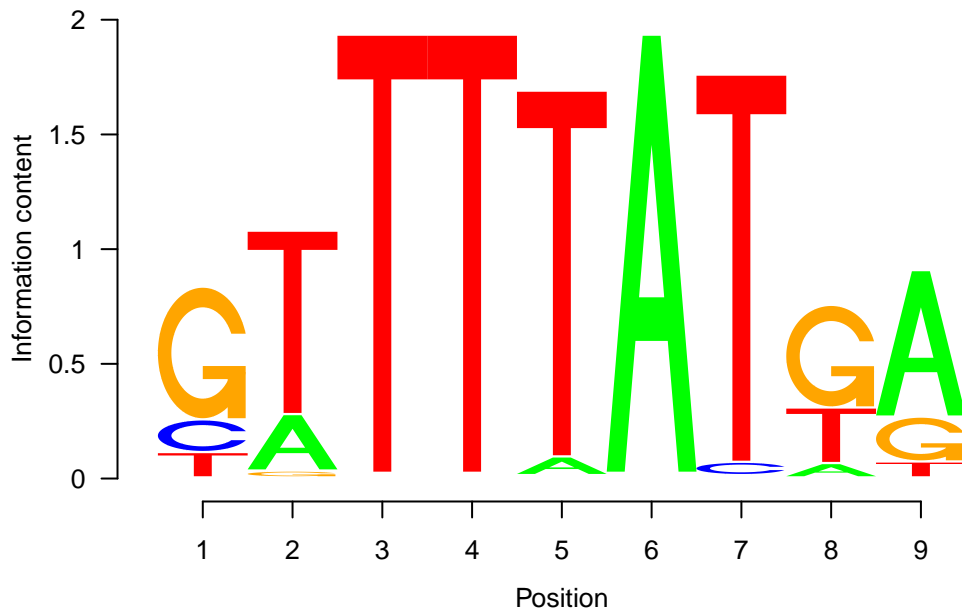
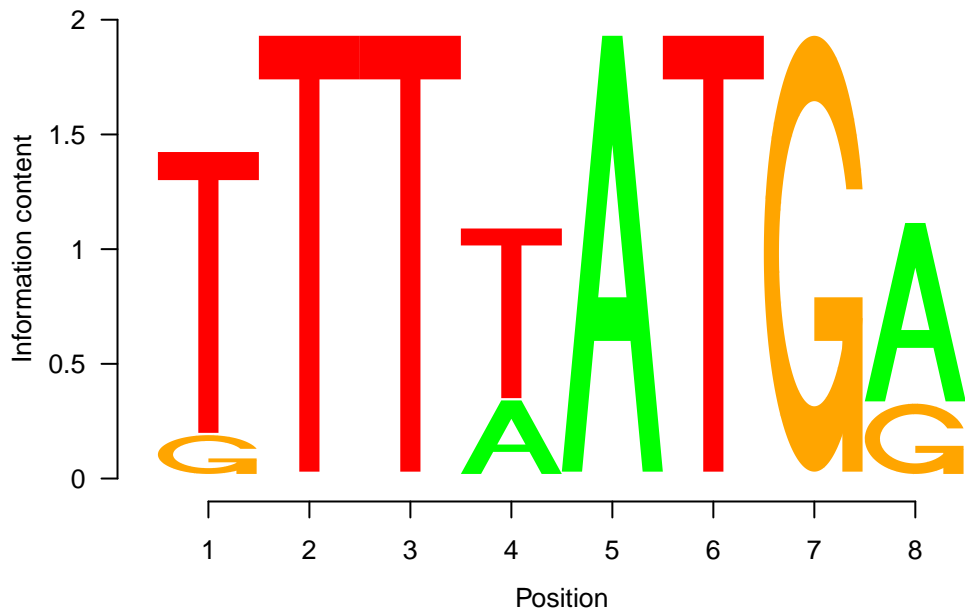


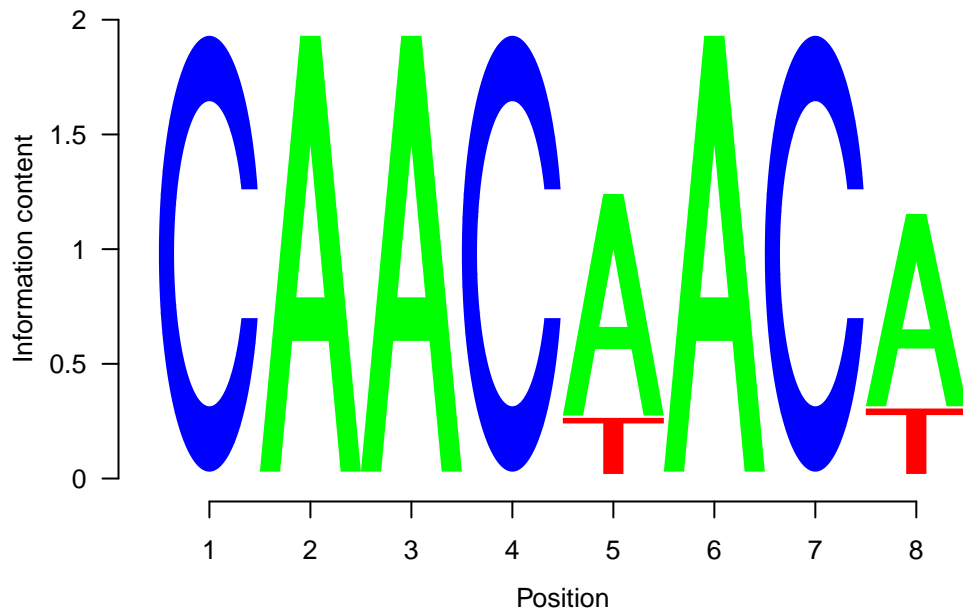
Genome: FlyFactorSurvey-AbdB_Cell_FBgn0000015



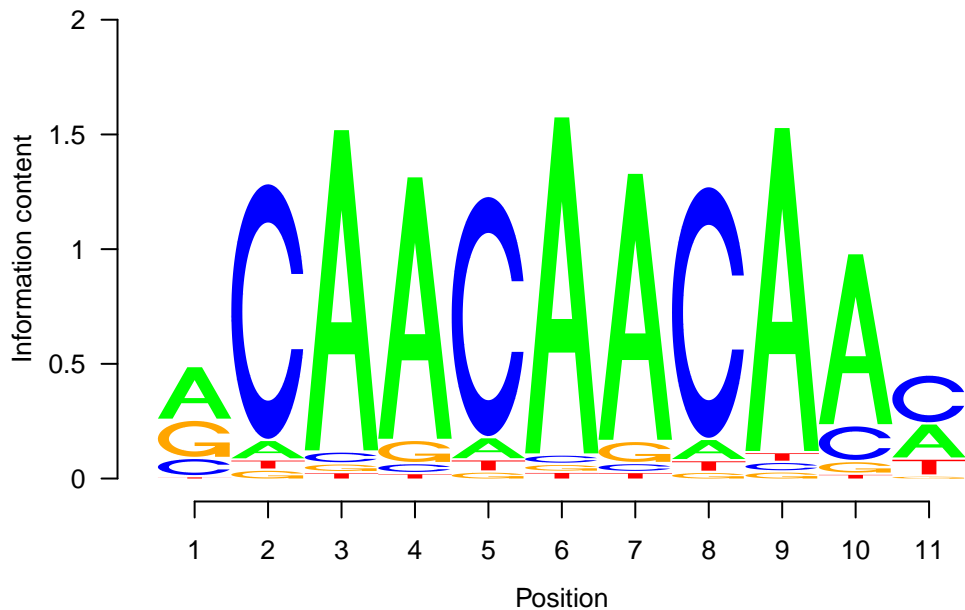
Genome: FlyFactorSurvey-AbdB_SOLEXA_FBgn0000015



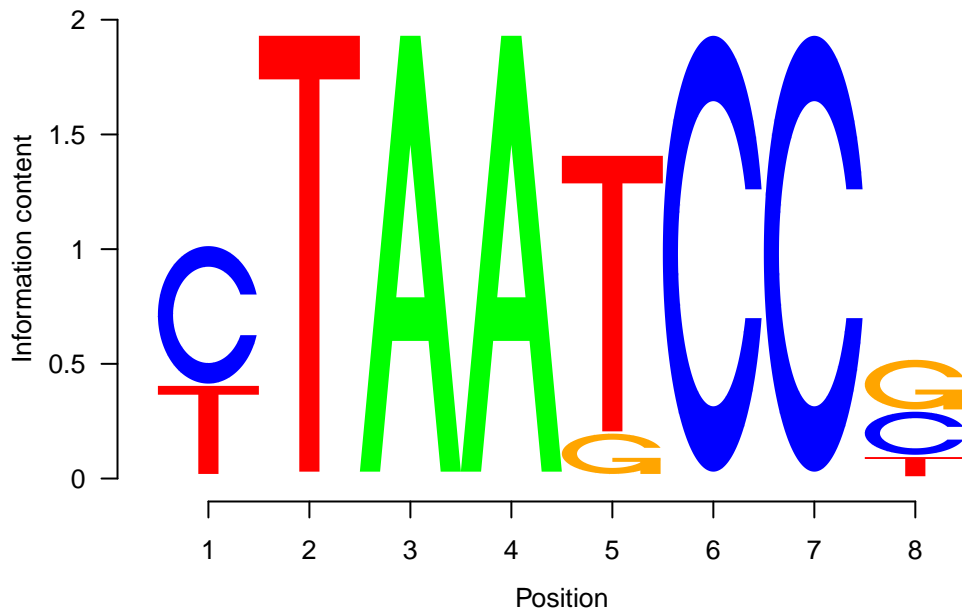
Genome: FlyFactorSurvey-Aef1_FlyReg_FBgn0005694



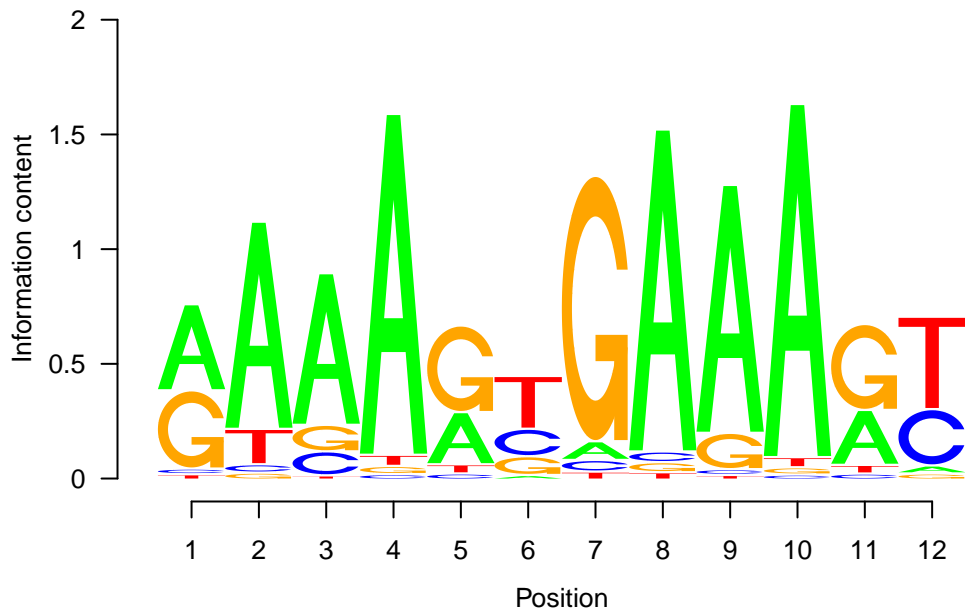
Genome: FlyFactorSurvey-Aef1_SANGER_5_FBgn000569



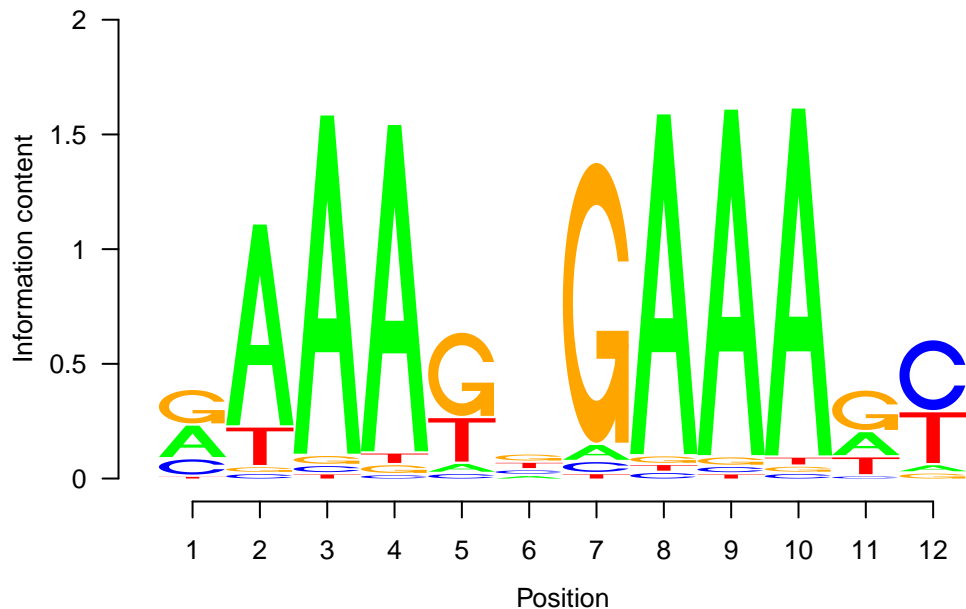
Genome: FlyFactorSurvey-bcd_FlyReg_FBgn0000166



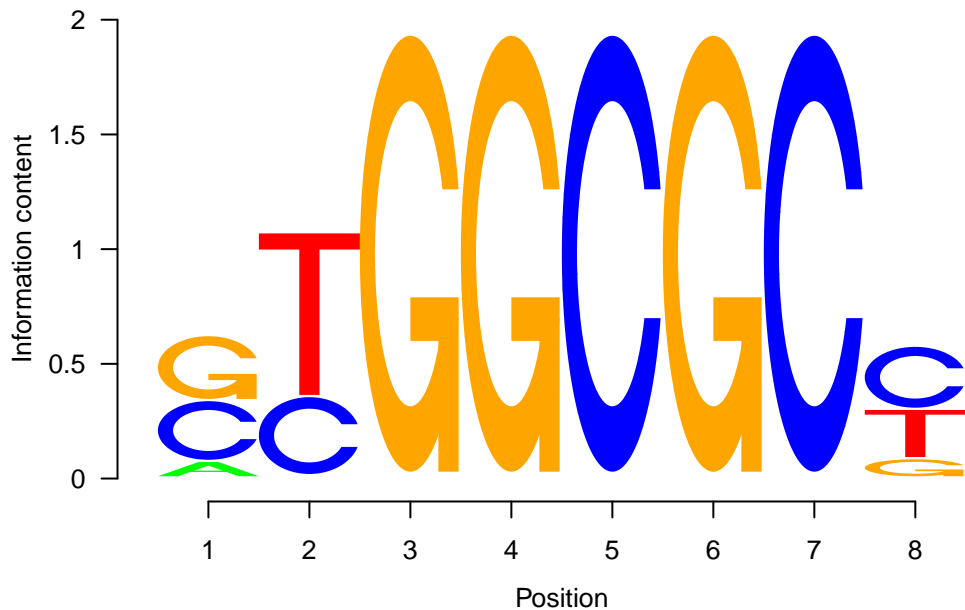
Genome: FlyFactorSurvey-Blimp.1_NAR_FBgn0035625



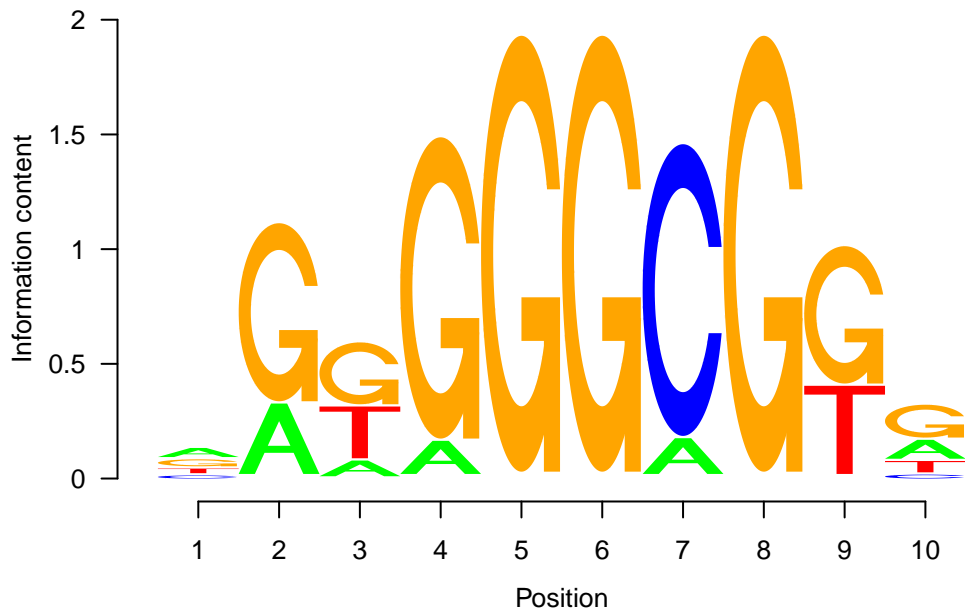
Genome: FlyFactorSurvey-Blimp.1_SANGER_5_FBgn00356



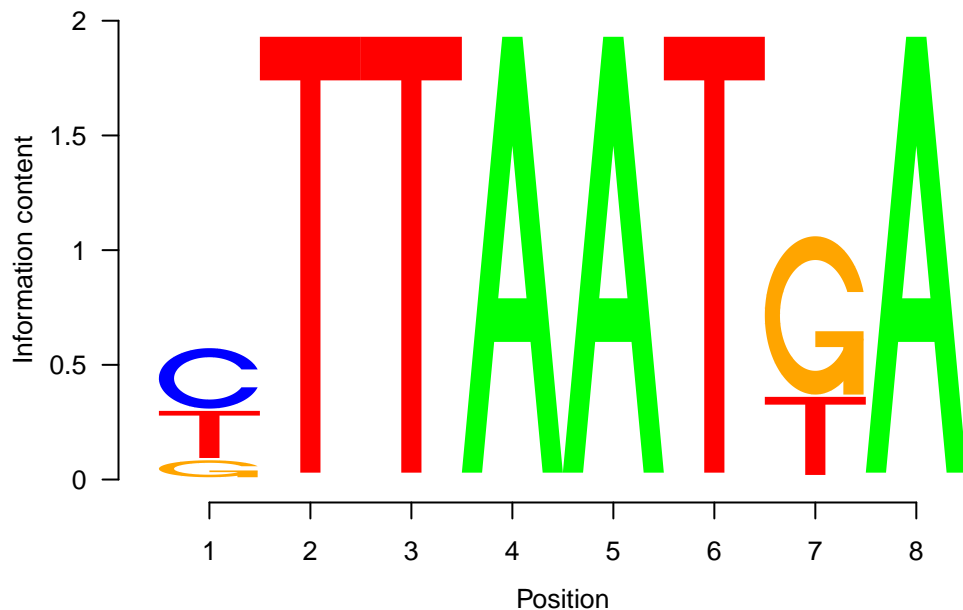
Genome: FlyFactorSurvey-brk_FlyReg_FBgn0024250



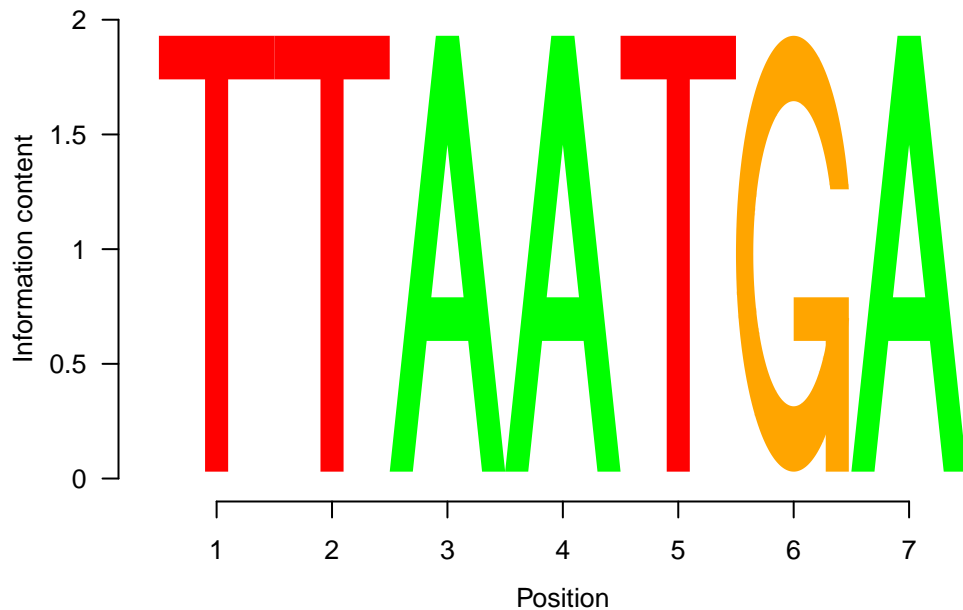
Genome: FlyFactorSurvey-btd_NAR_FBgn0000233



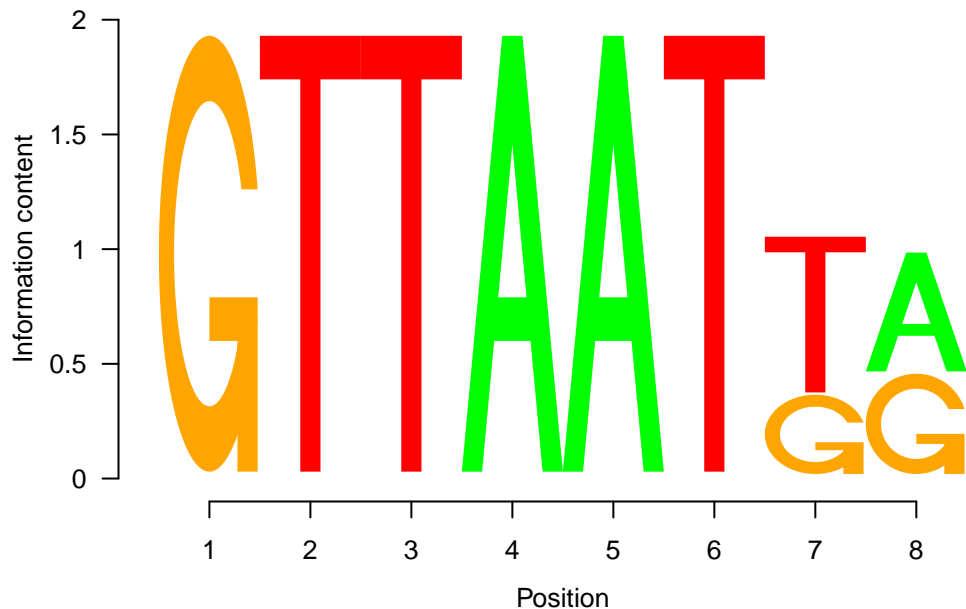
Genome: FlyFactorSurvey-Btn_Cell_FBgn0014949



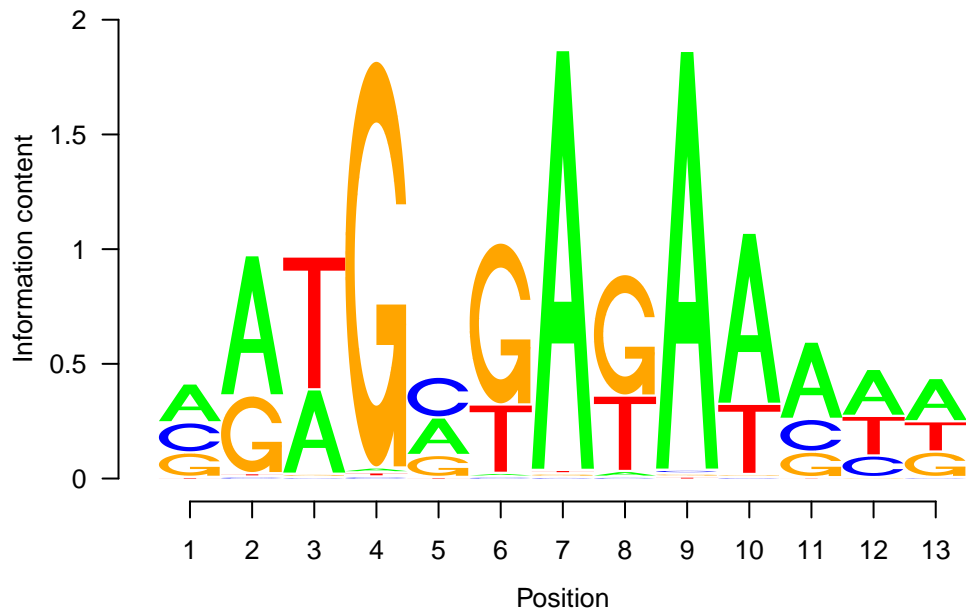
Genome: FlyFactorSurvey-Btn_SOLEXA_FBgn0014949



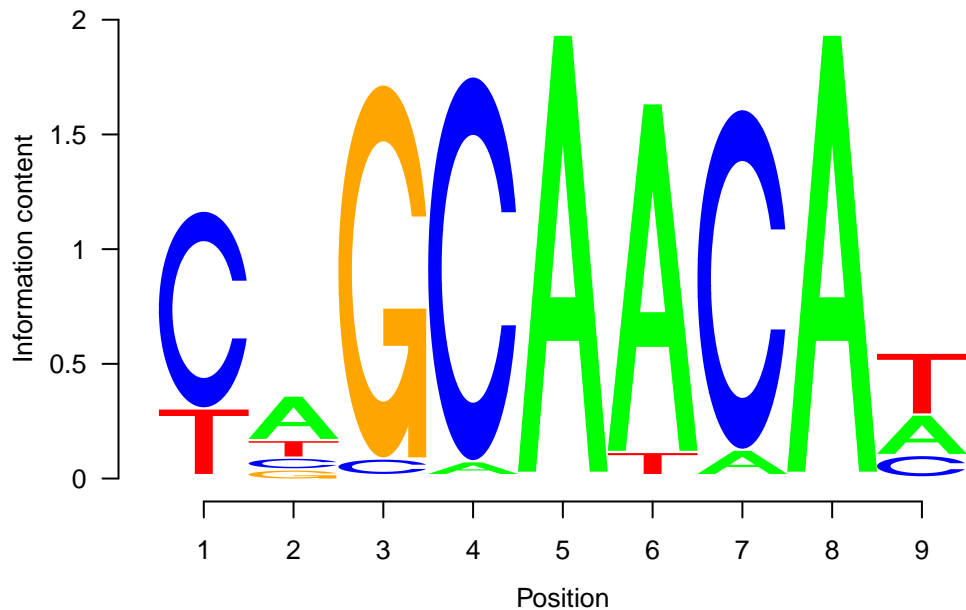
Genome: FlyFactorSurvey-C15_Cell_FBgn0004863



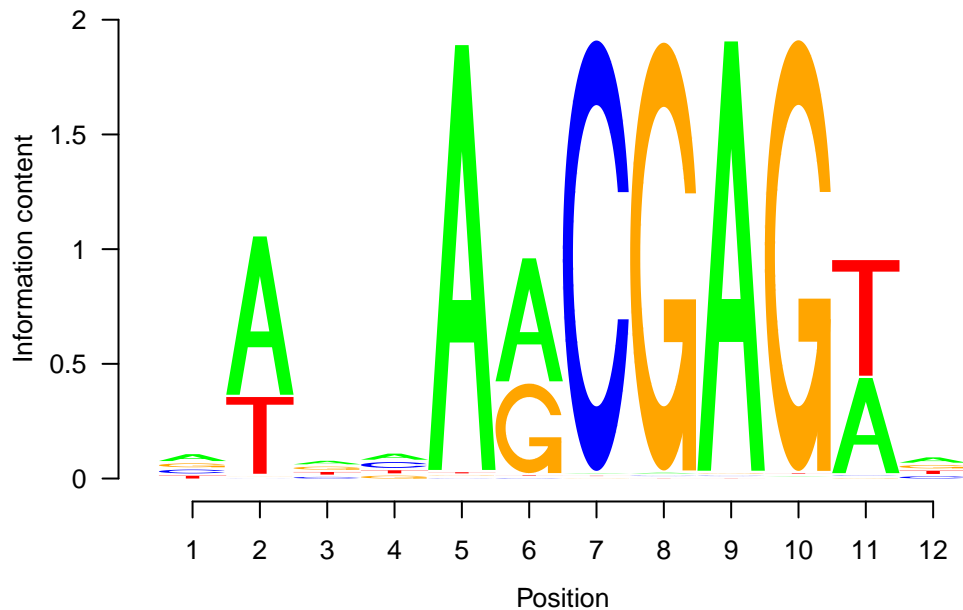
Genome: FlyFactorSurvey-CG10904_SANGER_5_FBgn0034



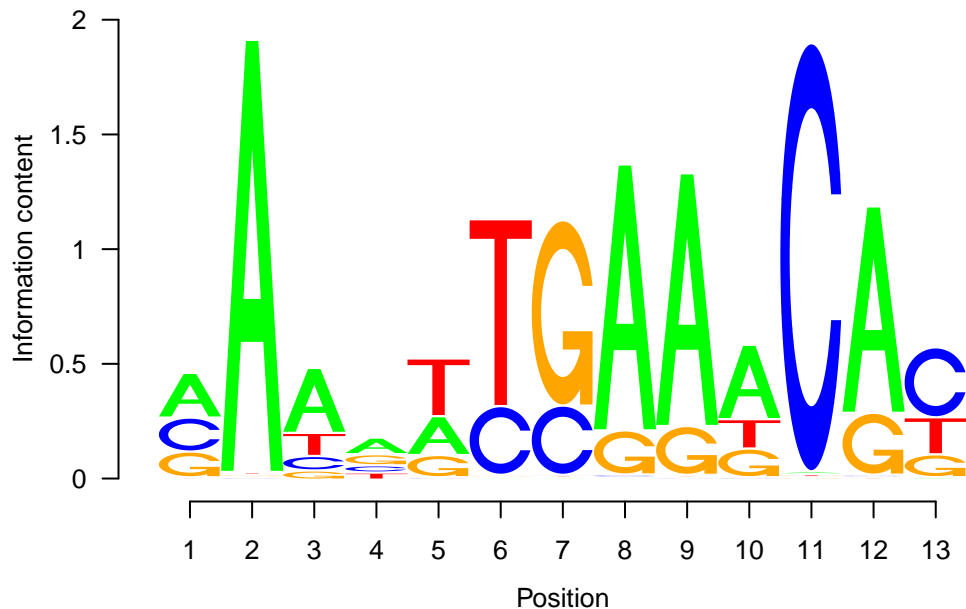
Genome: FlyFactorSurvey-CG11504_SANGER_5_FBgn0039



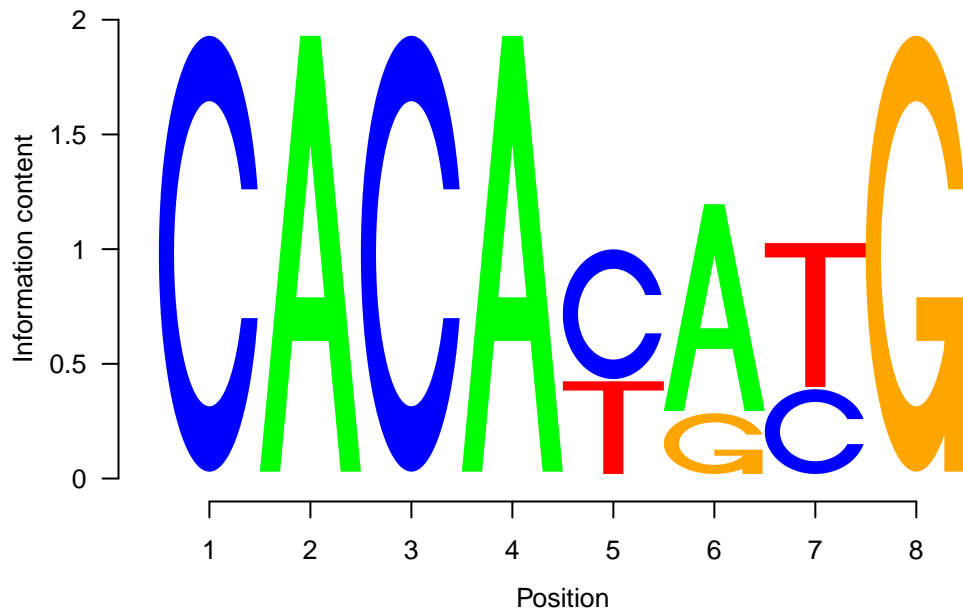
Genome: FlyFactorSurvey-CG12155_SANGER_5_FBgn0029



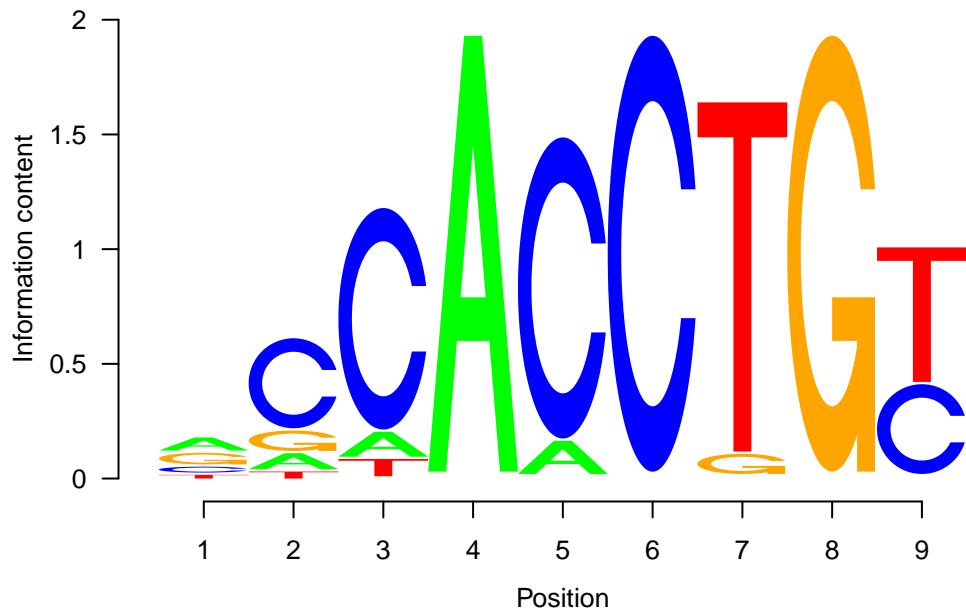
Genome: FlyFactorSurvey-CG14962_SANGER_5_FBgn0035



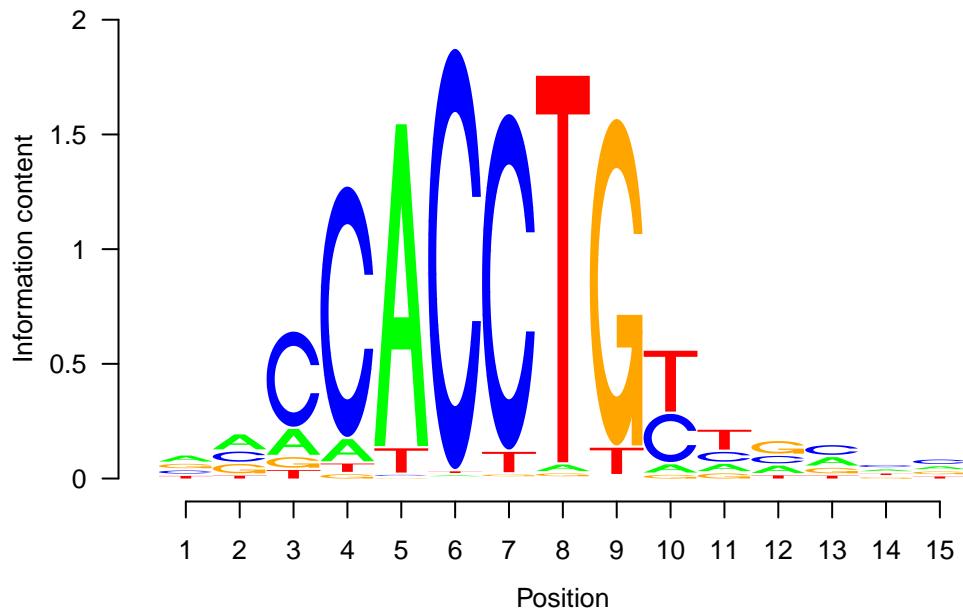
Genome: FlyFactorSurvey-CG16778_SANGER_5_FBgn0003



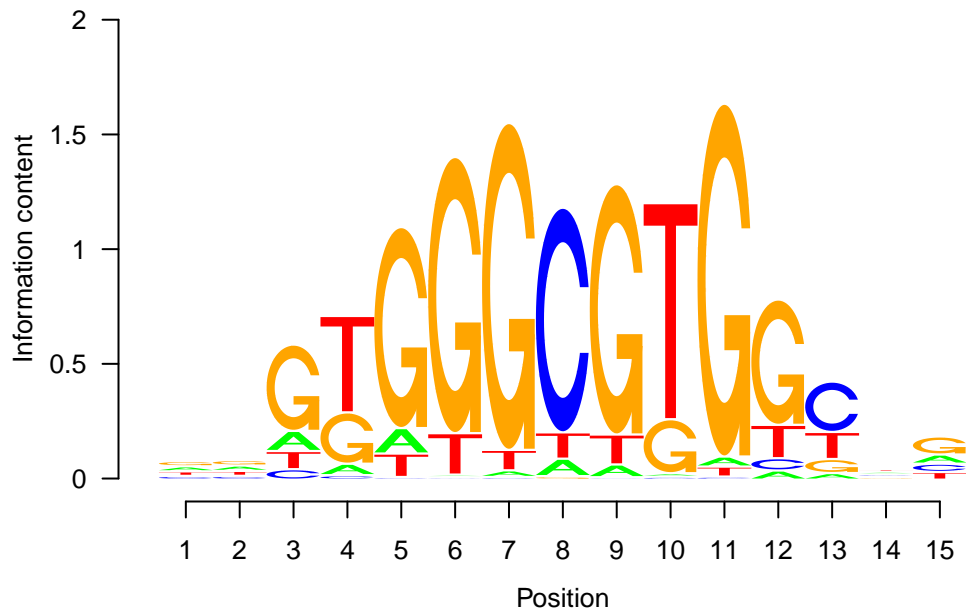
Genome: FlyFactorSurvey-CG17181_SANGER_5_FBgn0035



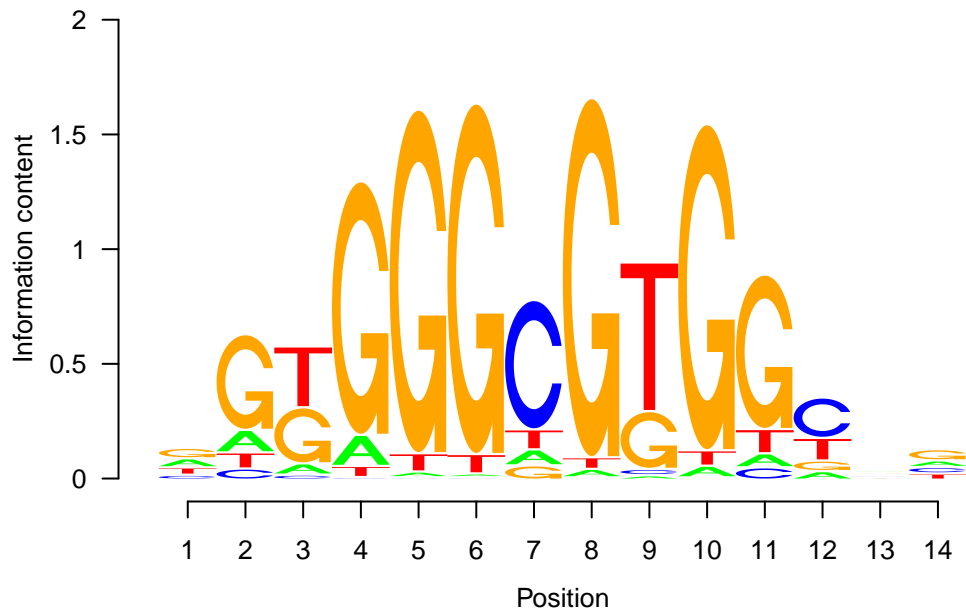
Genome: FlyFactorSurvey-CG17181_SOLEXA_5_FBgn00351



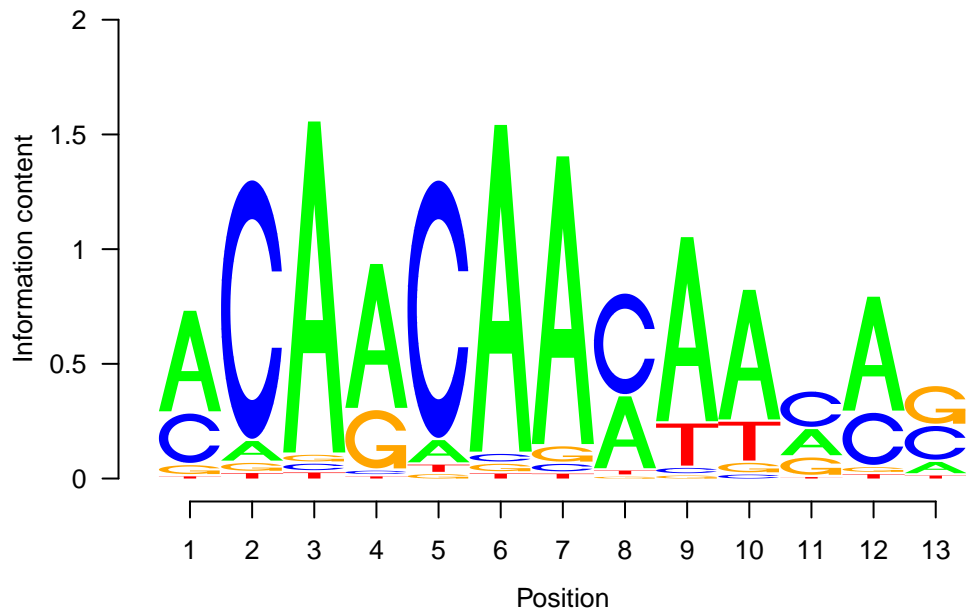
Genome: FlyFactorSurvey-CG3065_F1.3_SOLEXA_2.5_FBgn00



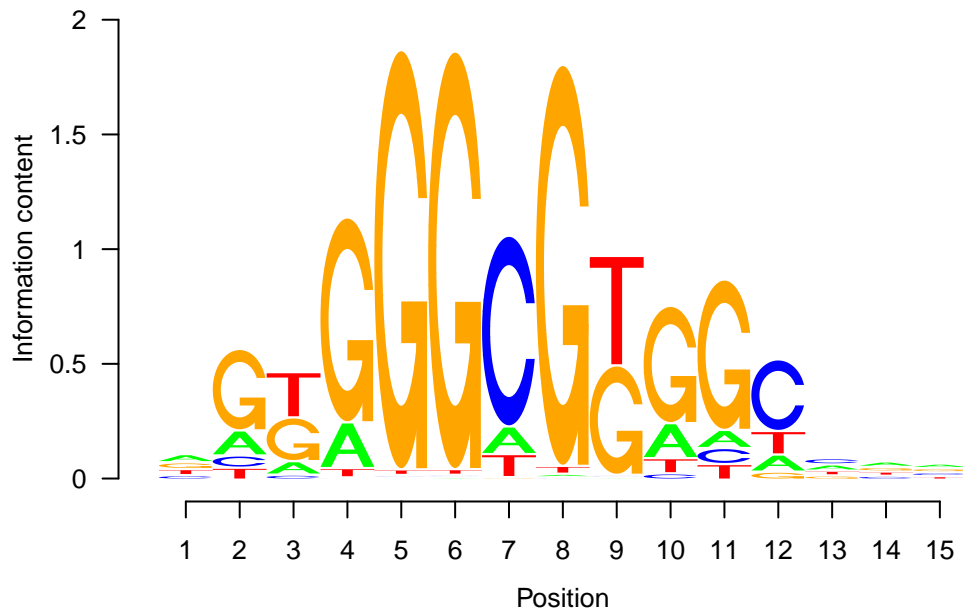
Genome: FlyFactorSurvey-CG3065_F1.5_SOLEXA_2.5_FBgn00



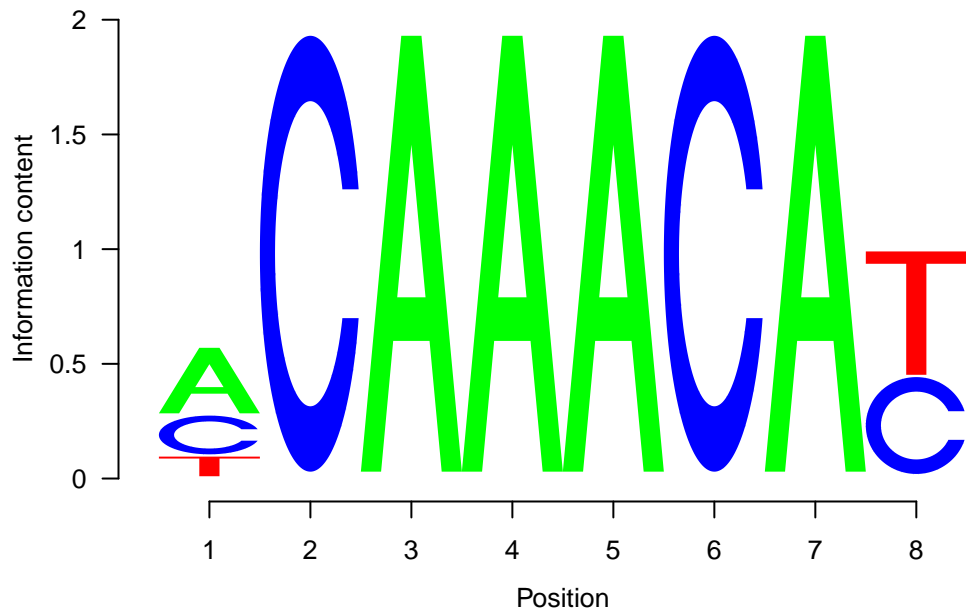
Genome: FlyFactorSurvey-CG4360_F1.3_SANGER_2.5_FBgn00



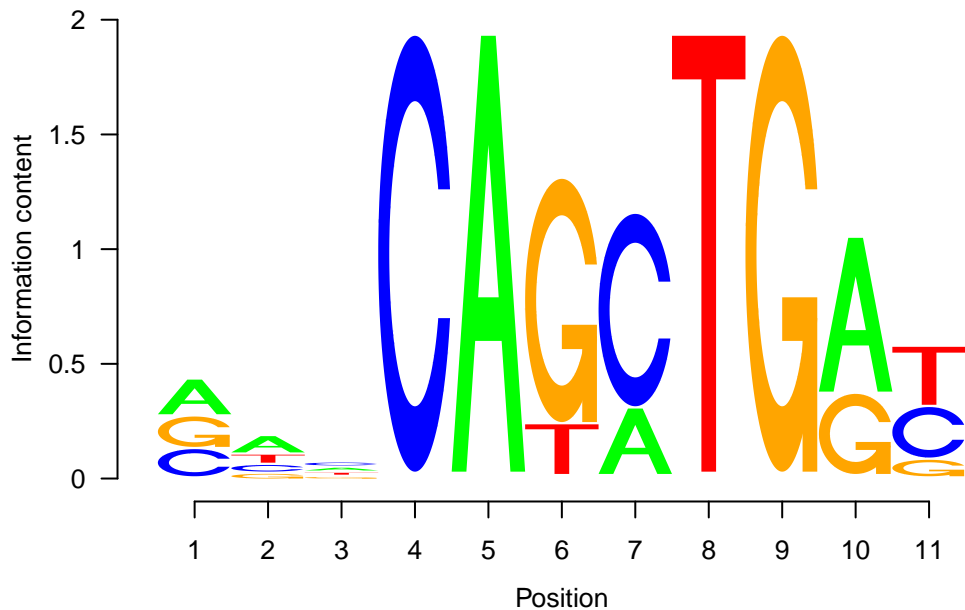
Genome: FlyFactorSurvey-CG5669_SOLEXA_5_FBgn00391



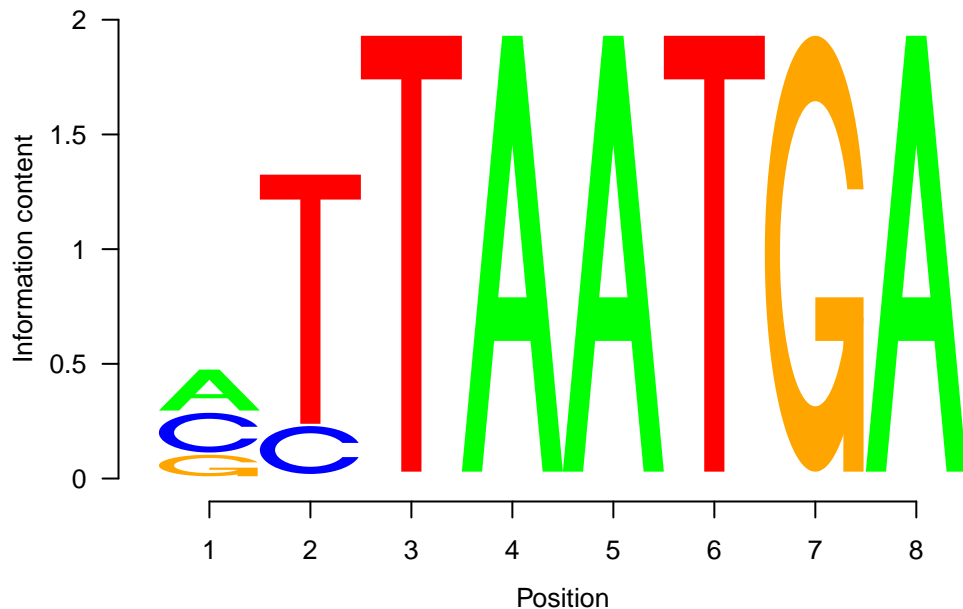
Genome: FlyFactorSurvey-CG5953_SANGER_5_FBgn00325



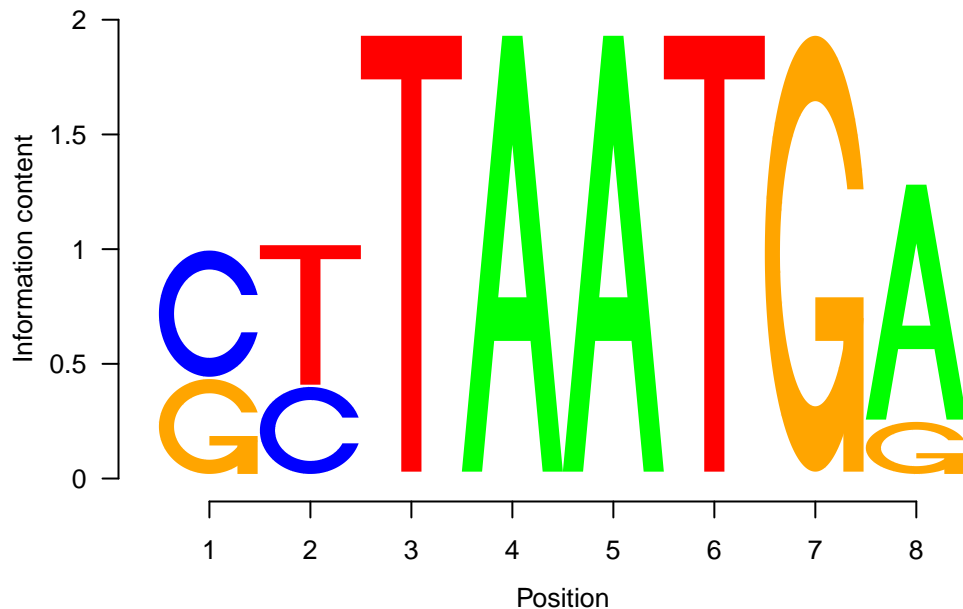
Genome: FlyFactorSurvey-crp_SANGER_10_FBgn0001994



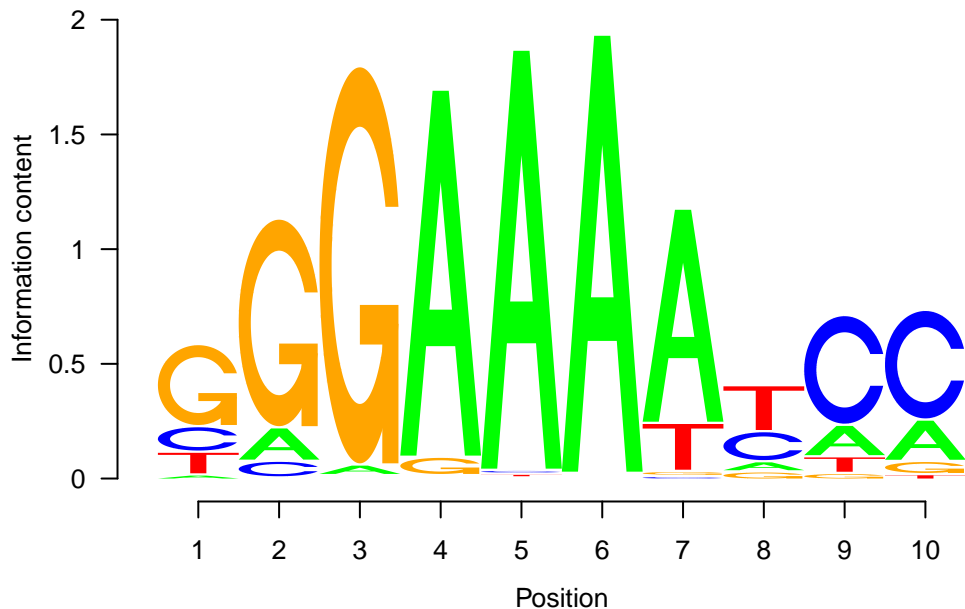
Genome: FlyFactorSurvey-Dfd_Cell_FBgn0000439



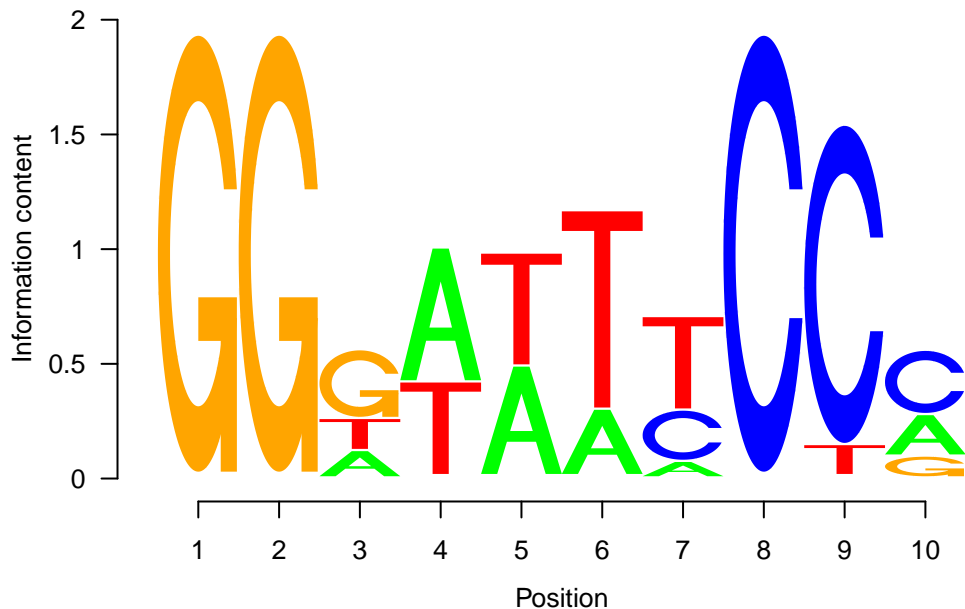
Genome: FlyFactorSurvey-Dfd_SOLEXA_FBgn0000439



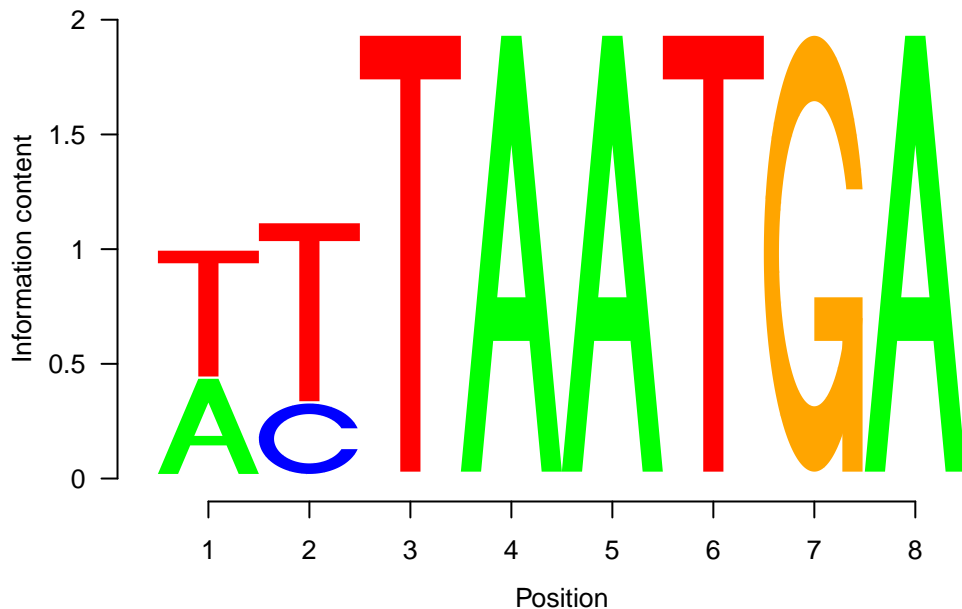
Genome: FlyFactorSurvey-dl_FlyReg_FBgn0000462



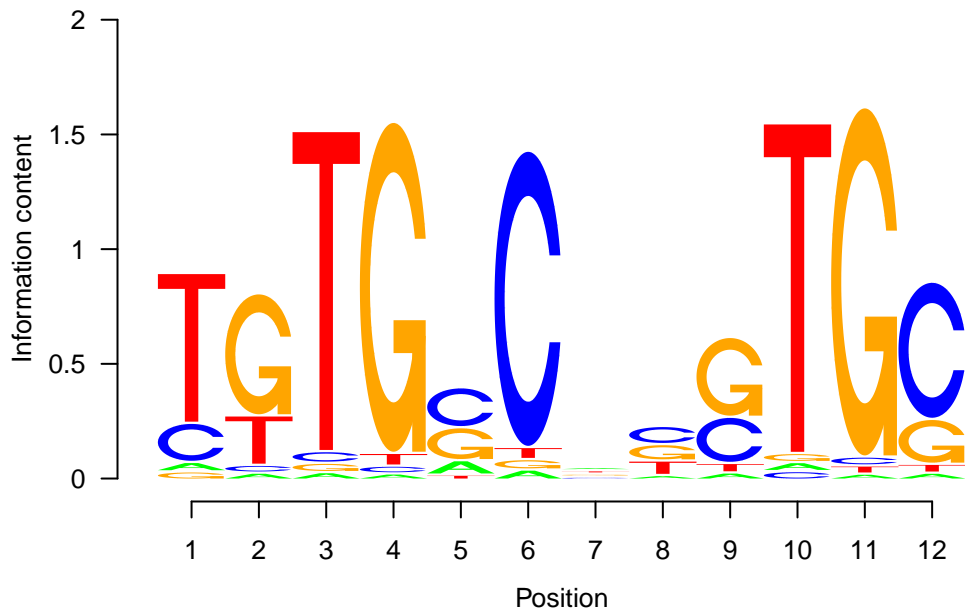
Genome: FlyFactorSurvey-dl_NBT_FBgn0000462



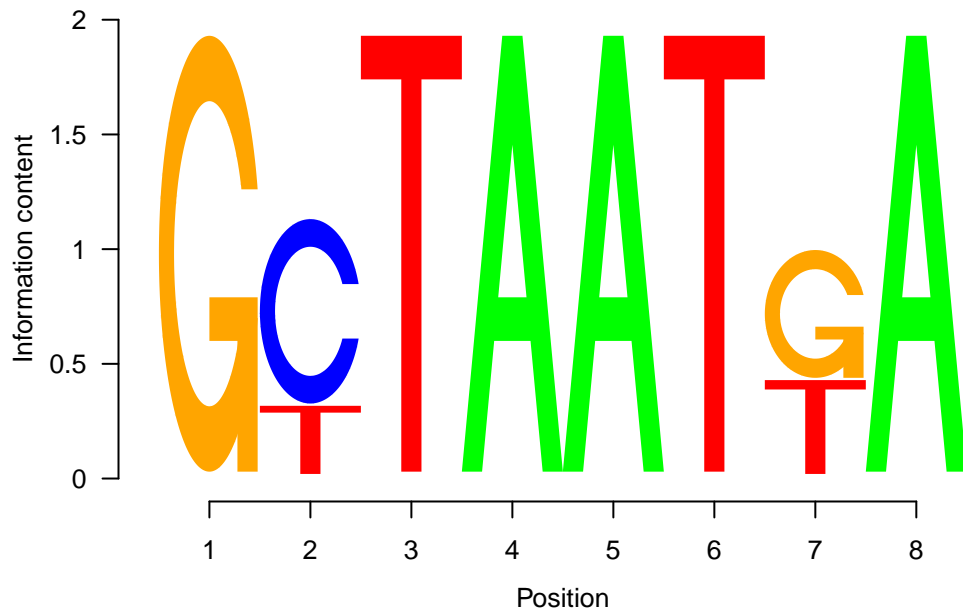
Genome: FlyFactorSurvey-Ems_Cell_FBgn0000576



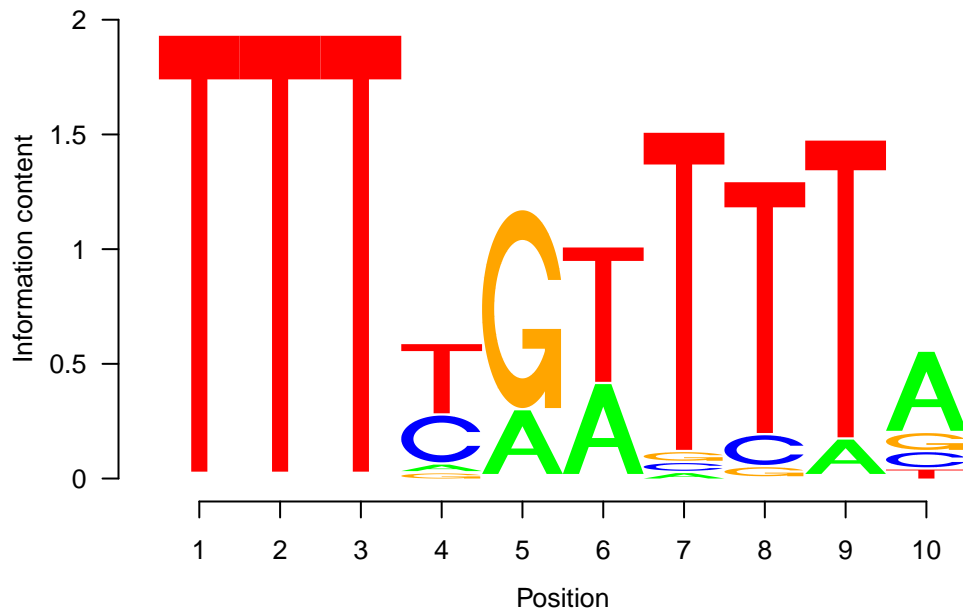
Genome: FlyFactorSurvey-Espl_FlyReg_FBgn0000591



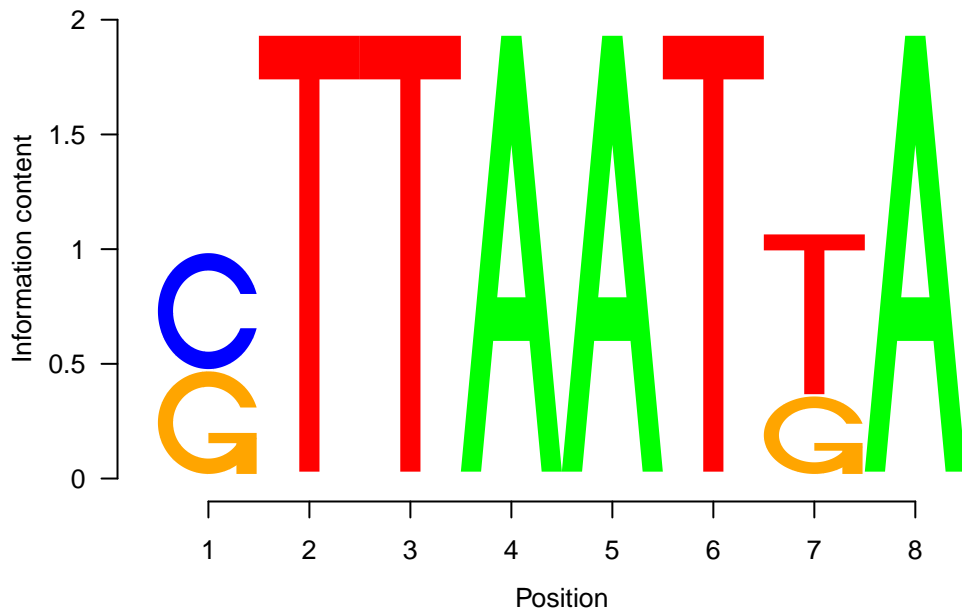
Genome: FlyFactorSurvey-Eve_Cell_FBgn0000606



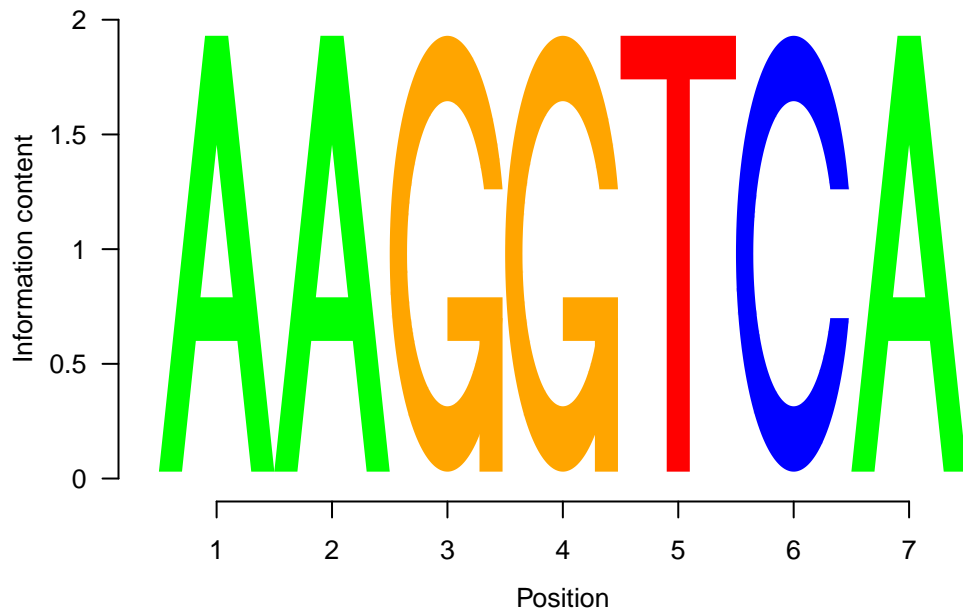
Genome: FlyFactorSurvey-exd_FlyReg_FBgn0000611



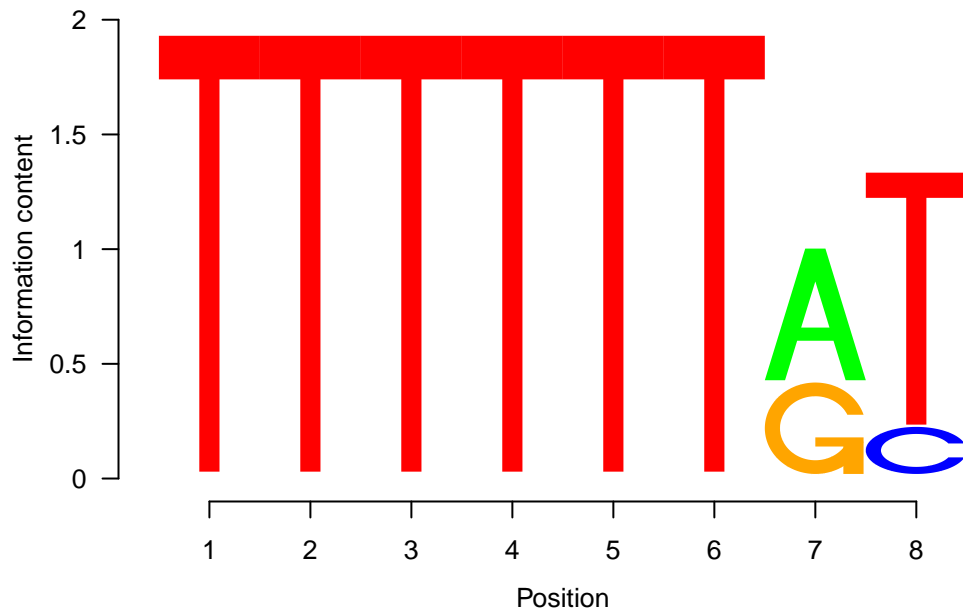
Genome: FlyFactorSurvey-Ftz_Cell_FBgn0001077



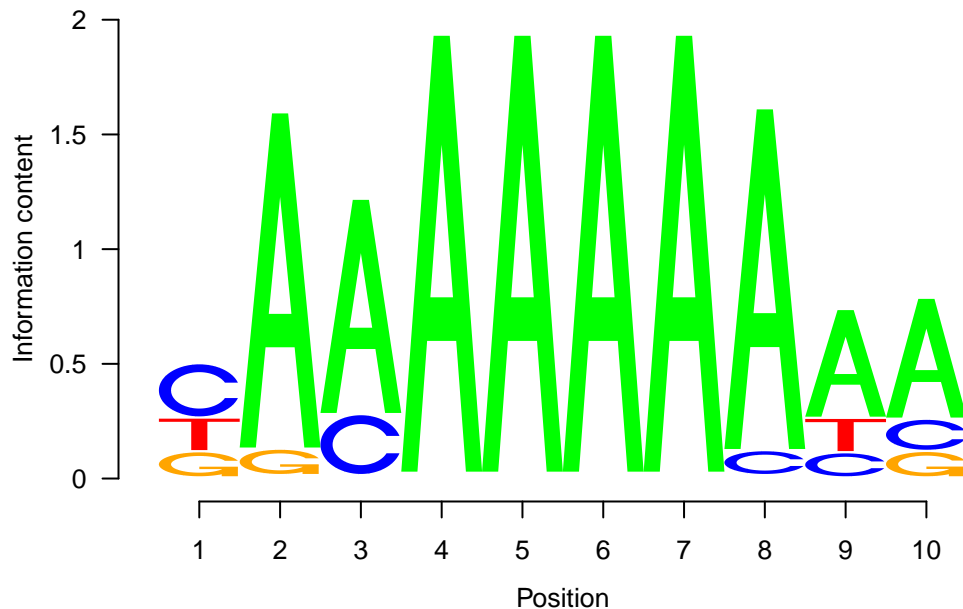
Genome: FlyFactorSurvey-ftz.f1_SANGER_5_FBgn000107



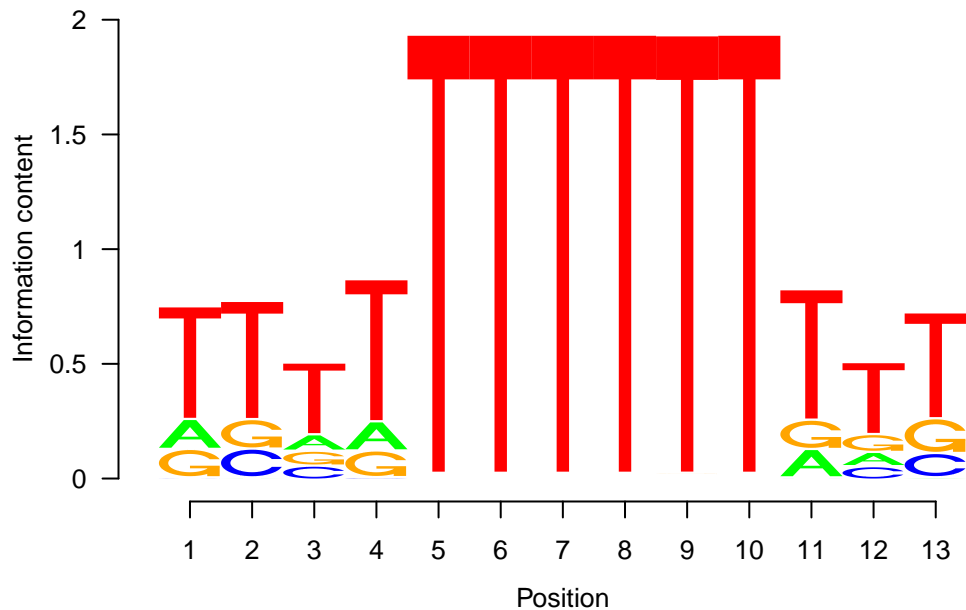
Genome: FlyFactorSurvey-hb_FlyReg_FBgn0001180



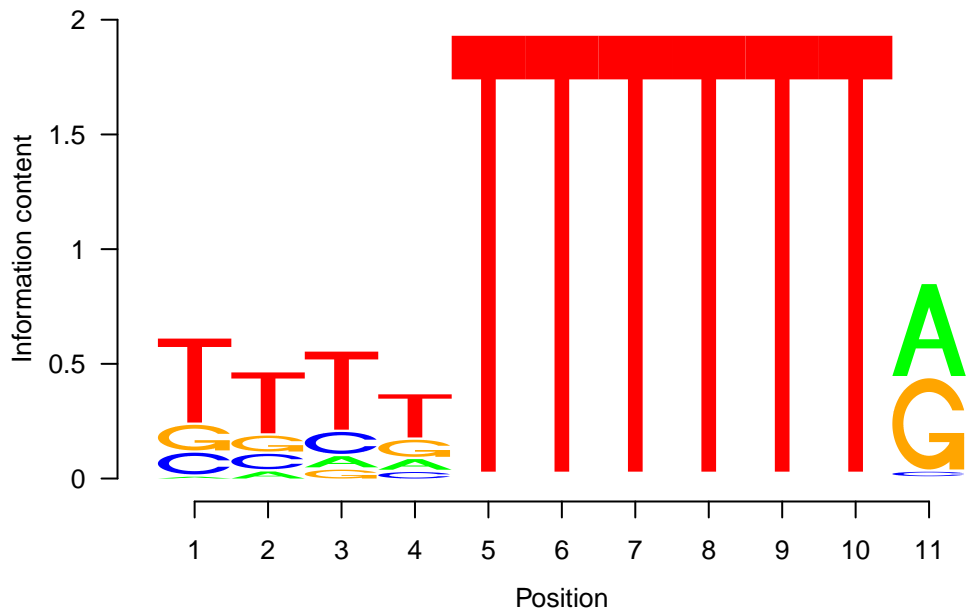
Genome: FlyFactorSurvey-hb_NAR_FBgn0001180



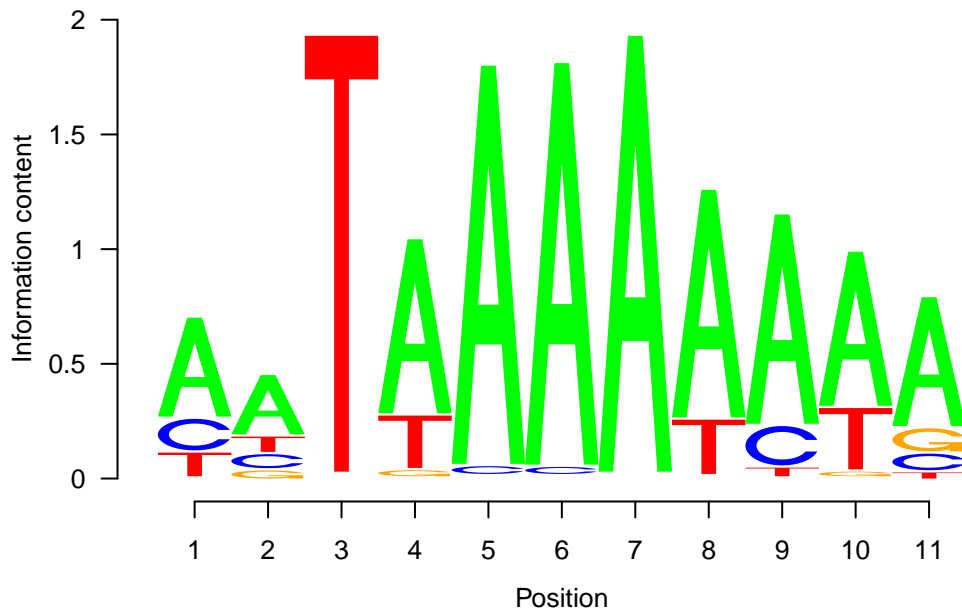
Genome: FlyFactorSurvey-hb_SANGER_2.5_FBgn0001180



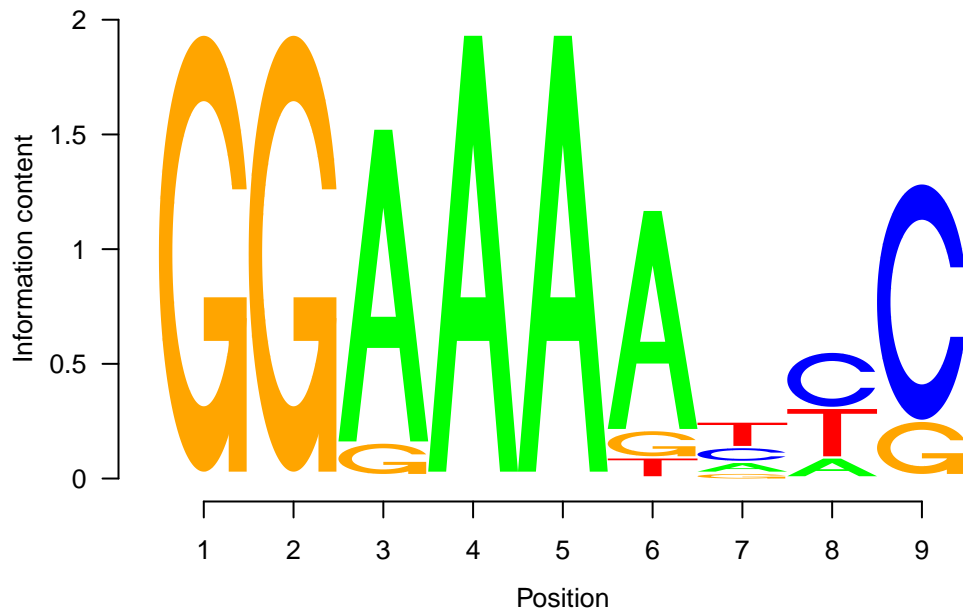
Genome: FlyFactorSurvey-hb_SOLEXA_5_FBgn0001180



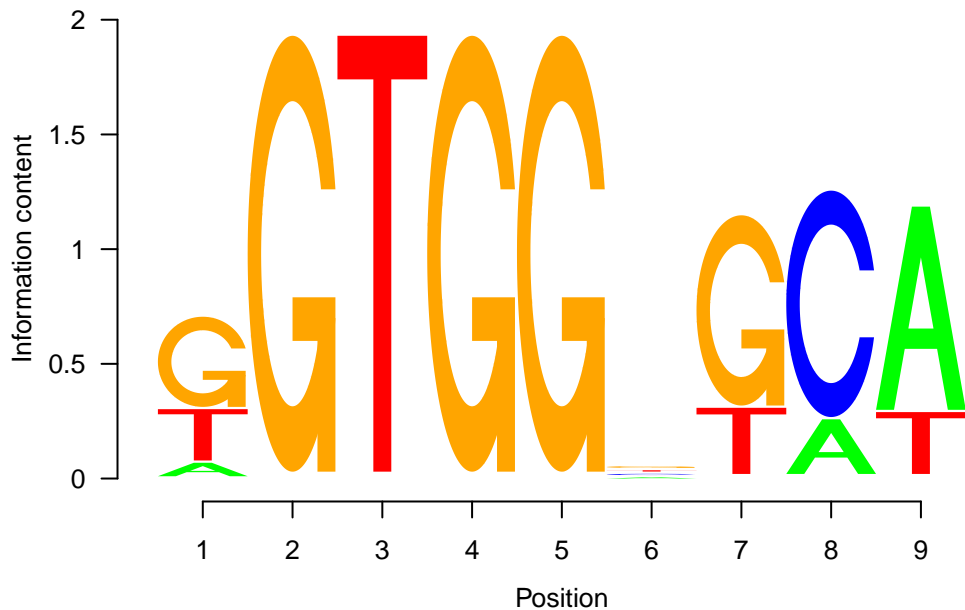
Genome: FlyFactorSurvey-jigr1_SANGER_5_FBgn0039350



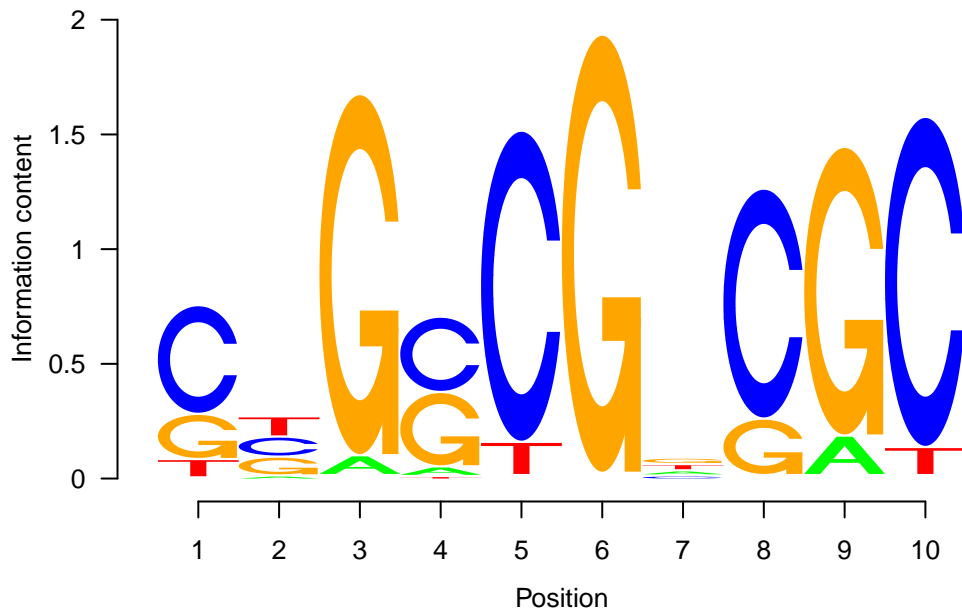
Genome: FlyFactorSurvey-lola.PC_SANGER_5_FBgn000563



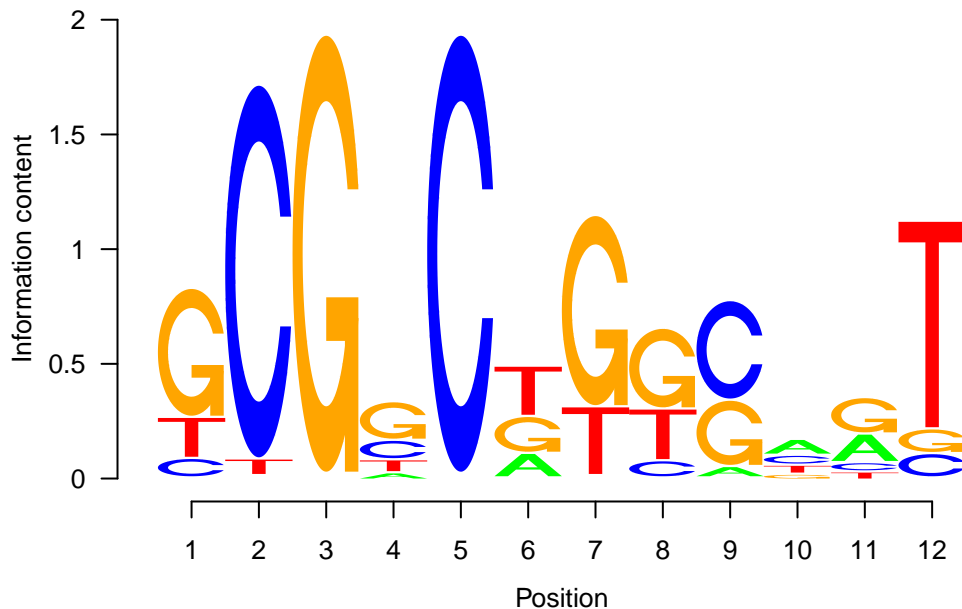
Genome: FlyFactorSurvey-lola.PL_SANGER_2.5_FBgn00056



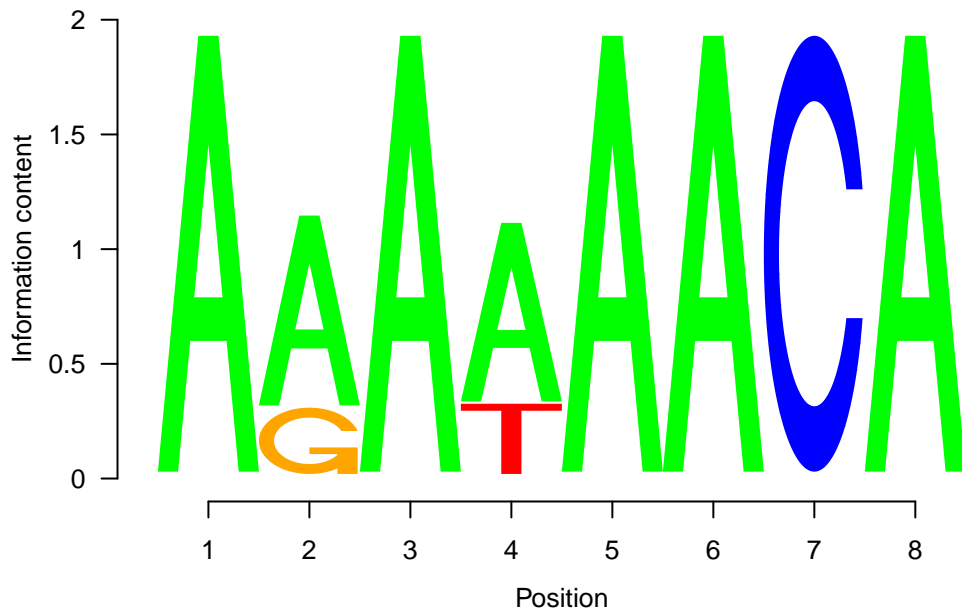
Genome: FlyFactorSurvey–Mad_FlyReg_FBgn0011648



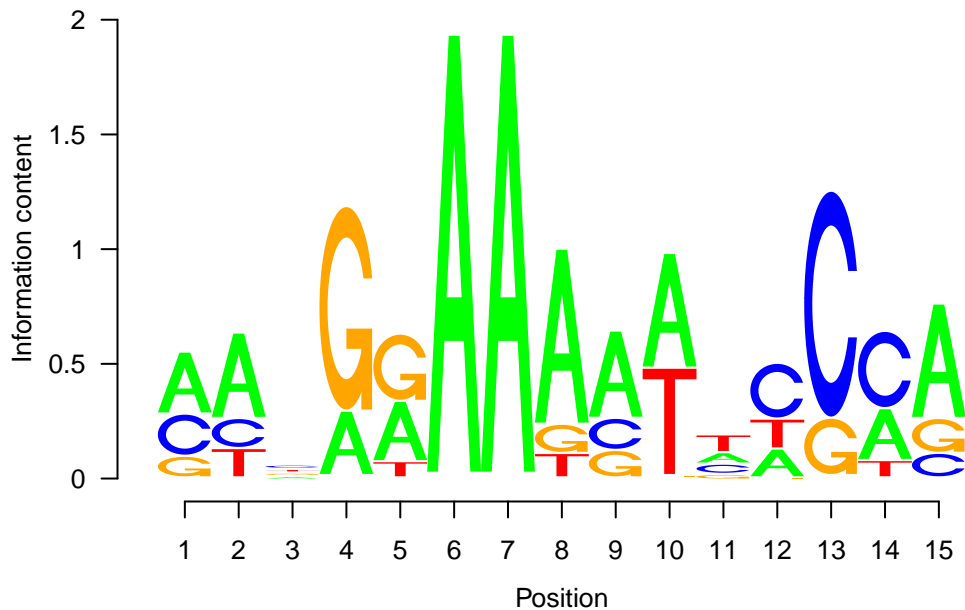
Genome: FlyFactorSurvey-Med_FlyReg_FBgn0011655



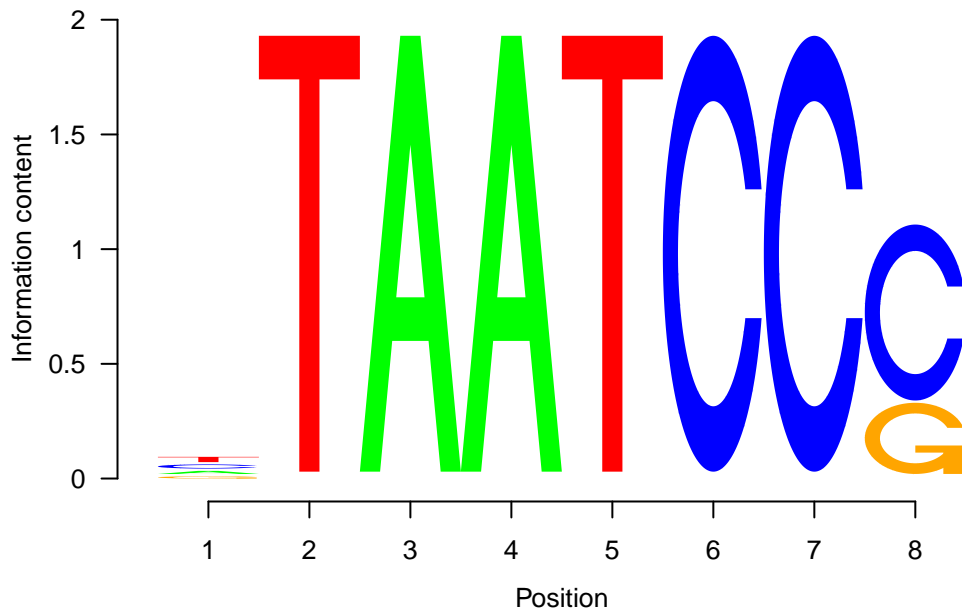
Genome: FlyFactorSurvey-Mirr_Cell_FBgn0014343



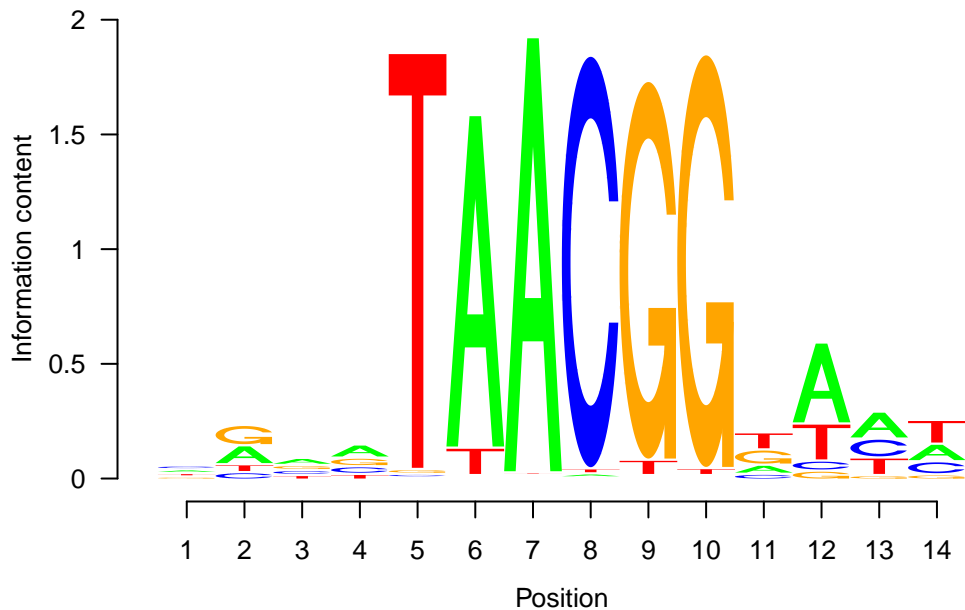
Genome: FlyFactorSurvey-NFAT_SANGER_5_FBgn003050



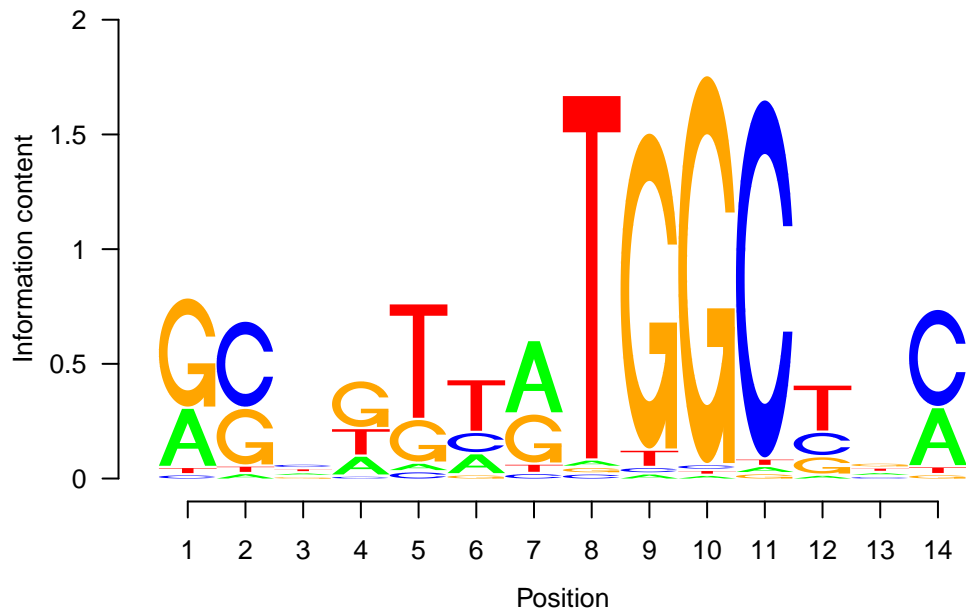
Genome: FlyFactorSurvey-Oc_SOLEXA_FBgn0004102



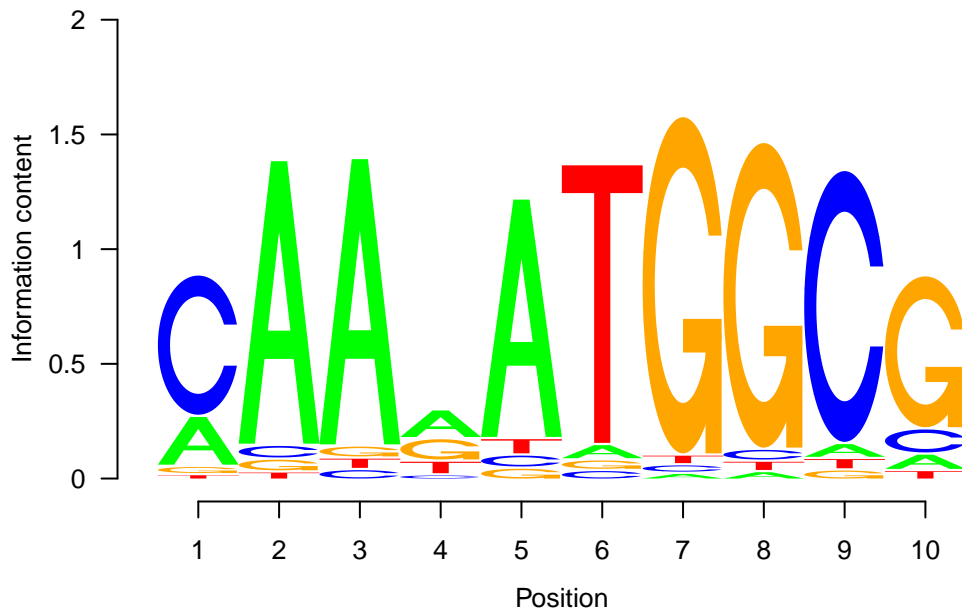
Genome: FlyFactorSurvey-ovo_SOLEXA_5_FBgn0003028



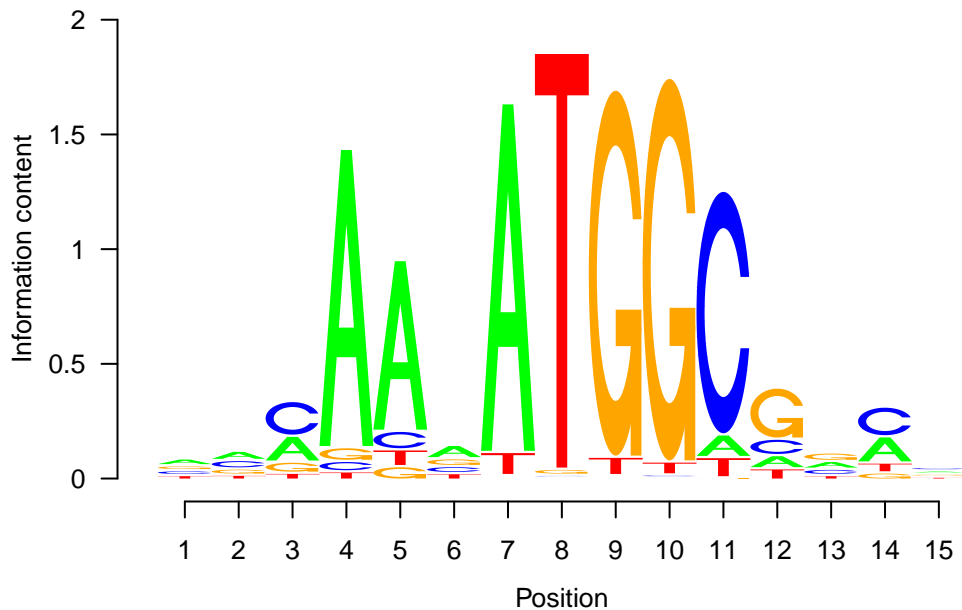
Genome: FlyFactorSurvey-pho_FlyReg_FBgn0002521



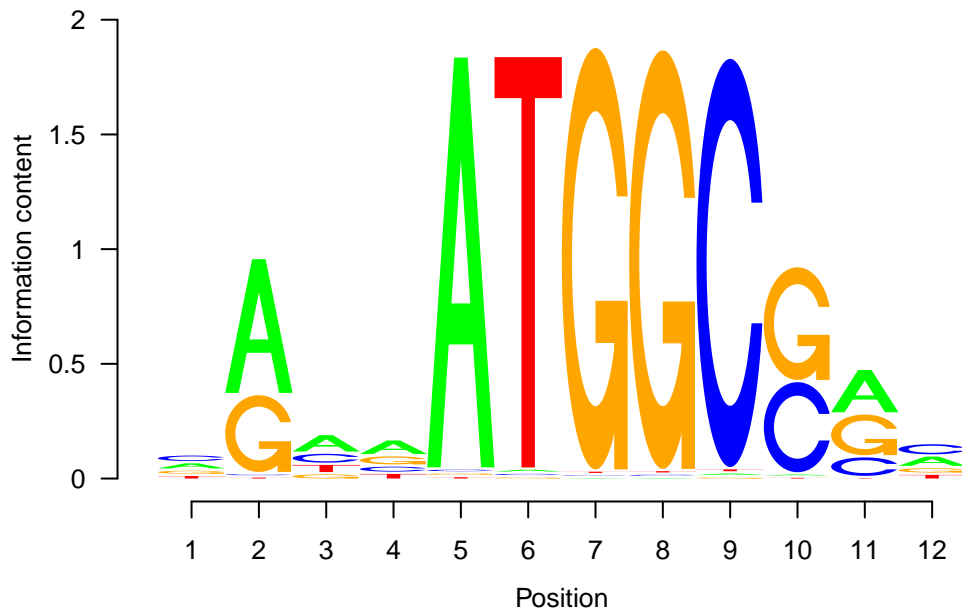
Genome: FlyFactorSurvey-phol_SANGER_5_FBgn0035997



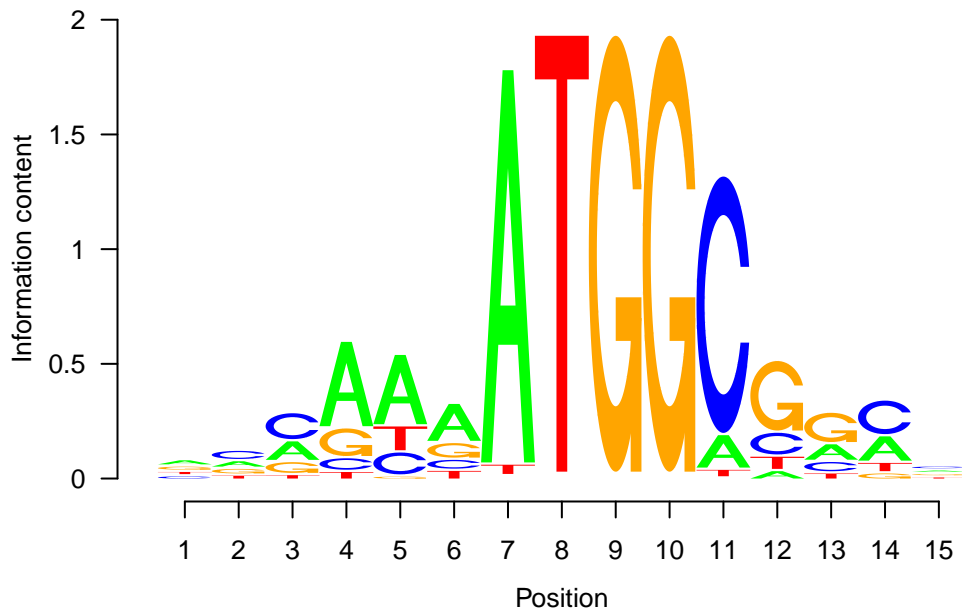
Genome: FlyFactorSurvey-phol_SOLEXA_5_FBgn0035997



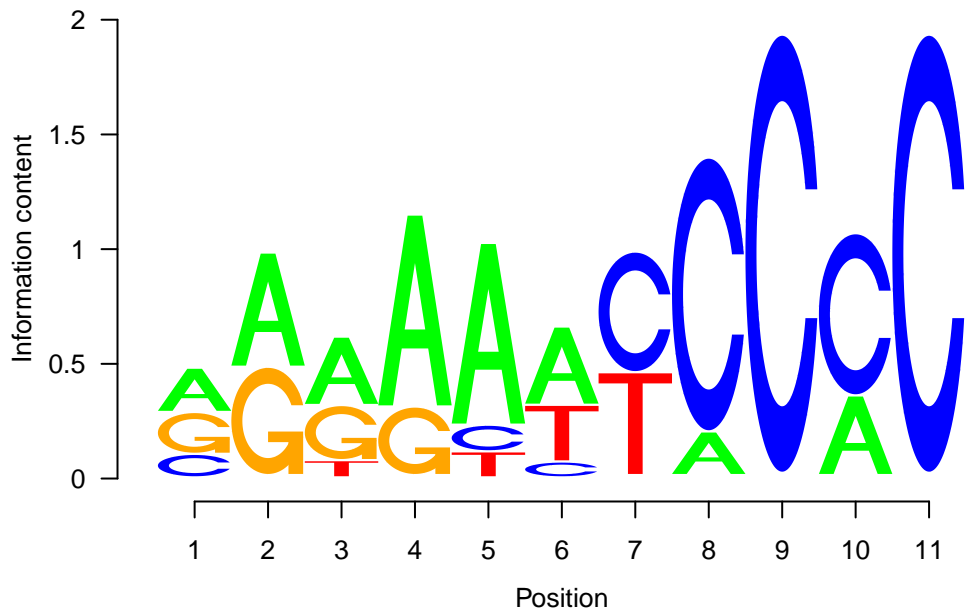
Genome: FlyFactorSurvey-pho_SANGER_10_FBgn000252



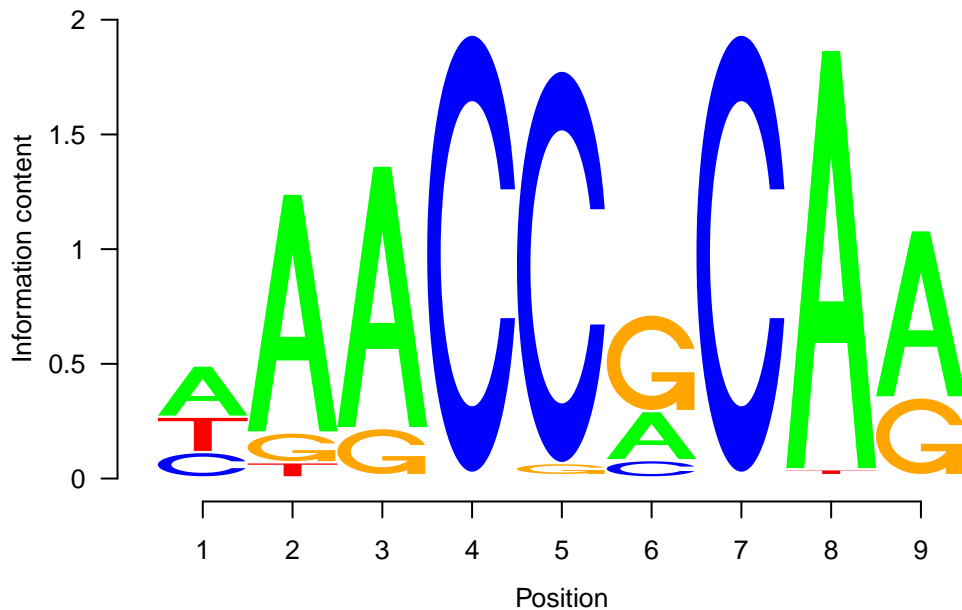
Genome: FlyFactorSurvey-pho_SOLEXA_5_FBgn0002521



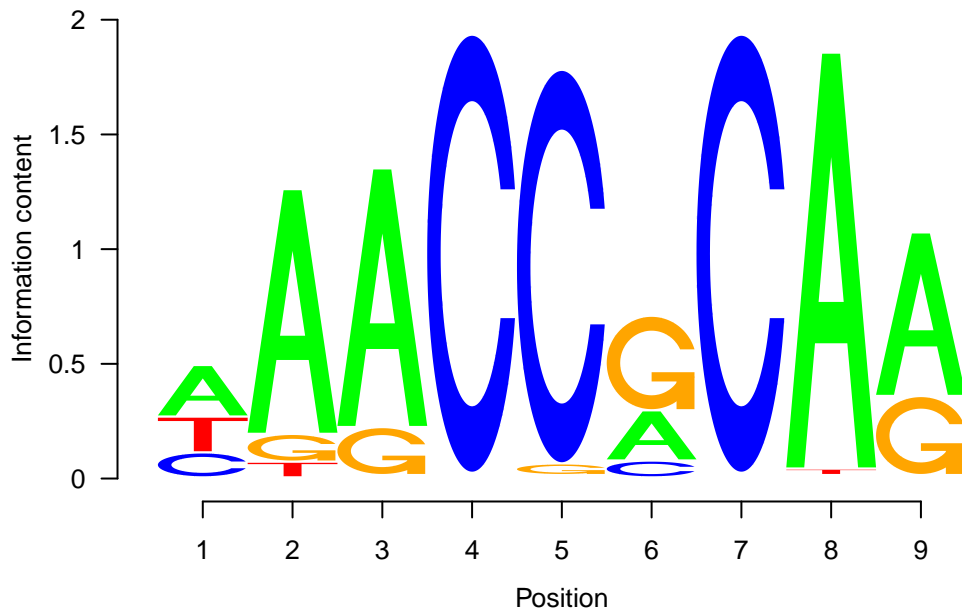
Genome: FlyFactorSurvey-Rel_SANGER_5_FBgn0014018



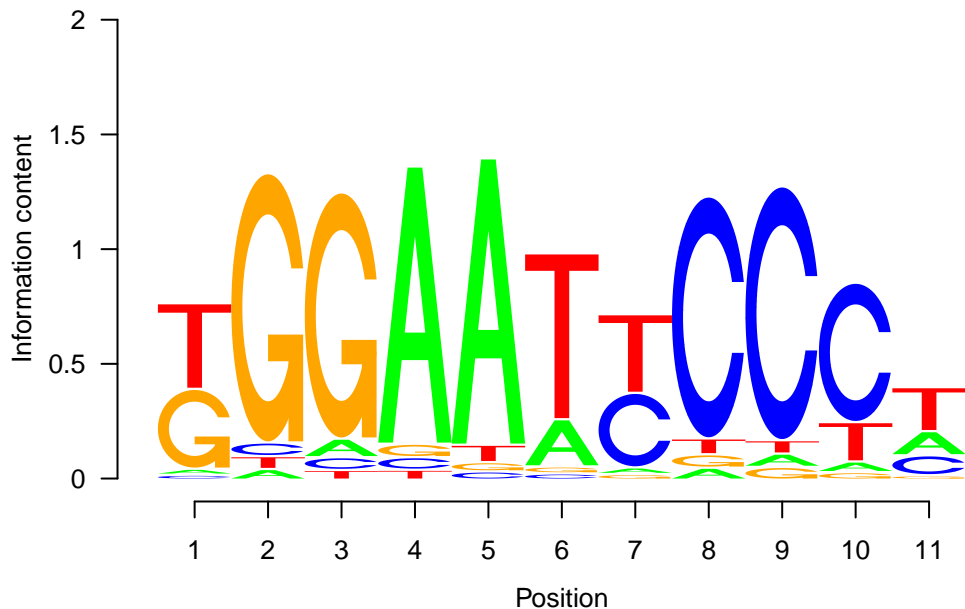
Genome: FlyFactorSurvey-run_Bgb_NBT_FBgn0003300



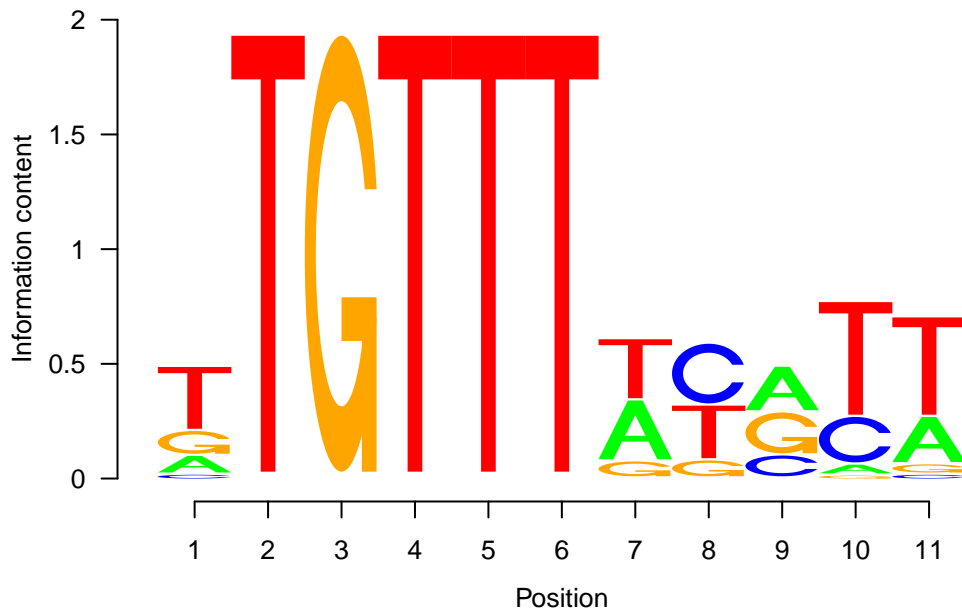
Genome: FlyFactorSurvey-run_Bgb_NBT_FBgn0013753



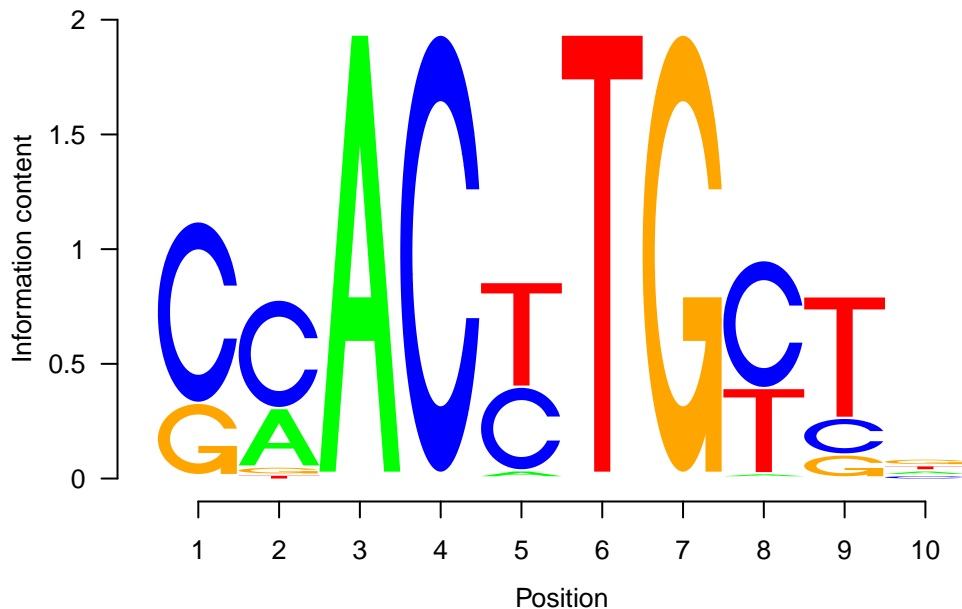
Genome: FlyFactorSurvey-shn.F1.2_SANGER_5_FBgn00033



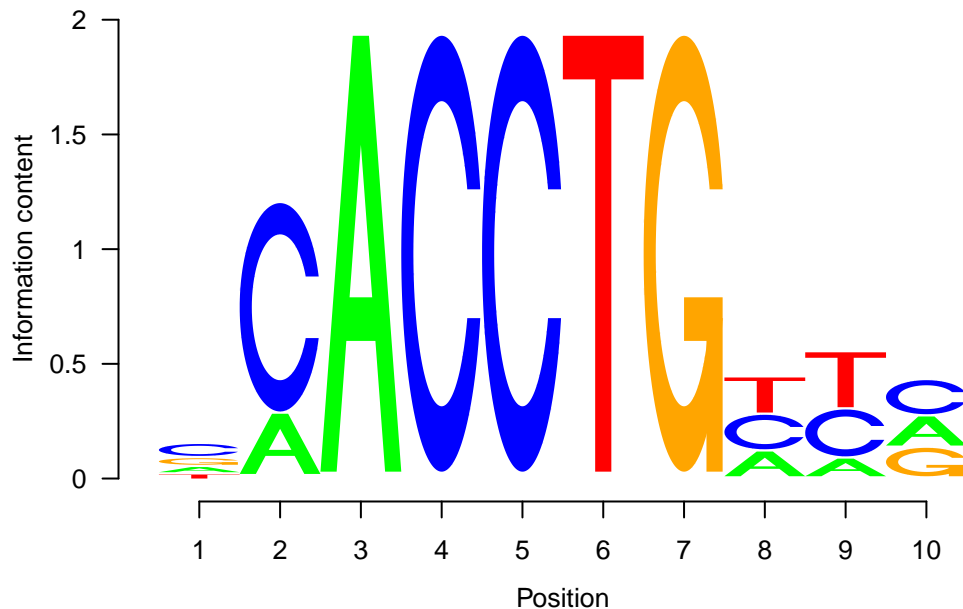
Genome: FlyFactorSurvey-slp1_NAR_FBgn0003430



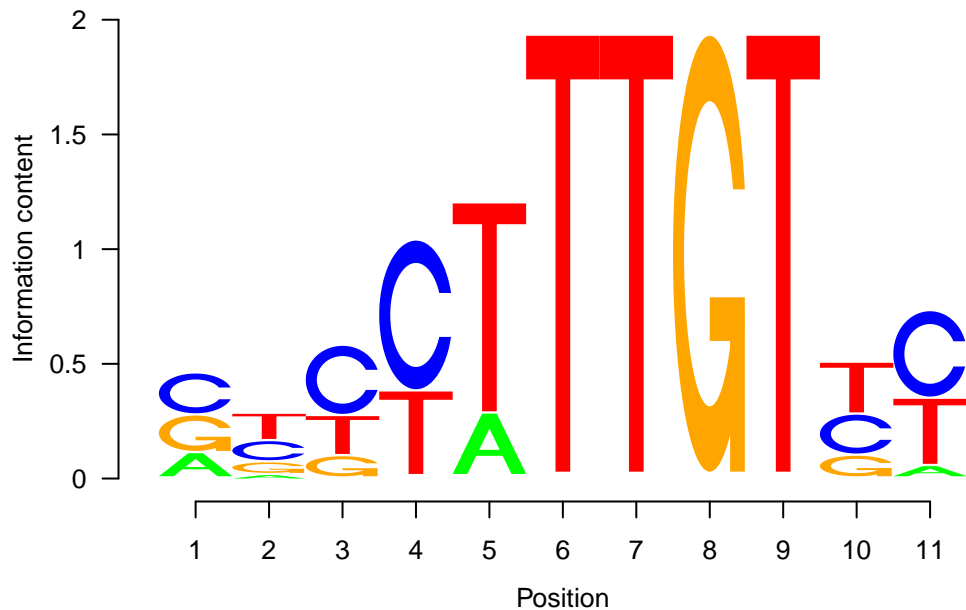
Genome: FlyFactorSurvey-sna_FlyReg_FBgn0003448



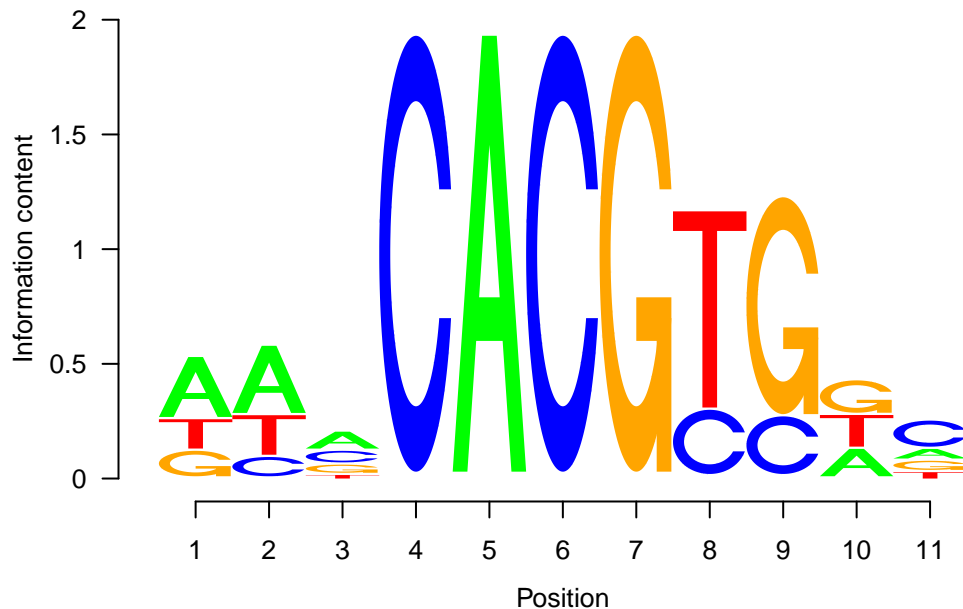
Genome: FlyFactorSurvey-sna_SANGER_10_FBgn000344



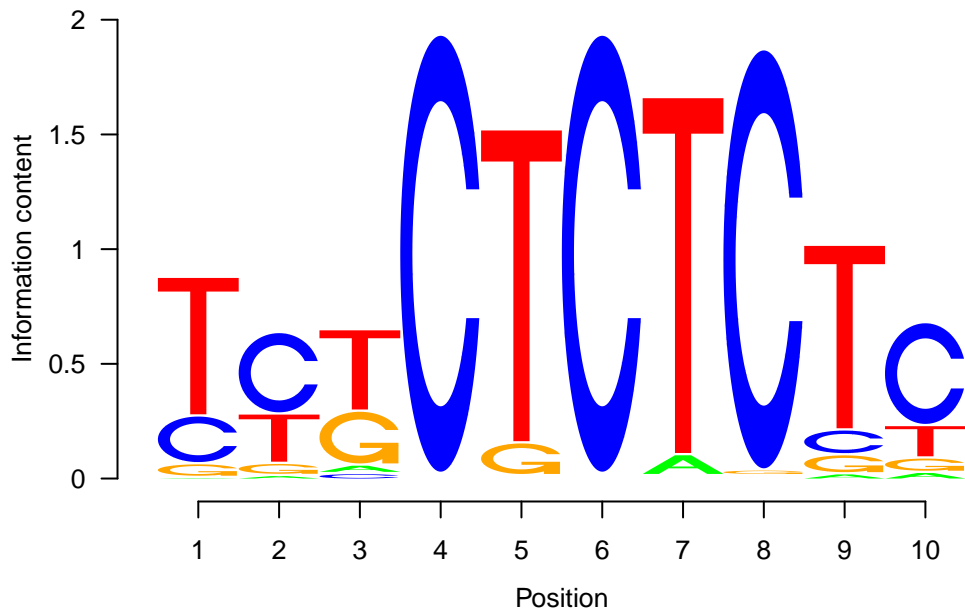
Genome: FlyFactorSurvey-Sox14_SANGER_10_FBgn00056



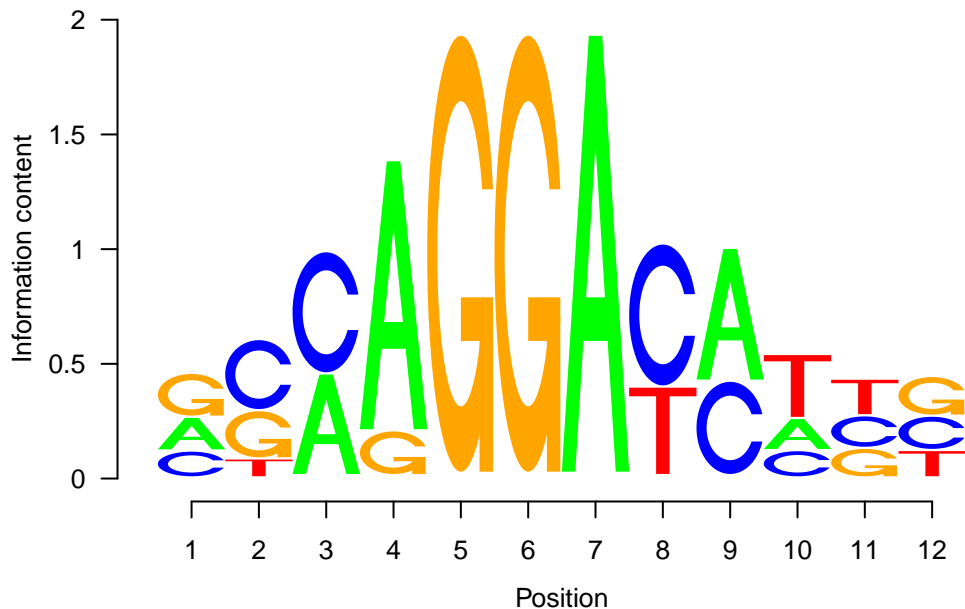
Genome: FlyFactorSurvey-tai_SANGER_5_FBgn0041092



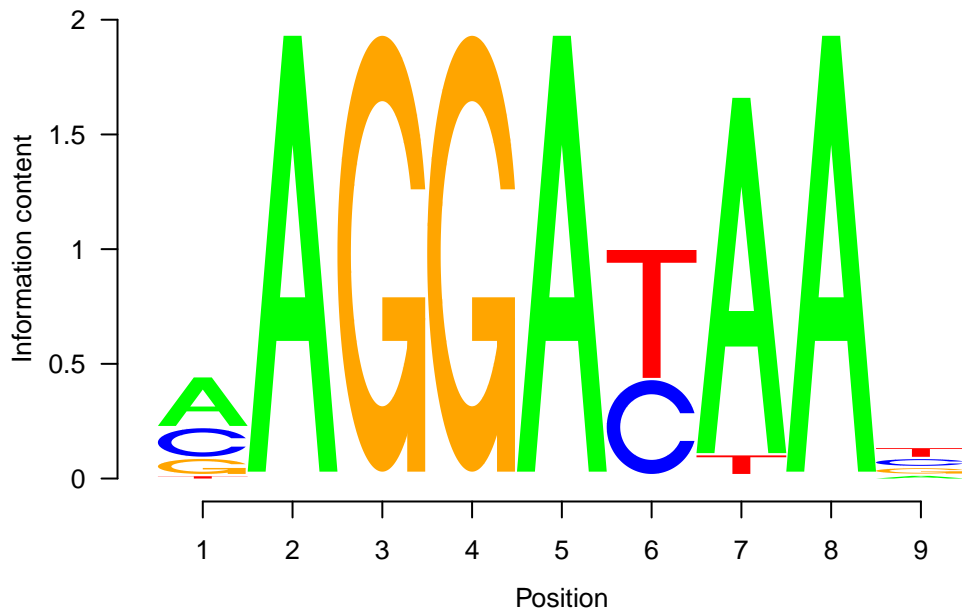
Genome: FlyFactorSurvey-Trl_FlyReg_FBgn0013263



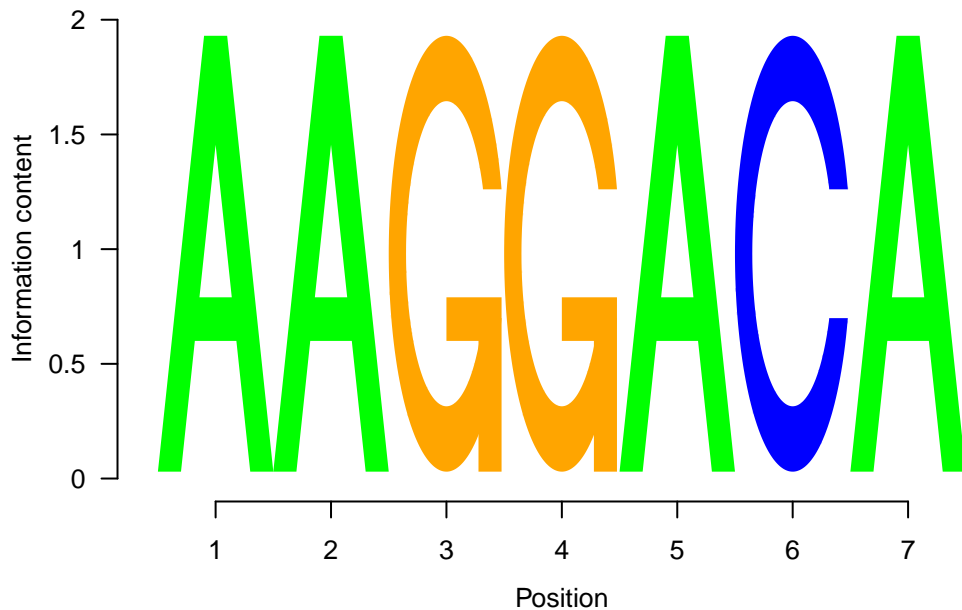
Genome: FlyFactorSurvey-ttk_FlyReg_FBgn0003870



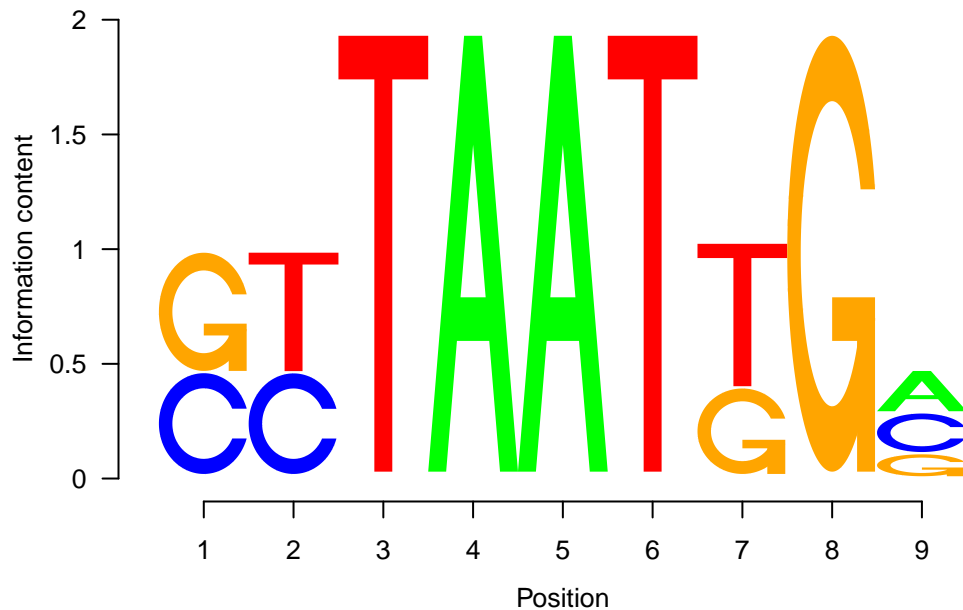
Genome: FlyFactorSurvey-ttk_NAR_FBgn0003870



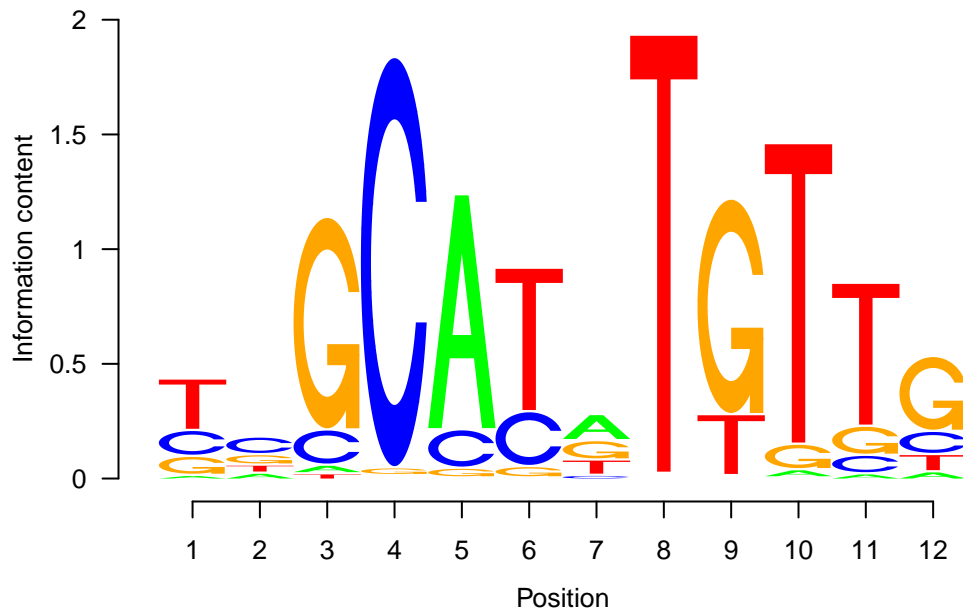
Genome: FlyFactorSurvey-ttk.PF_SANGER_5_FBgn000387



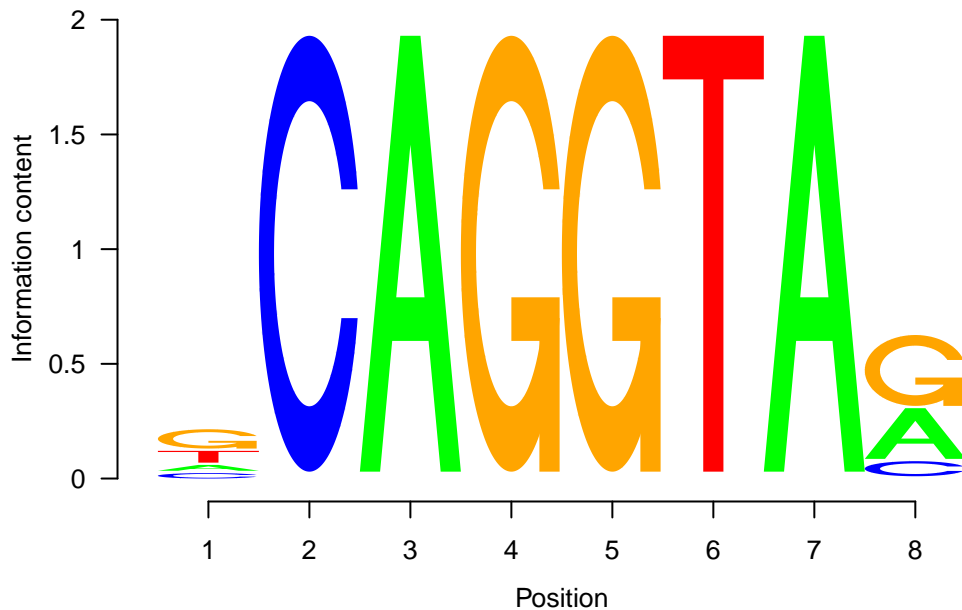
Genome: FlyFactorSurvey-tup_SOLEXA_10_FBgn0003896



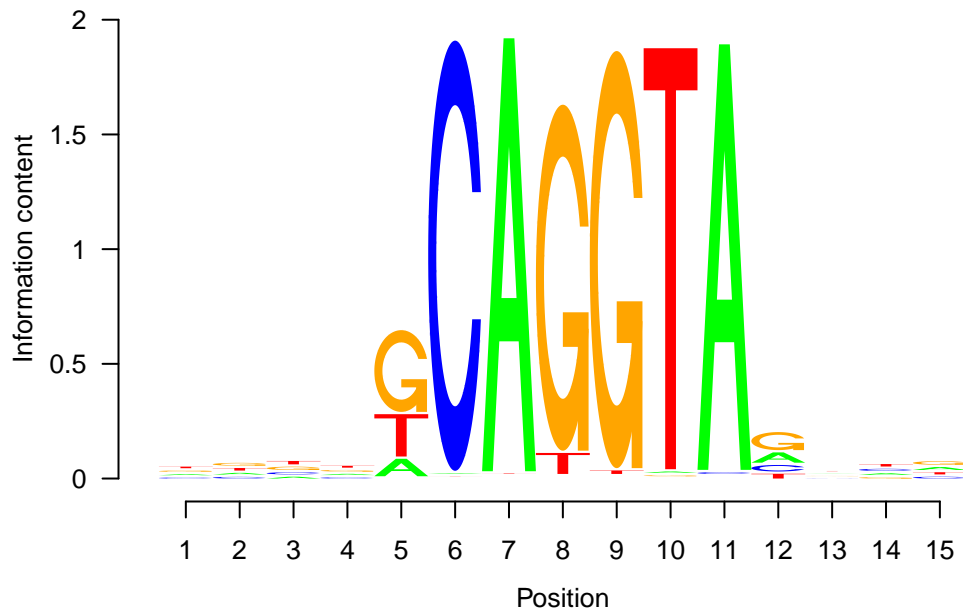
Genome: FlyFactorSurvey-twi_FlyReg_FBgn0003900



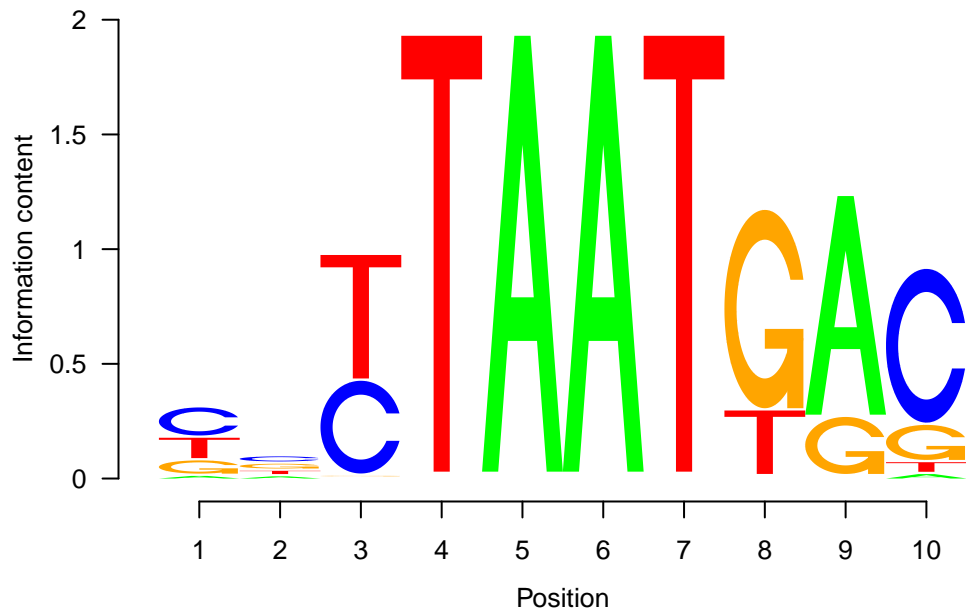
Genome: FlyFactorSurvey-vfl_SANGER_5_FBgn0259789



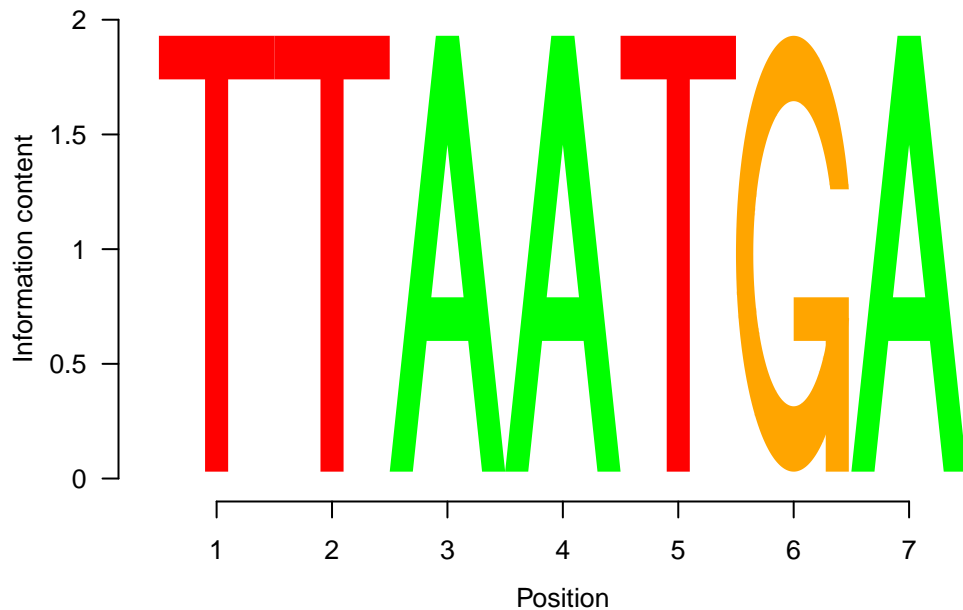
Genome: FlyFactorSurvey-vfl_SOLEXA_5_FBgn0259789



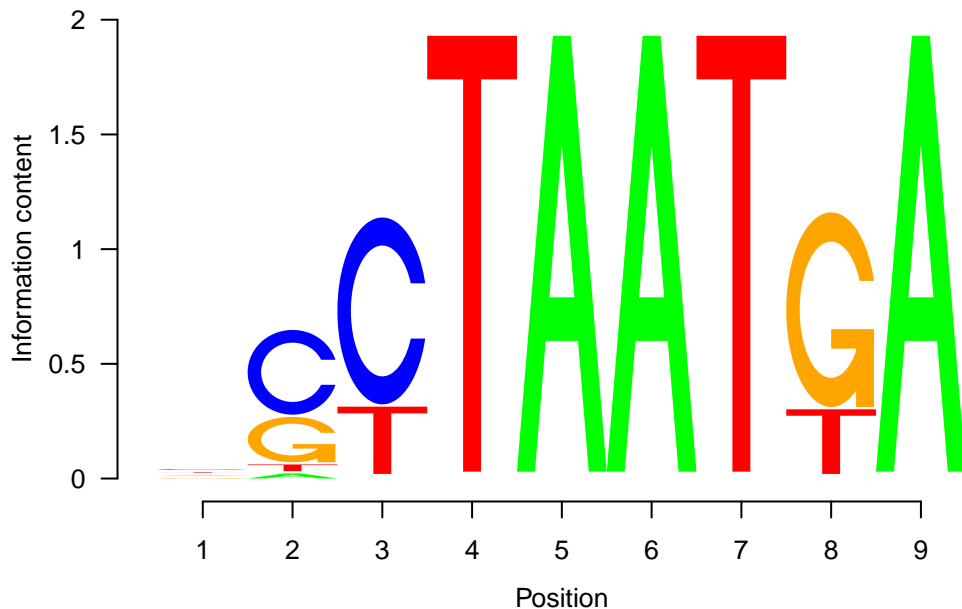
Genome: FlyFactorSurvey-zen2_SOLEXA_2_FBgn0004054



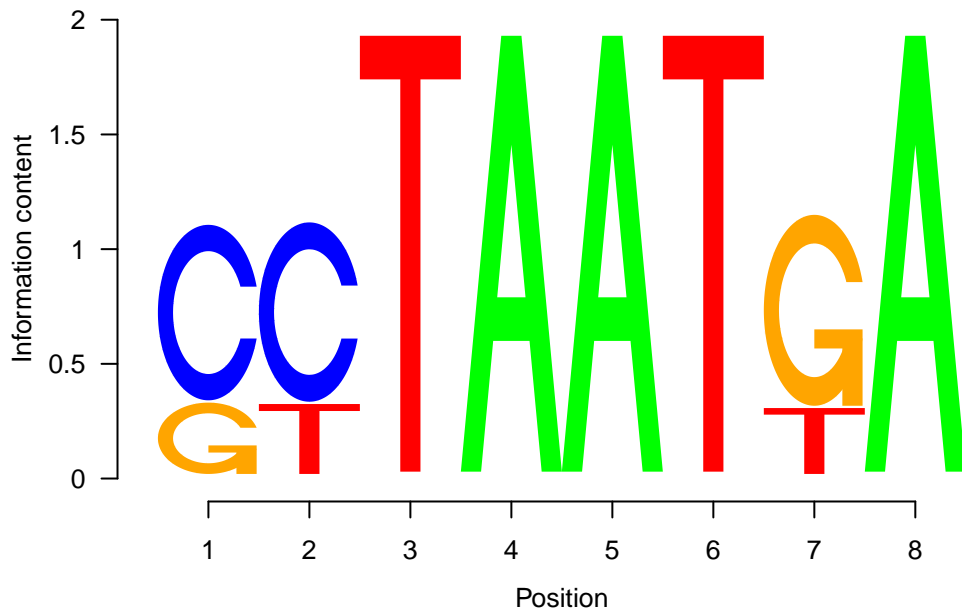
Genome: FlyFactorSurvey-Zen2_SOLEXA_FBgn0004054



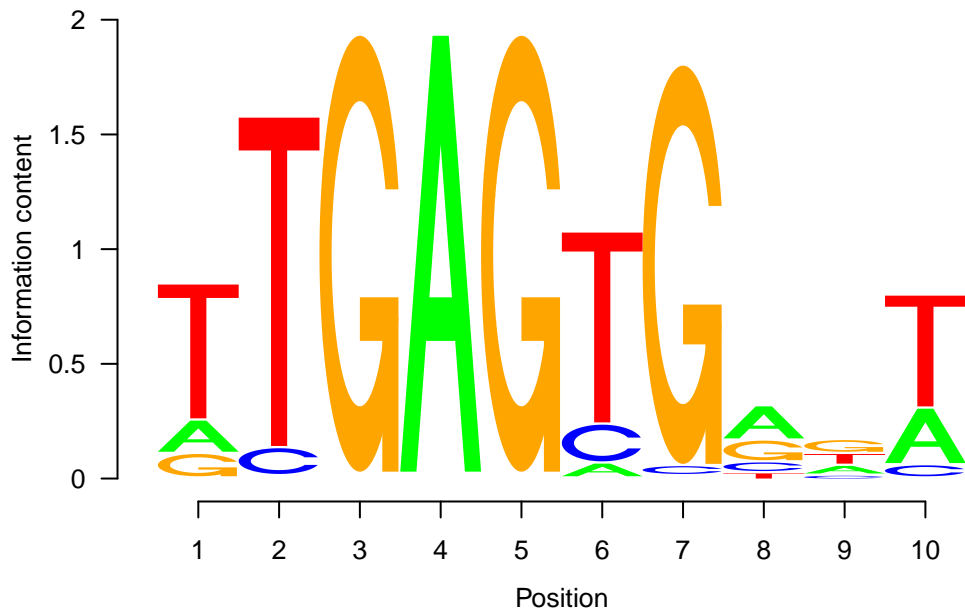
Genome: FlyFactorSurvey-Zen_Cell_FBgn0004053



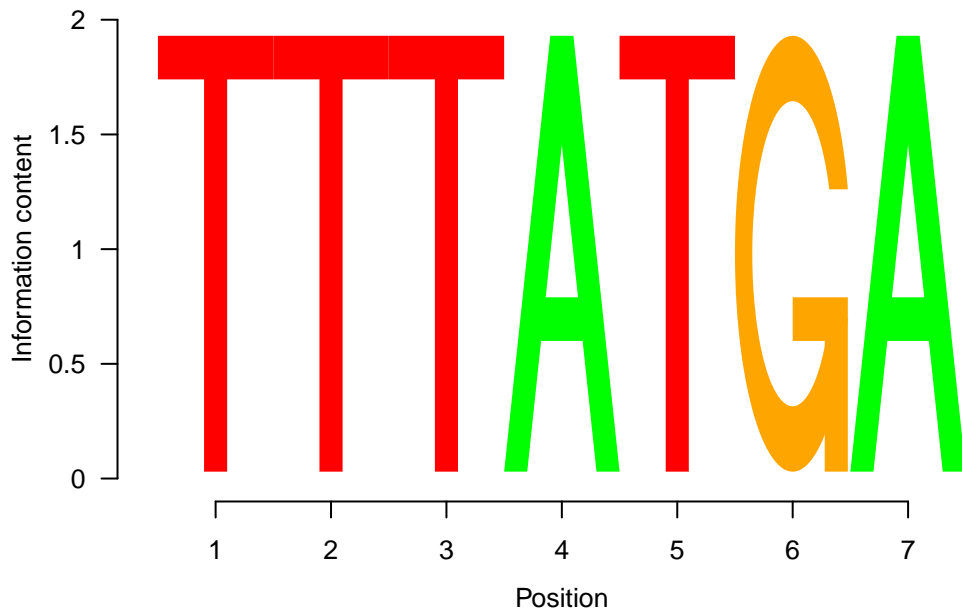
Genome: FlyFactorSurvey-Zen_SOLEXA_FBgn0004053



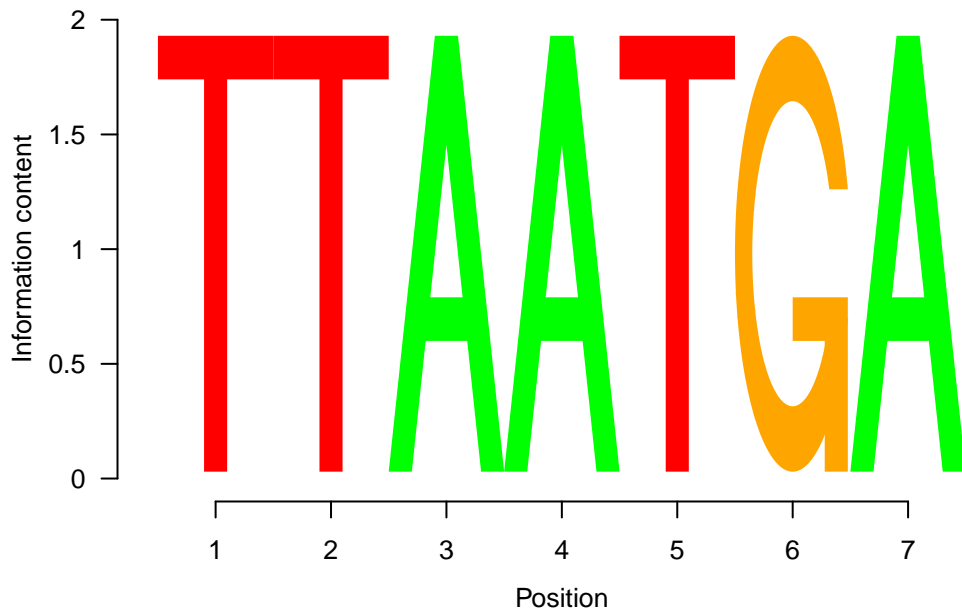
Genome: FlyFactorSurvey-z_FlyReg_FBgn0004050



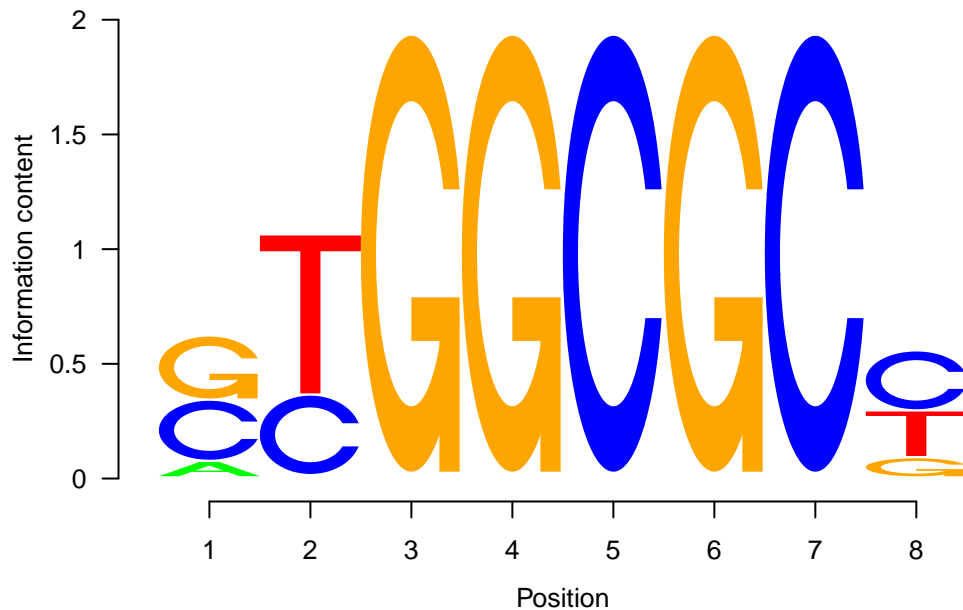
Genome: JASPAR_2014-Abd-B-MA0165.1



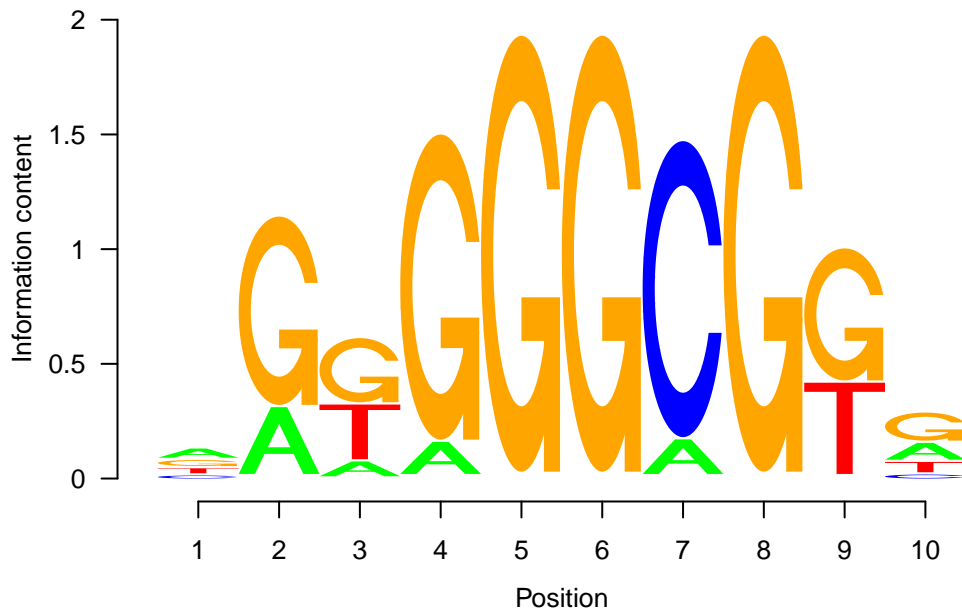
Genome: JASPAR_2014-Antp-MA0166.1



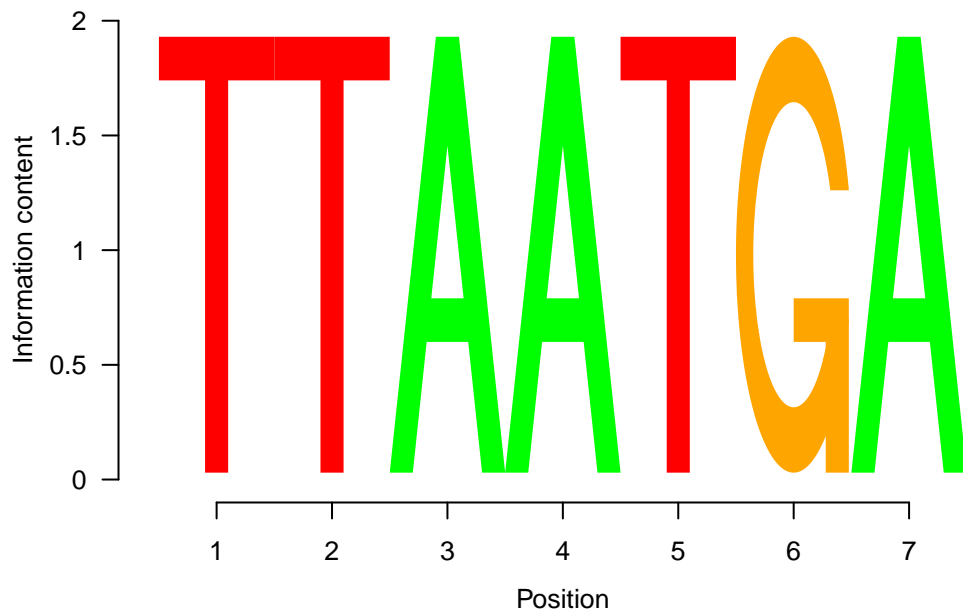
Genome: JASPAR_2014-brk-MA0213.1



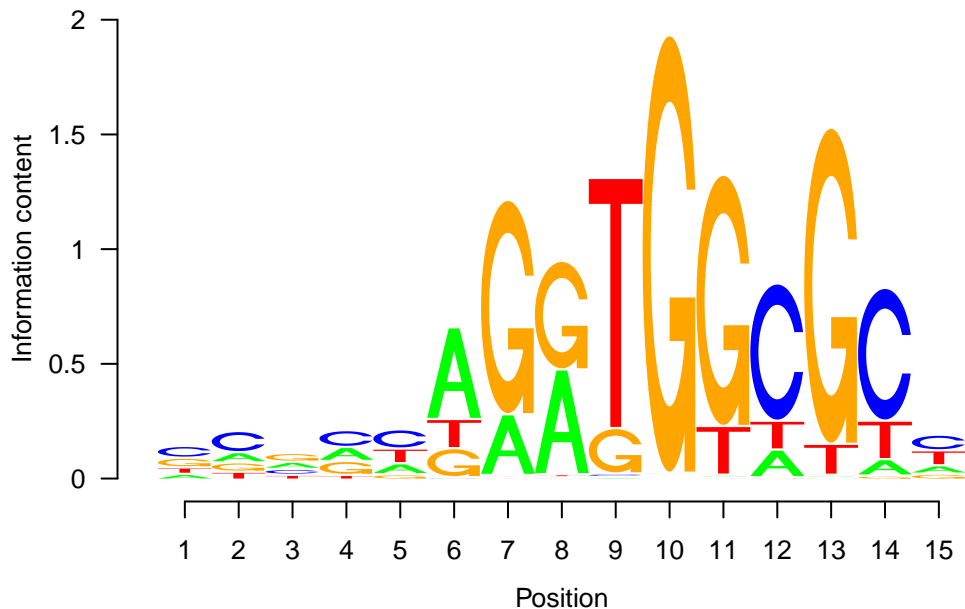
Genome: JASPAR_2014-btd-MA0443.1



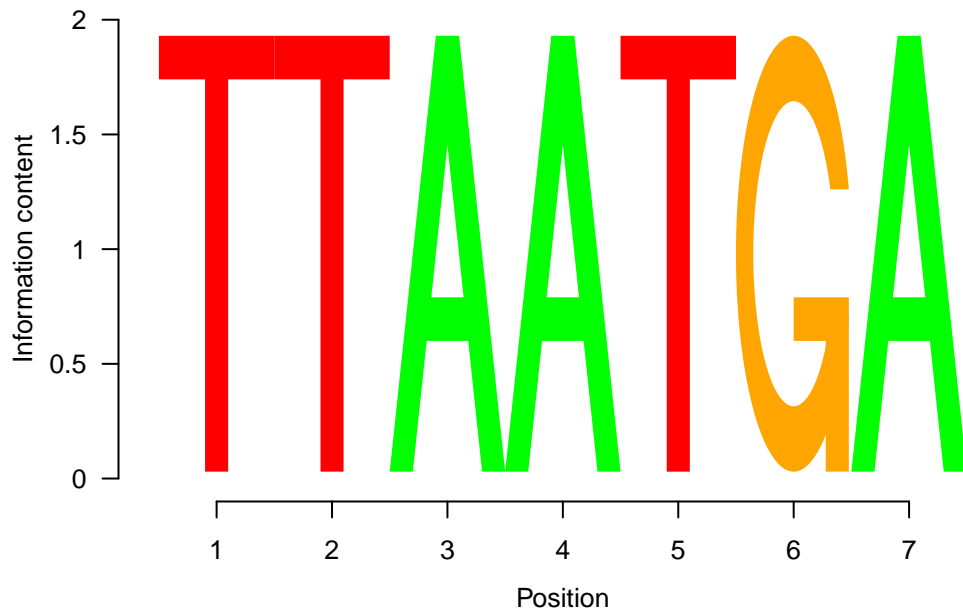
Genome: JASPAR_2014-btn-MA0215.1



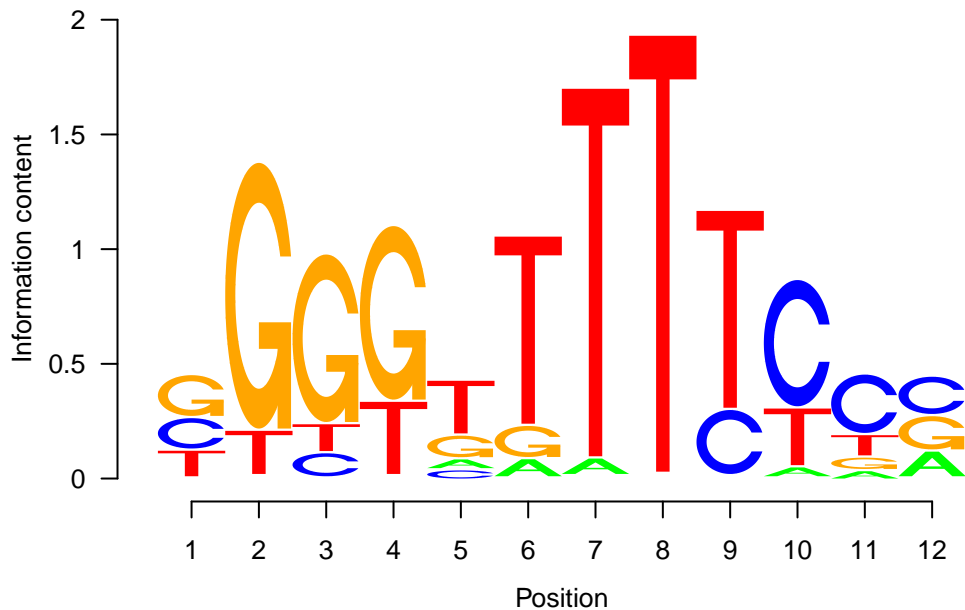
Genome: JASPAR_2014-CTCF-MA0531.1



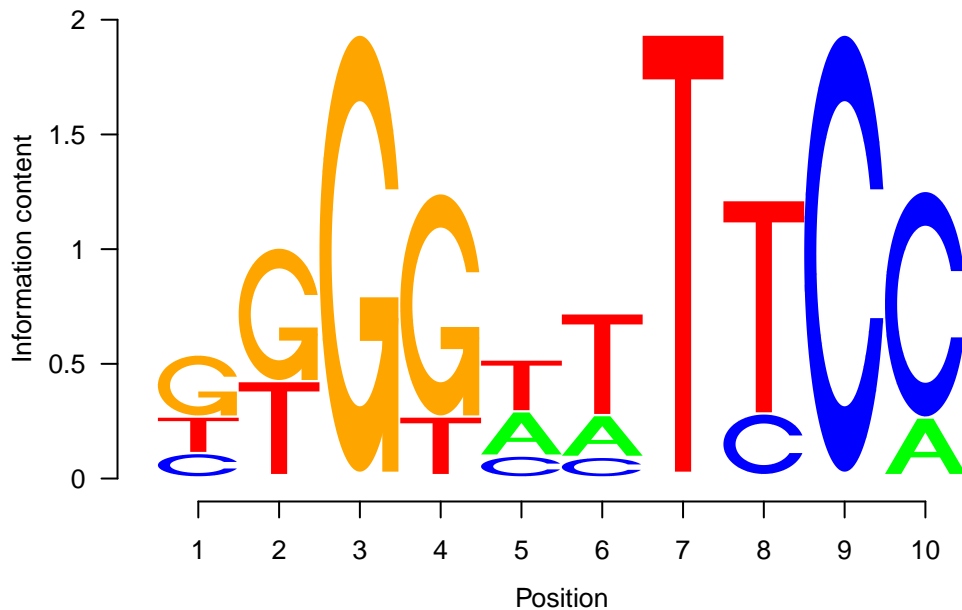
Genome: JASPAR_2014-Dfd-MA0186.1



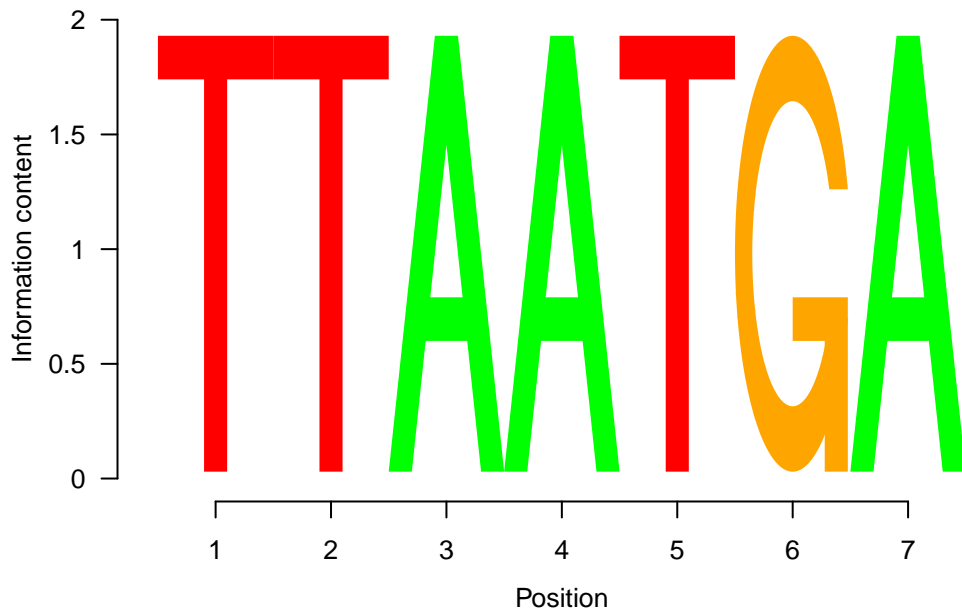
Genome: JASPAR_2014-dl_1-MA0022.1



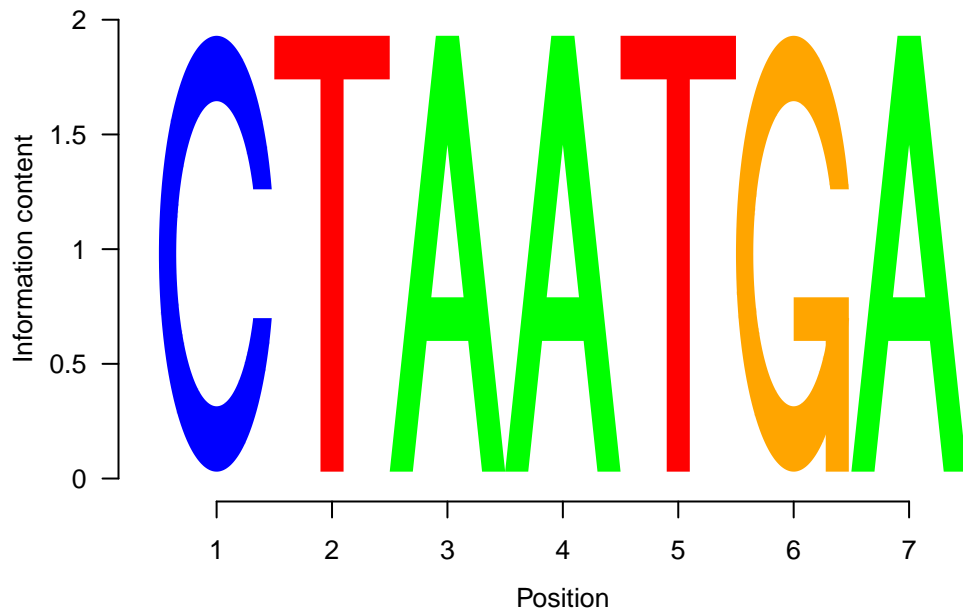
Genome: JASPAR_2014-dl_2-MA0023.1



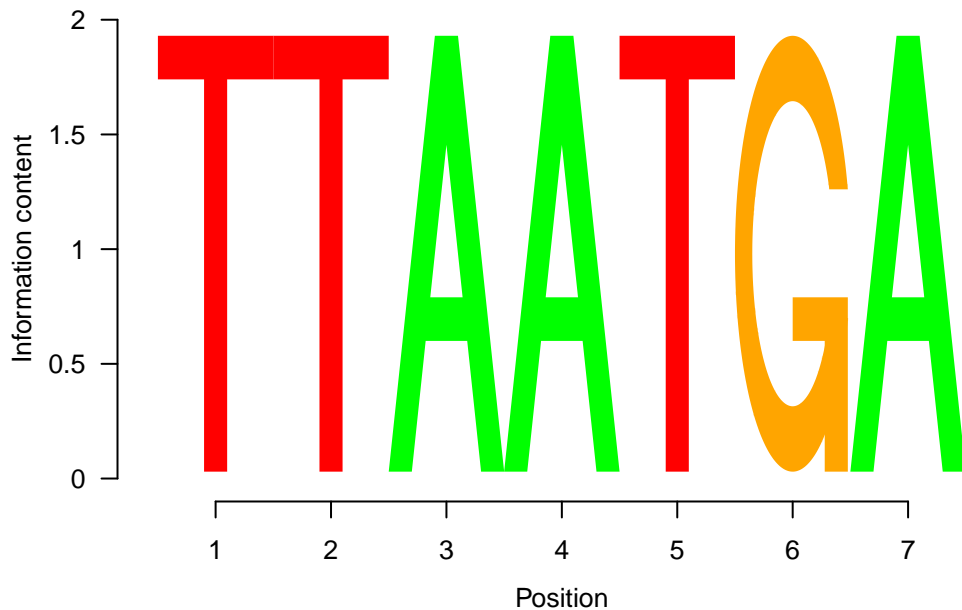
Genome: JASPAR_2014-ems-MA0219.1



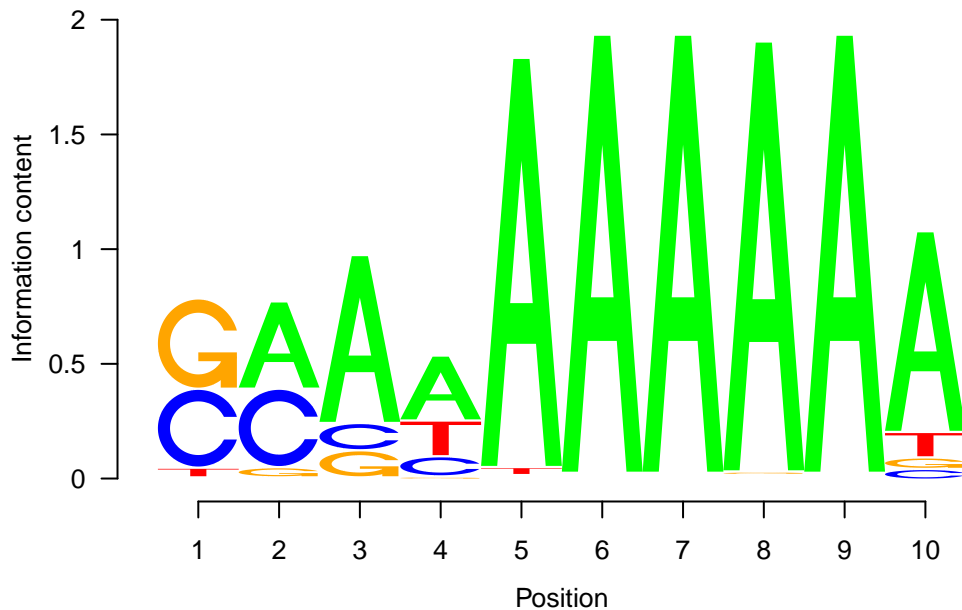
Genome: JASPAR_2014-eve-MA0221.1



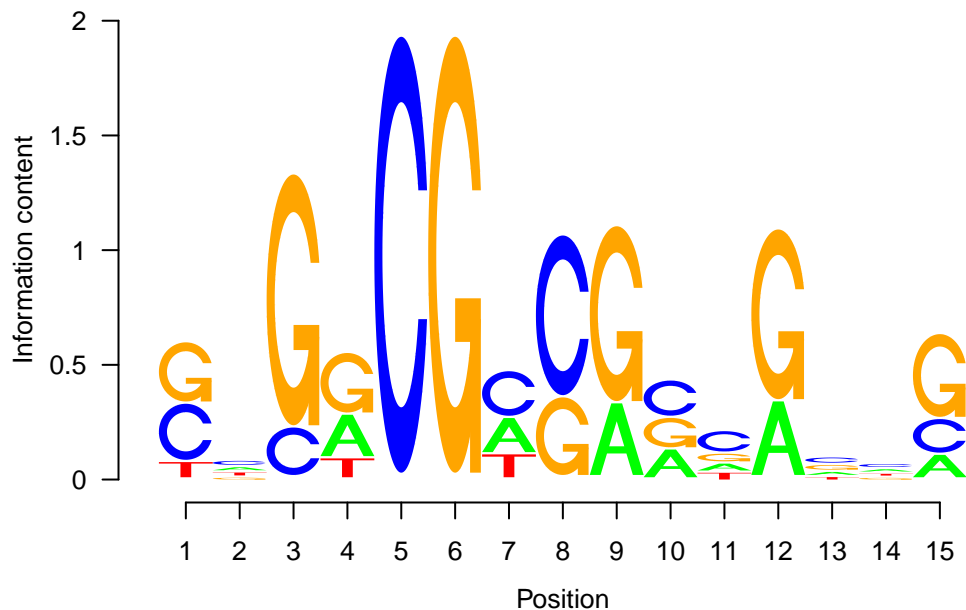
Genome: JASPAR_2014-ftz-MA0225.1



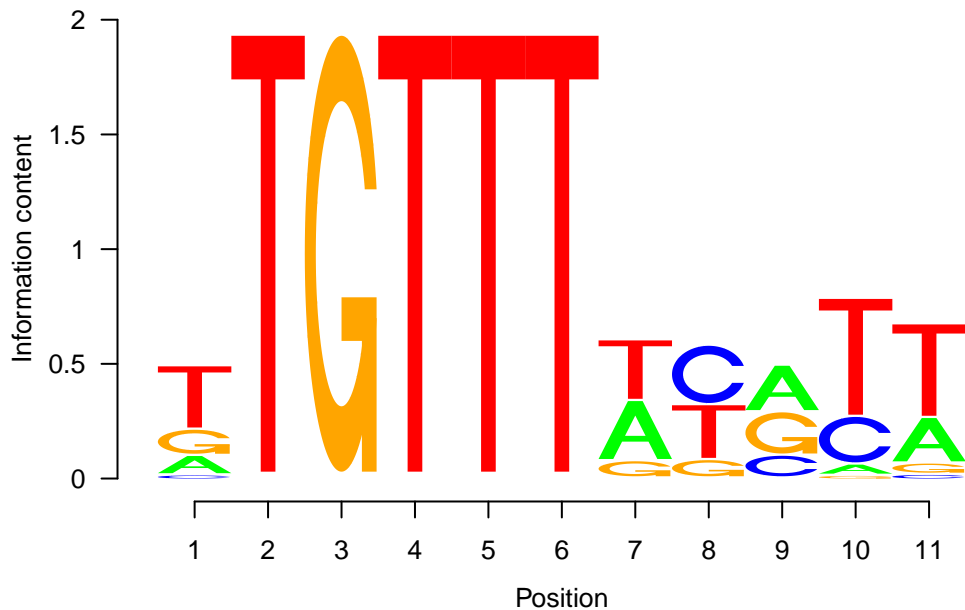
Genome: JASPAR_2014-hb-MA0049.1



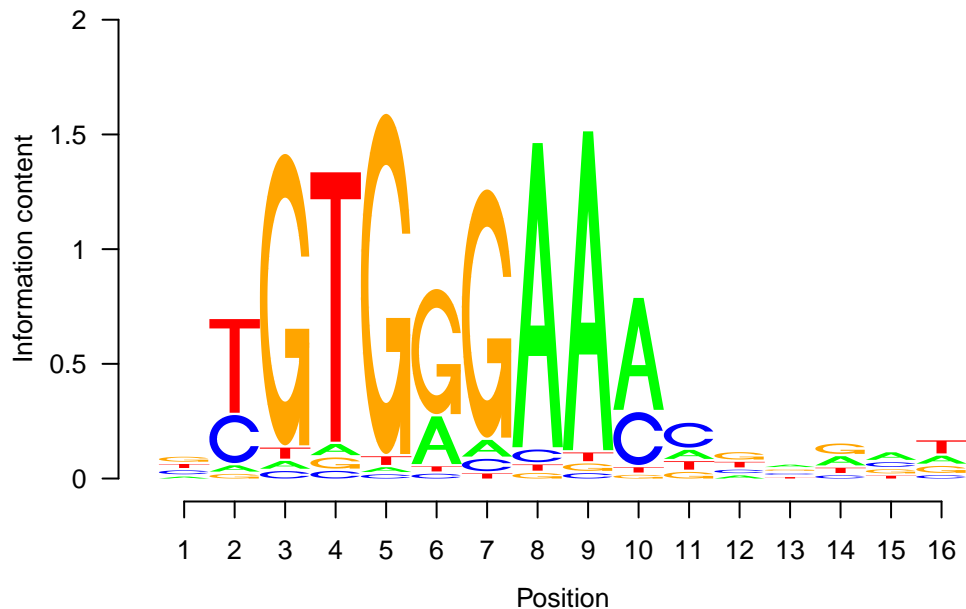
Genome: JASPAR_2014-Mad-MA0535.1



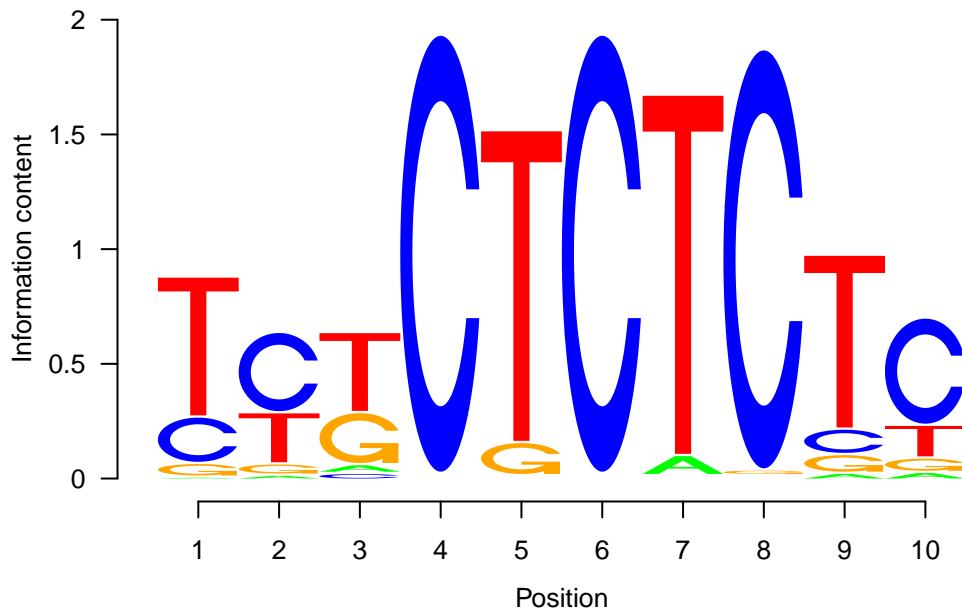
Genome: JASPAR_2014-slp1-MA0458.1



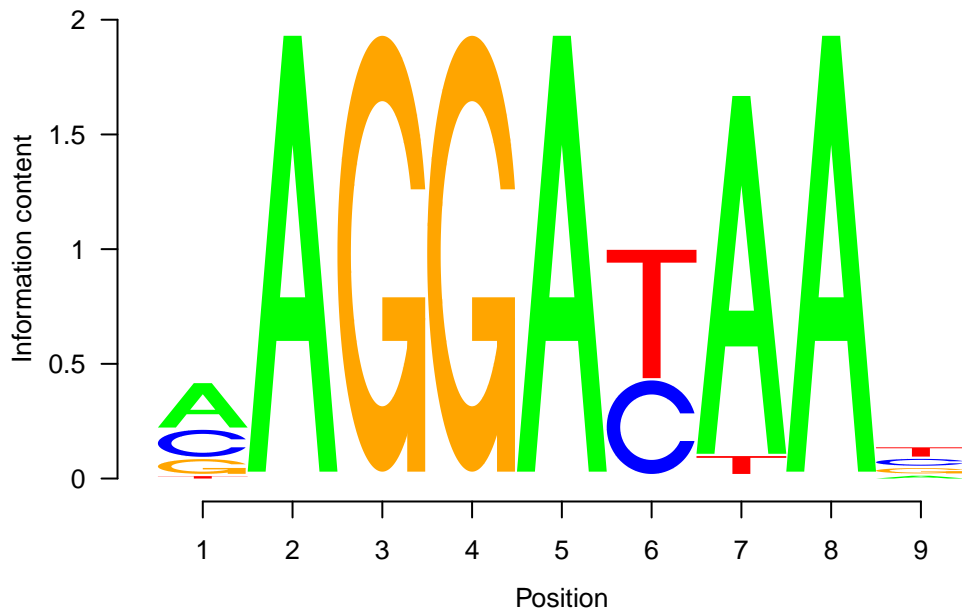
Genome: JASPAR_2014-Su(H)-MA0085.1



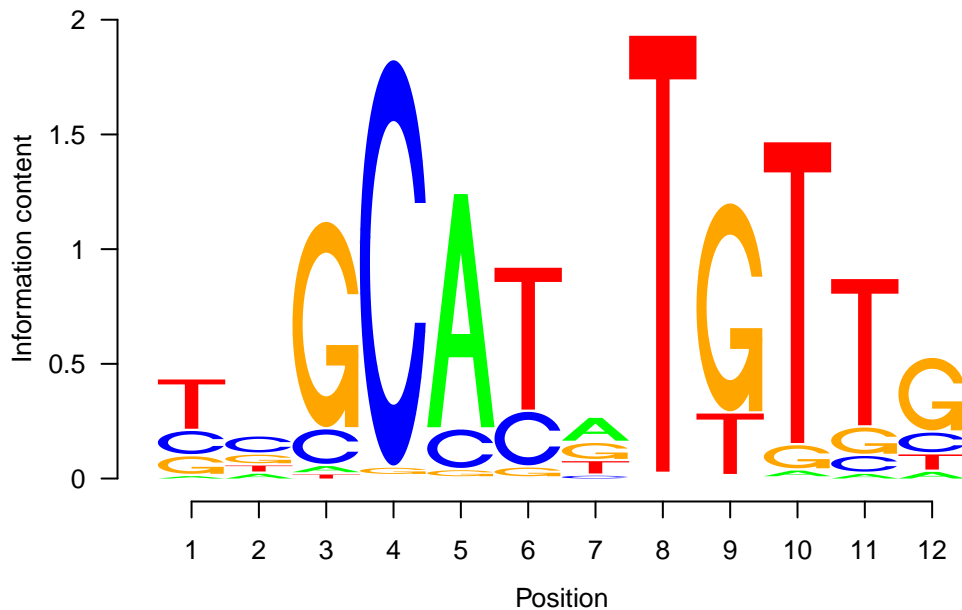
Genome: JASPAR_2014-Trl-MA0205.1



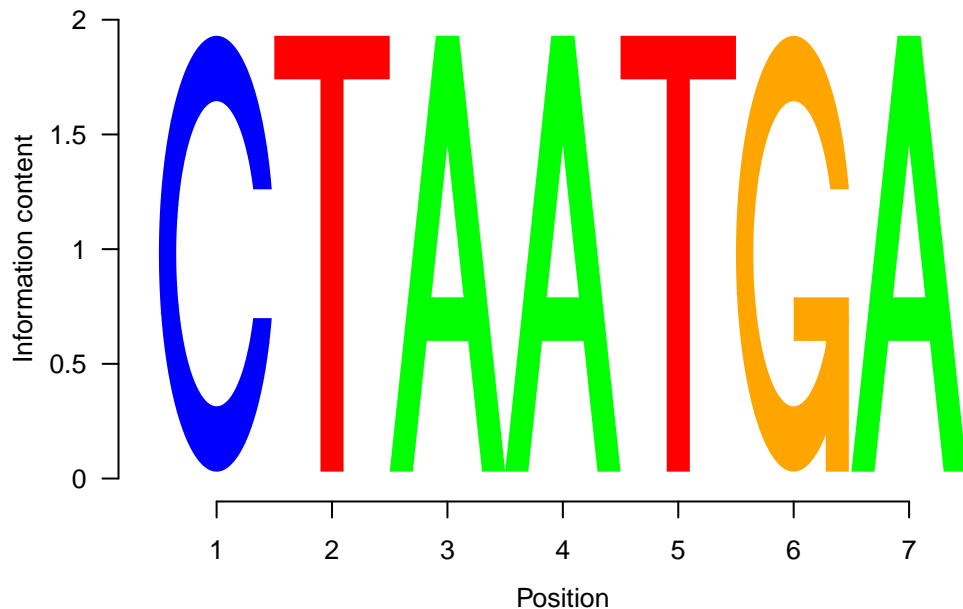
Genome: JASPAR_2014-ttk-MA0460.1



Genome: JASPAR_2014-twi-MA0249.1



Genome: JASPAR_2014-zen-MA0256.1



Genome: JASPAR_2014-z-MA0255.1

