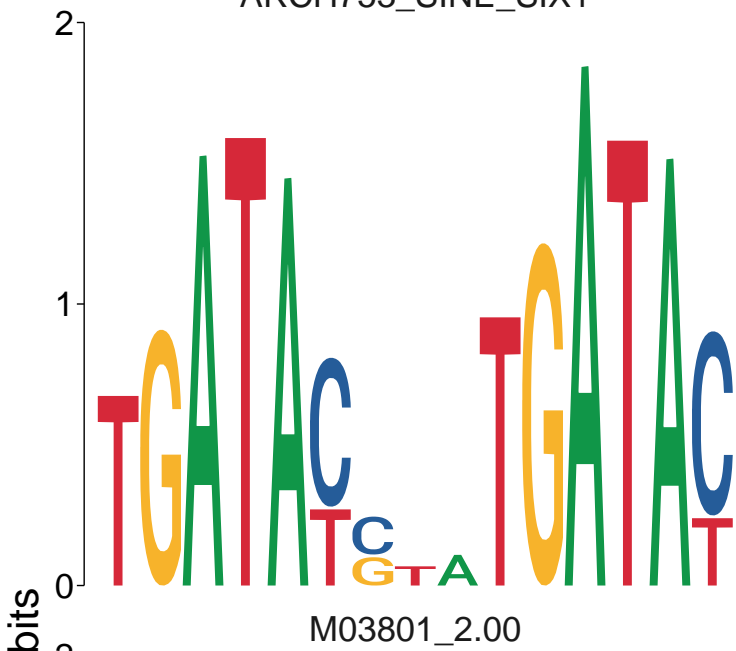
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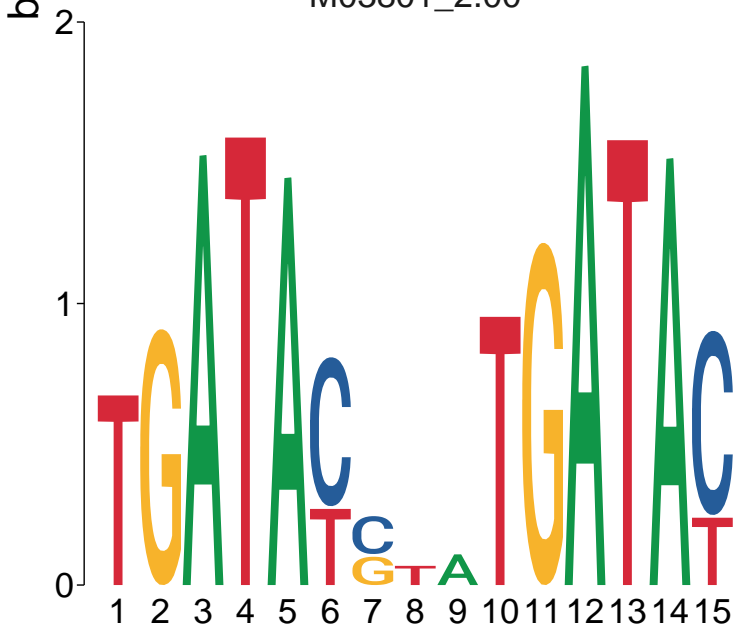
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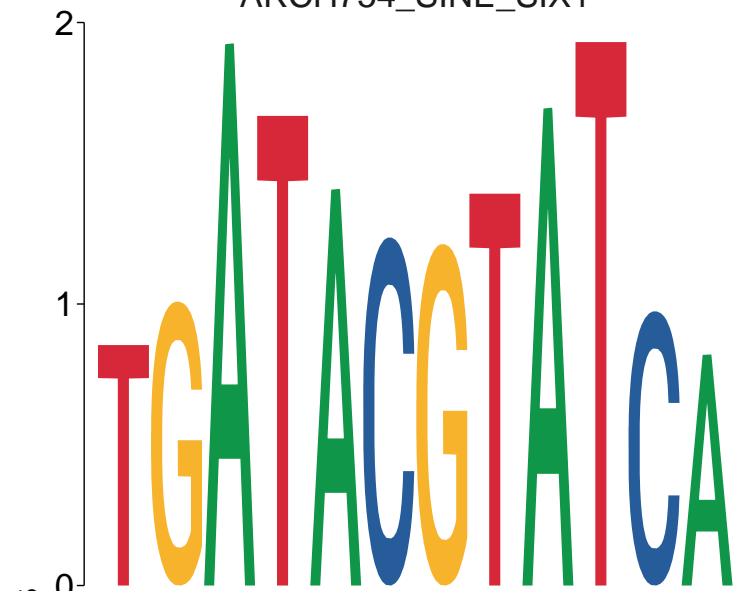
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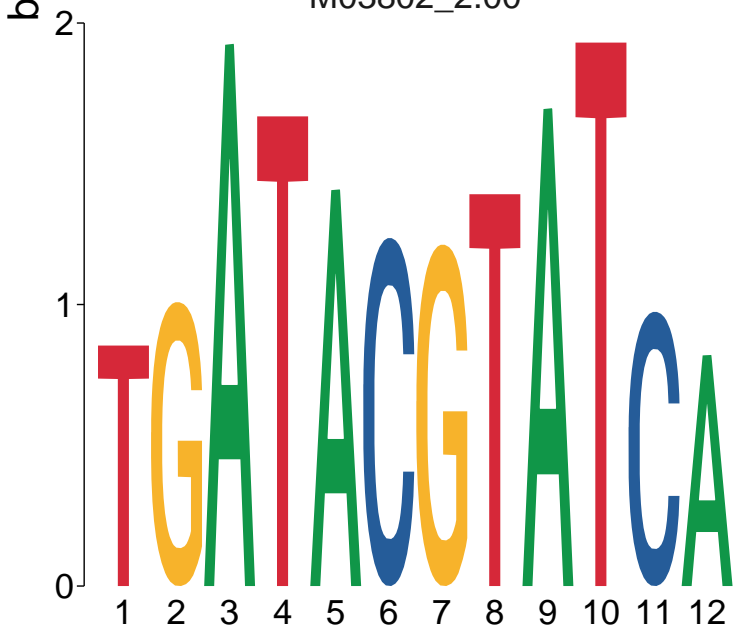
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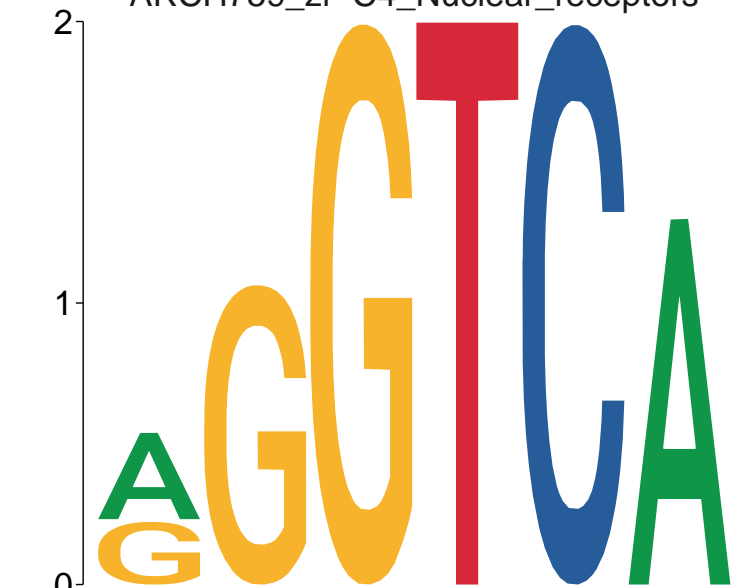
ARCH754_SINE_SIX1



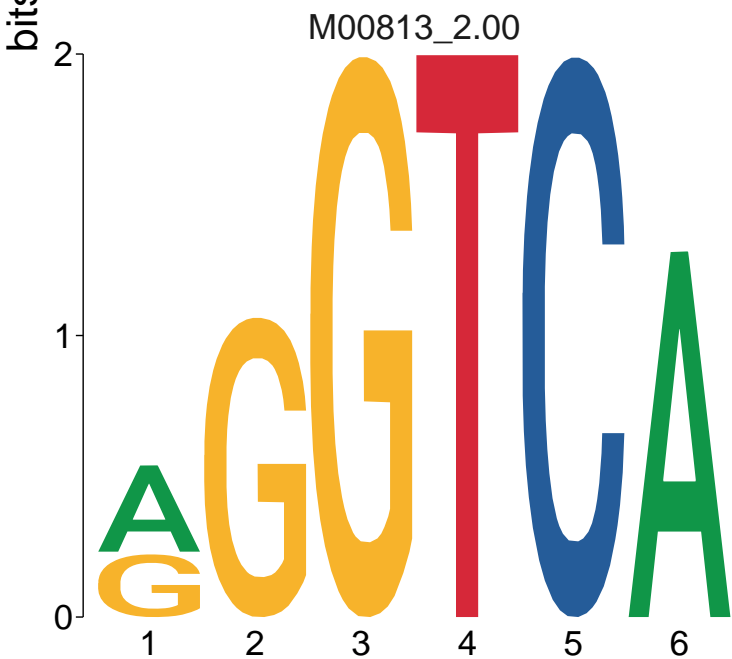
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ARCH759_zf-C4_Nuclear_receptors



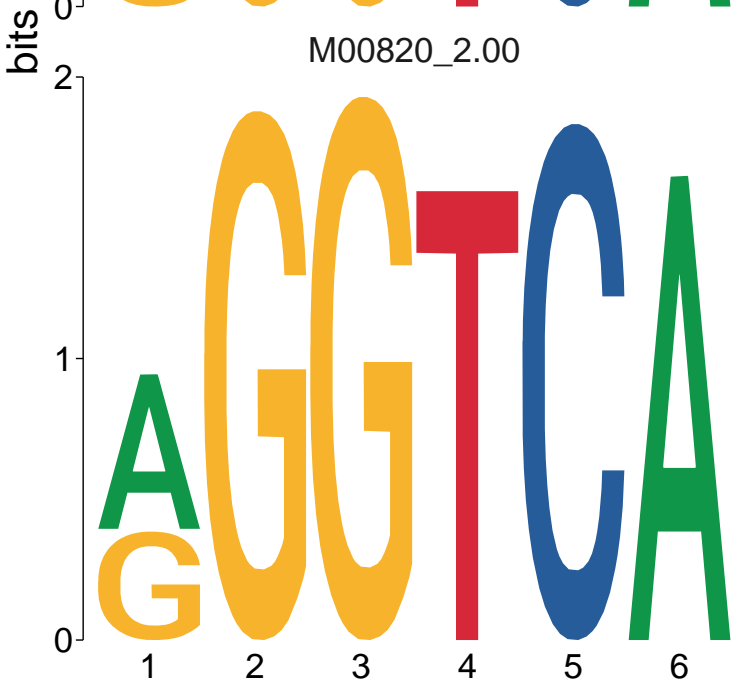
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ARCH760_zf-C4_Nuclear_receptors



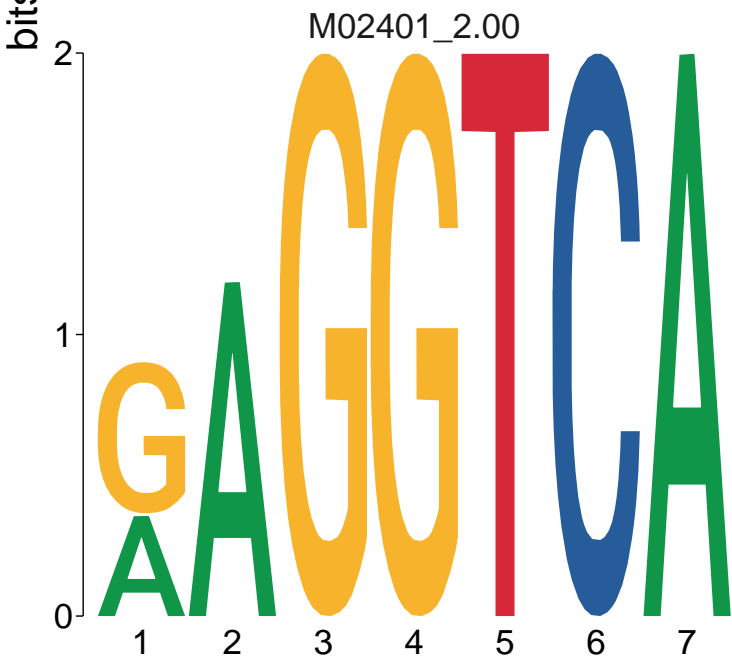
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ARCH761_zf-C4_Nuclear_receptors



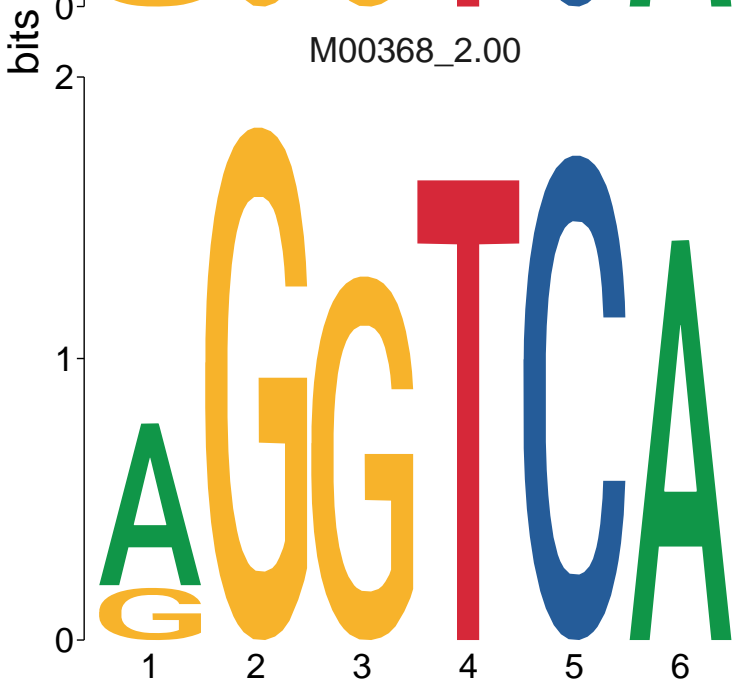
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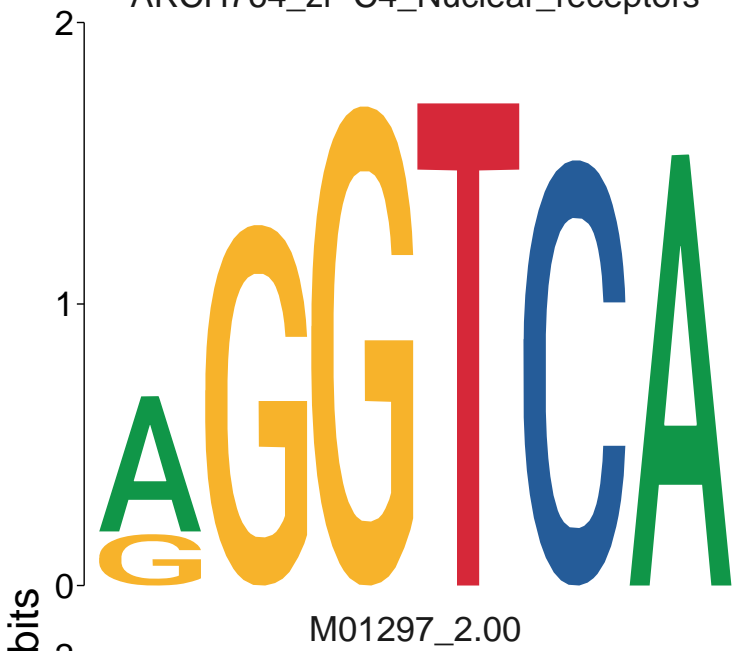
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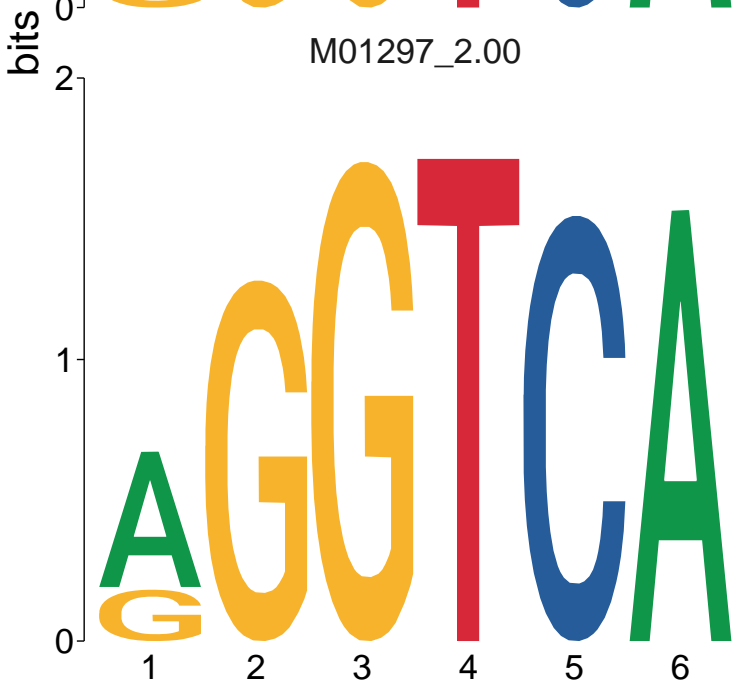
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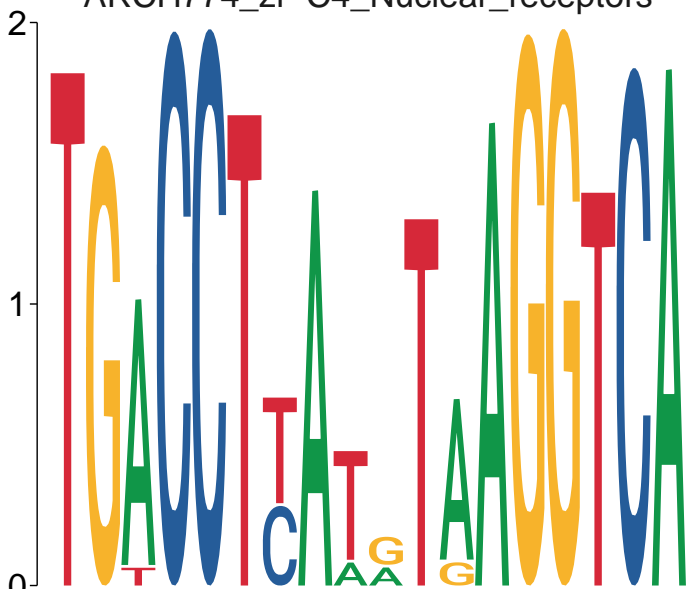
ARCH764_zf-C4_Nuclear_receptors



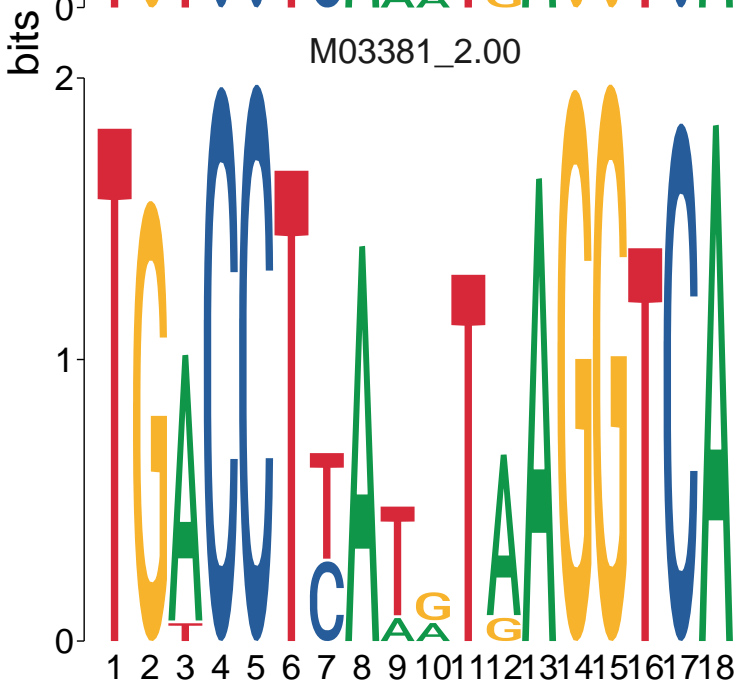
M01297_2.00



ARCH774_zf-C4_Nuclear_receptors



M03381_2.00



ARCH786_zf-C4_Nuclear_receptors



M05632_2.00



M05630_2.00



M05584_2.00



M05582_2.00

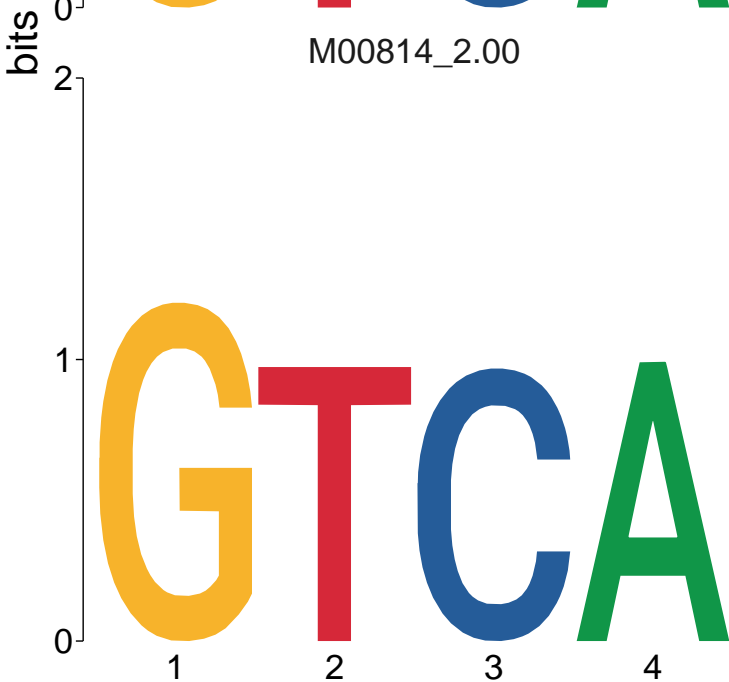


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ARCH791_NR2E1



M00814_2.00



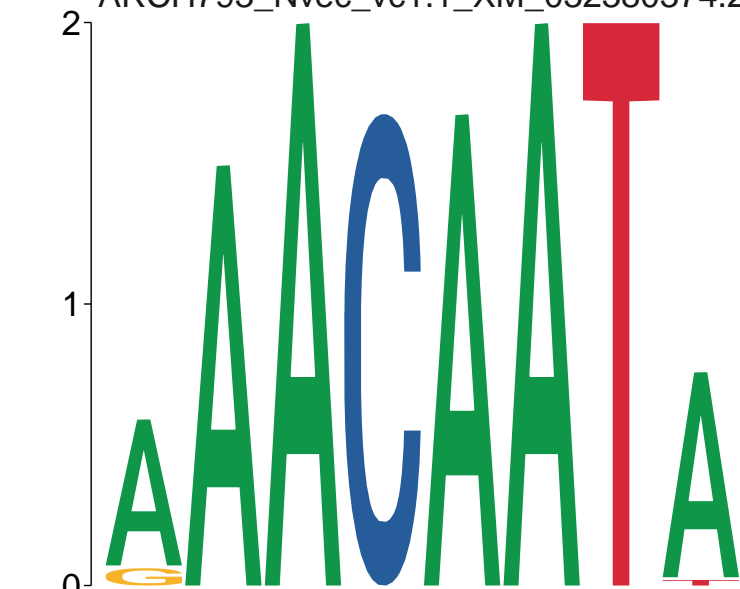
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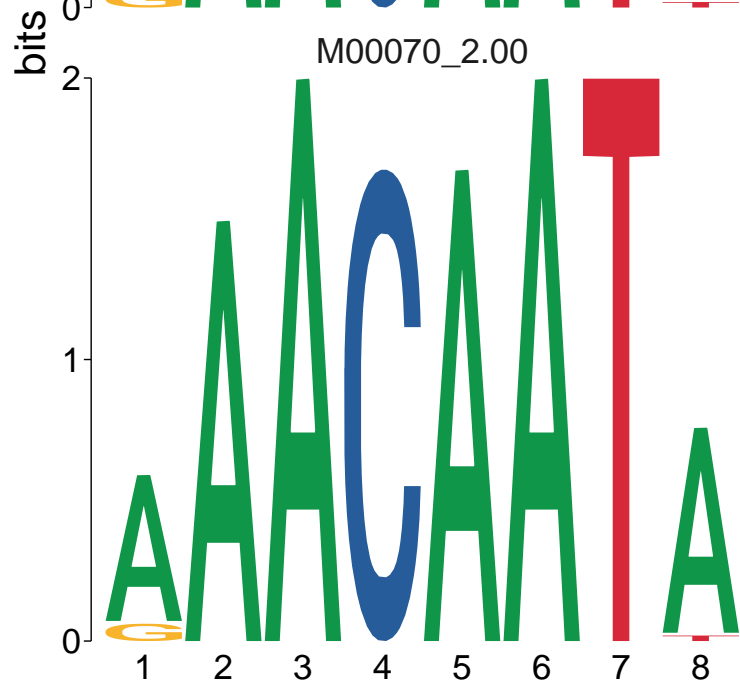
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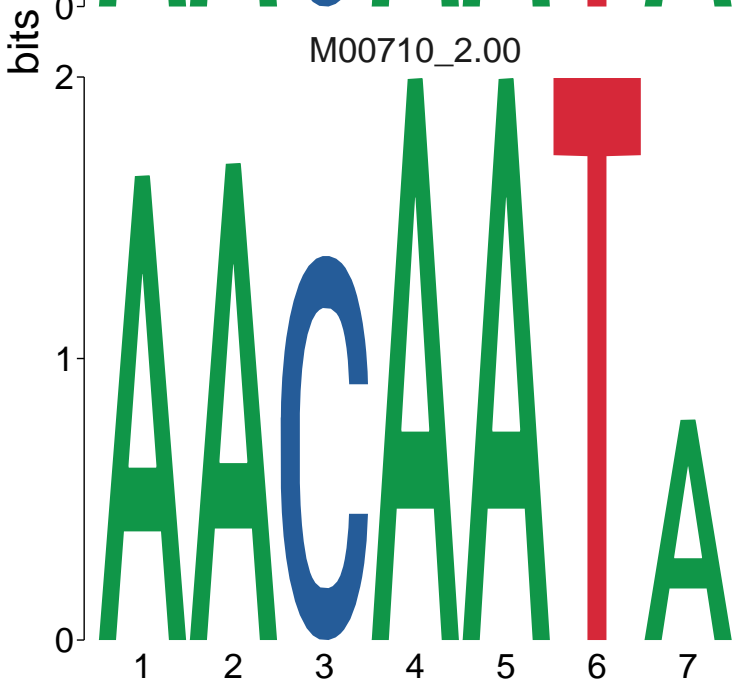
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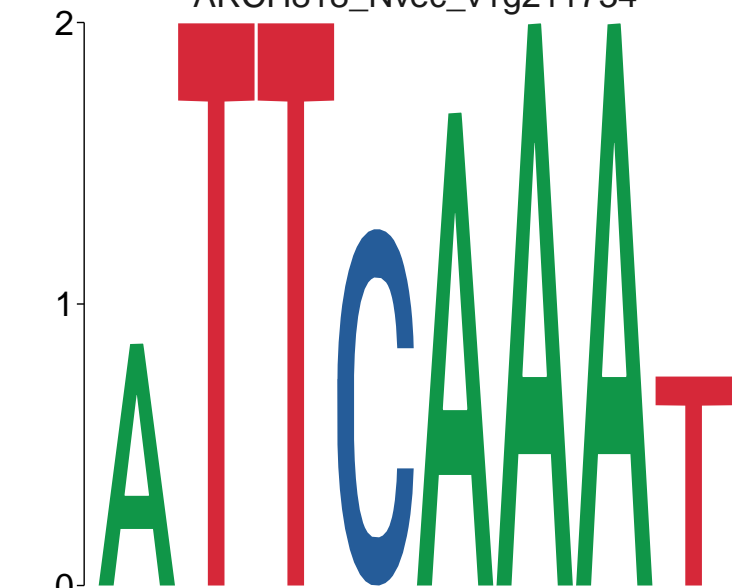
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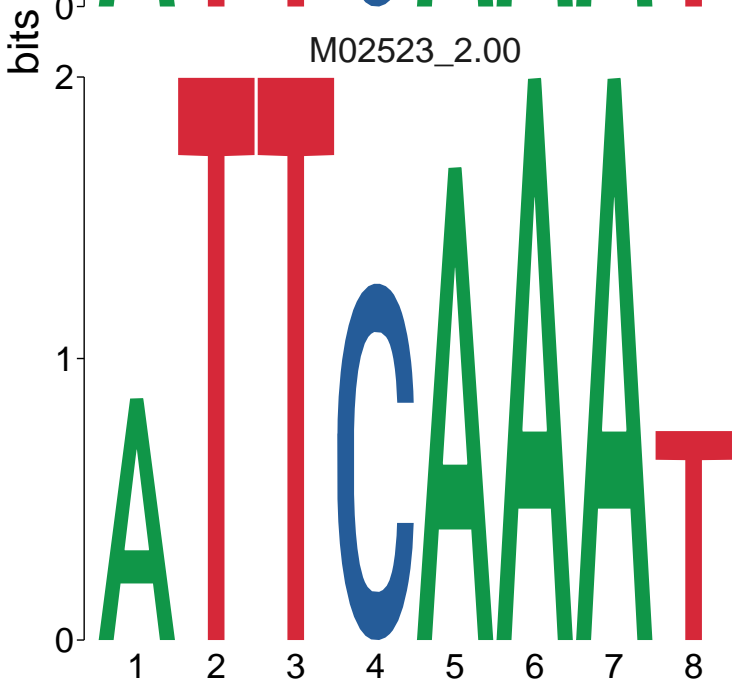
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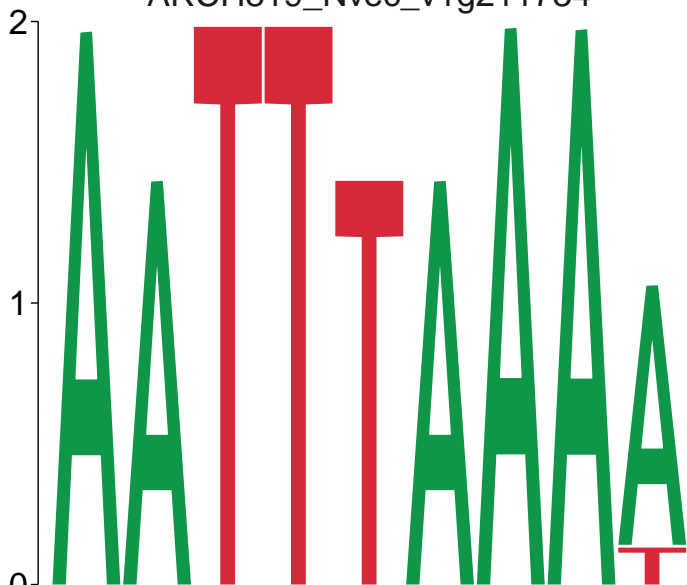
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M02523_2.00



ARCH819_Nvec_v1g211734



M02524_2.00

