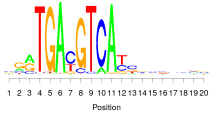


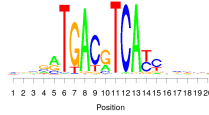
1 Evidence for (at least) one motif

ATF2 (1193 ChIP-Seq positives)

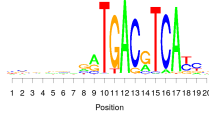
PWM



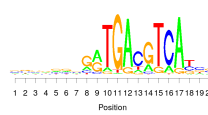
PMM1



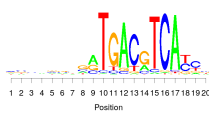
PMM2



PMM3

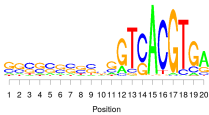


PMM4

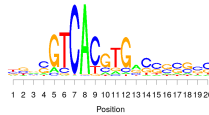


ATF3 (962 ChIP-Seq positives)

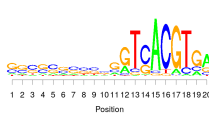
PWM



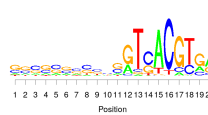
PMM1



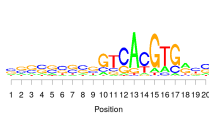
PMM2



PMM3

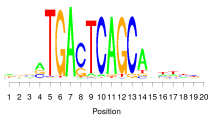


PMM4

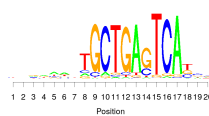


BACH1 (2292 ChIP-Seq positives)

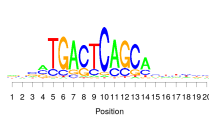
PWM



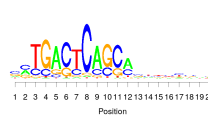
PMM1



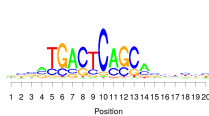
PMM2



PMM3

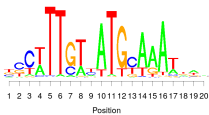


PMM4

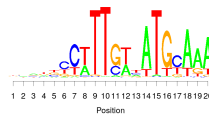


BCL11A (504 ChIP-Seq positives)

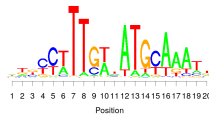
PWM



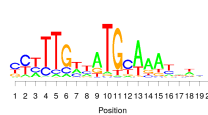
PMM1



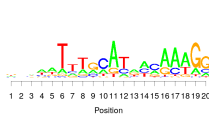
PMM2



PMM3

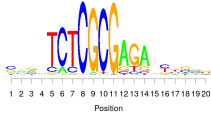


PMM4

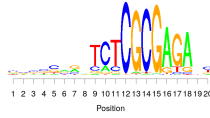


BRCA1 (405 ChIP-Seq positives)

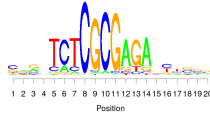
PWM



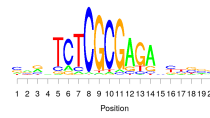
PMM1



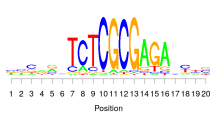
PMM2



PMM3

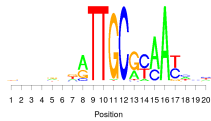


PMM4

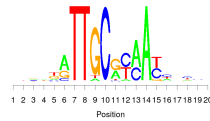


CEBPB (3112 ChIP-Seq positives)

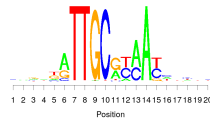
PWM



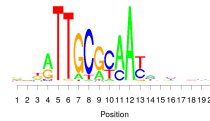
PMM1



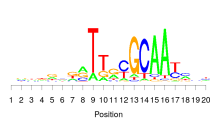
PMM2



PMM3

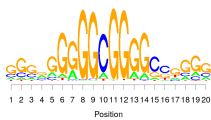


PMM4

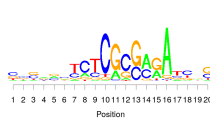


CHD2 (1368 ChIP-Seq positives)

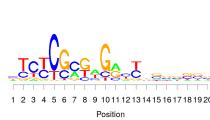
PWM



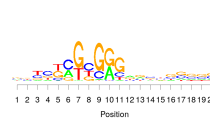
PMM1



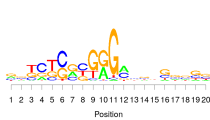
PMM2



PMM3

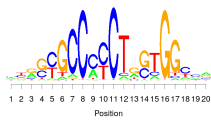


PMM4

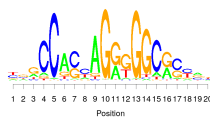


CTCF (10822 ChIP-Seq positives)

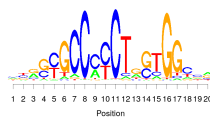
PWM



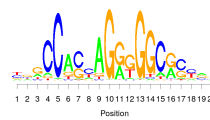
PMM1



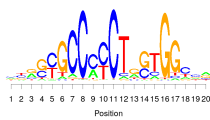
PMM2



PMM3

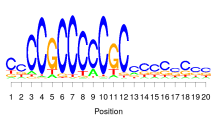


PMM4

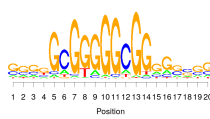


EGR1 (1749 ChIP-Seq positives)

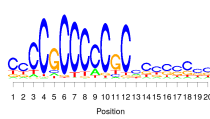
PWM



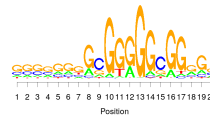
PMM1



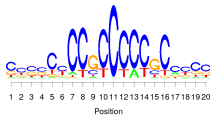
PMM2



PMM3

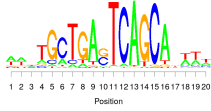


PMM4

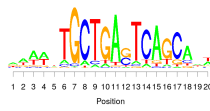


MAFK (2284 ChIP-Seq positives)

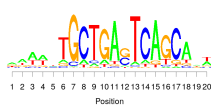
PWM



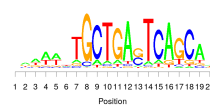
PMM1



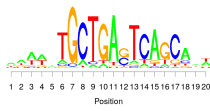
PMM2



PMM3

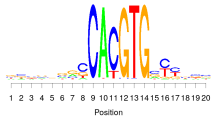


PMM4

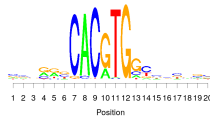


MAX (2225 ChIP-Seq positives)

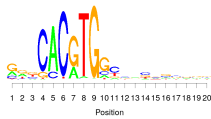
PWM



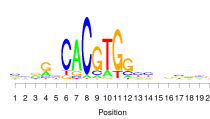
PMM1



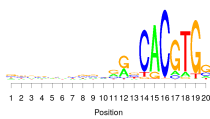
PMM2



PMM3

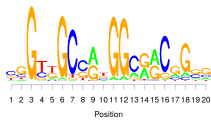


PMM4

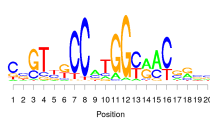


MXI1 (1271 ChIP-Seq positives)

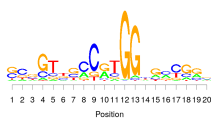
PWM



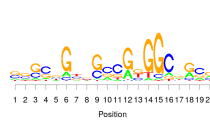
PMM1



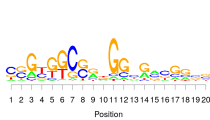
PMM2



PMM3

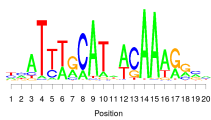


PMM4

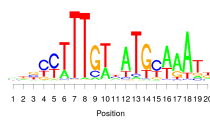


NANOG (1095 ChIP-Seq positives)

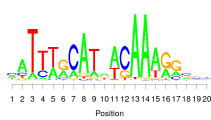
PWM



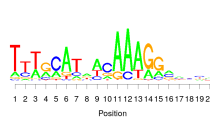
PMM1



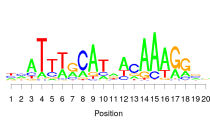
PMM2



PMM3

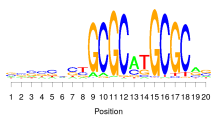


PMM4

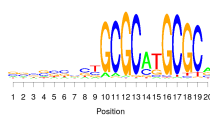


NRF1 (903 ChIP-Seq positives)

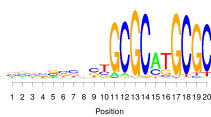
PWM



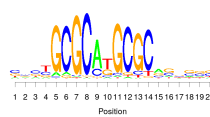
PMM1



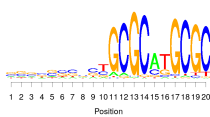
PMM2



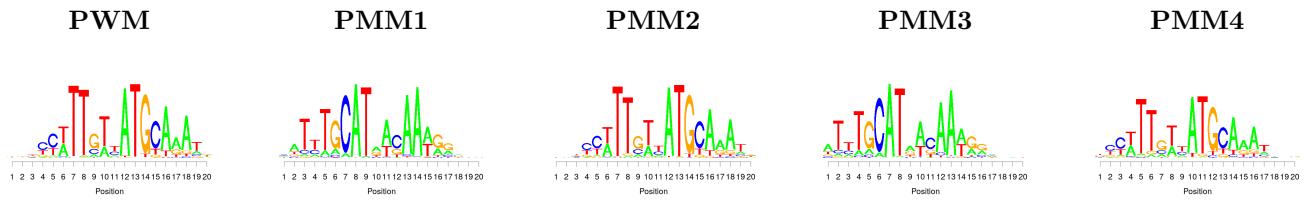
PMM3



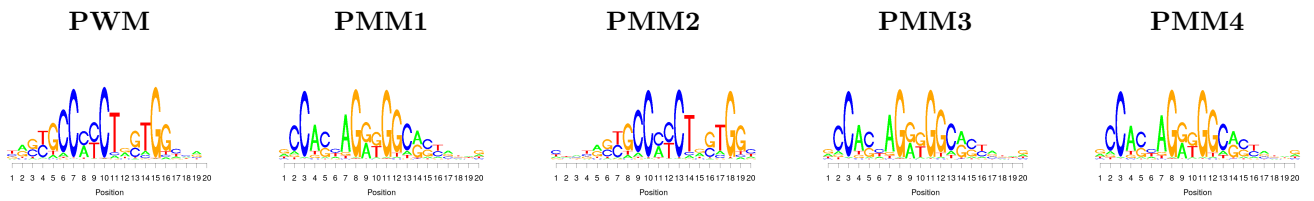
PMM4



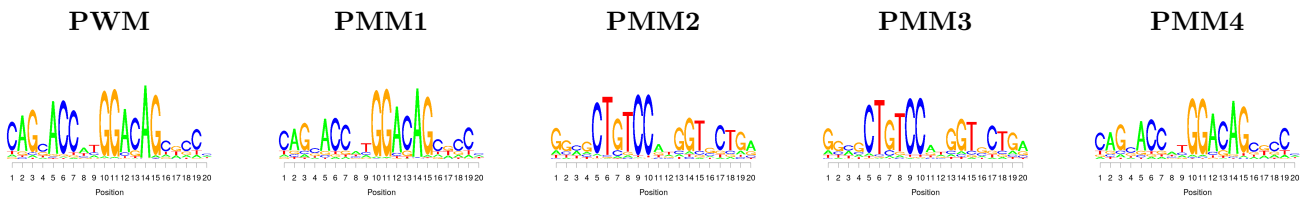
POU5F1 (800 ChIP-Seq positives)



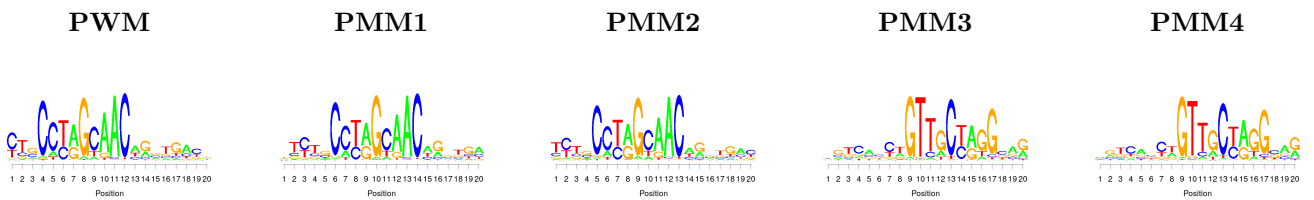
RAD21 (15136 ChIP-Seq positives)



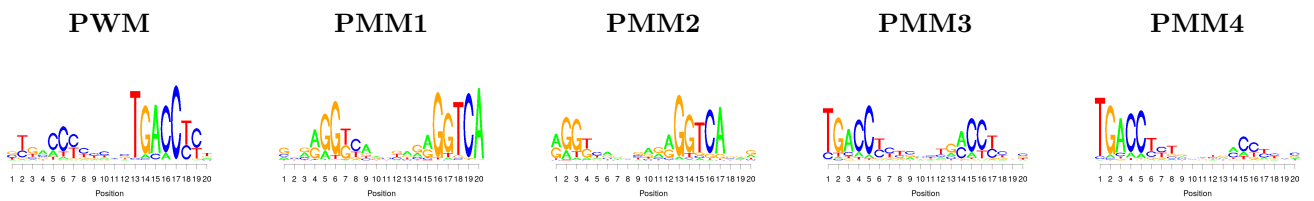
REST (2656 ChIP-Seq positives)



RFX5 (338 ChIP-Seq positives)

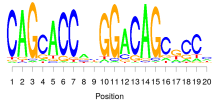


RXRA (262 ChIP-Seq positives)

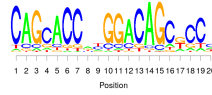


SIN3AK20 (4260 ChIP-Seq positives)

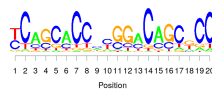
PWM



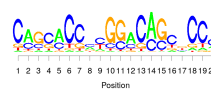
PMM1



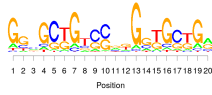
PMM2



PMM3

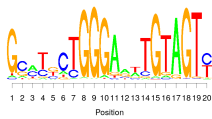


PMM4

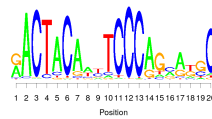


SIX5 (684 ChIP-Seq positives)

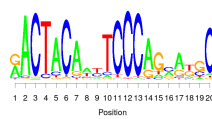
PWM



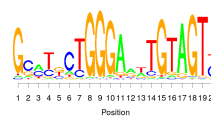
PMM1



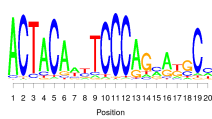
PMM2



PMM3

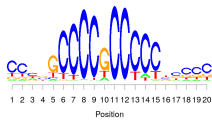


PMM4

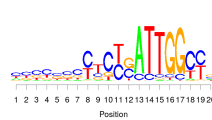


SP1 (3009 ChIP-Seq positives)

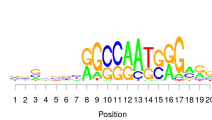
PWM



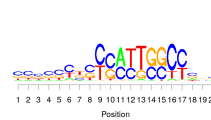
PMM1



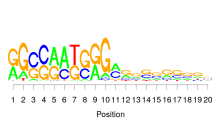
PMM2



PMM3

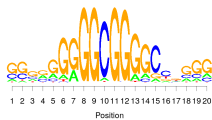


PMM4

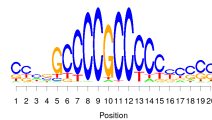


SP2 (492 ChIP-Seq positives)

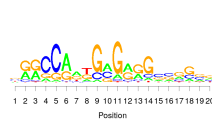
PWM



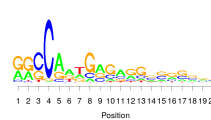
PMM1



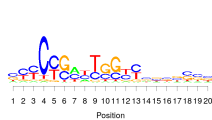
PMM2



PMM3

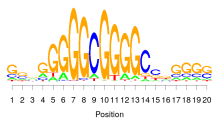


PMM4

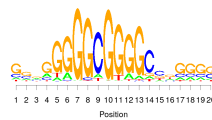


SP4 (1150 ChIP-Seq positives)

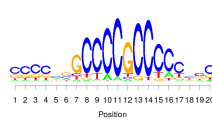
PWM



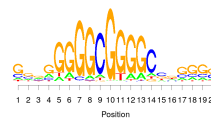
PMM1



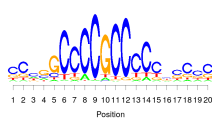
PMM2



PMM3

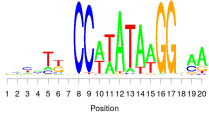


PMM4

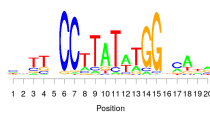


SRF (1020 ChIP-Seq positives)

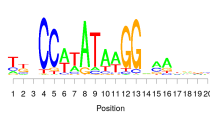
PWM



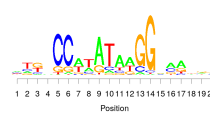
PMM1



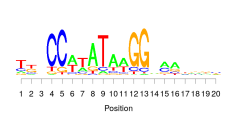
PMM2



PMM3

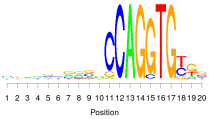


PMM4

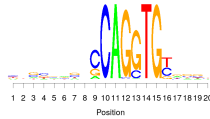


TCF12 (1567 ChIP-Seq positives)

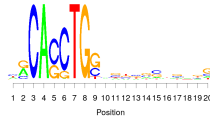
PWM



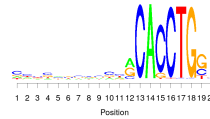
PMM1



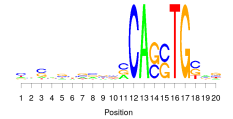
PMM2



PMM3

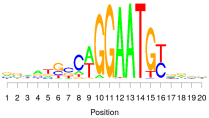


PMM4

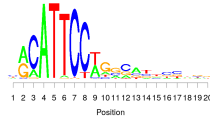


TEAD4 (3972 ChIP-Seq positives)

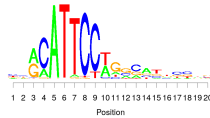
PWM



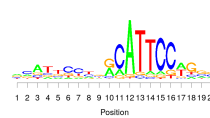
PMM1



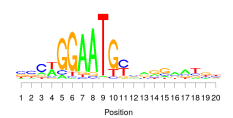
PMM2



PMM3

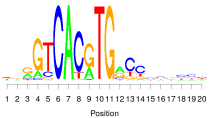


PMM4

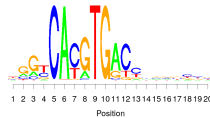


USF1 (5208 ChIP-Seq positives)

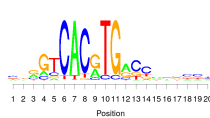
PWM



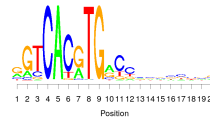
PMM1



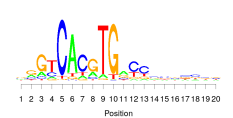
PMM2



PMM3

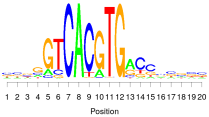


PMM4

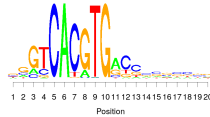


USF2 (1391 ChIP-Seq positives)

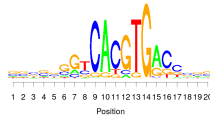
PWM



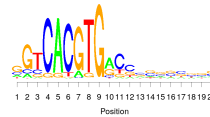
PMM1



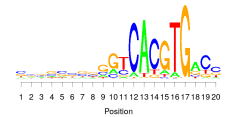
PMM2



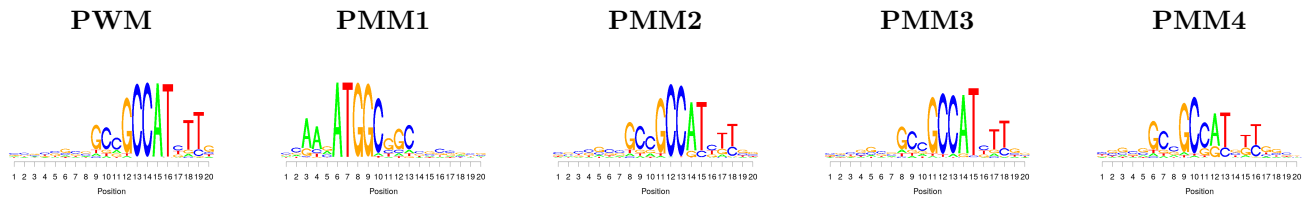
PMM3



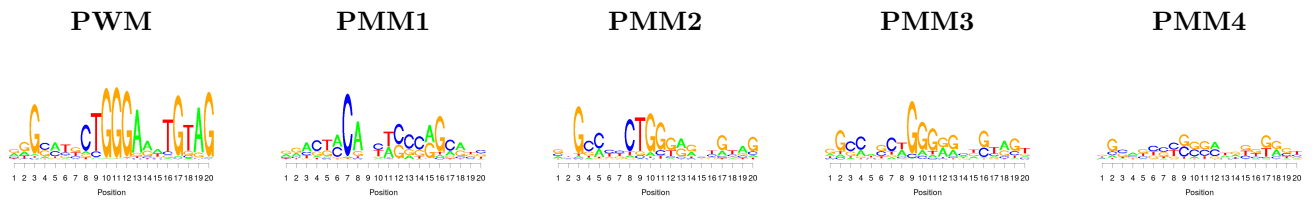
PMM4



YY1 (3646 ChIP-Seq positives)

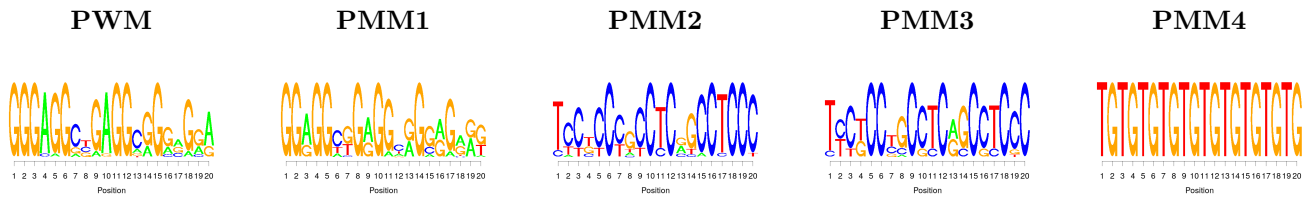


ZNF143 (6134 ChIP-Seq positives)

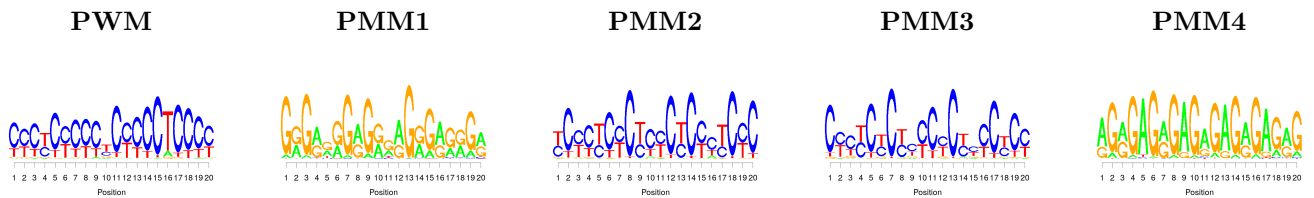


2 No meaningful motif identified

CHD1 (1380 ChIP-Seq positives)

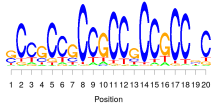


CTBP2 (1418 ChIP-Seq positives)

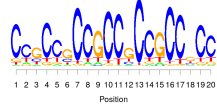


TAF7 (2072 ChIP-Seq positives)

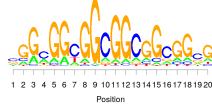
PWM



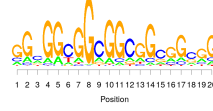
PMM1



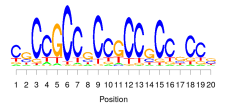
PMM2



PMM3

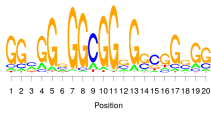


PMM4

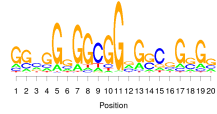


TBP (3429 ChIP-Seq positives)

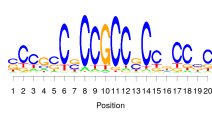
PWM



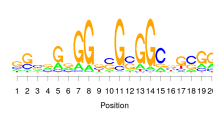
PMM1



PMM2



PMM3



PMM4

