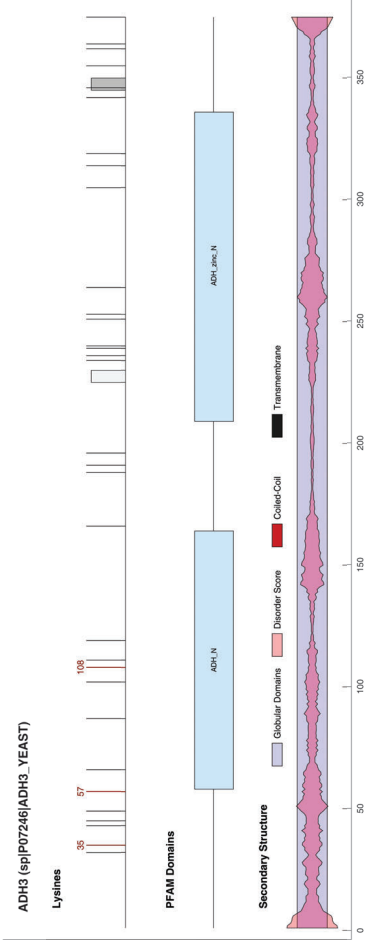
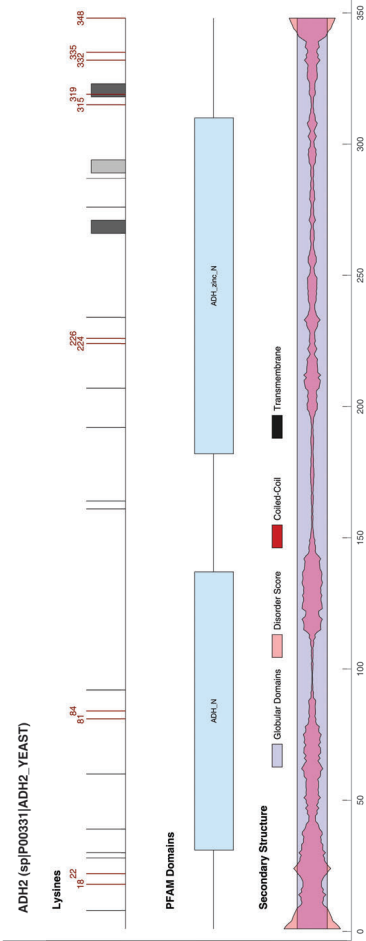
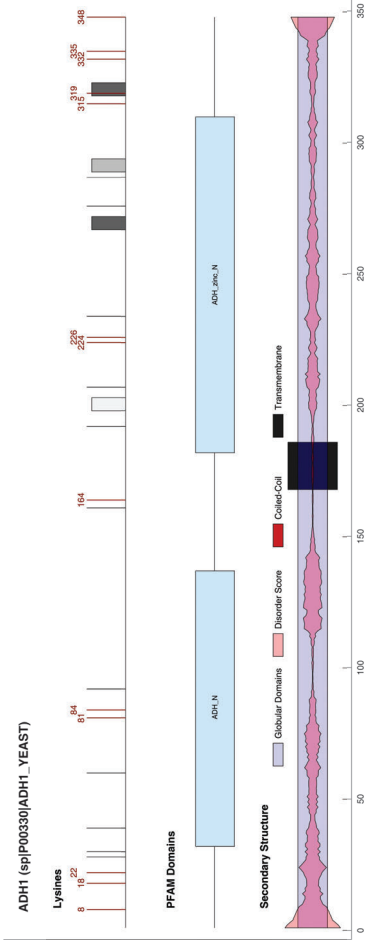
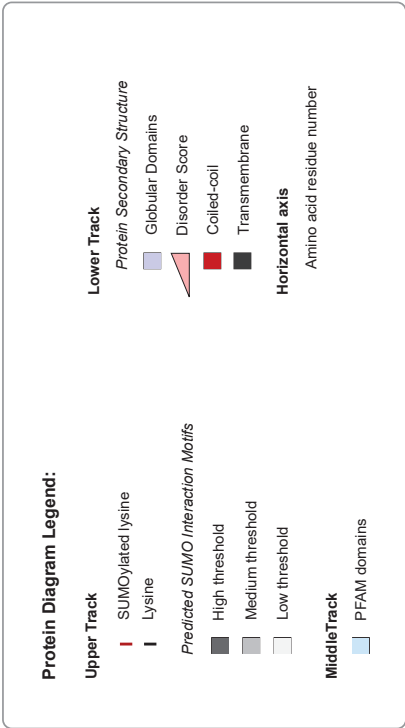
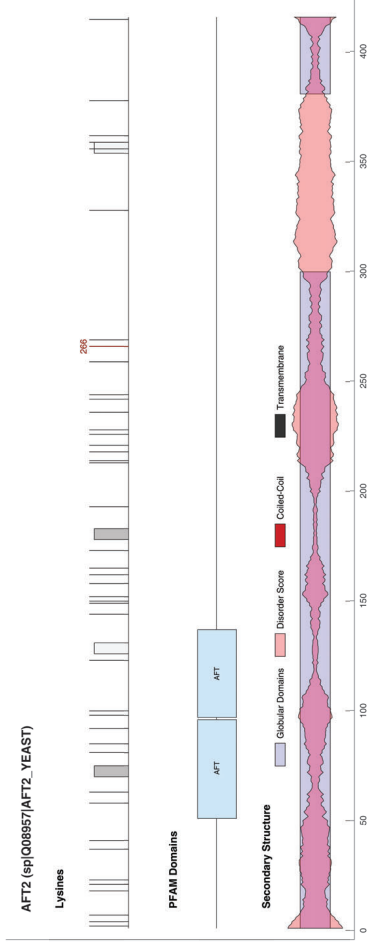
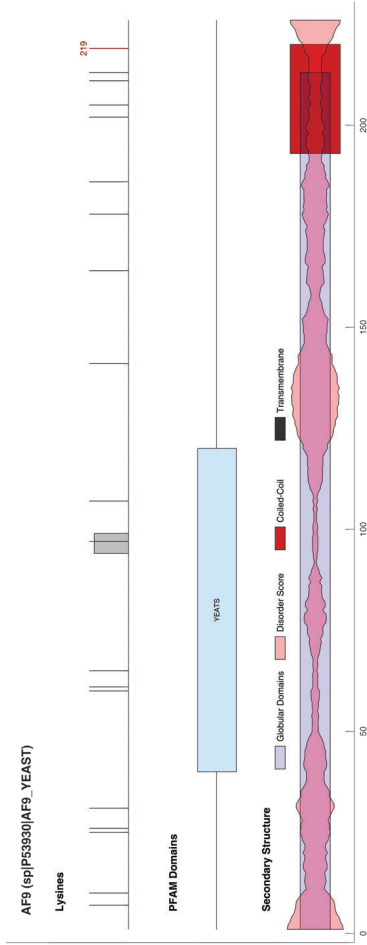
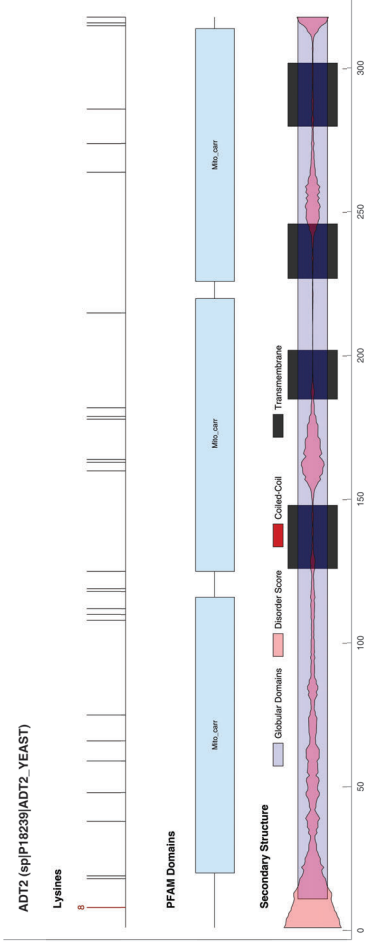
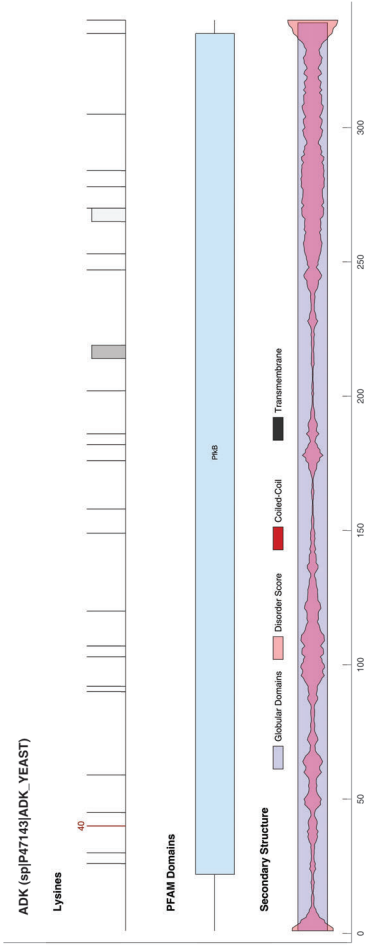
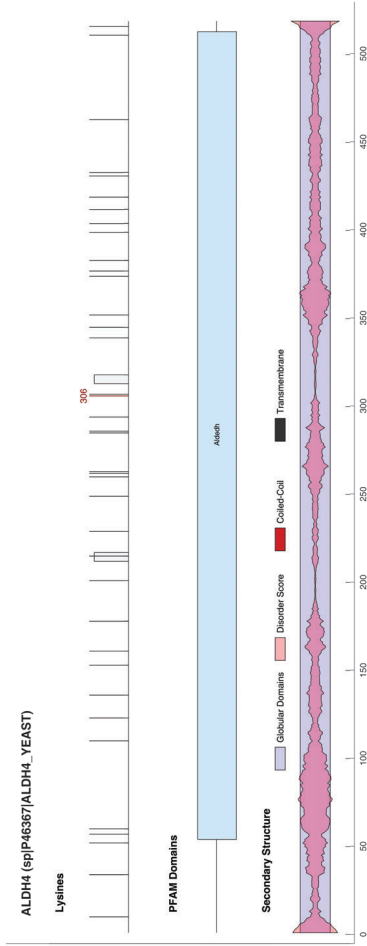
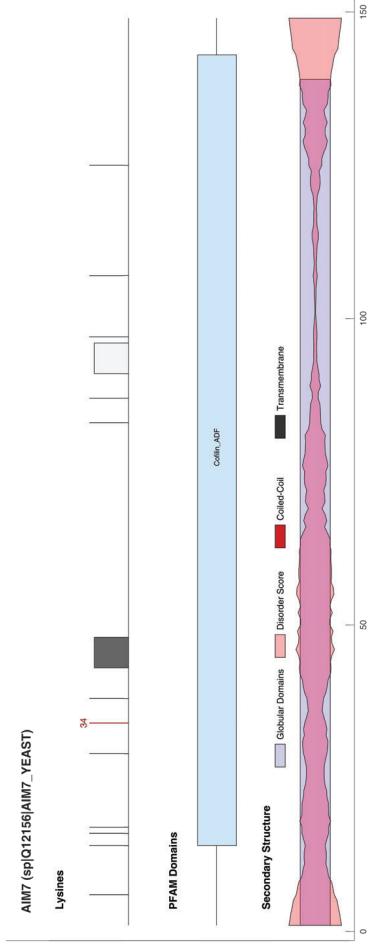
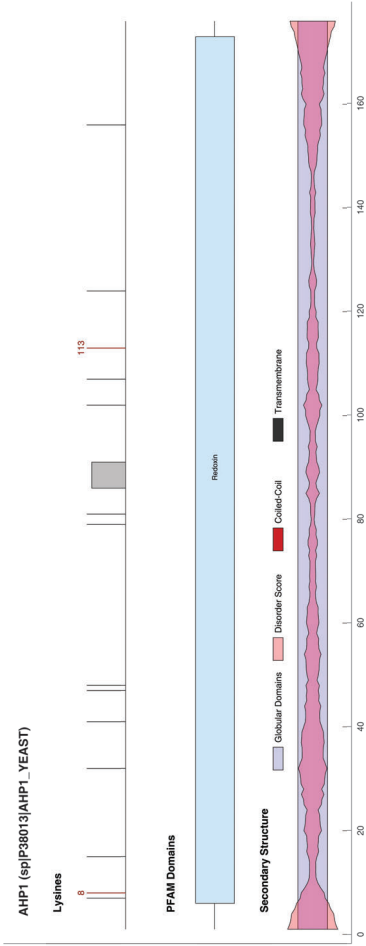
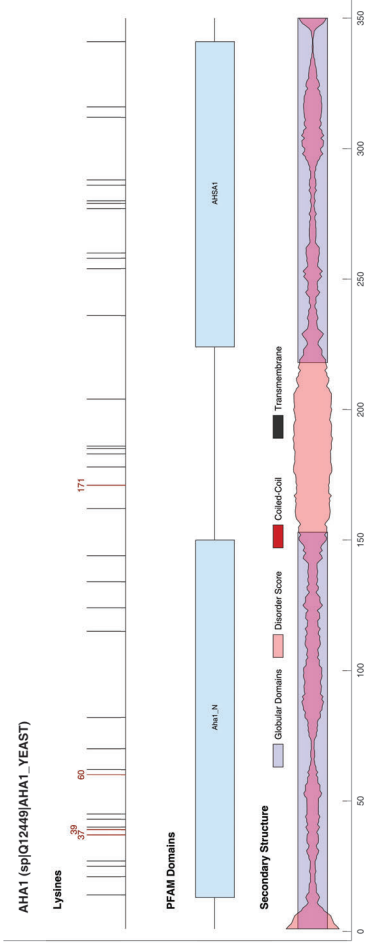
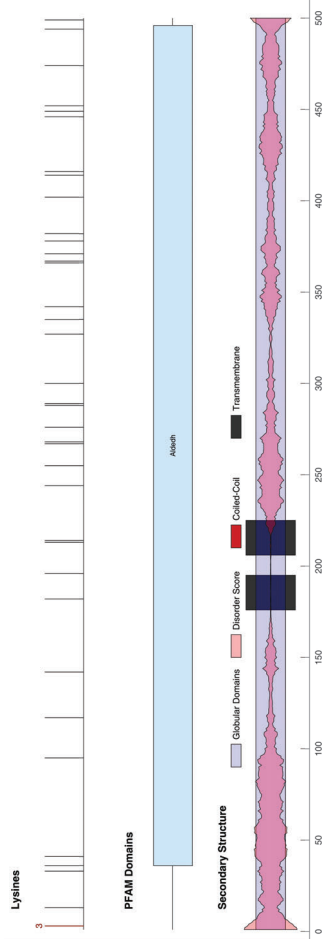




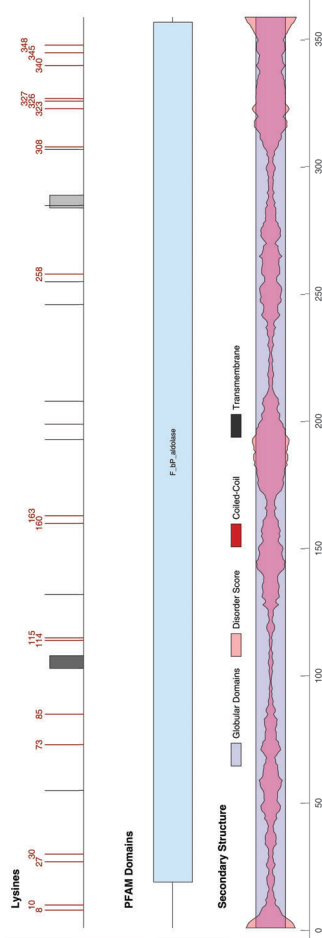




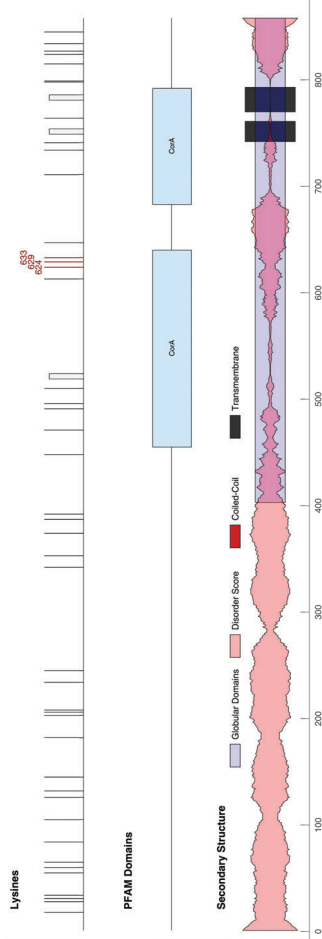
ALDH6 (sp|P54115|ALDH6_YEAST)



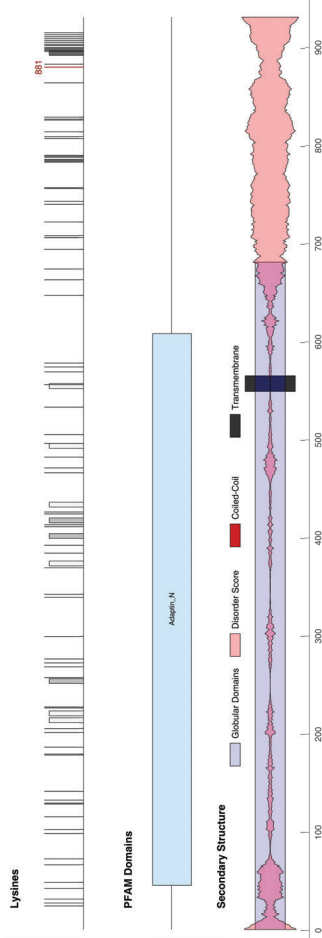
ALF (sp|P14540|ALF_YEAST)

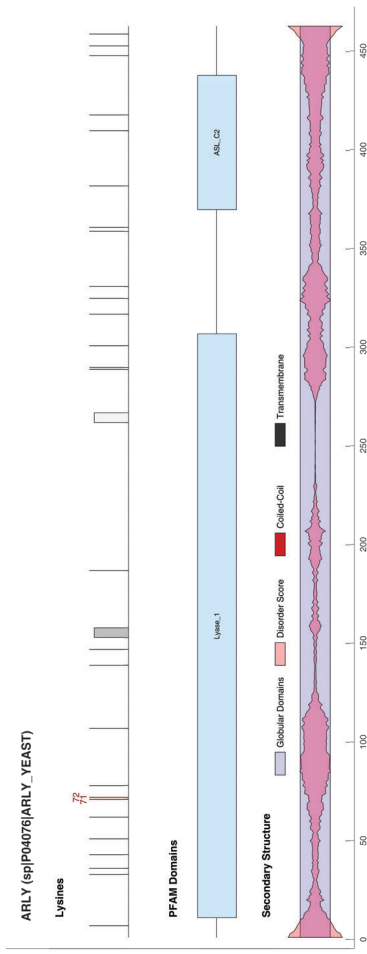
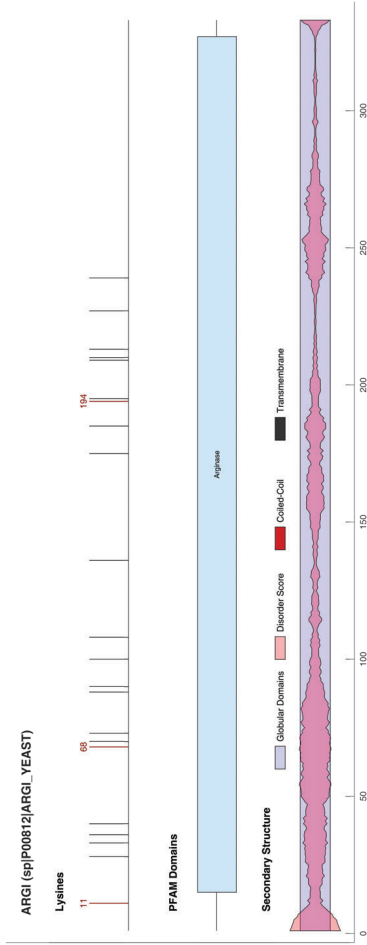
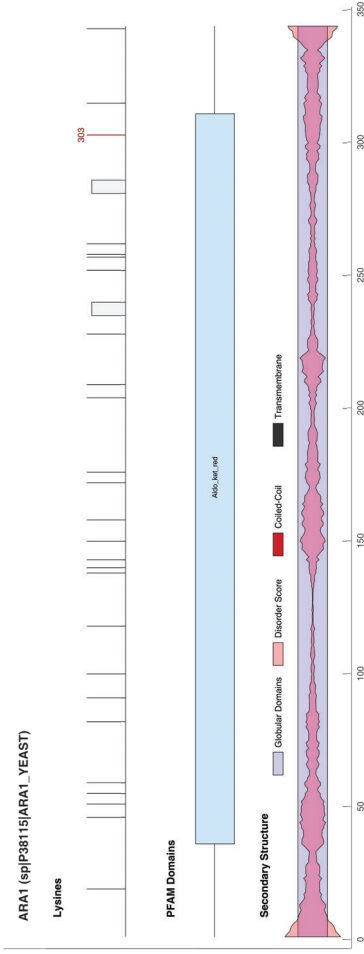
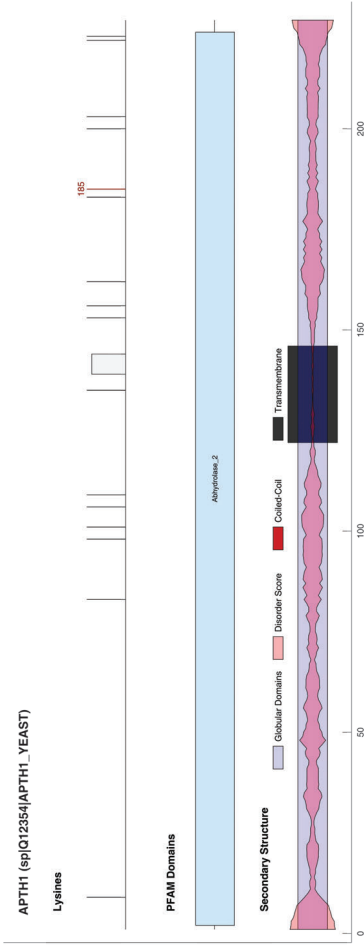


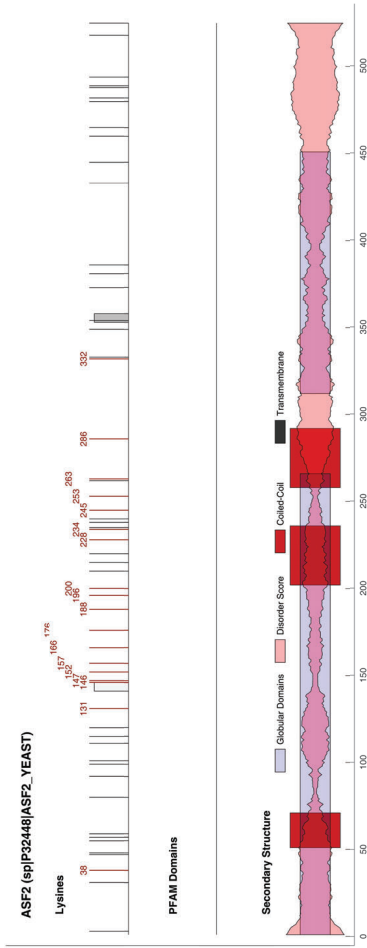
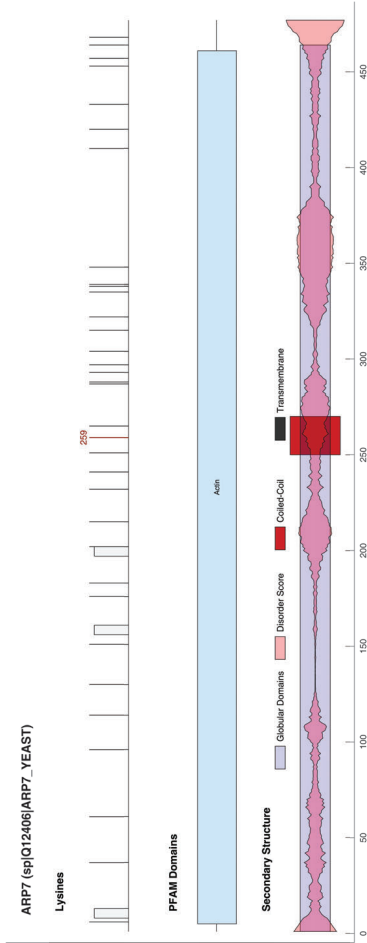
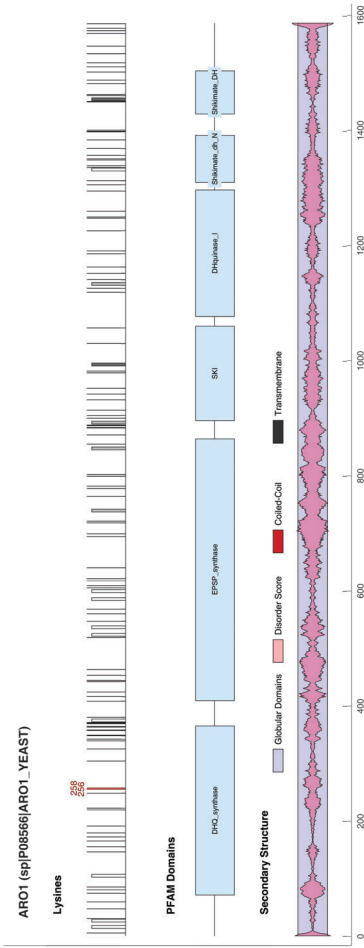
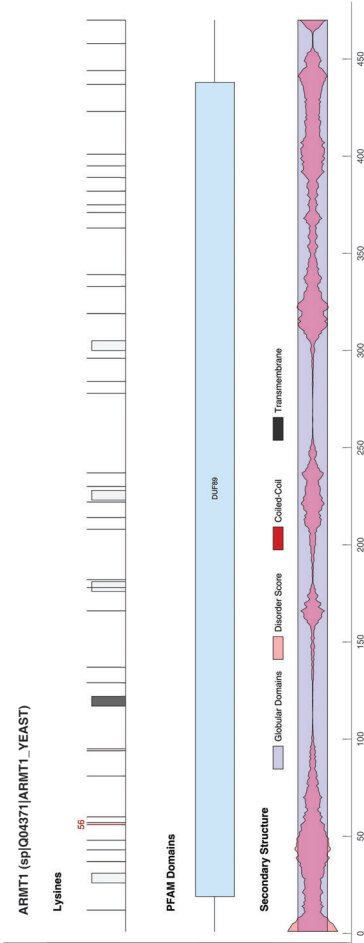
ALR2 (sp|P43553|ALR2_YEAST)

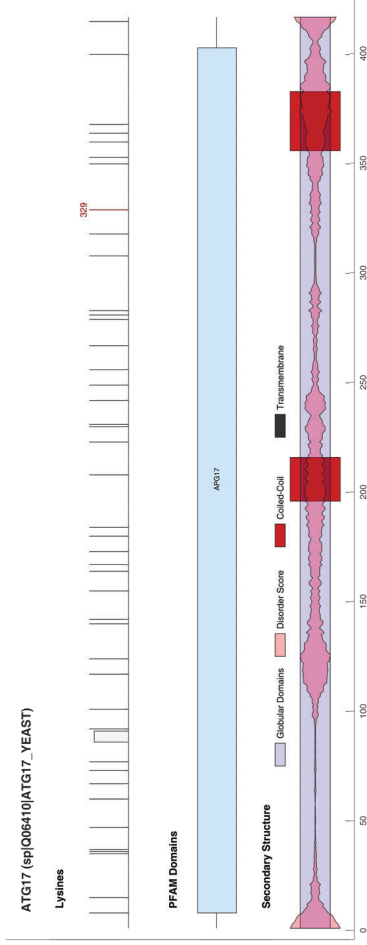
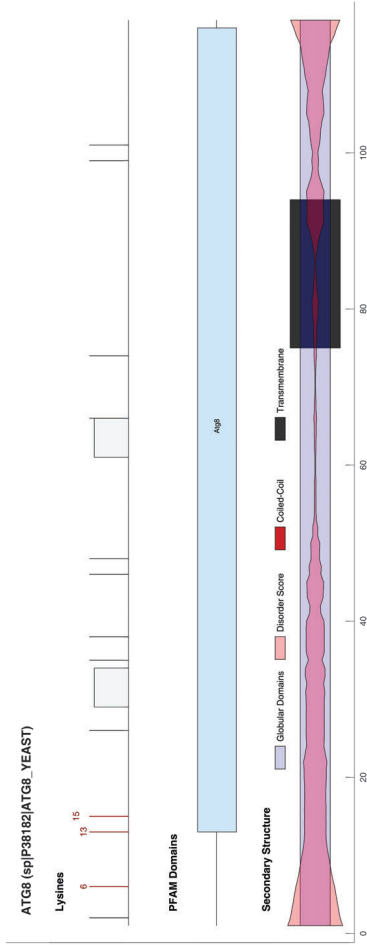
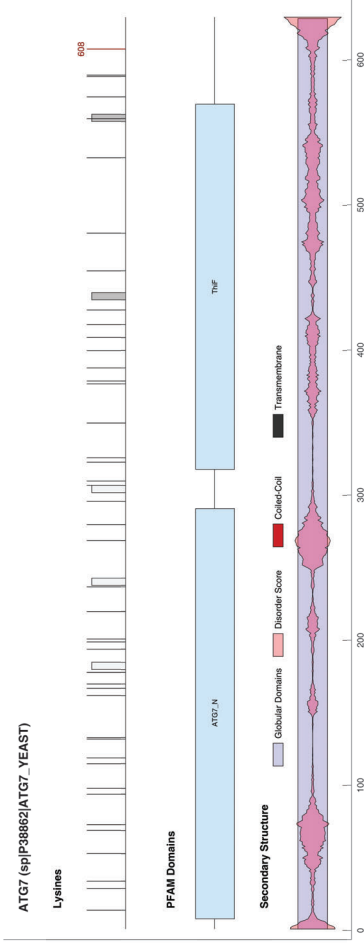
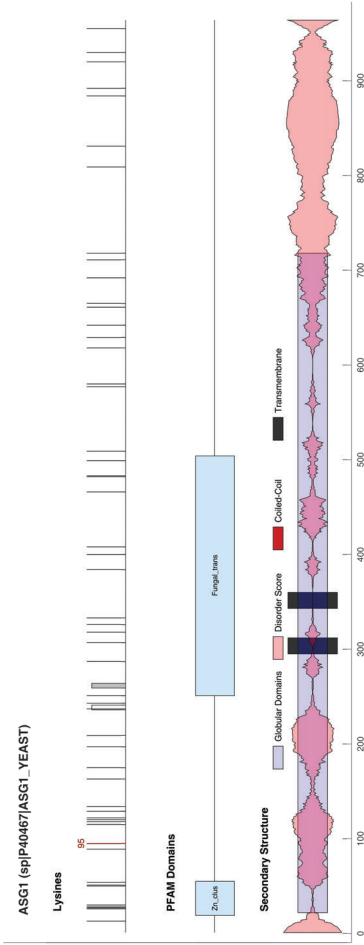


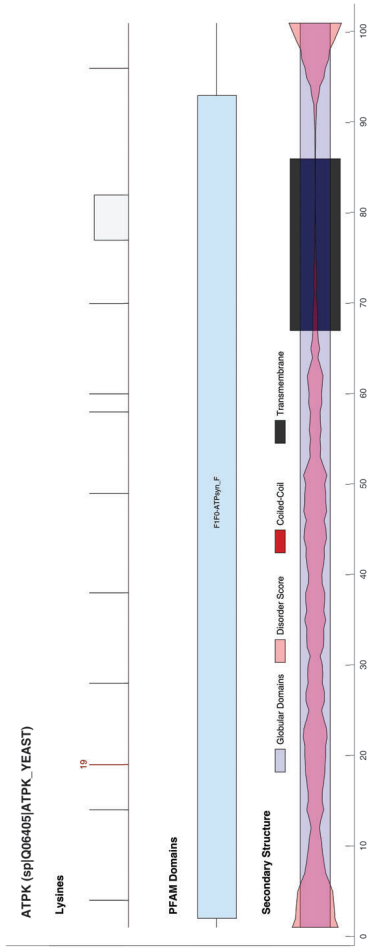
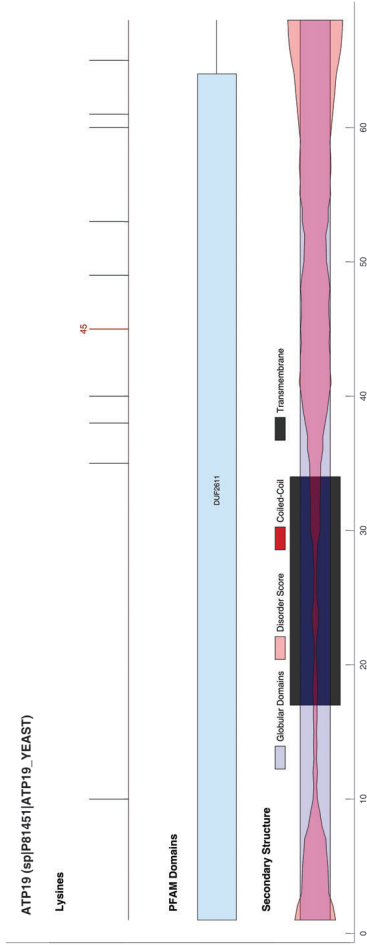
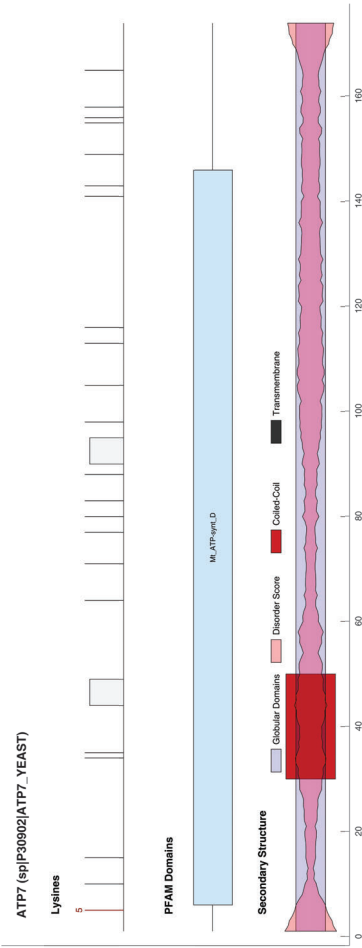
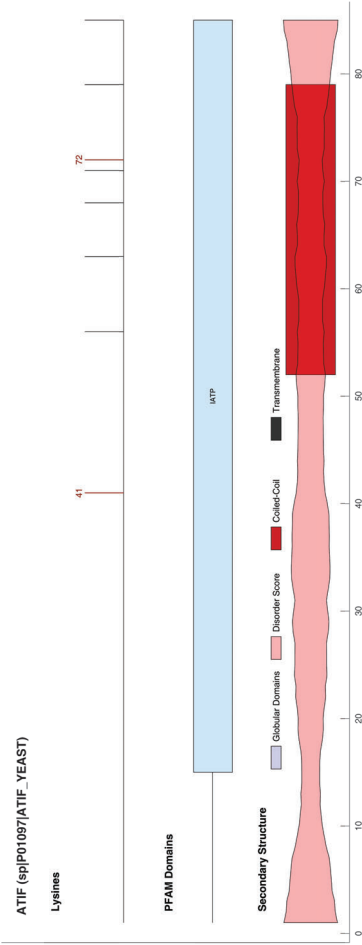
AP3D (sp|Q08951|AP3D_YEAST)

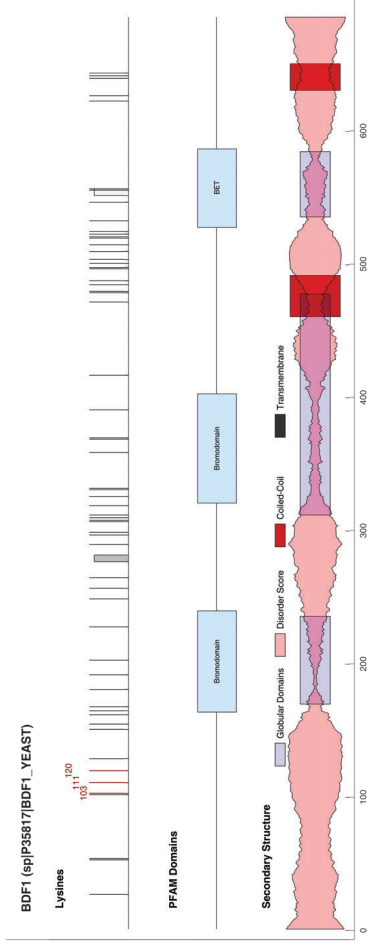
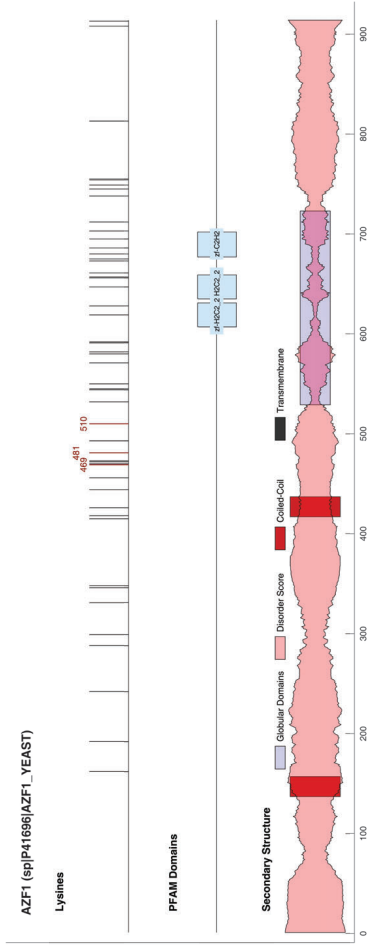
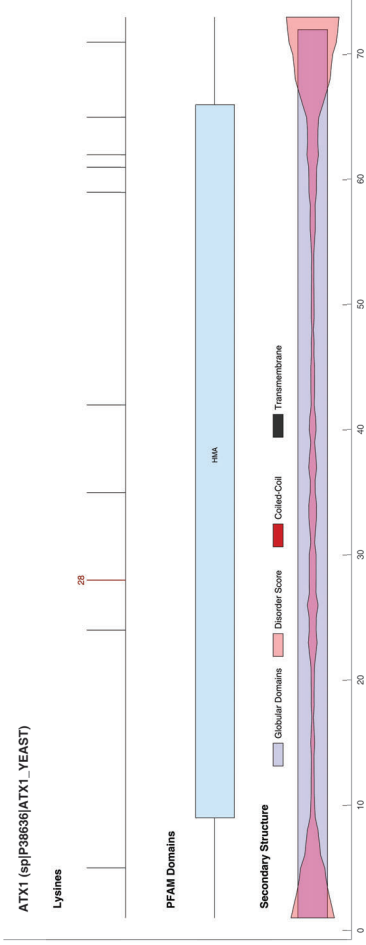
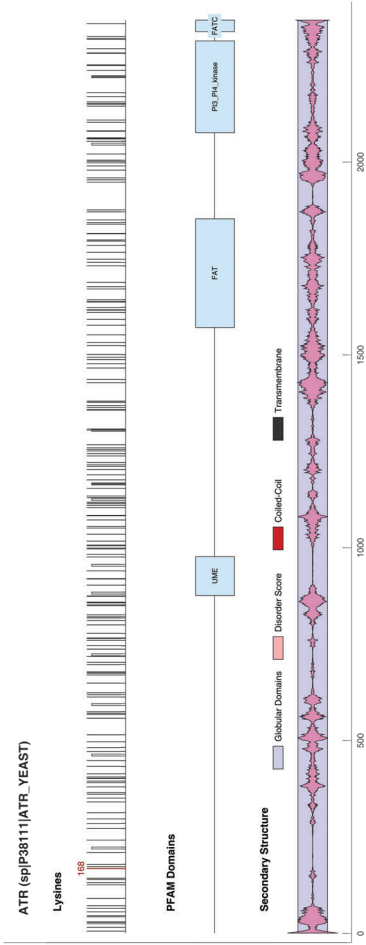




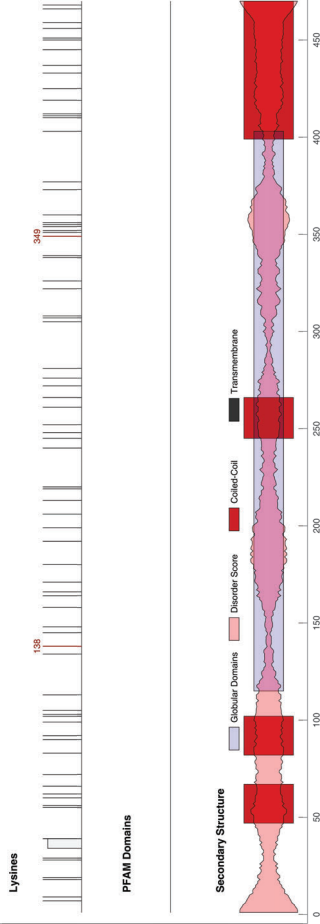




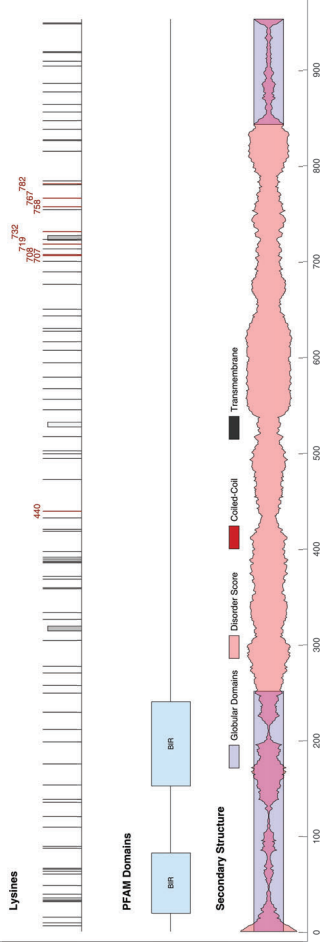




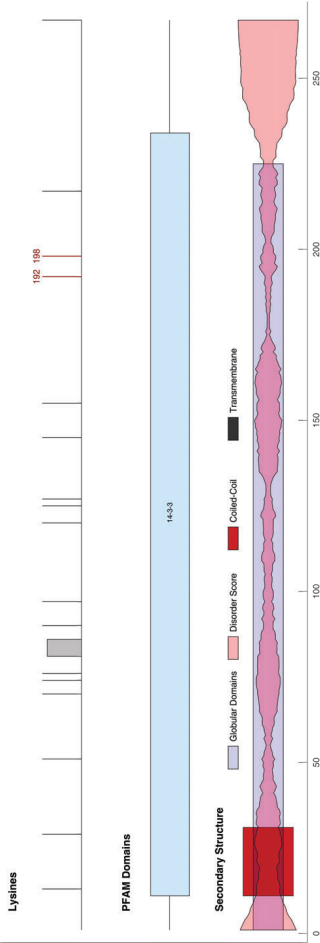
BFR1 (sp|P38934|BFR1_YEAST)



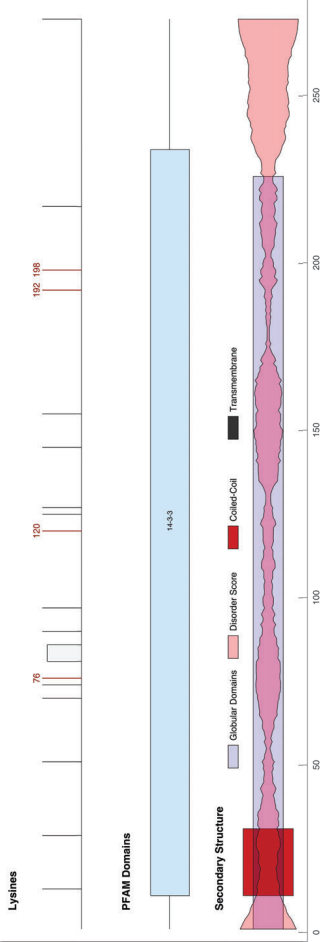
BIR1 (sp|P47134|BIR1_YEAST)

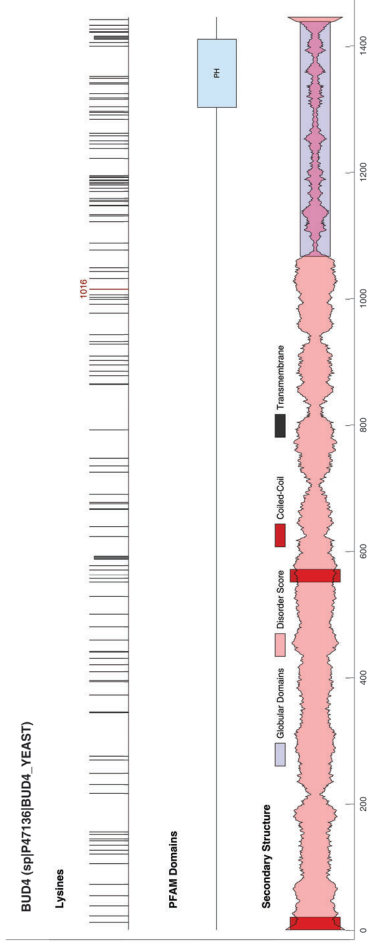
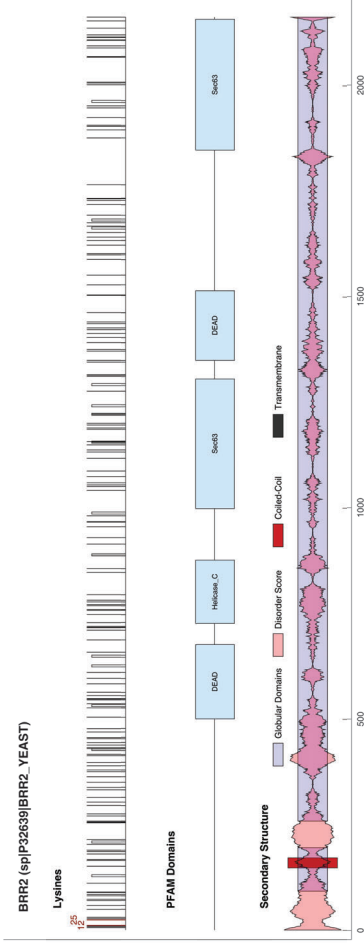
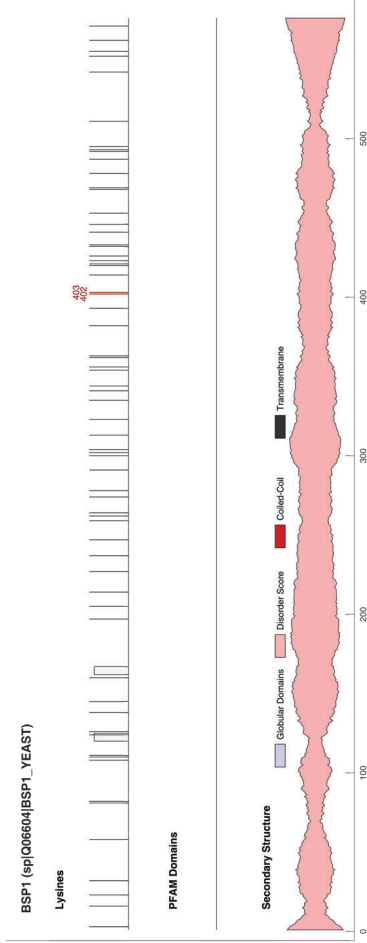
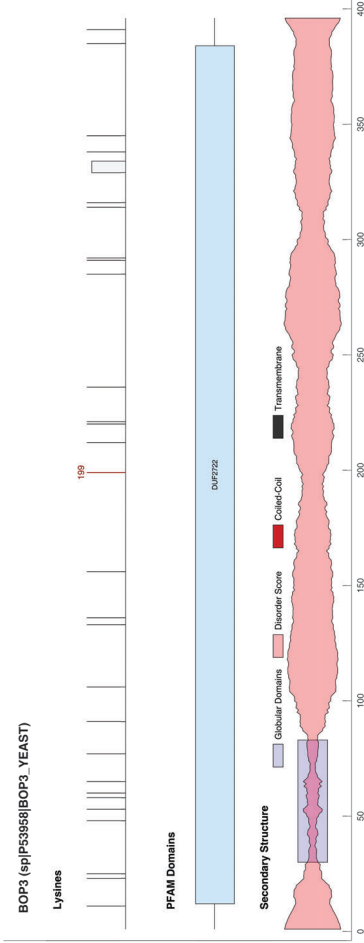


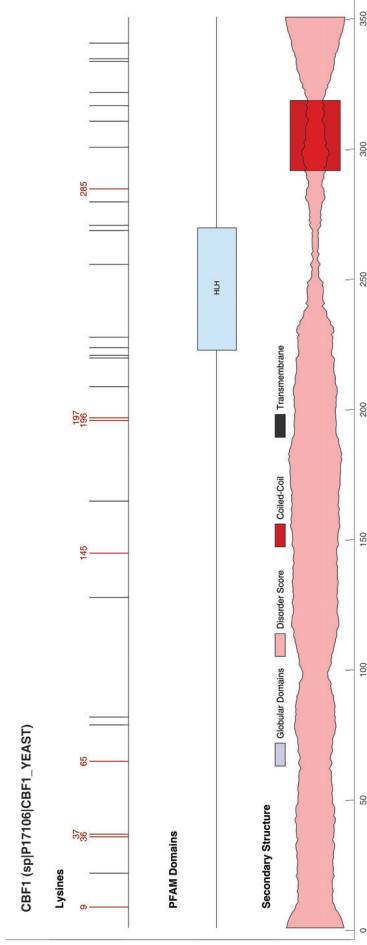
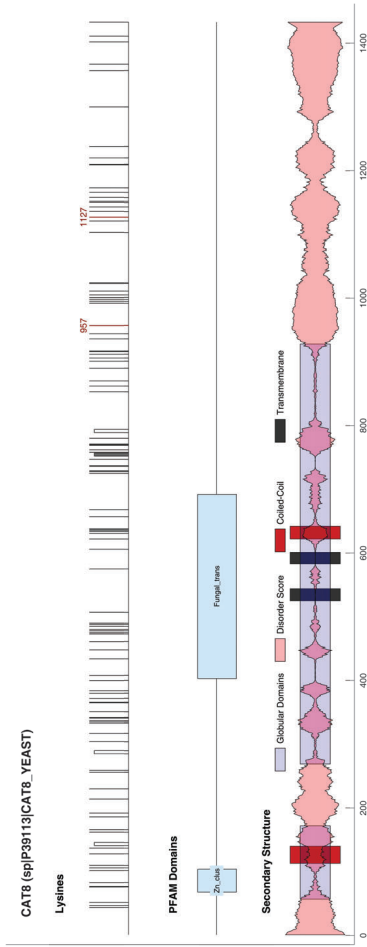
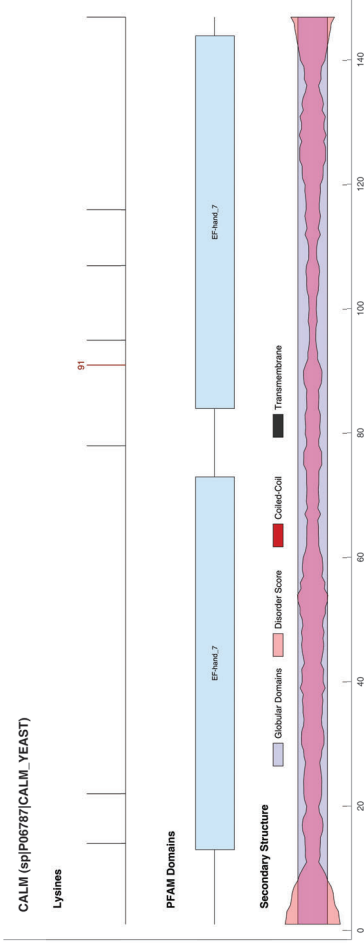
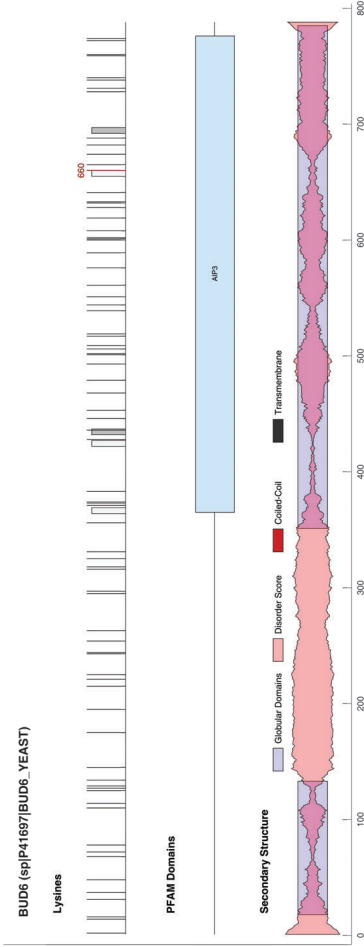
BMH1 (sp|P29311|BMH1_YEAST)

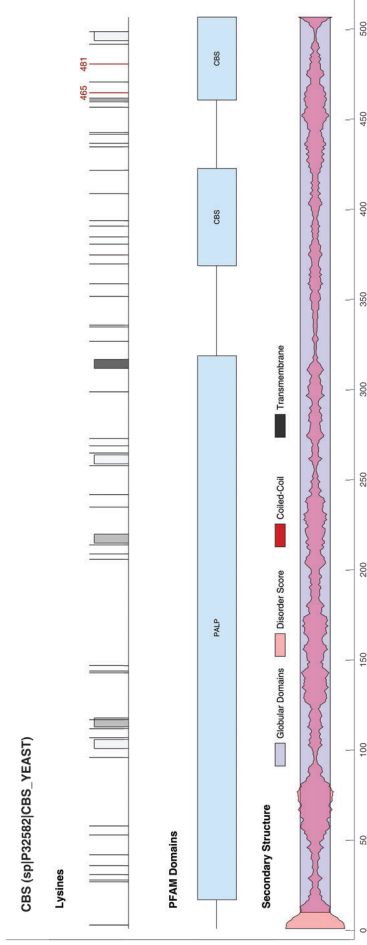
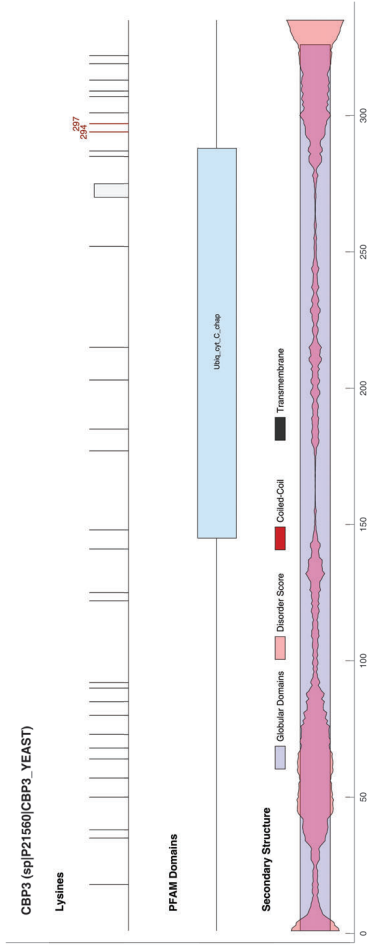
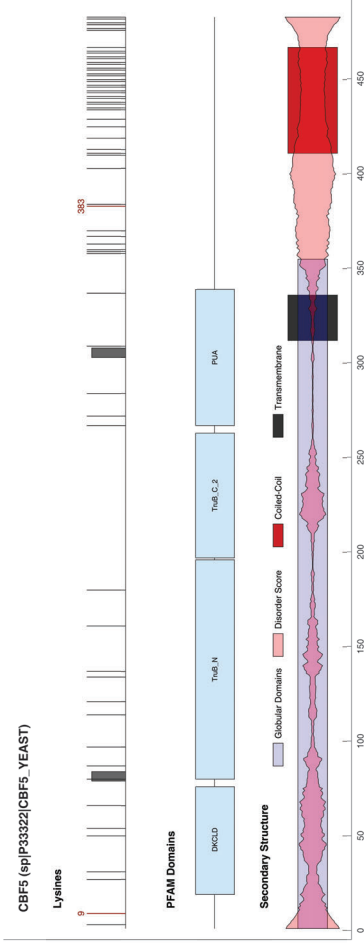
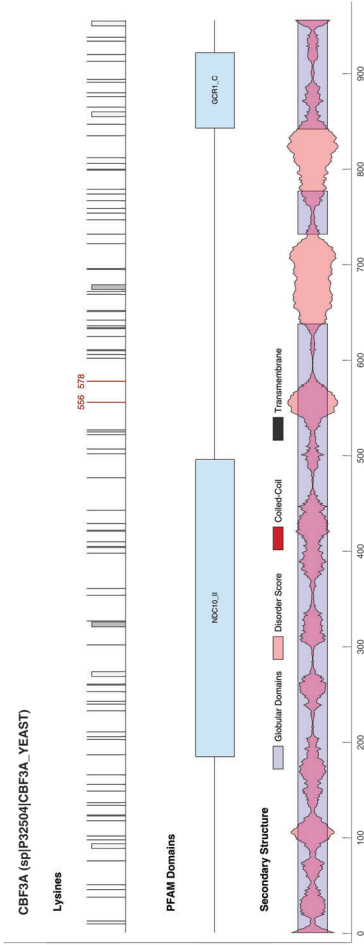


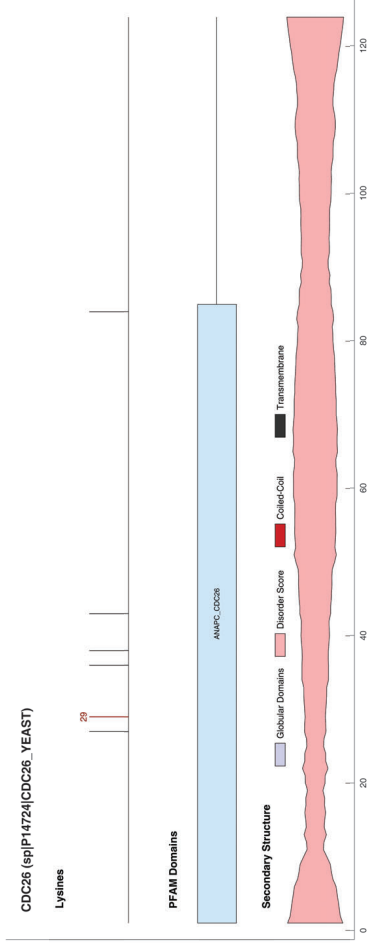
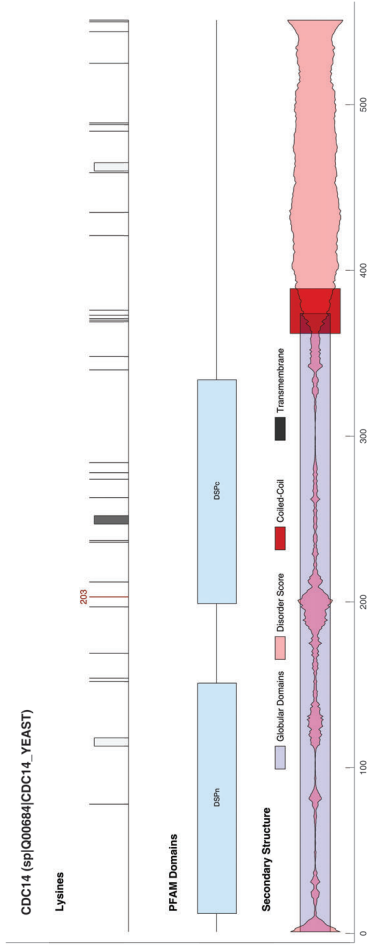
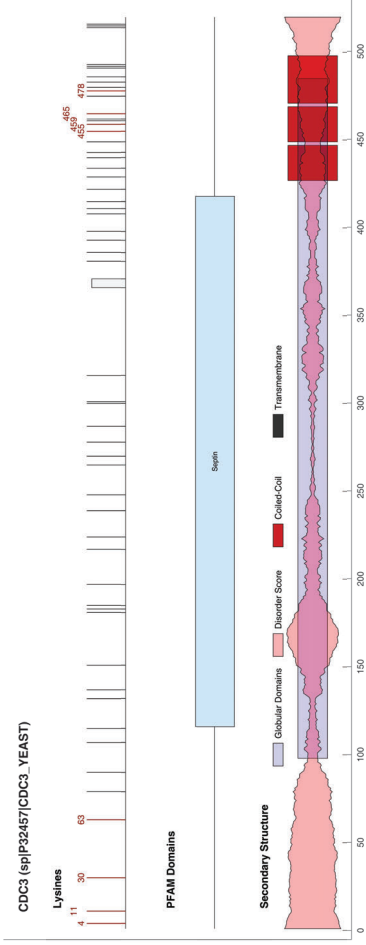
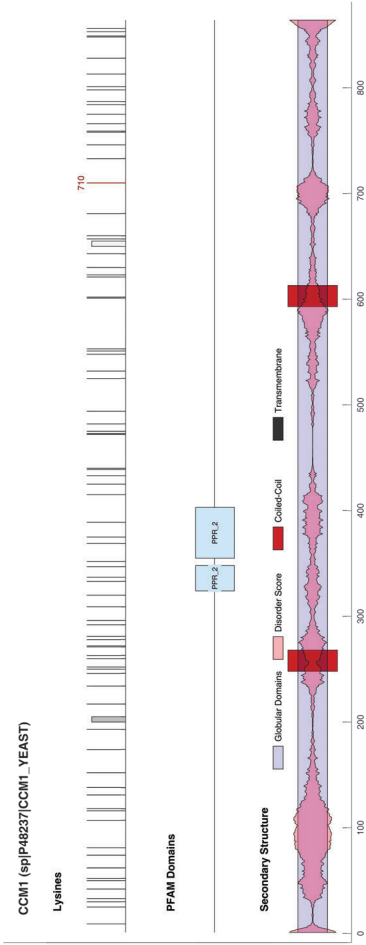
BMH2 (sp|P34730|BMH2_YEAST)

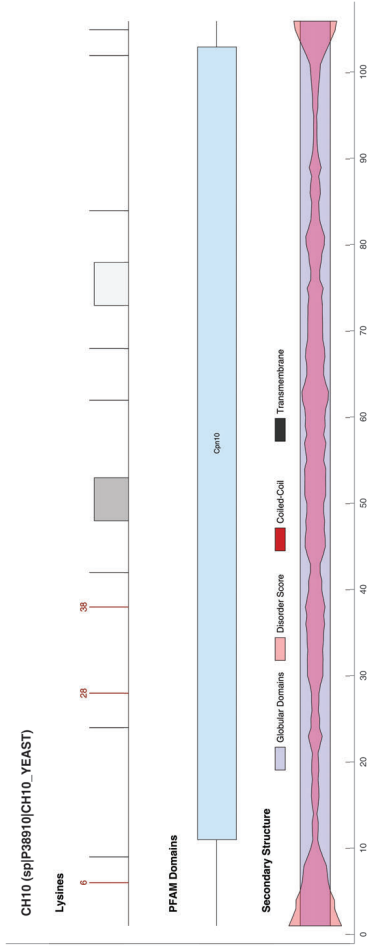
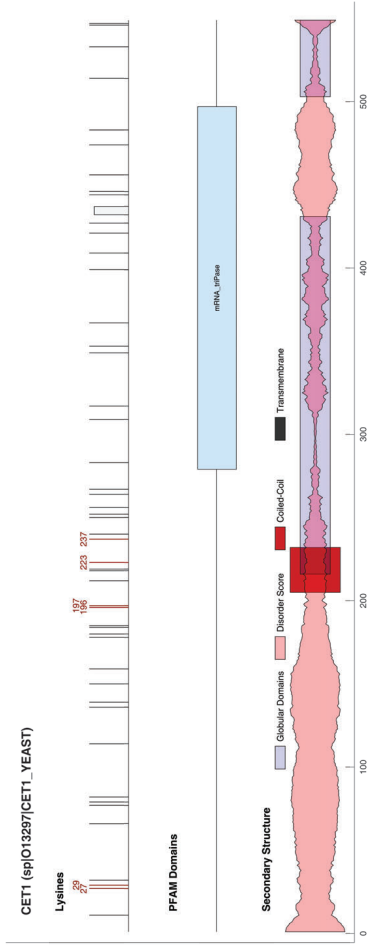
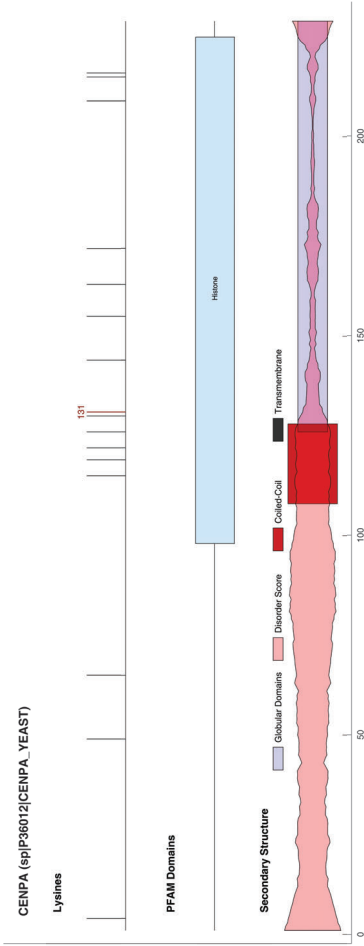
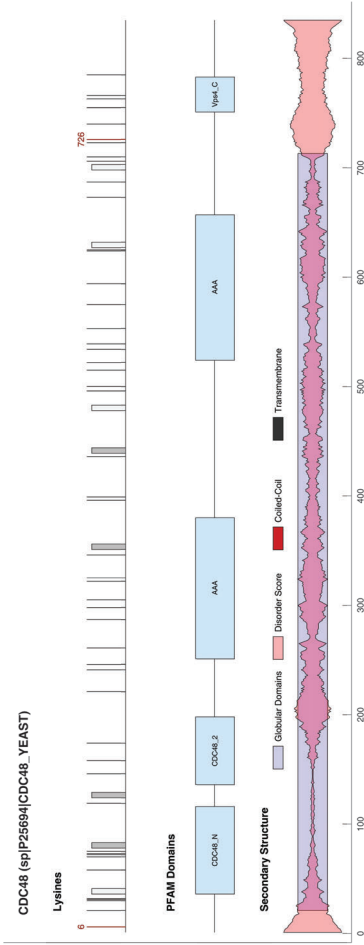












CHA4 (sp|P43634|CHA4_YEAST)

Lysines

ZNF132

PFAM Domains

- Zn_Zf
- Fingrid_Lines

Secondary Structure

- Globular Domains
- Disorder Score
- Coiled-Coil
- Transmembrane

CHD1 (sp|P32657|CHD1_YEAST)

Lysines 122

PFAM Domains

DUF3038

Helix_C

SWP2_N

Coiled-Coil

Transmembrane

Secondary Structure

Alpha-Helix (red)

Beta-Strand (blue)

Loop (grey)

CISY1 (sp|P00890|CISY1_YEAST)

Lysines

CLC1 (sp|P17891|CLC1_YEAST)

Lysines

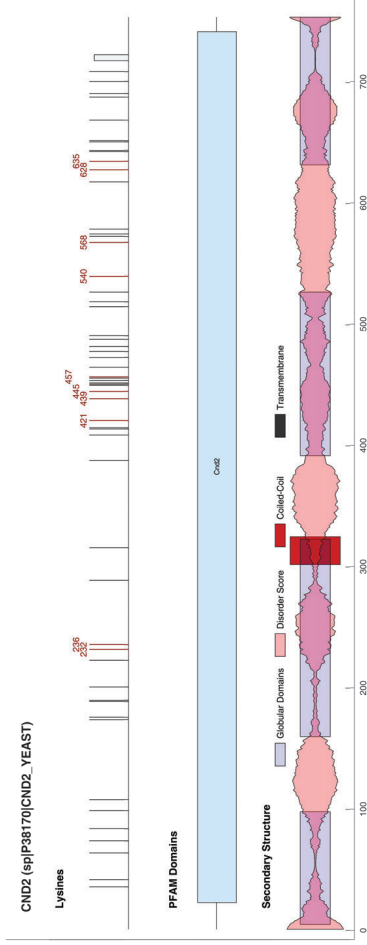
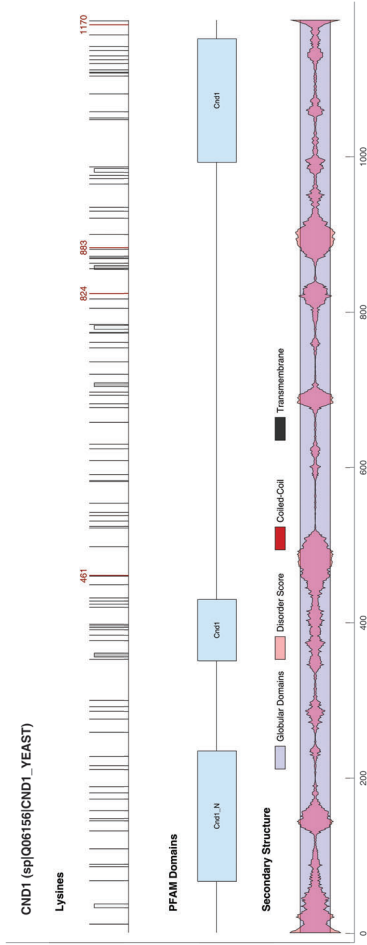
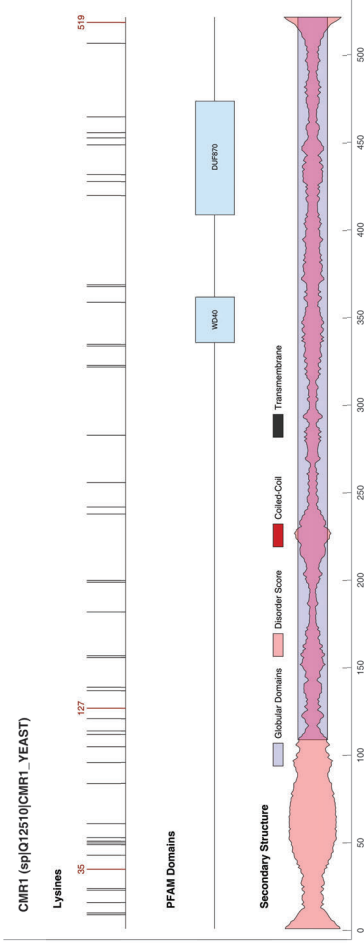
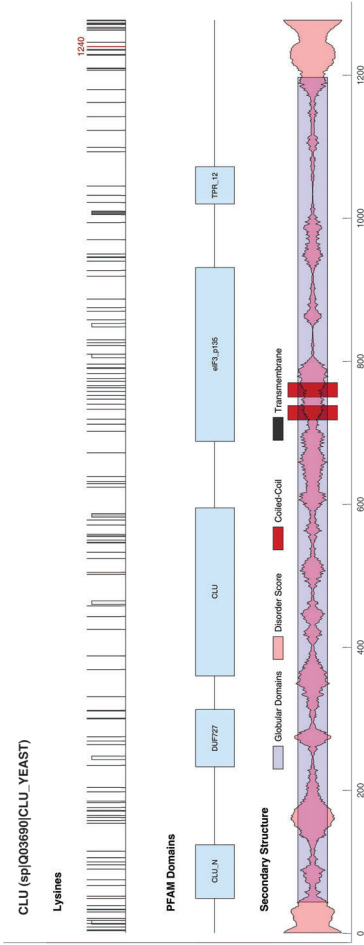
PFAM Domains

Ctrm1p_ch

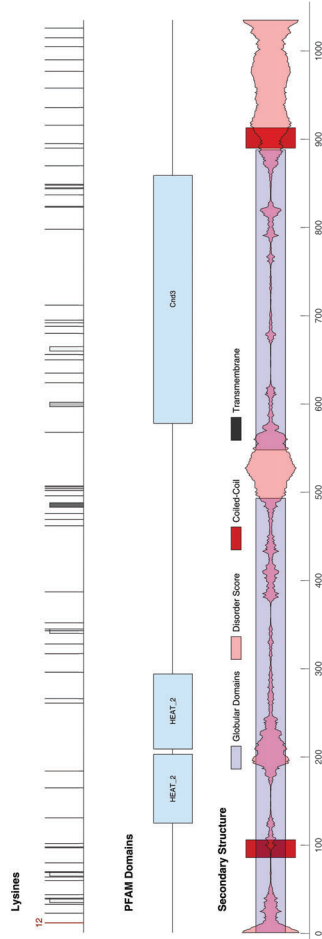
Secondary Structure

Disorder Score

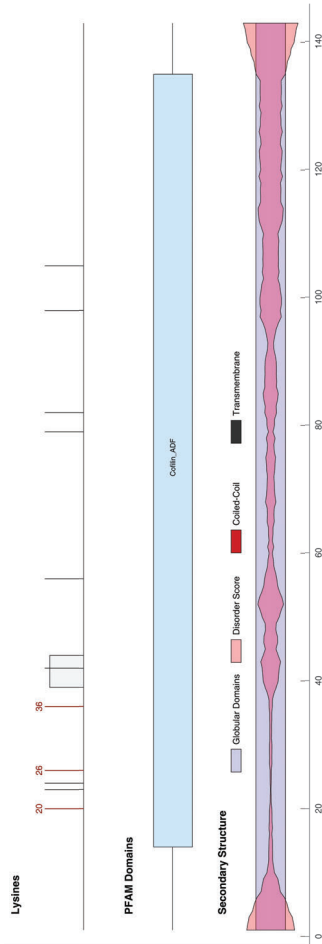
Transmembrane



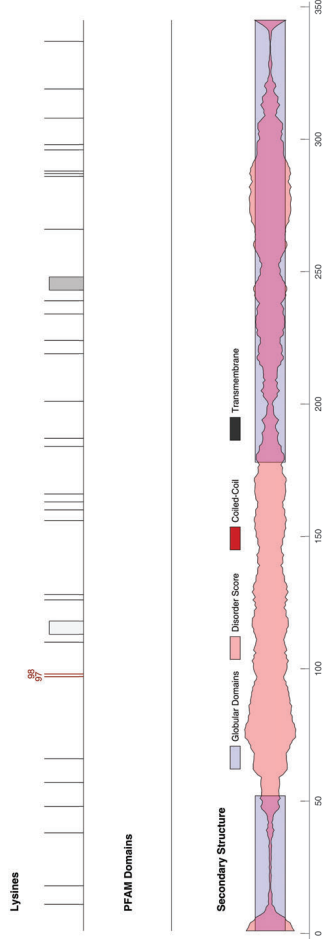
CND3 (sp|Q06680|CND3_YEAST)



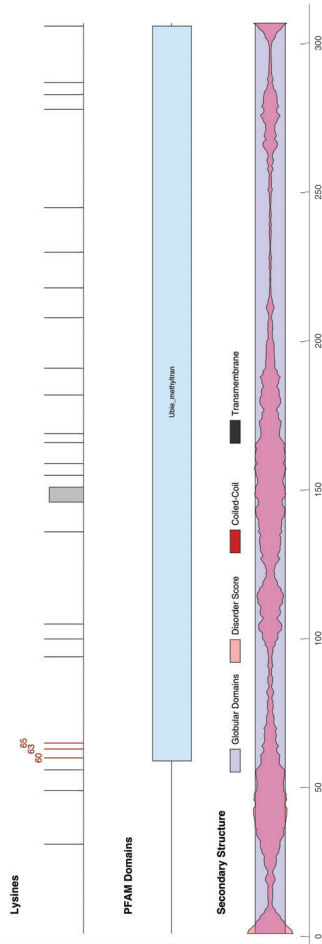
COF1 (sp|Q03048|COF1_YEAST)

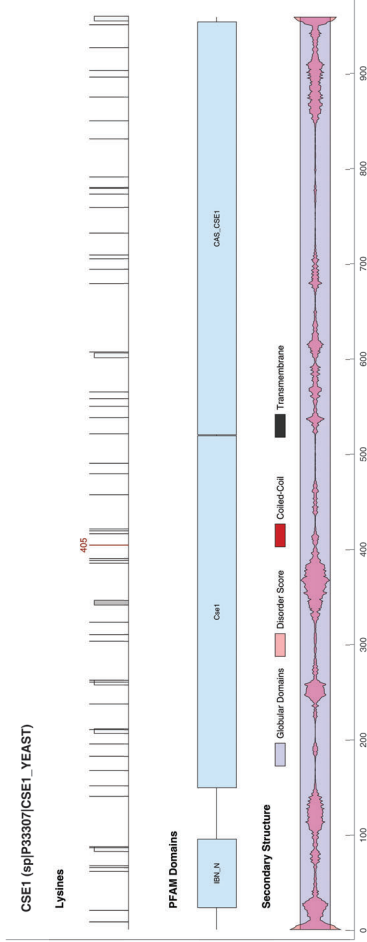
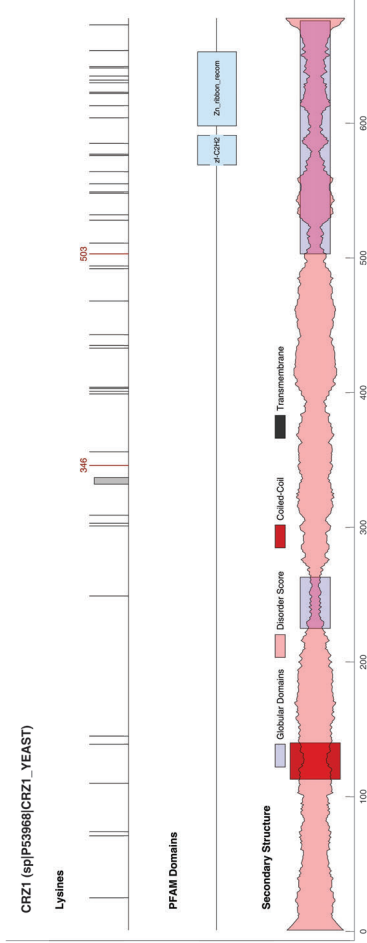
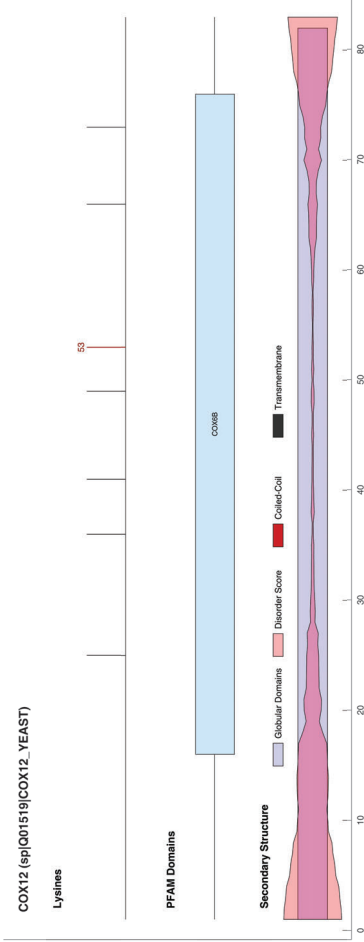
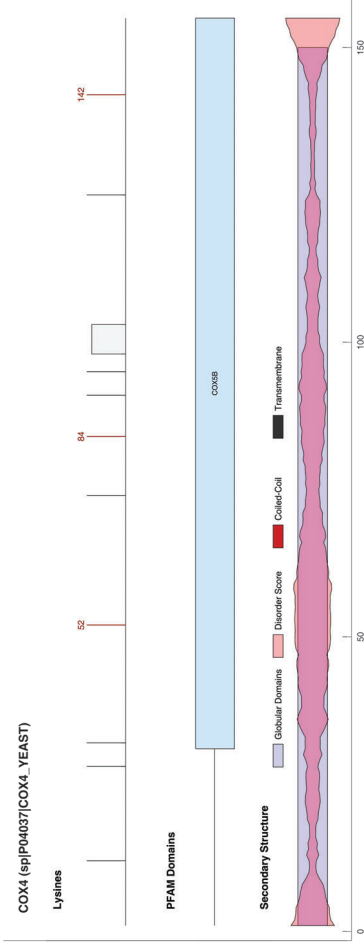


COM1 (sp|P46946|COM1_YEAST)

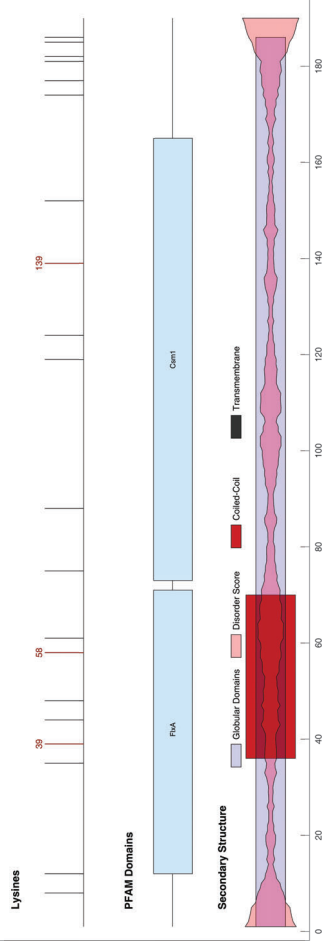


COO5 (sp|P49017|COO5_YEAST)

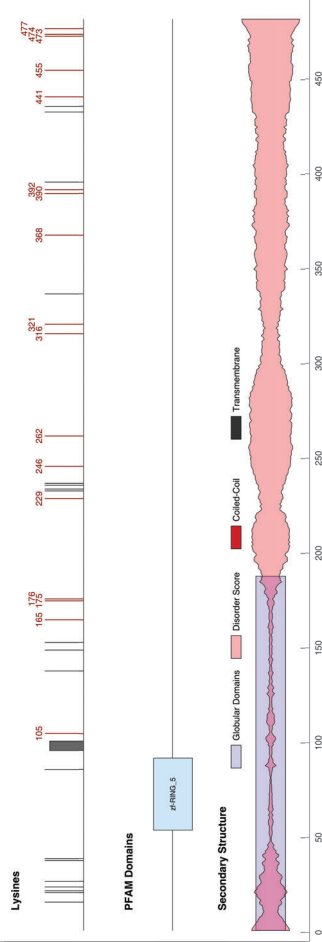




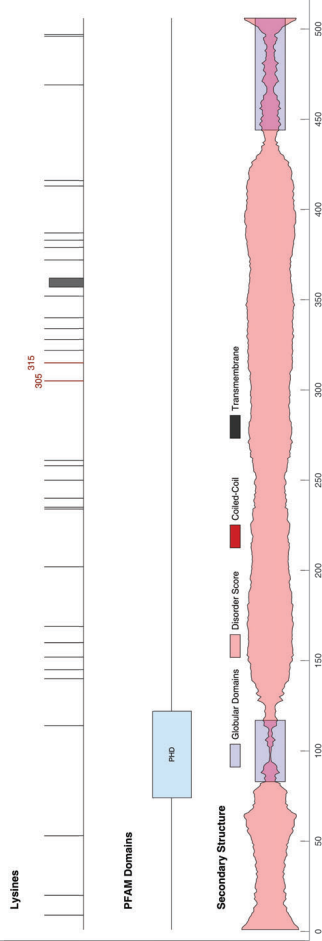
CSM1 (sp|P25651|CSM1_YEAST)



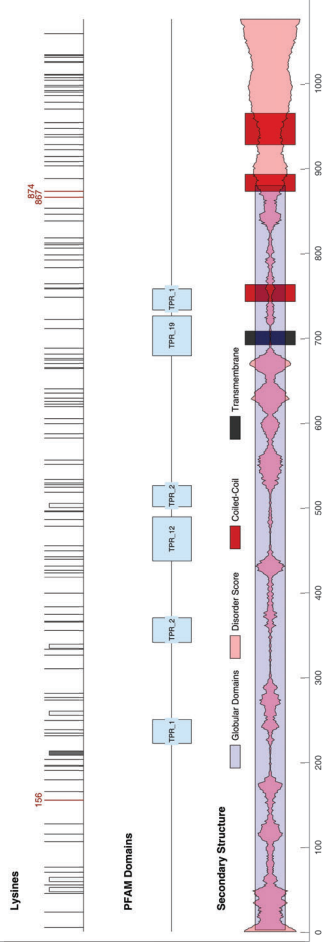
CST9 (sp|Q6032|CST9_YEAST)



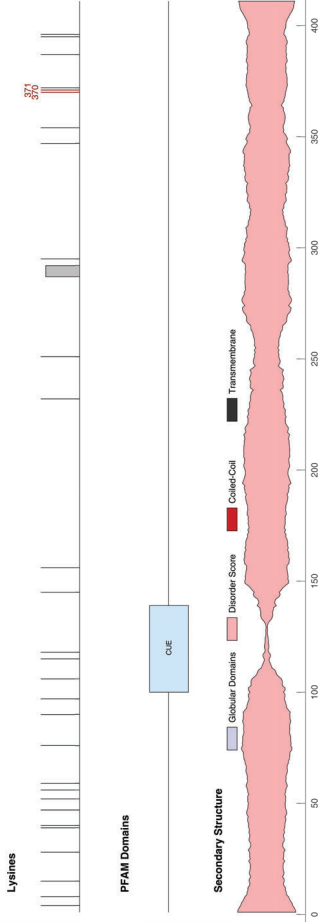
CT16 (sp|Q08923|CT16_YEAST)



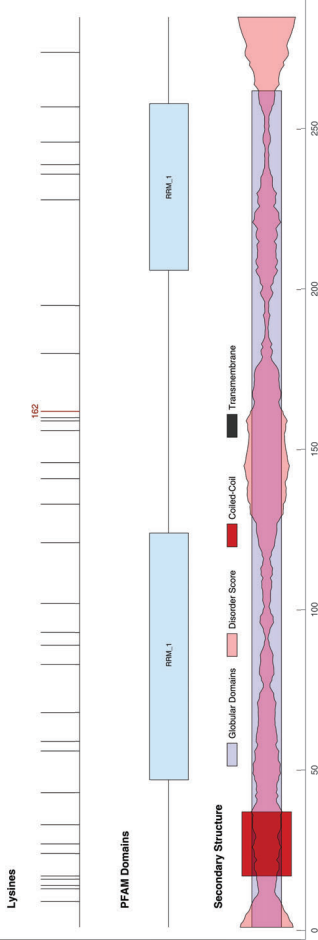
CTR9 (sp|P89105|CTR9_YEAST)



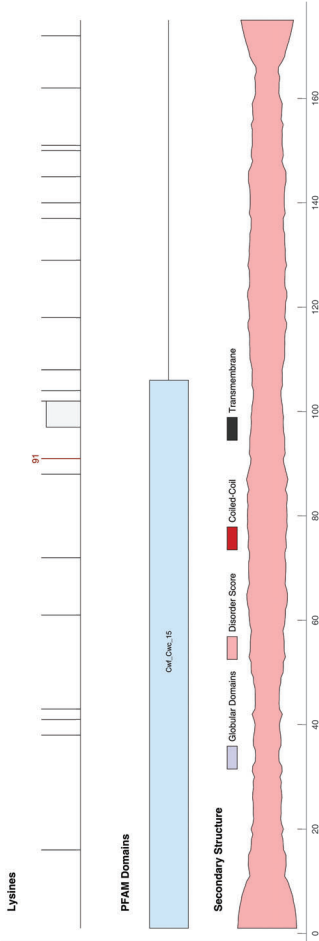
CUE5 (sp|Q08412|CUE5_YEAST)



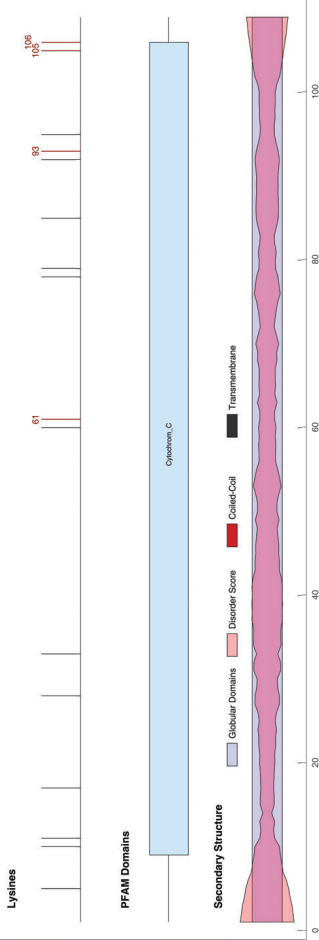
CUS2 (sp|P53830|CUS2_YEAST)

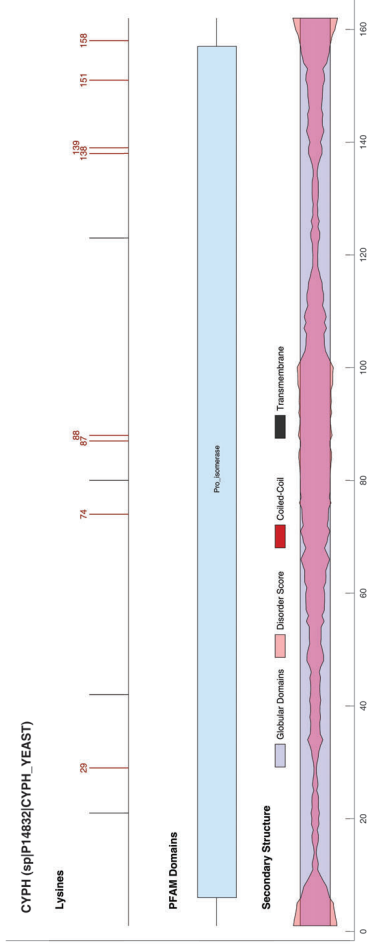
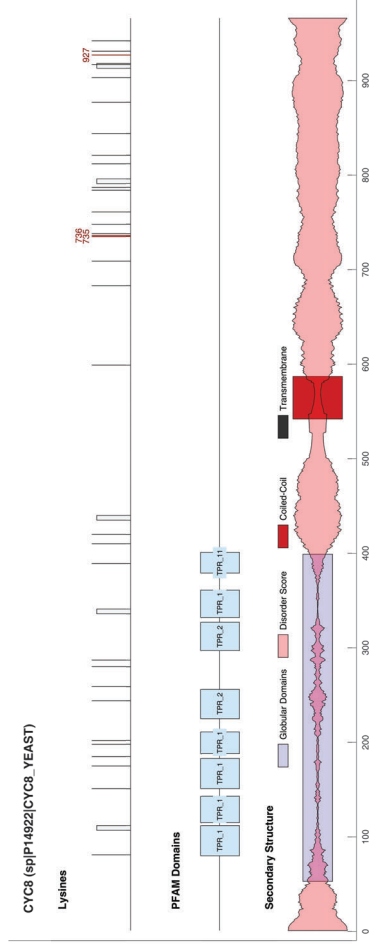


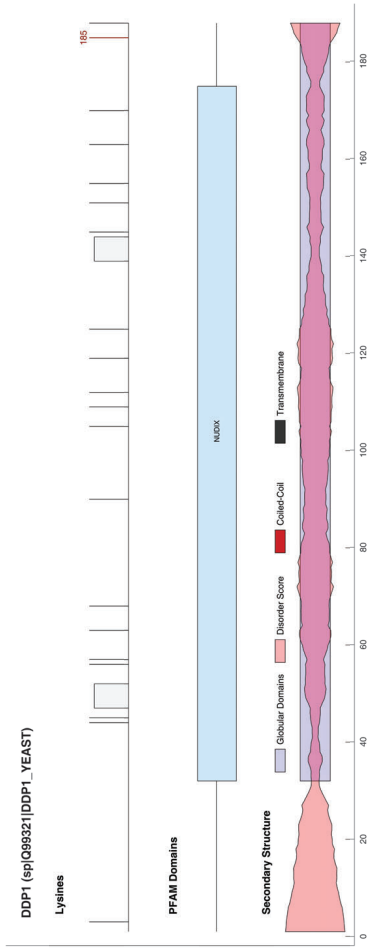
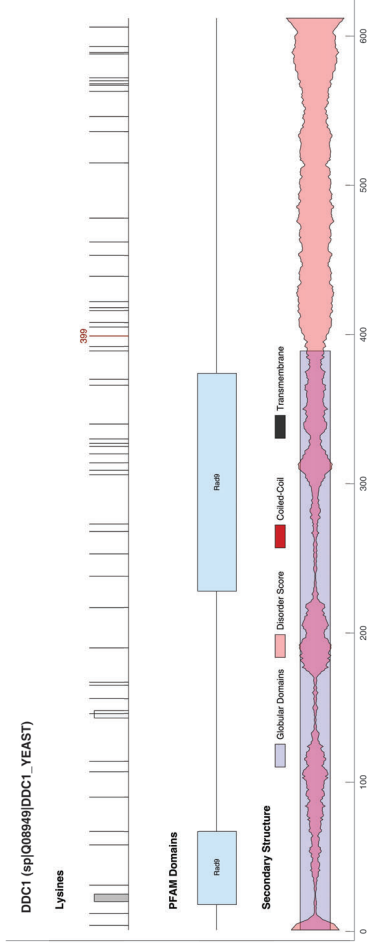
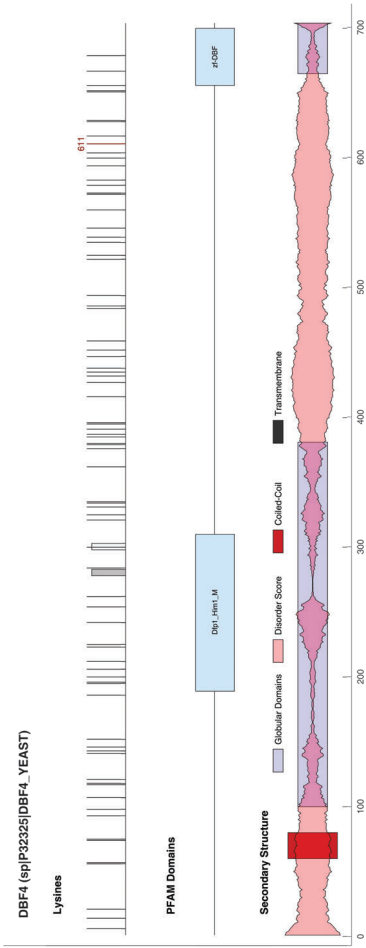
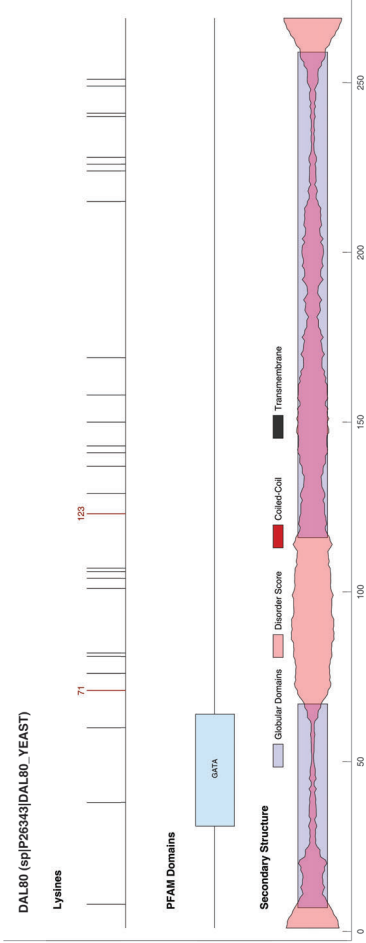
CWC15 (sp|Q03772|CWC15_YEAST)

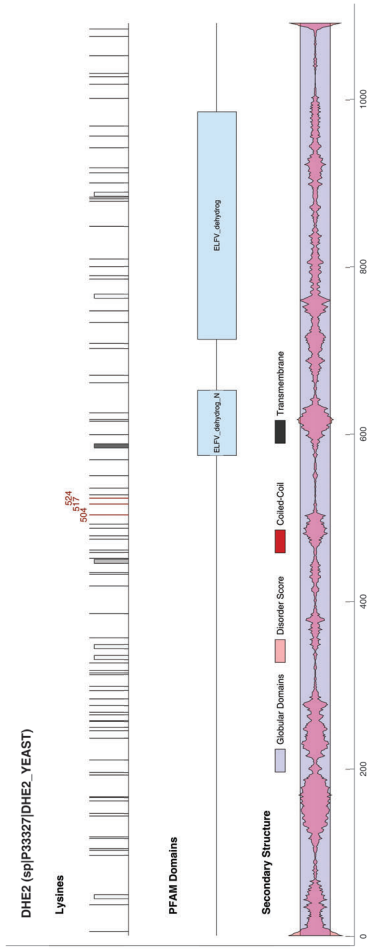
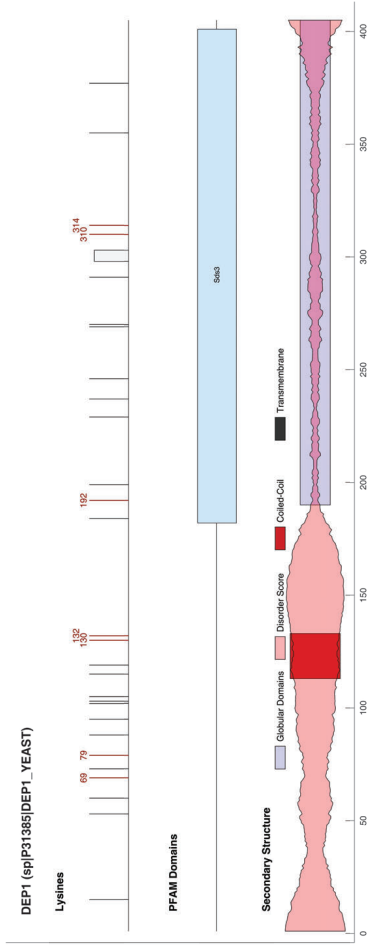
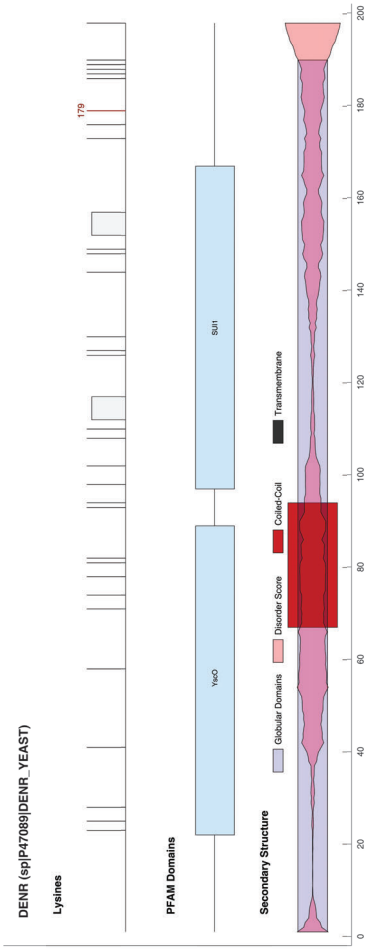
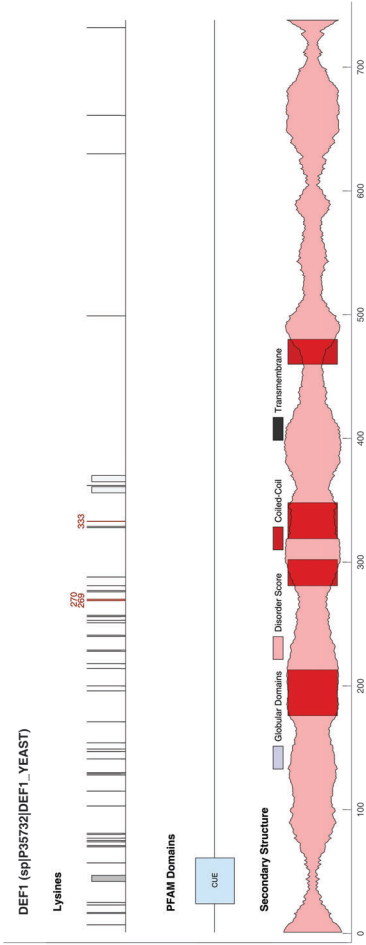


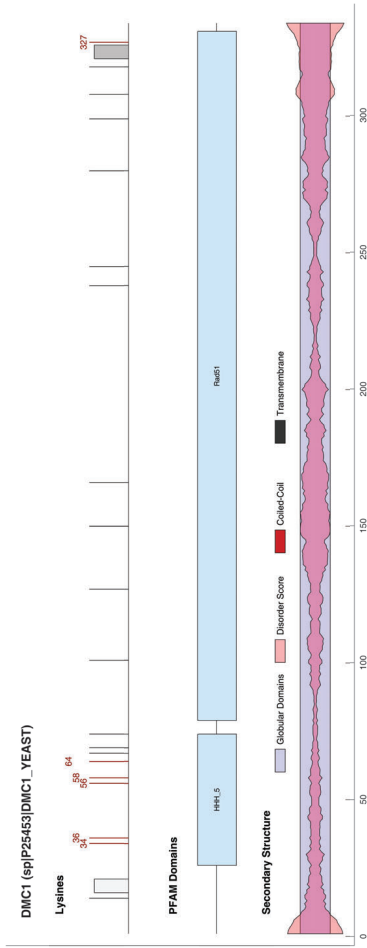
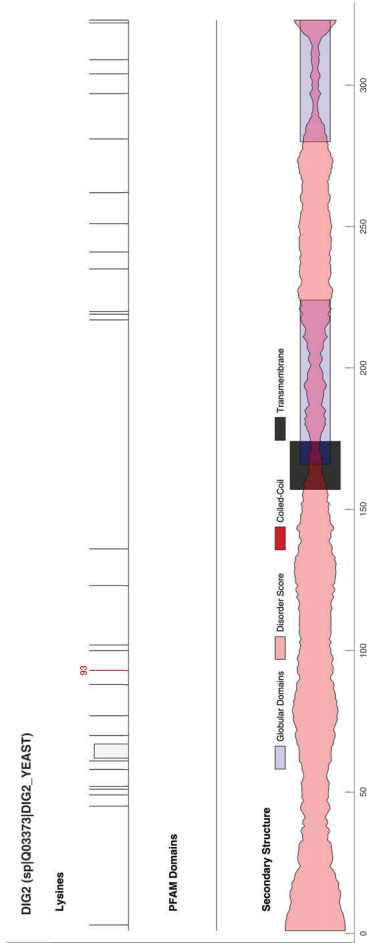
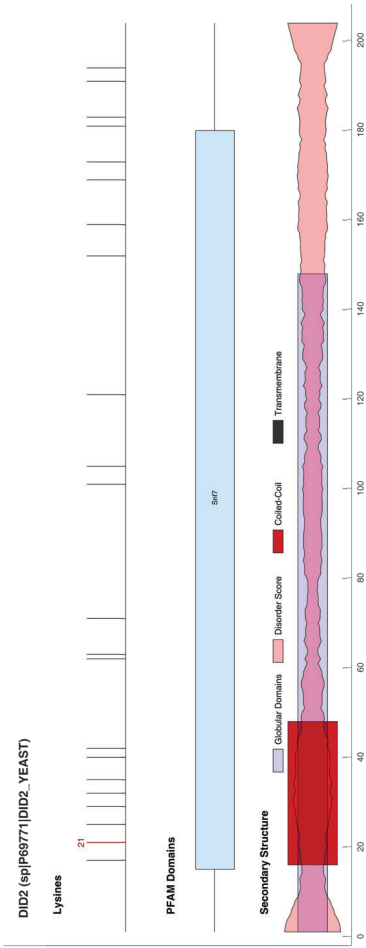
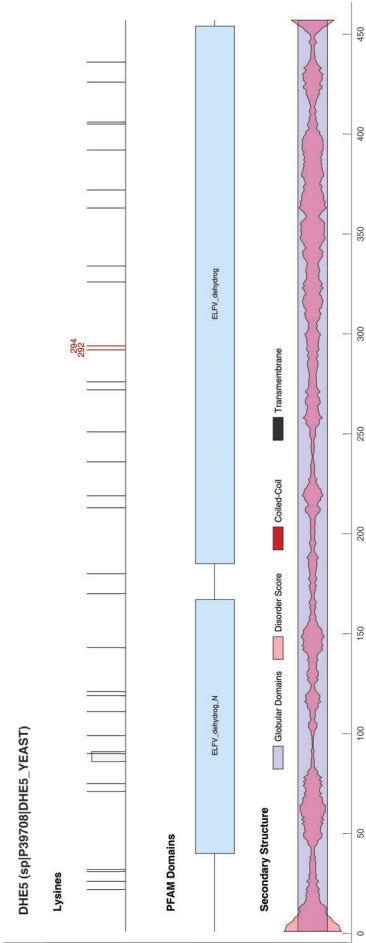
CYC1 (sp|P00044|CYC1_YEAST)

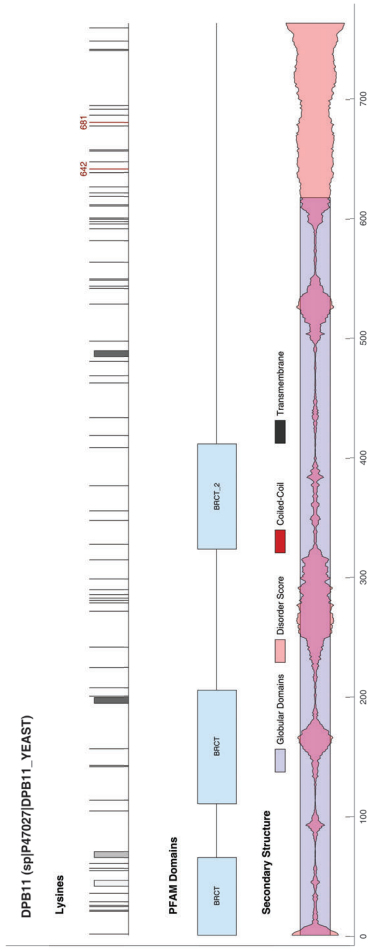
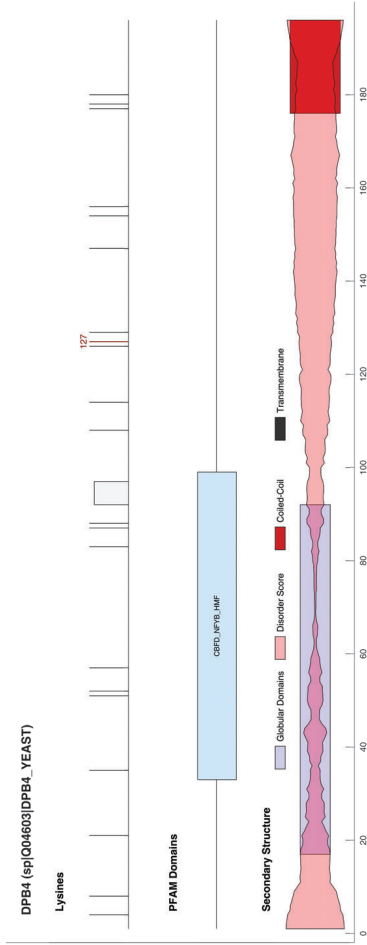
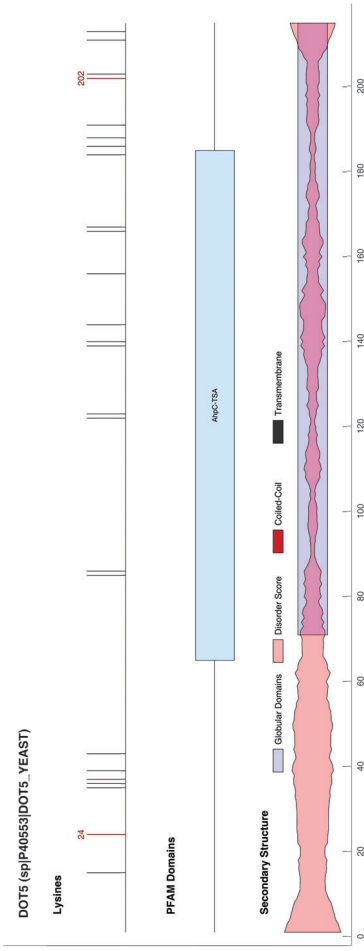
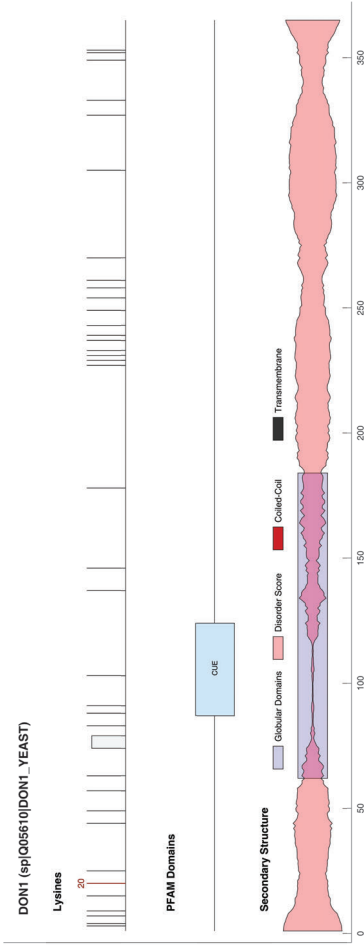


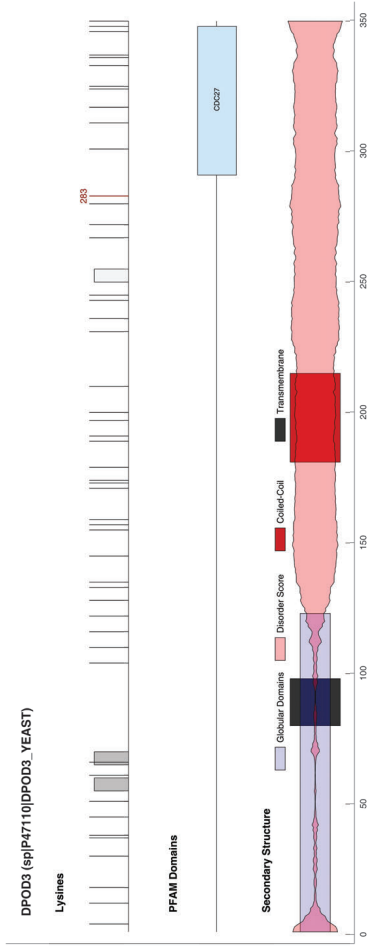
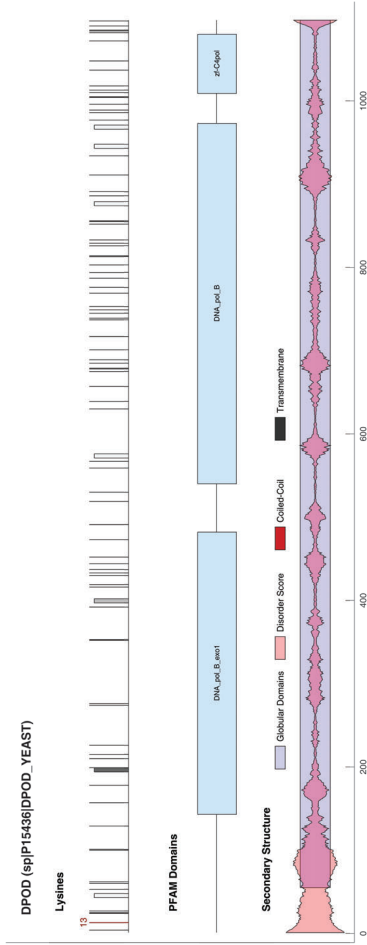
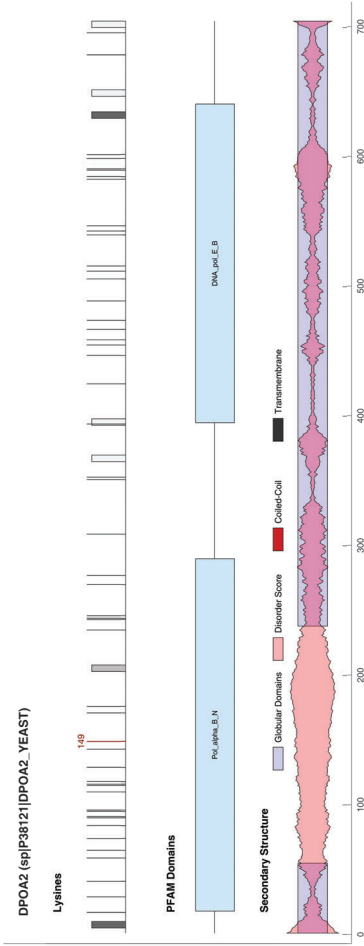
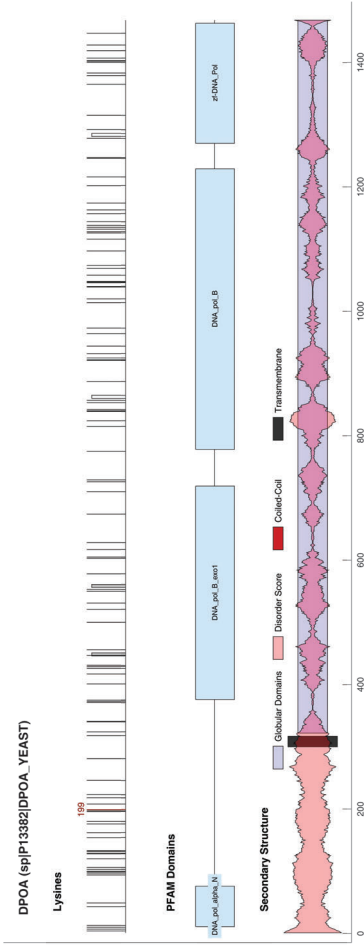


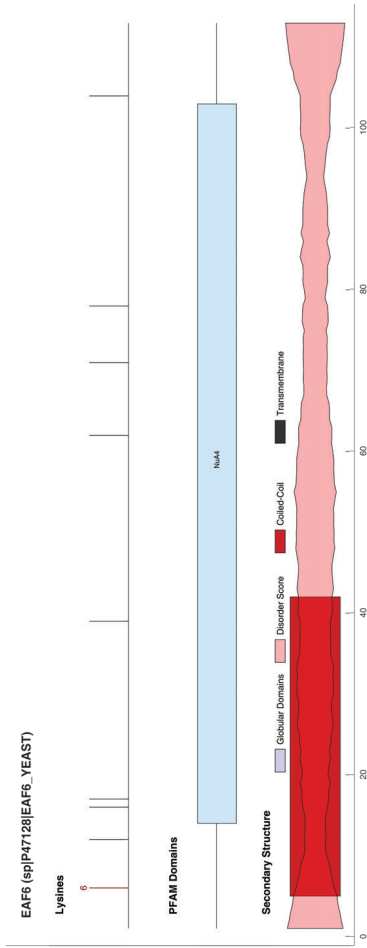
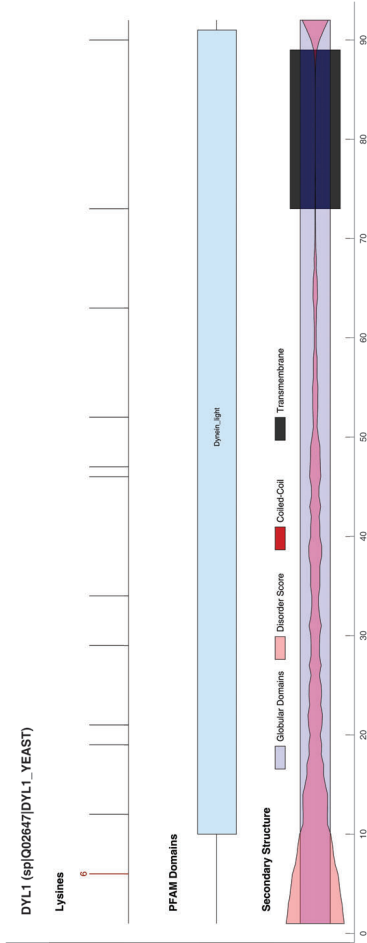
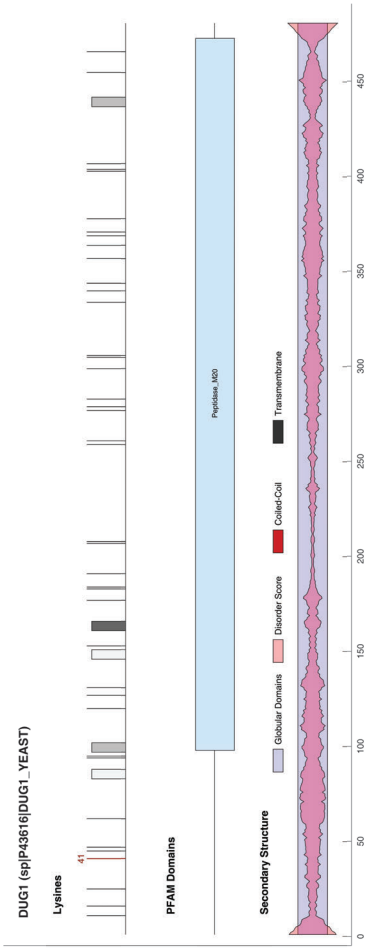
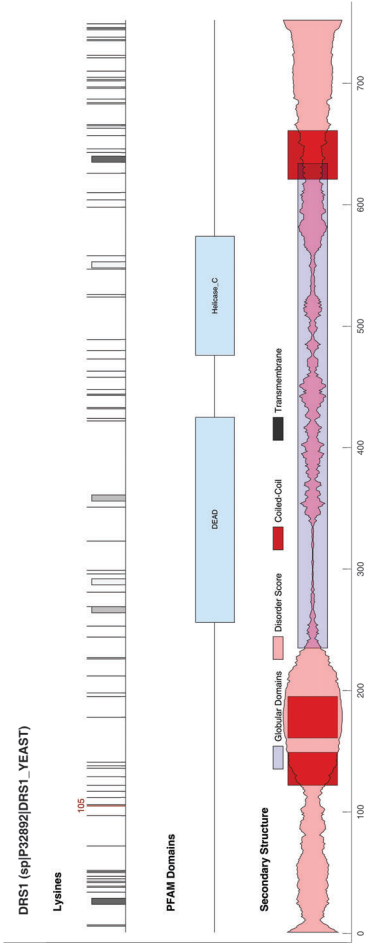


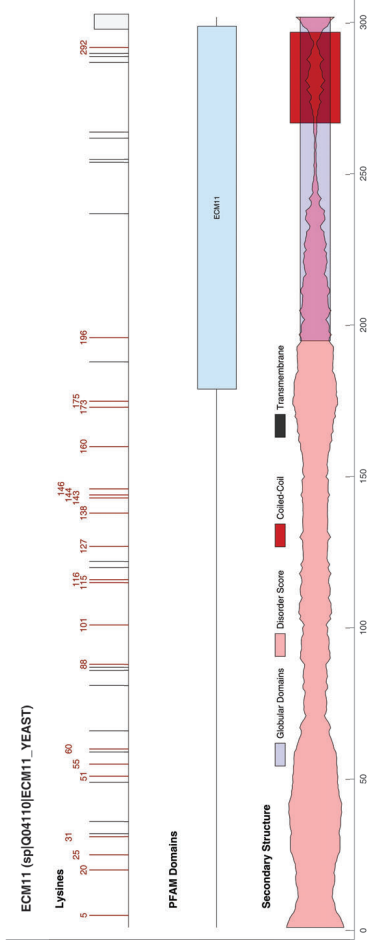
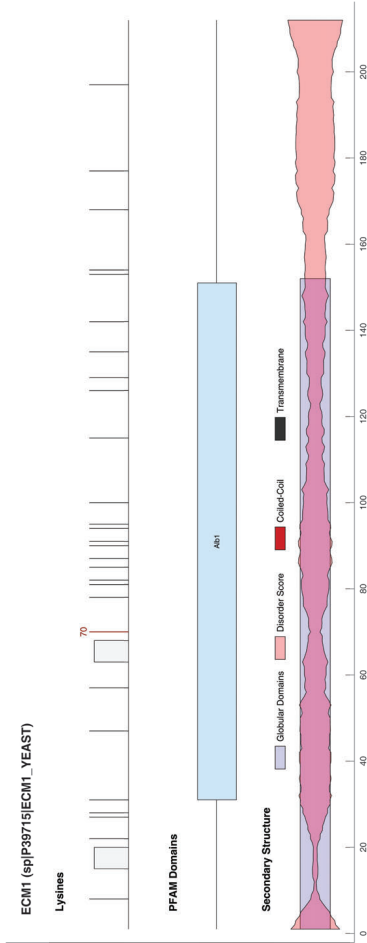
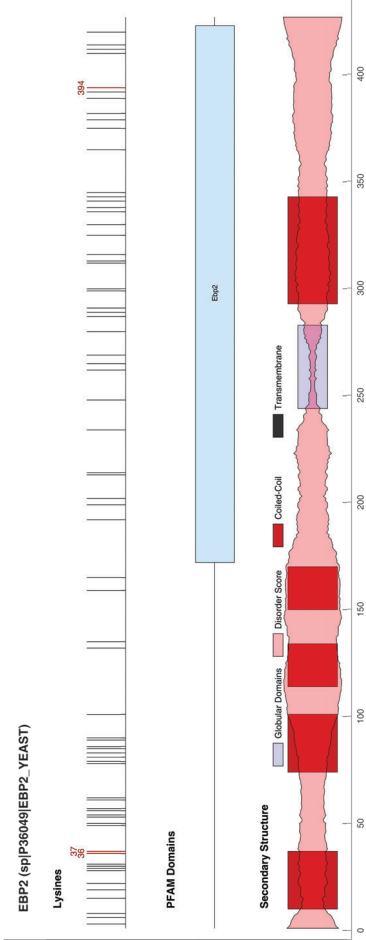
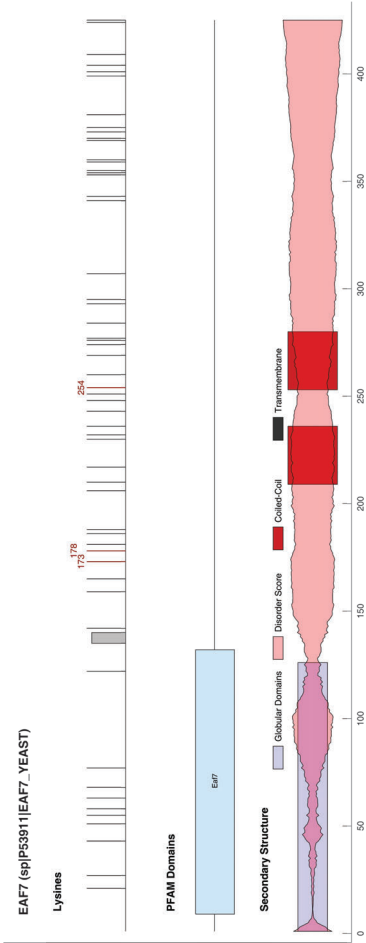


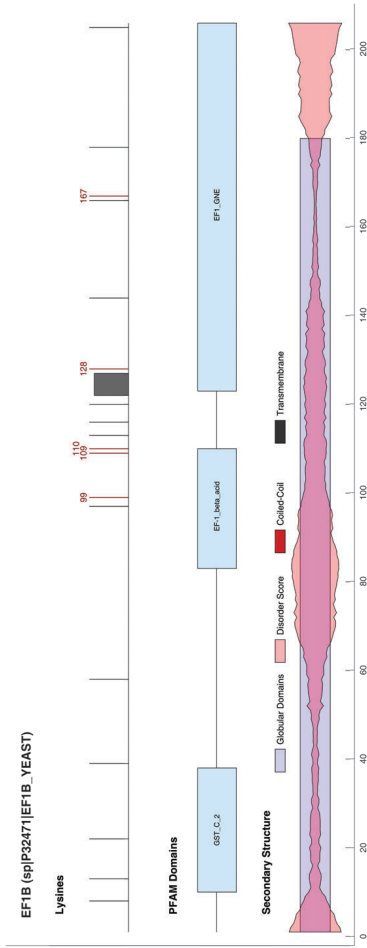
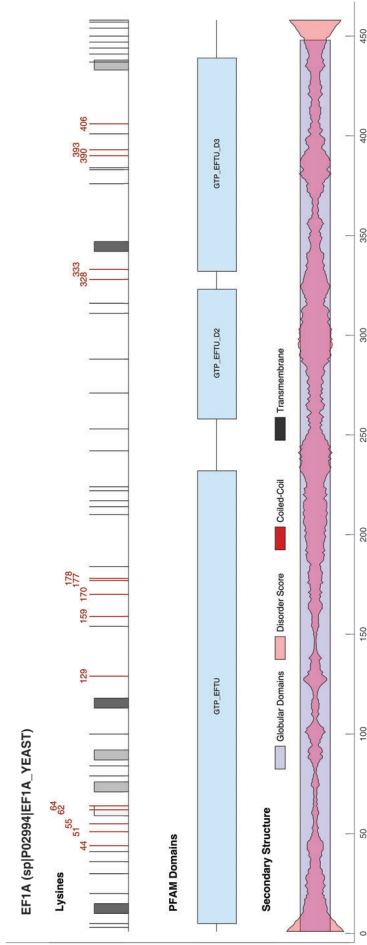
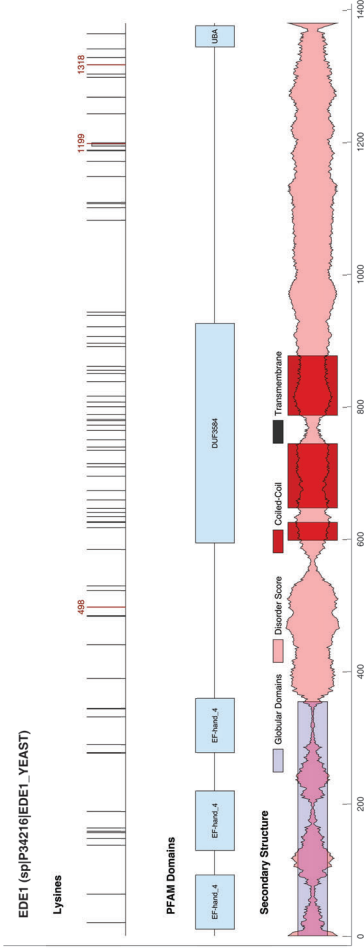
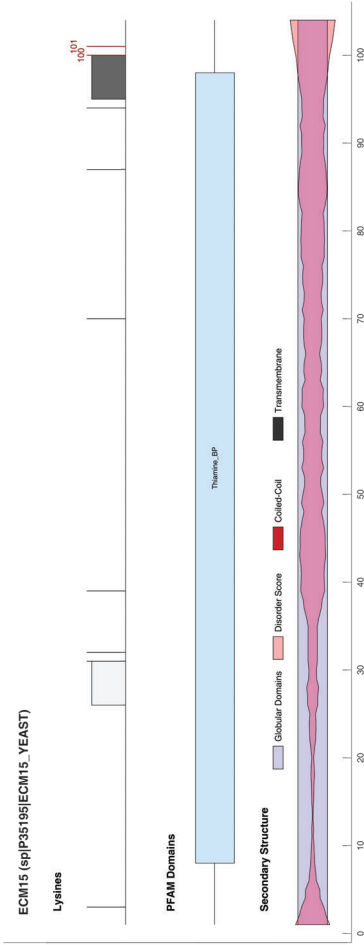


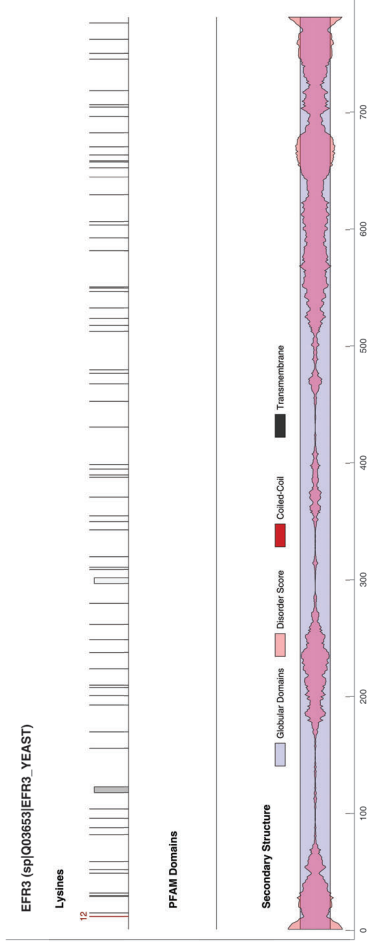
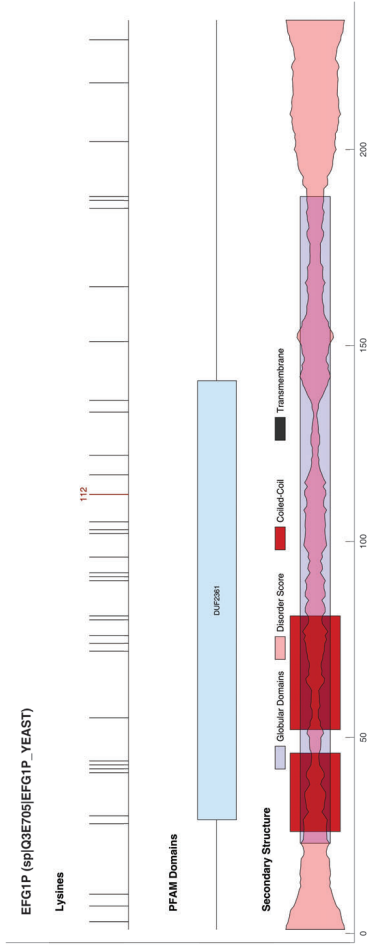
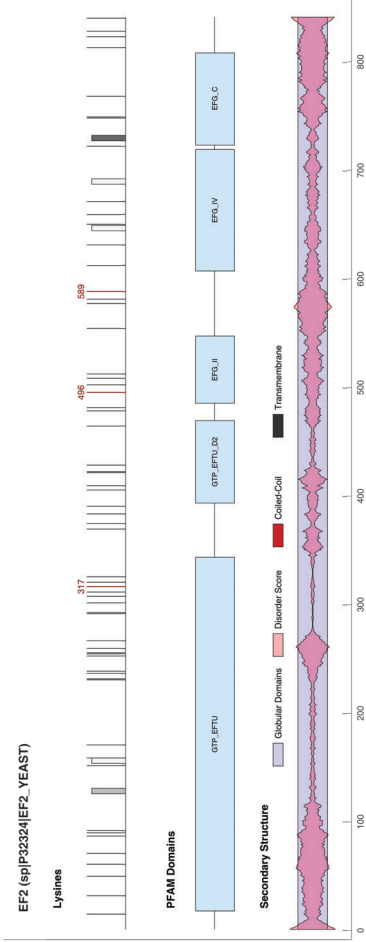
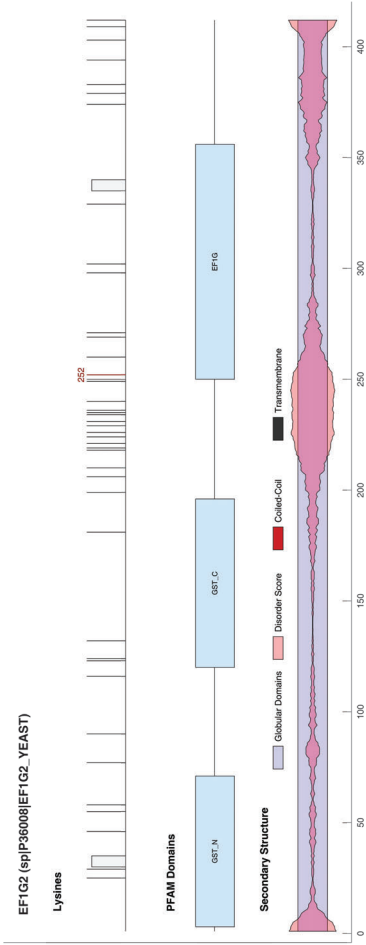




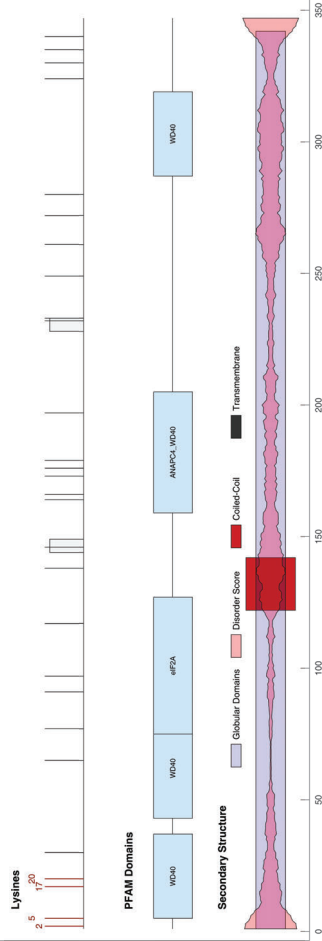




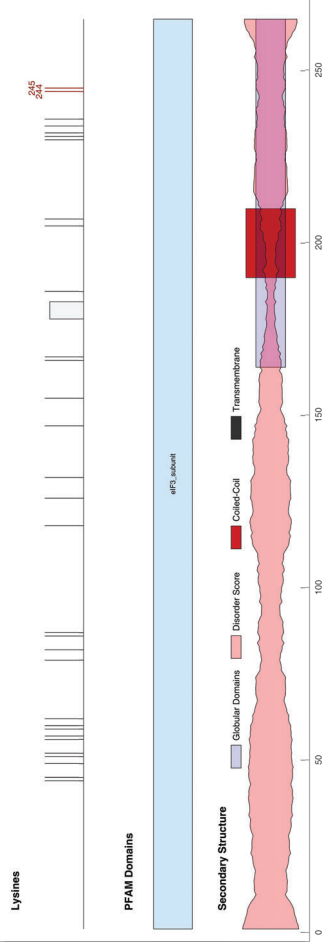




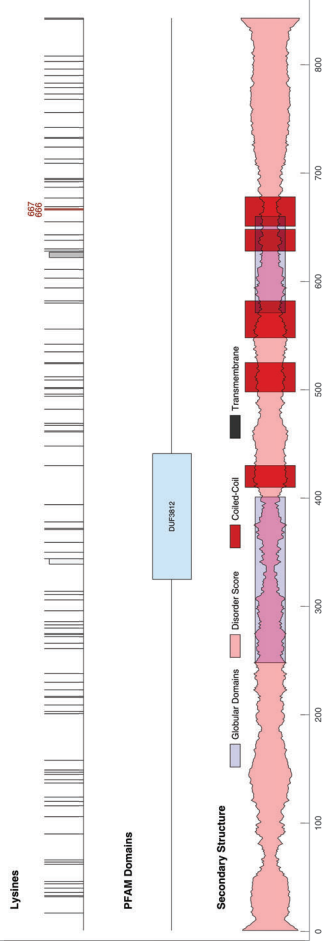
EIF3I (sp|P40217|EIF3I_YEAST)



EIF3J (sp|Q05775|EIF3J_YEAST)

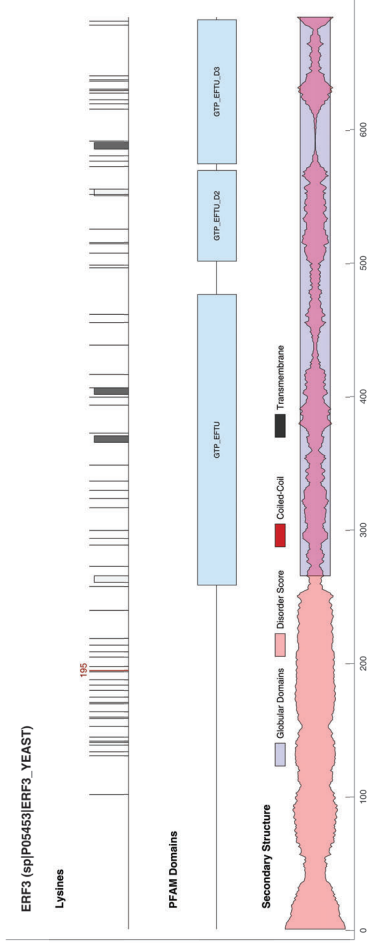
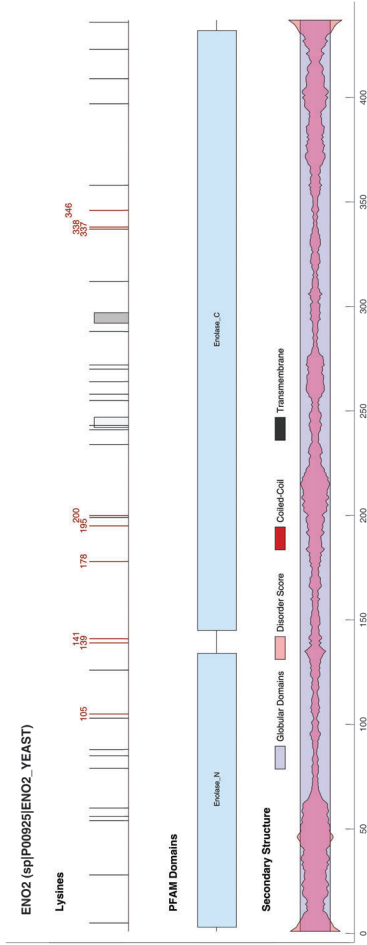
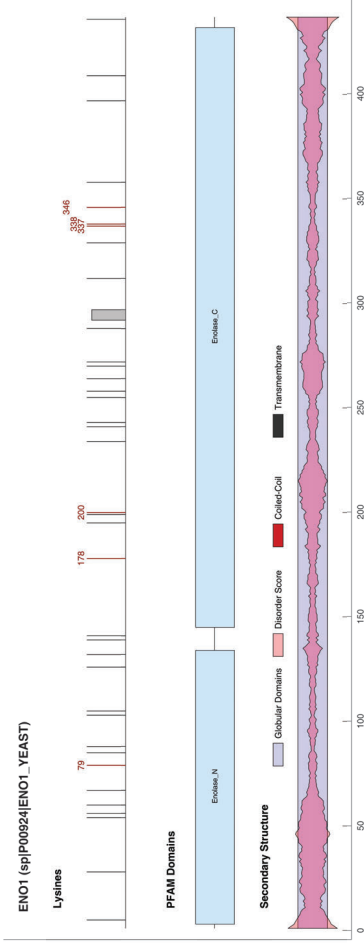
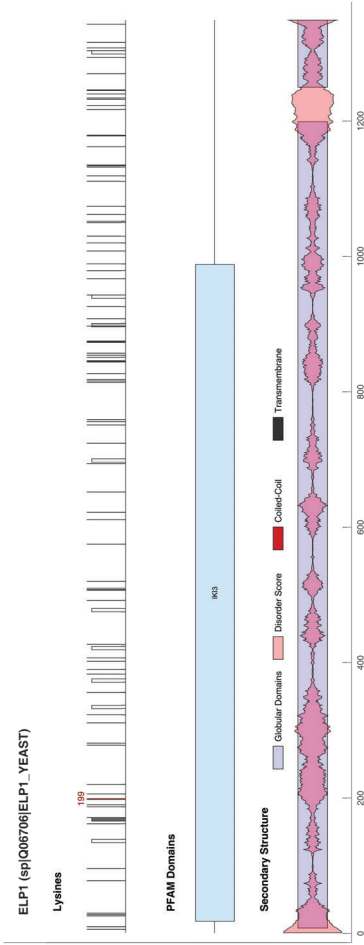


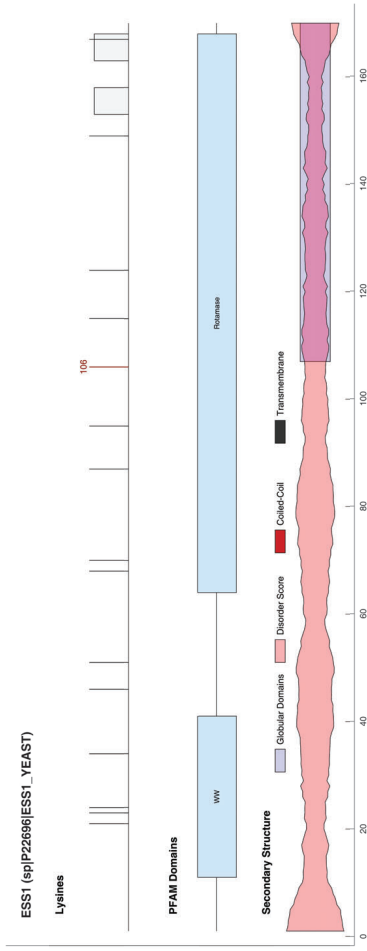
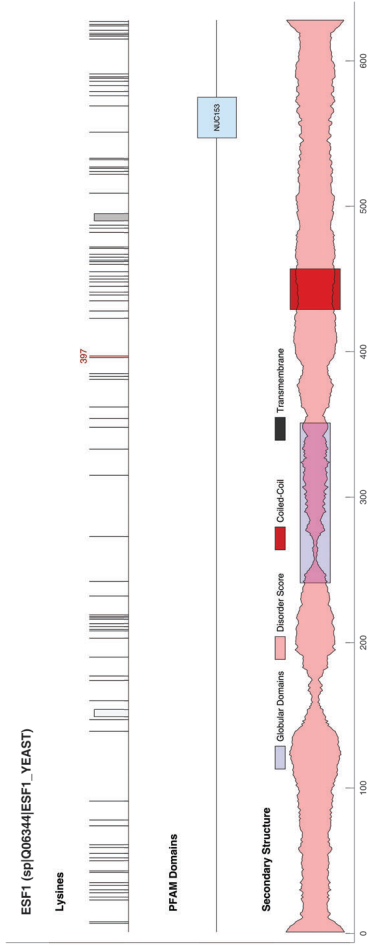
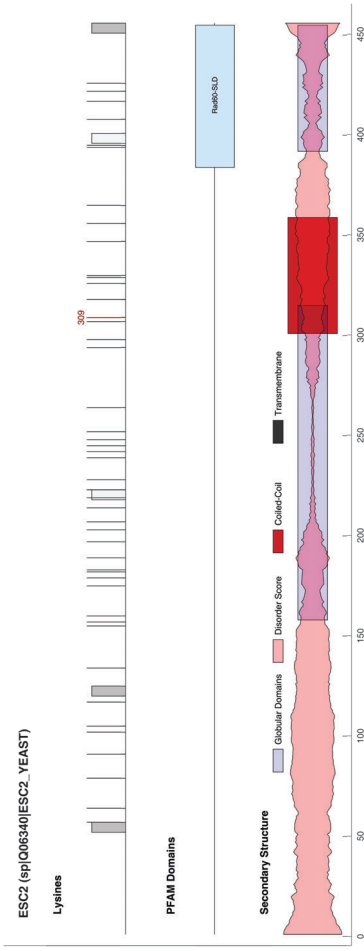
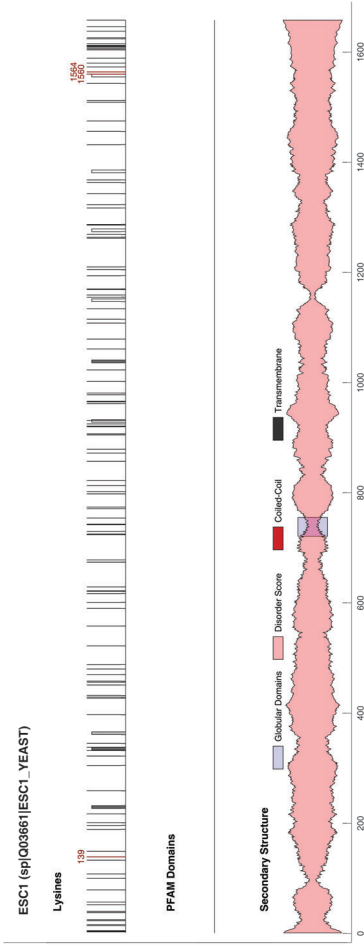
EIS1 (sp|Q05050|EIS1_YEAST)

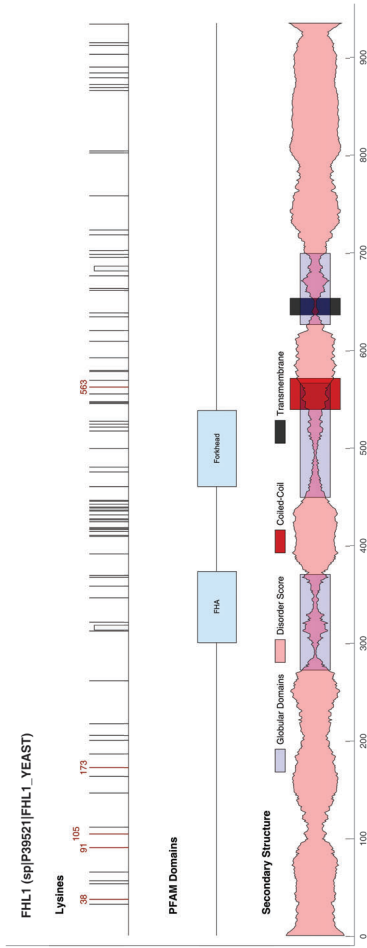
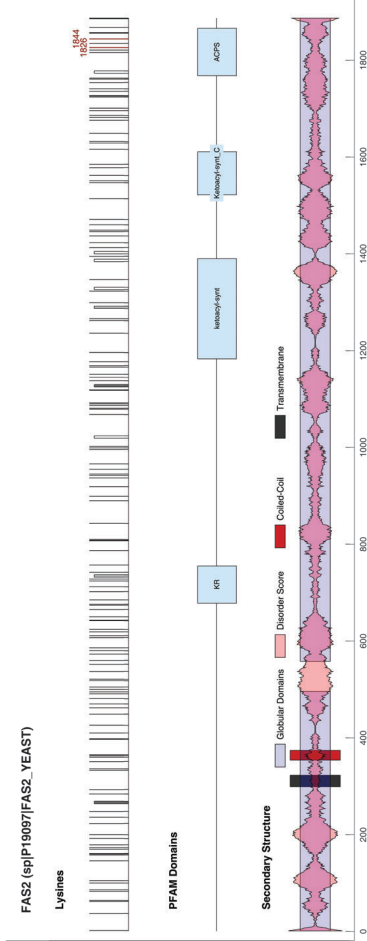
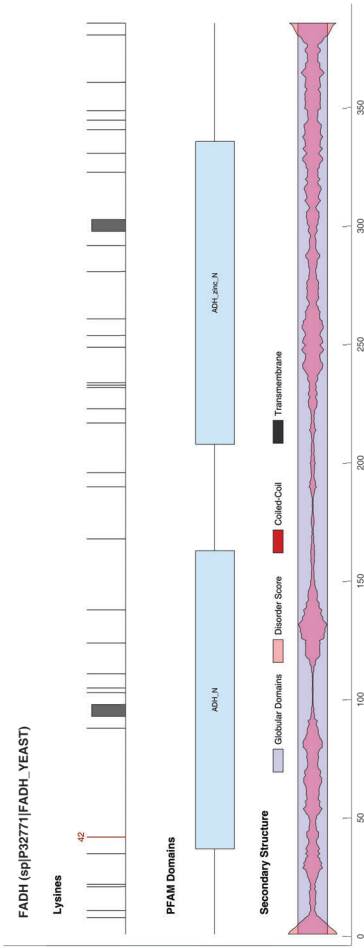
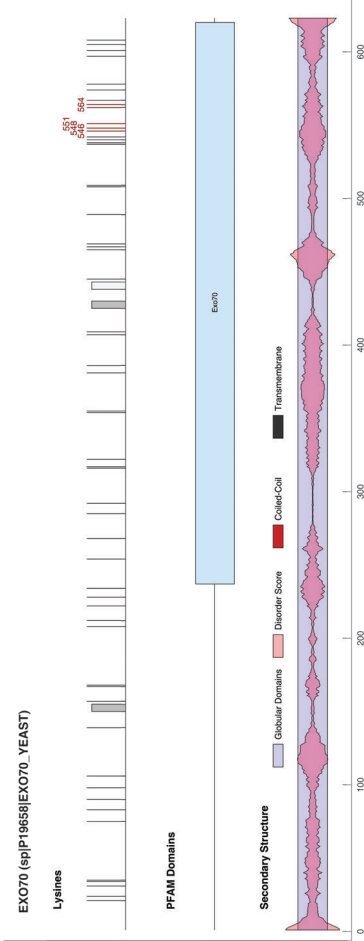


ELG1 (sp|Q12050|ELG1_YEAST)

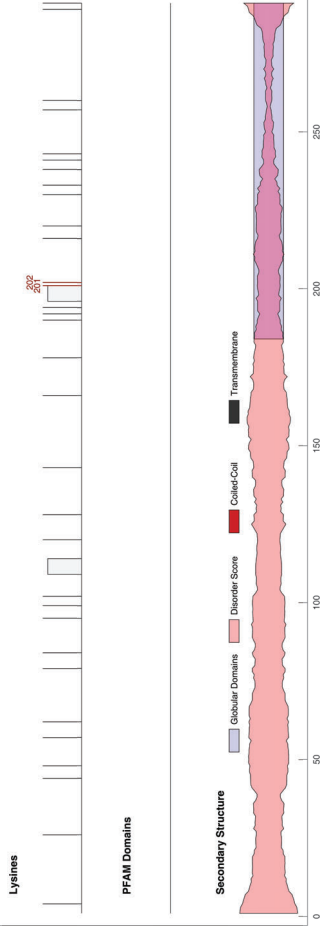




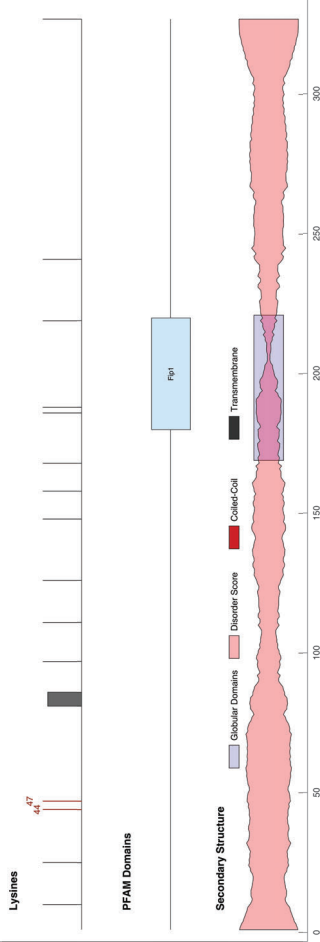




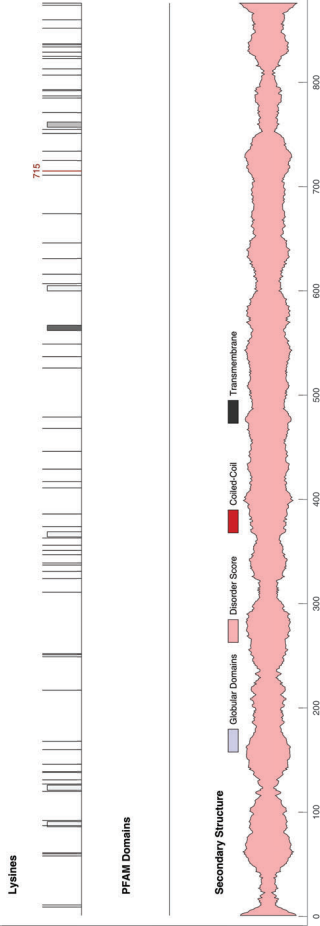
FIN1 (sp|Q03898|FIN1_YEAST)



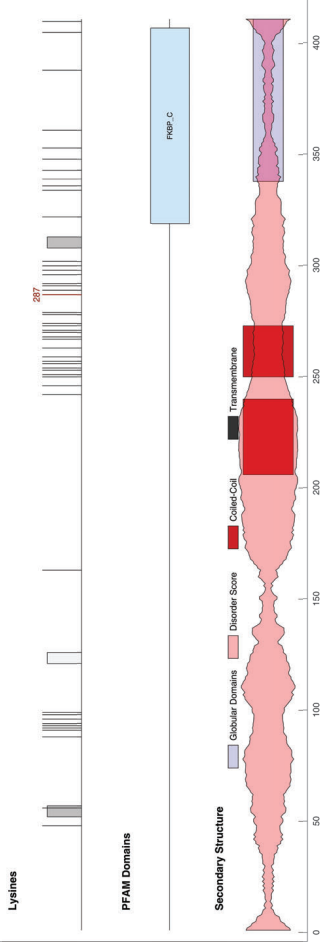
FIP1 (sp|P45976|FIP1_YEAST)

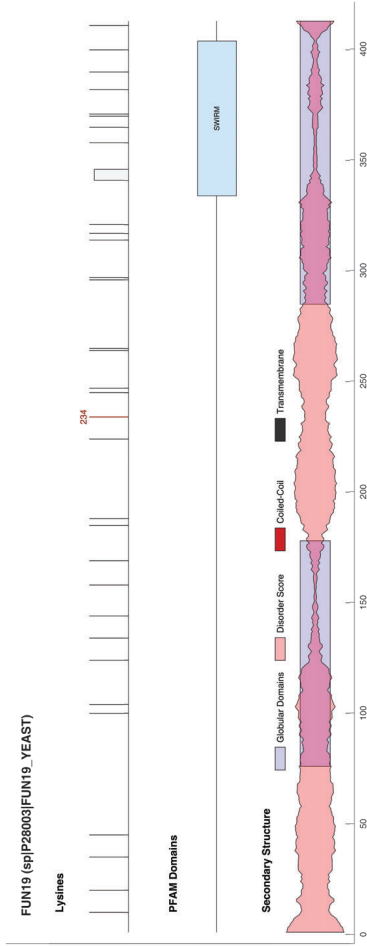
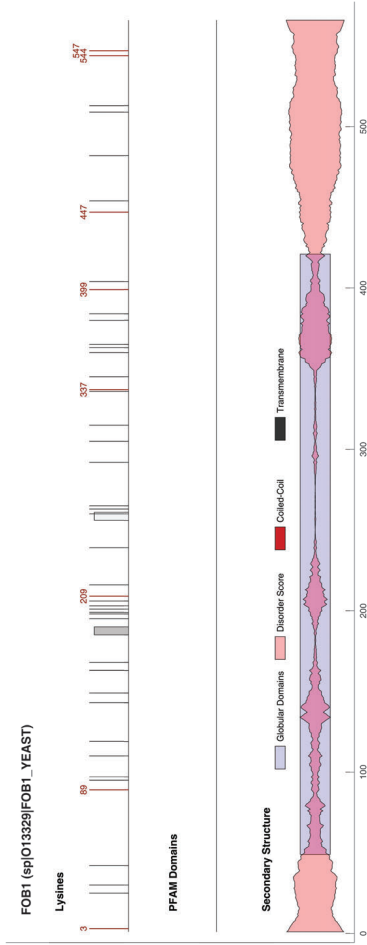
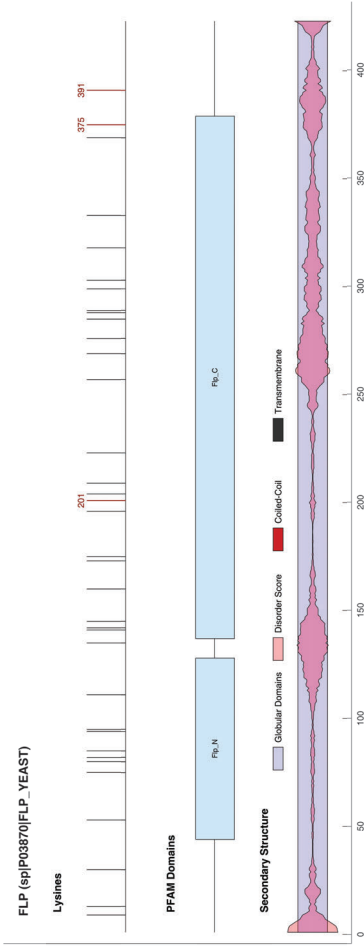
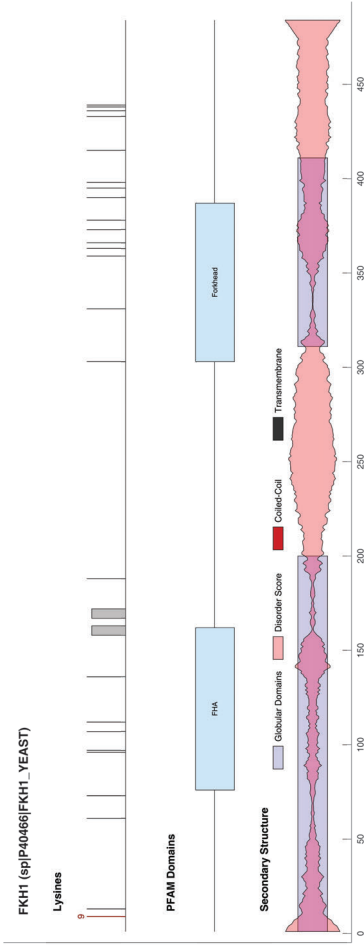


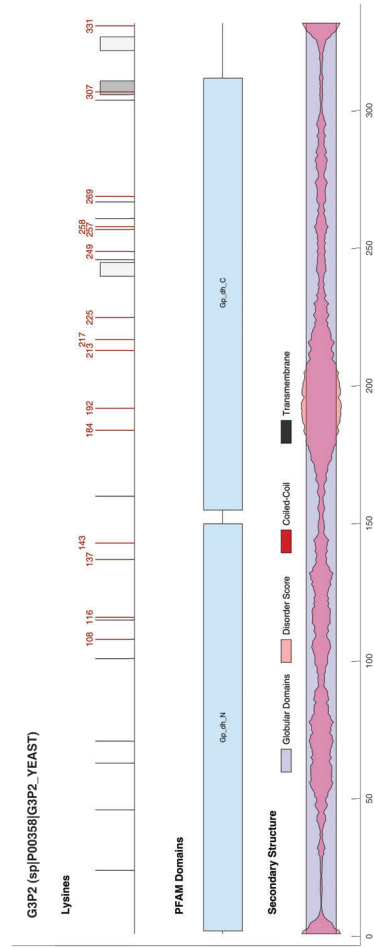
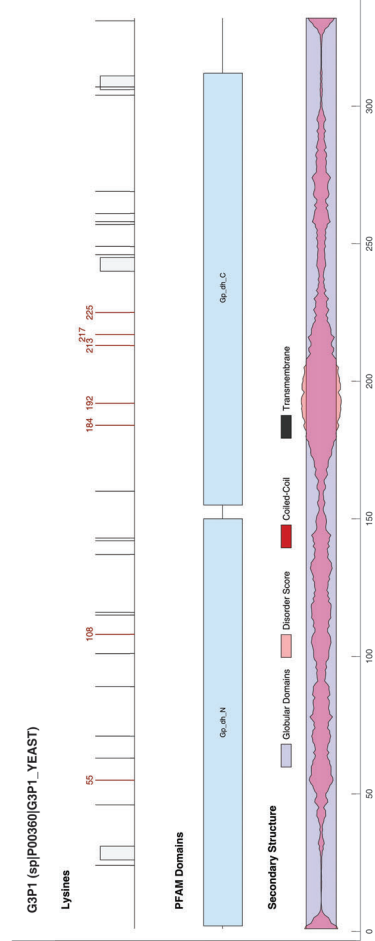
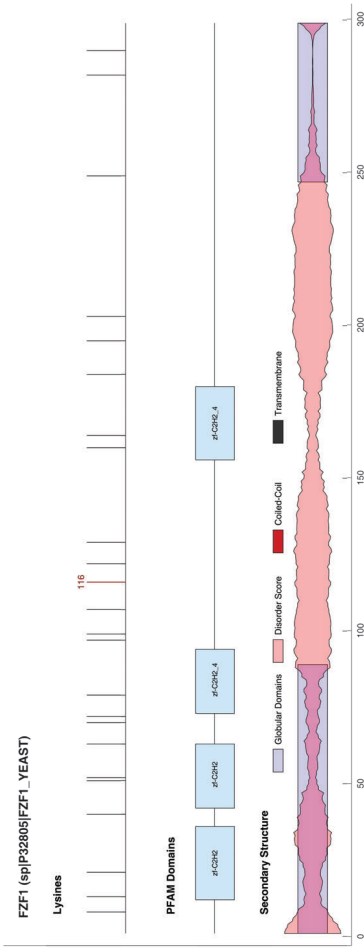
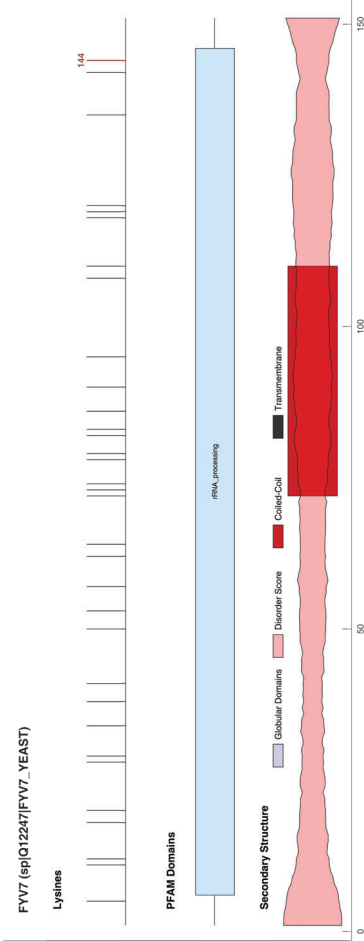
FIR1 (sp|P40020|FIR1_YEAST)

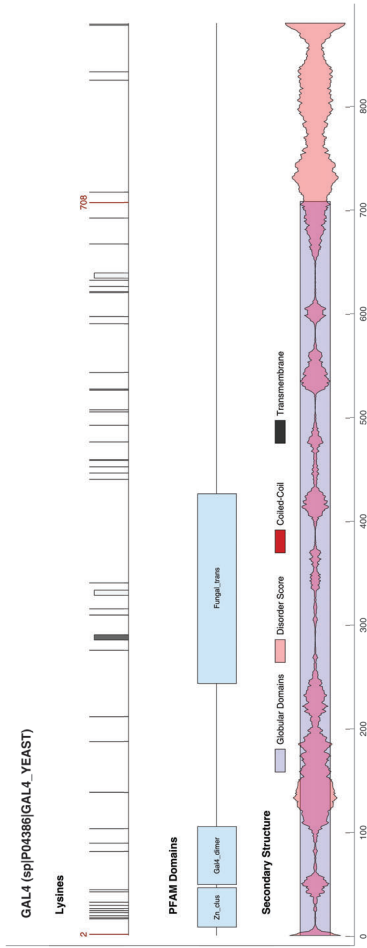
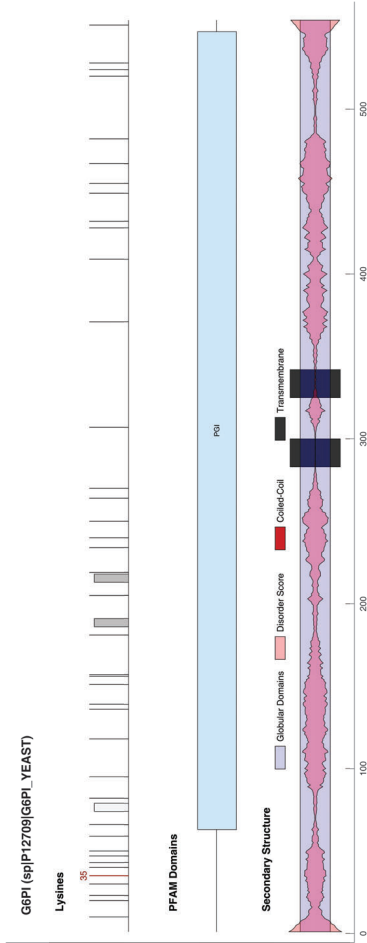
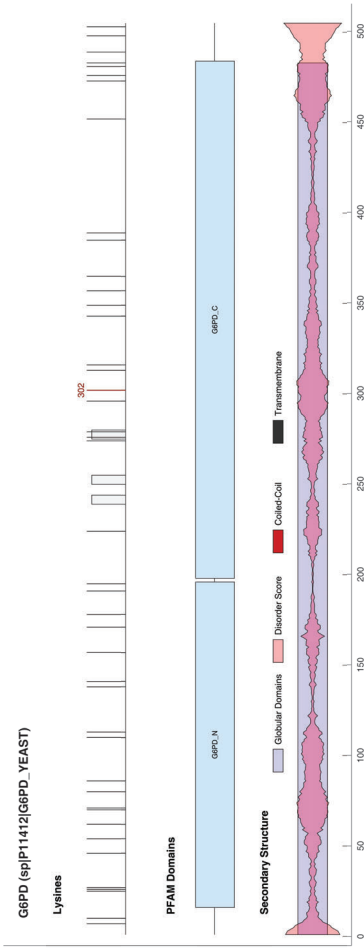
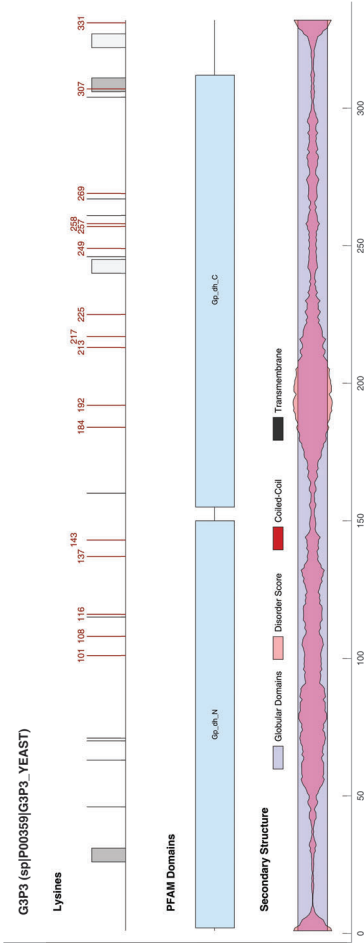


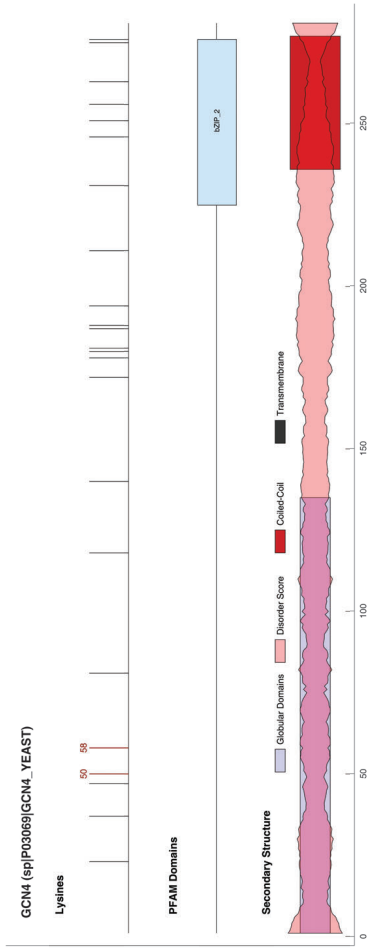
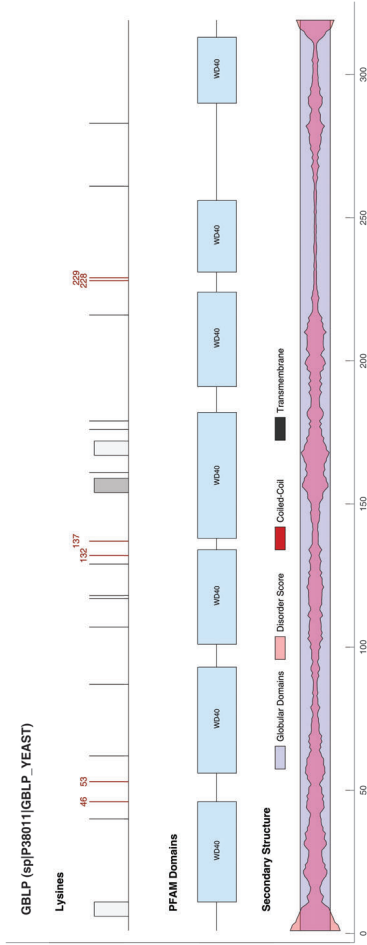
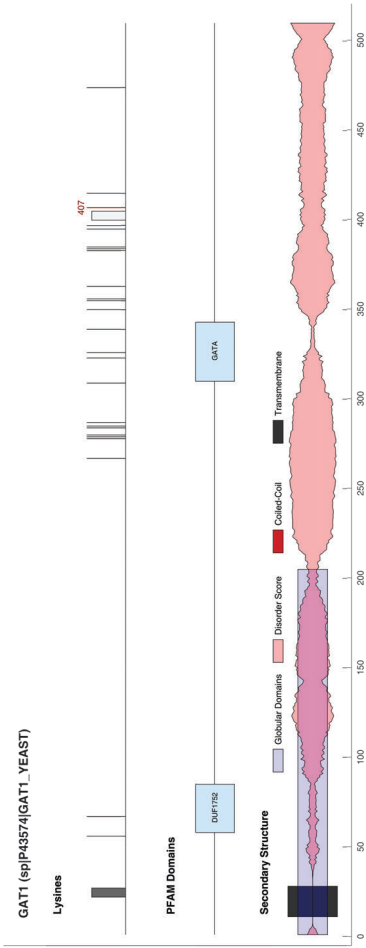
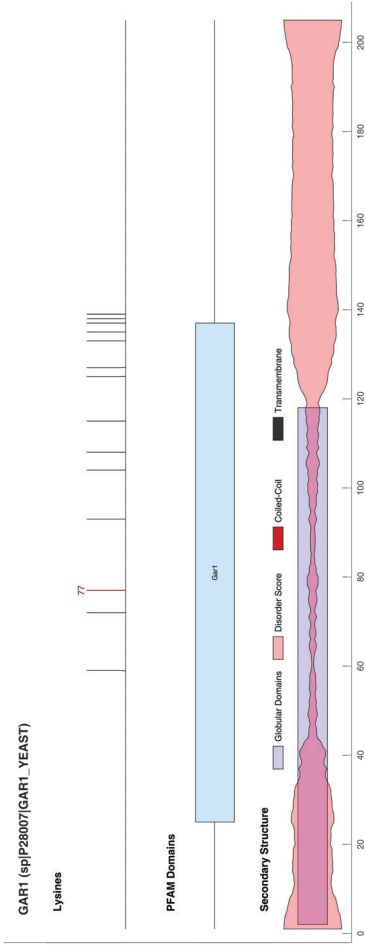
FKBP3 (sp|P3891|FKBP3_YEAST)

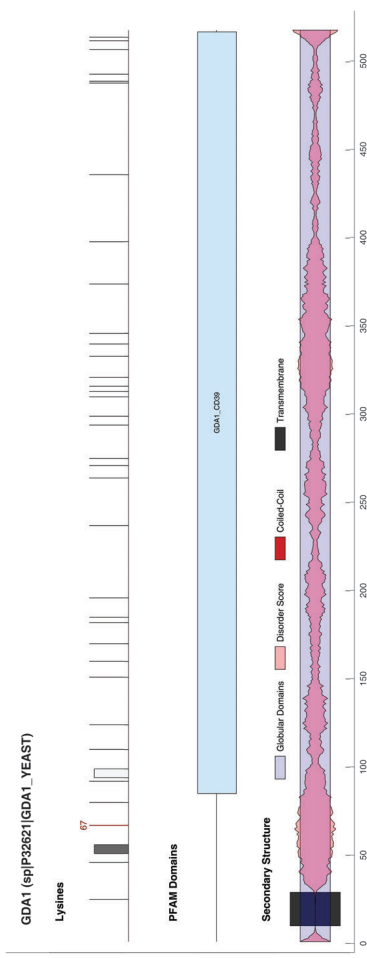
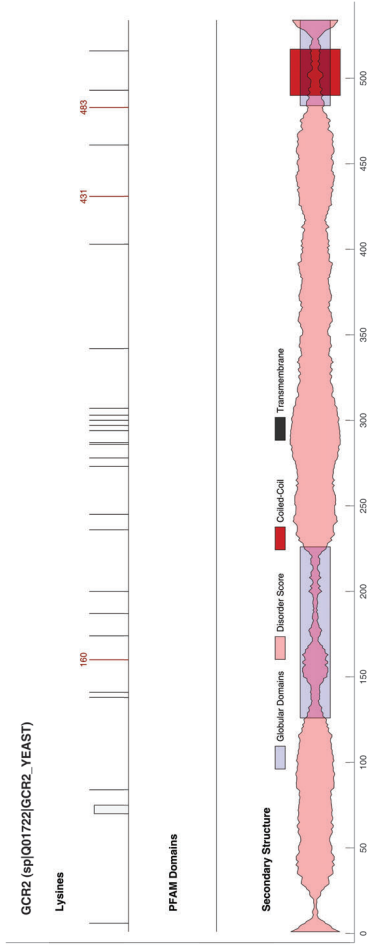
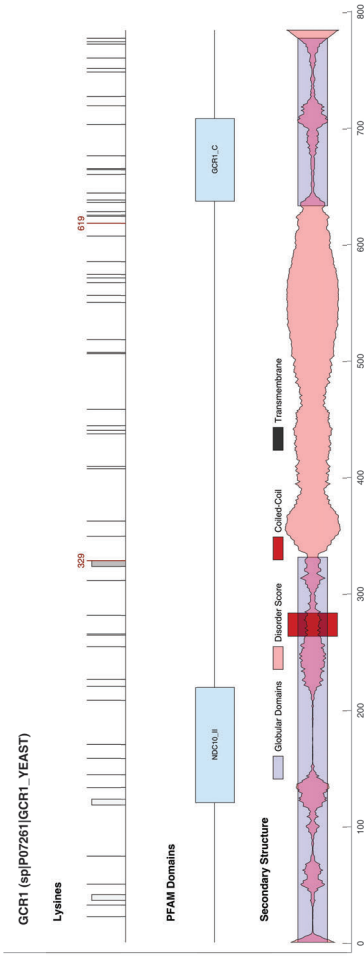
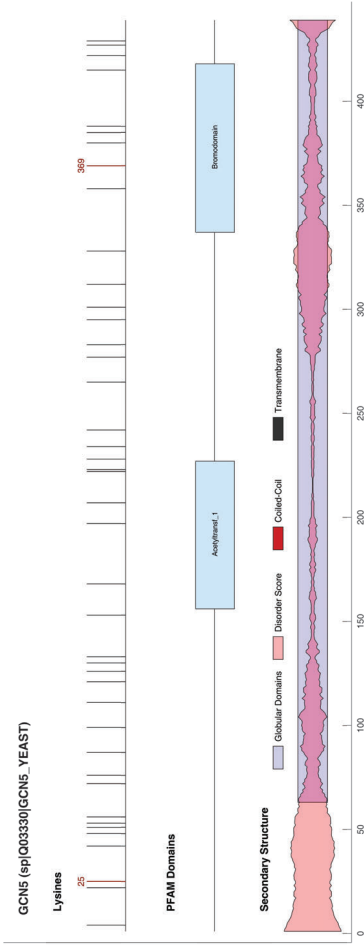




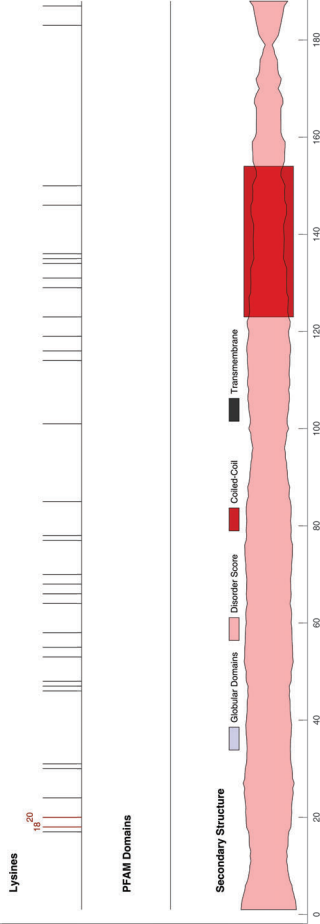




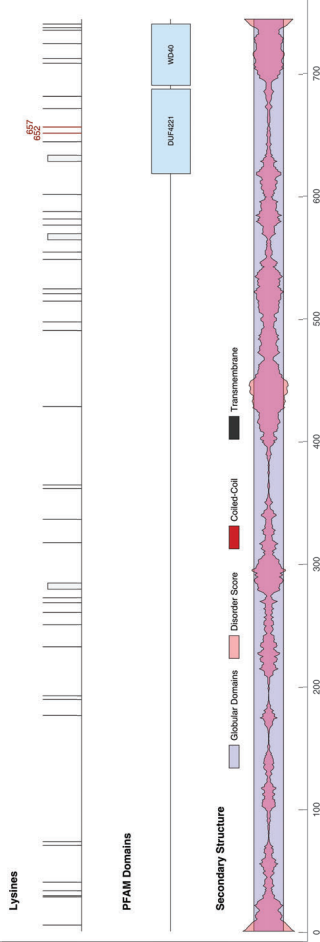




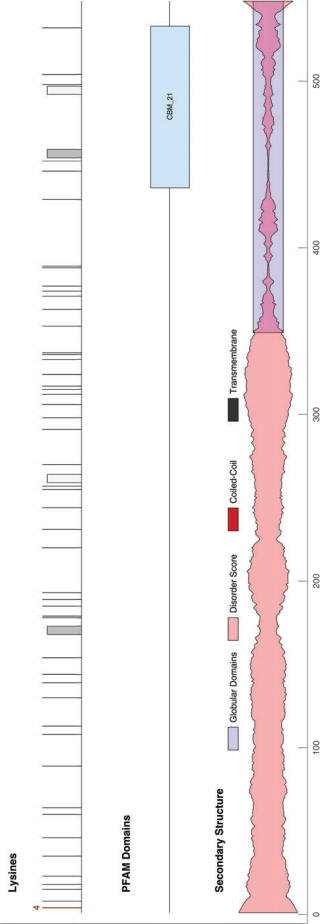
GFD1 (sp|Q04839|GFD1_YEAST)



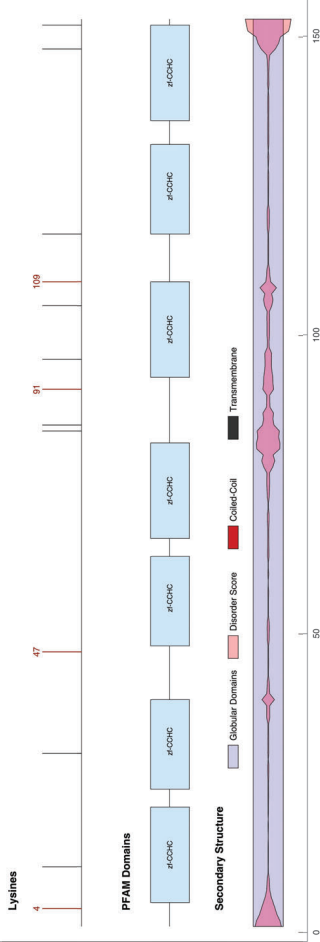
GID7 (sp|P25569|GID7_YEAST)

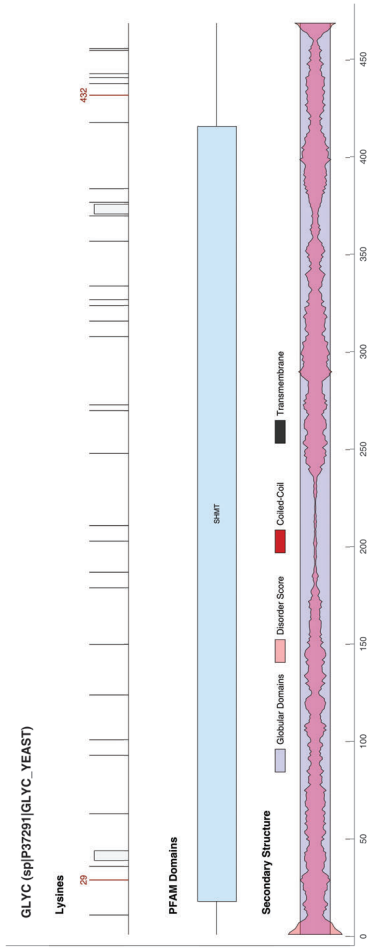
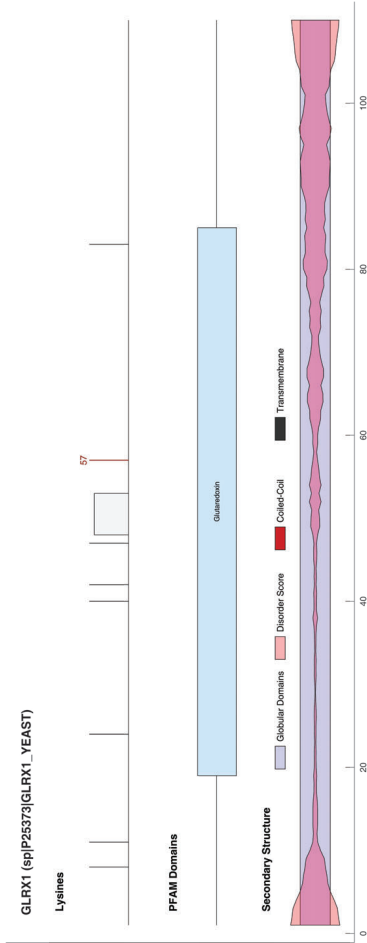
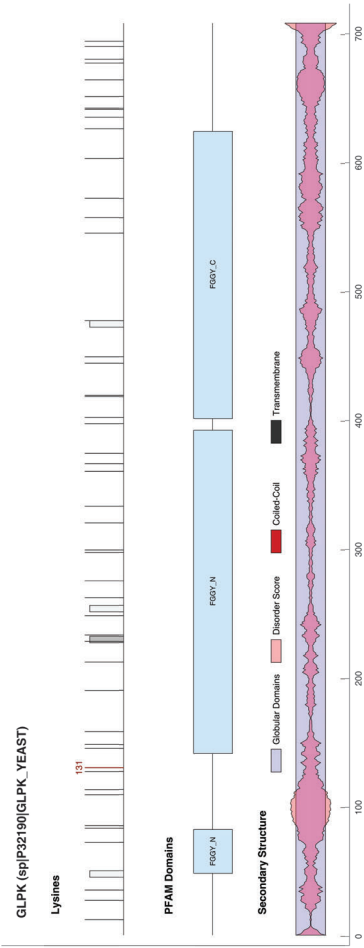
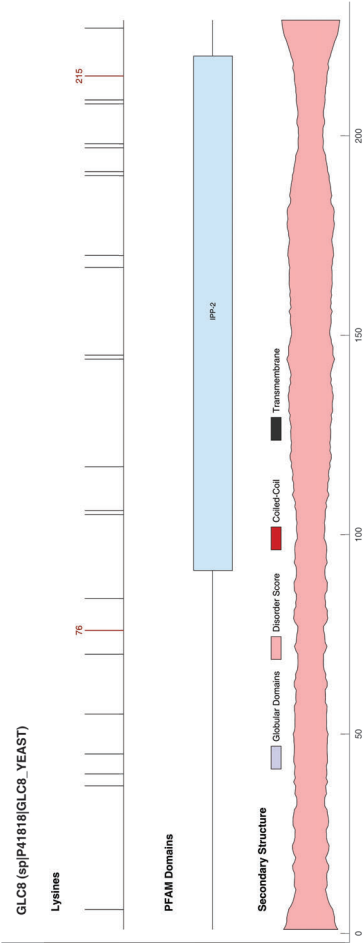


GIP2 (sp|P40036|GIP2_YEAST)

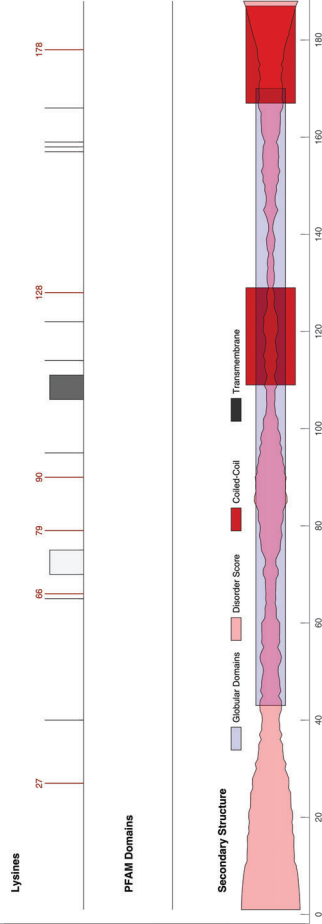


GIS2 (sp|P53849|GIS2_YEAST)

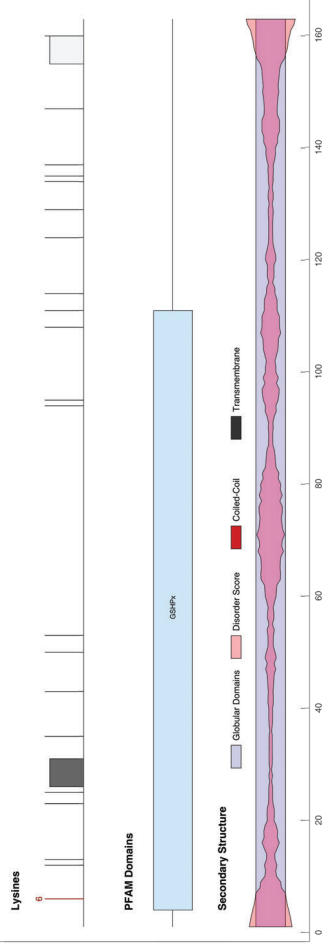




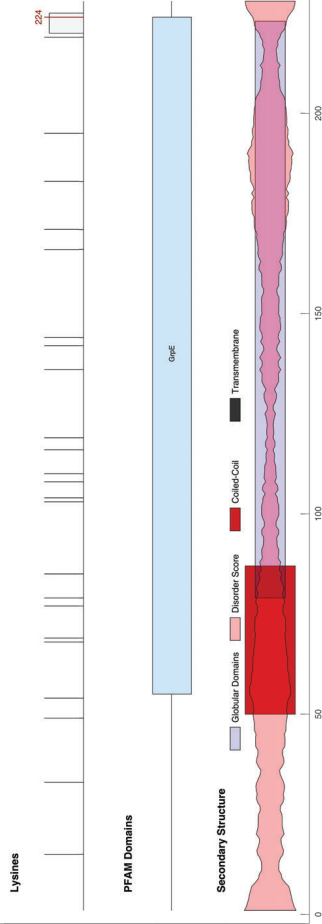
GMC2 (sp|Q06201|GMC2_YEAST)



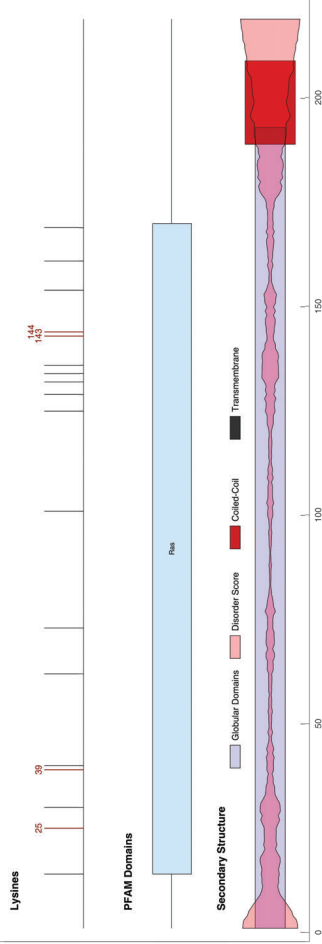
GPX3 (sp|P40581|GPX3_YEAST)

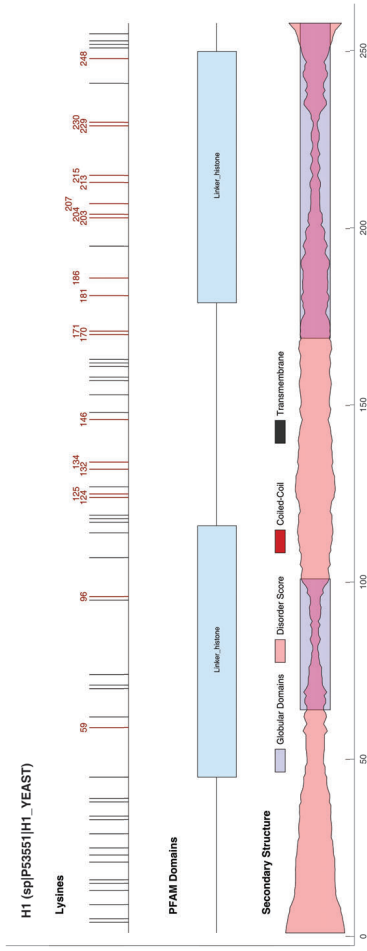
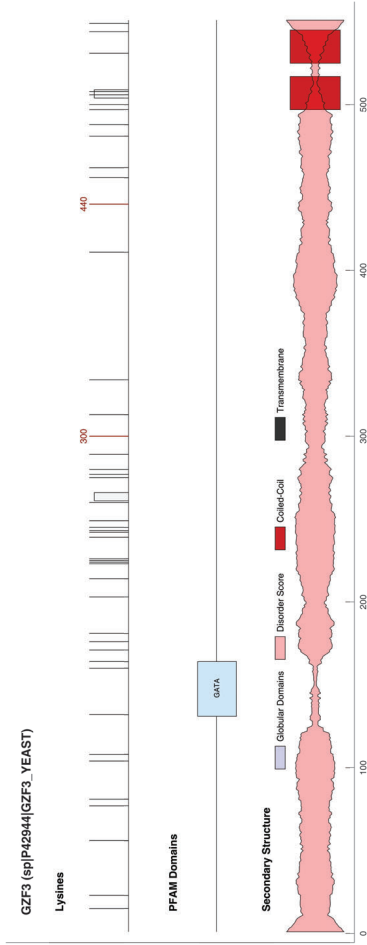
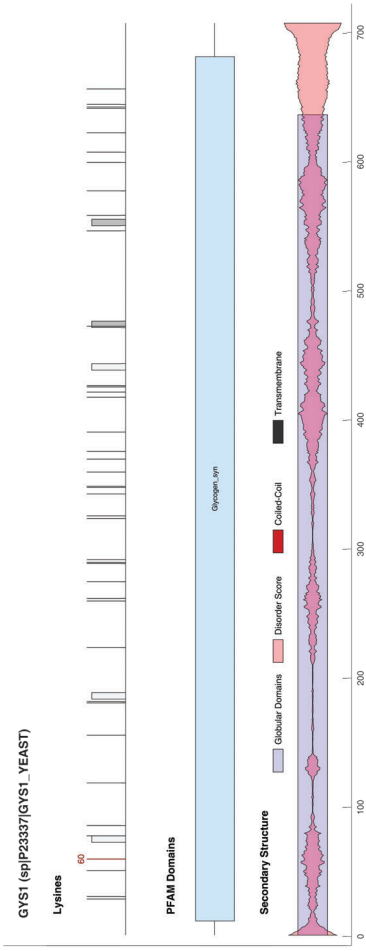
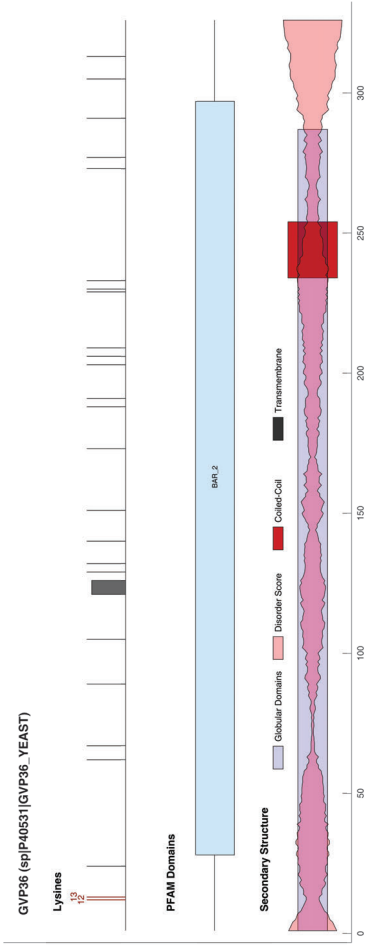


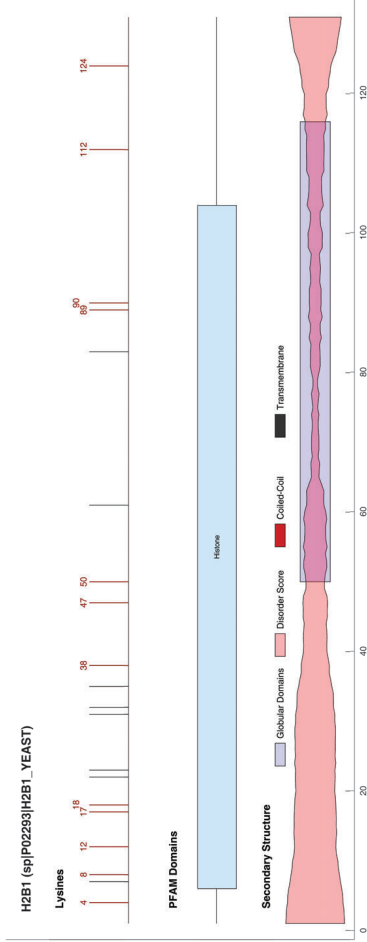
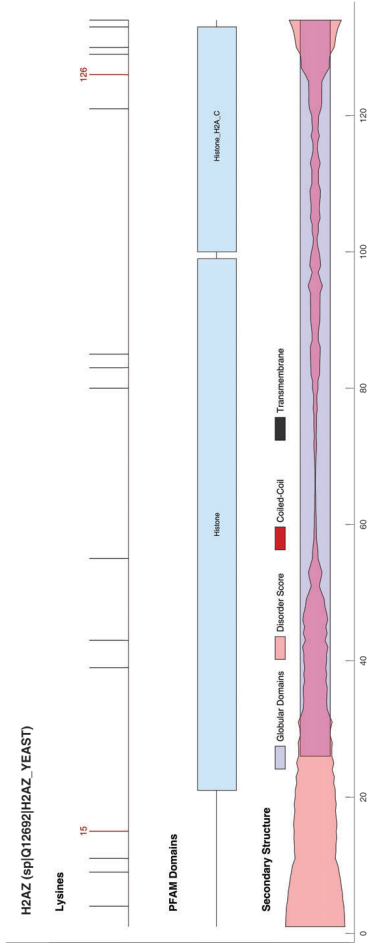
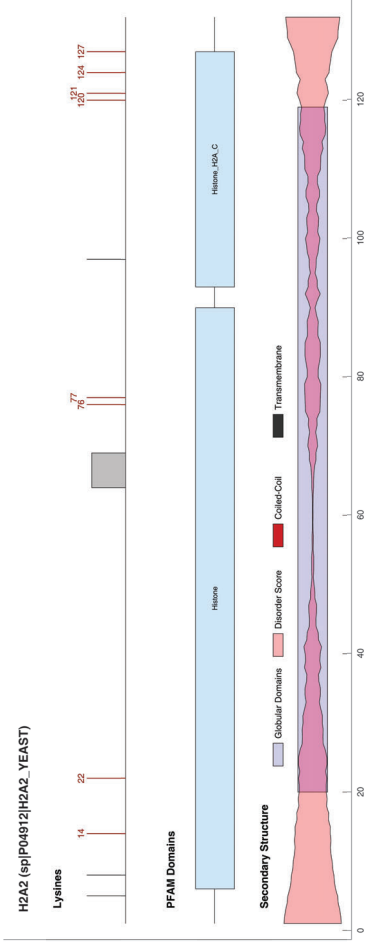
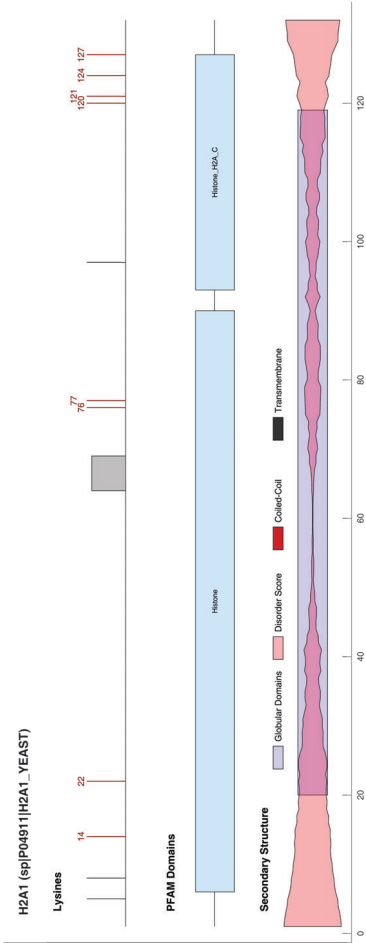
GRPE (sp|P38523|GRPE_YEAST)

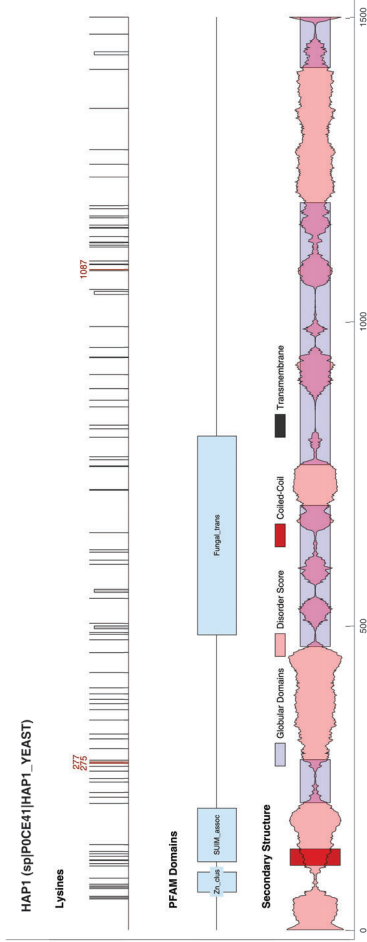
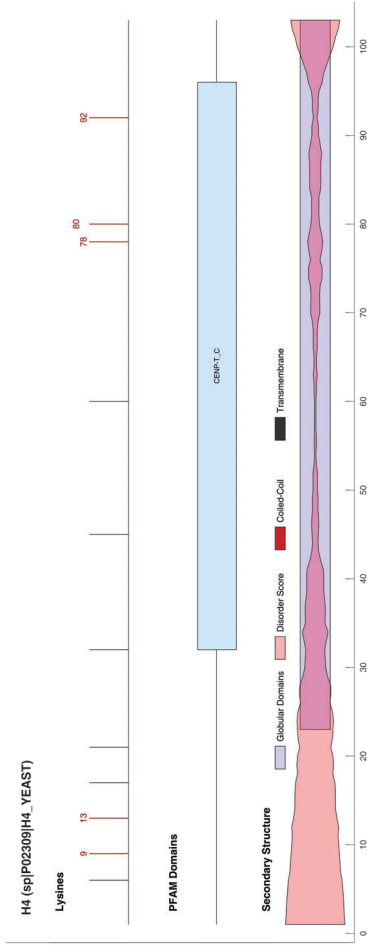
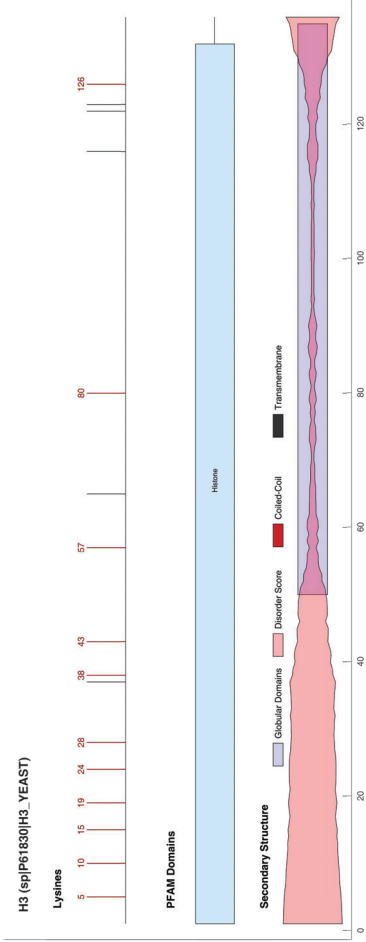
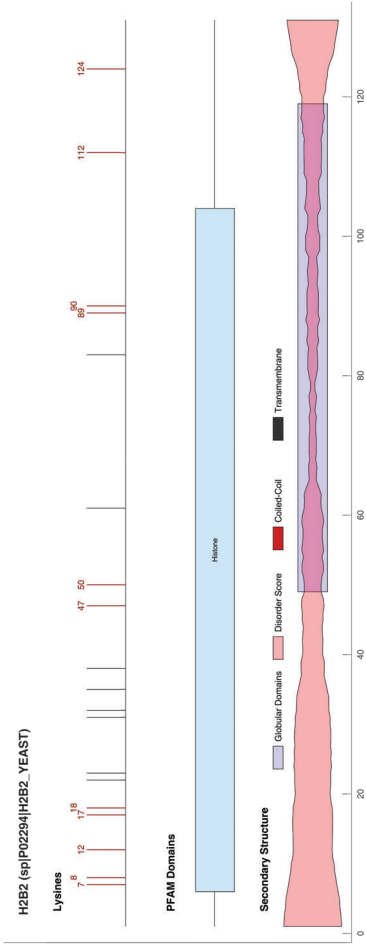


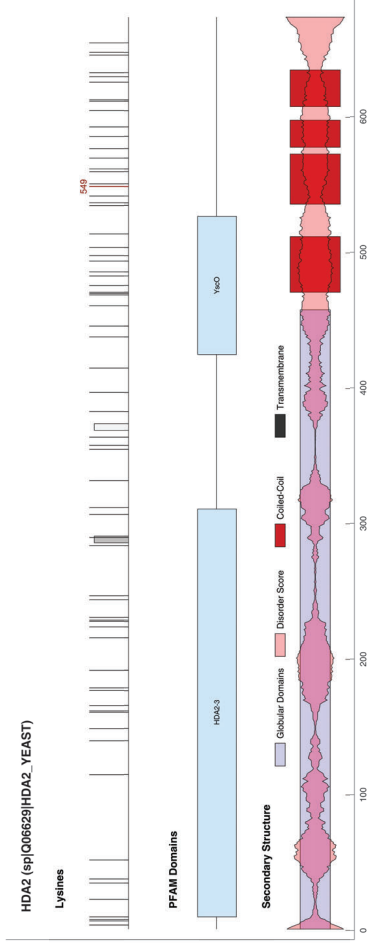
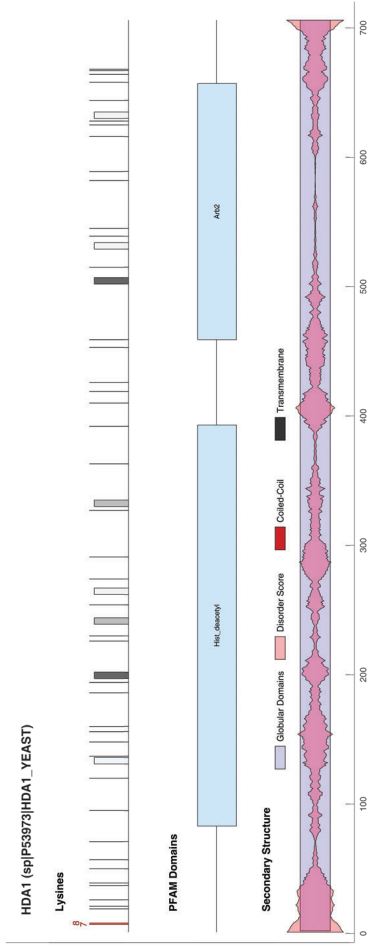
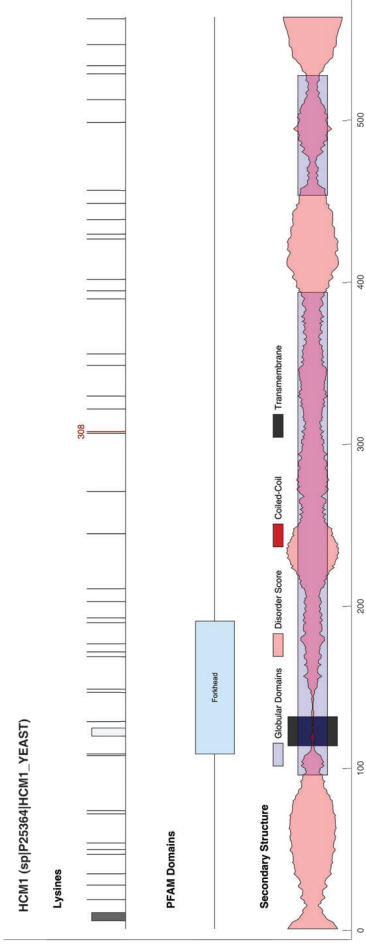
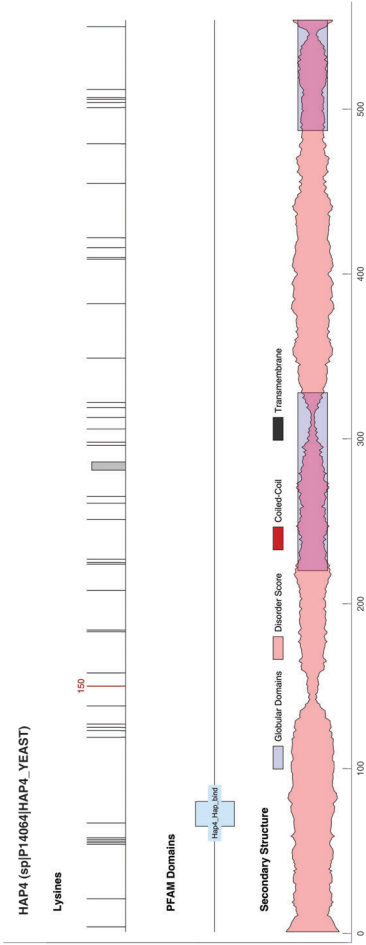
GSP1 (sp|P32835|GSP1_YEAST)

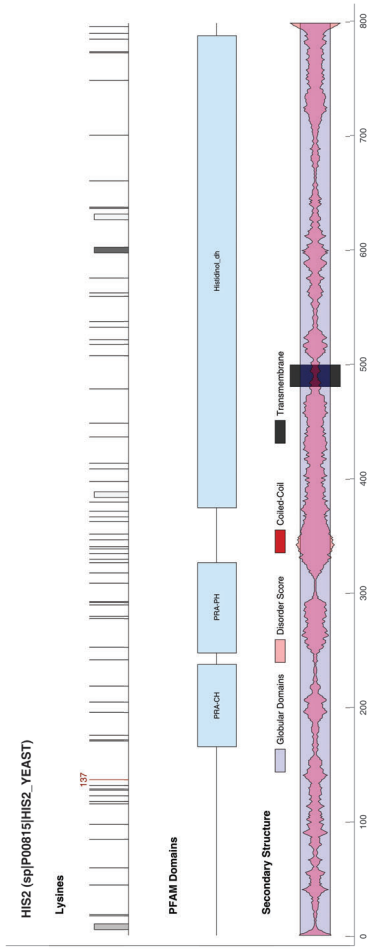
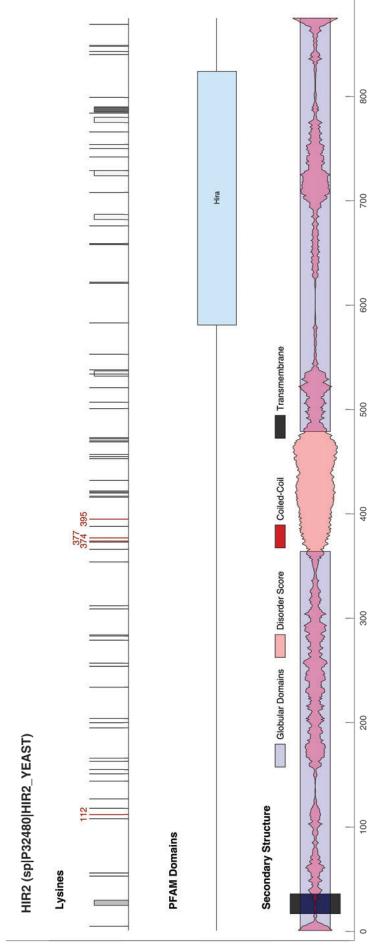
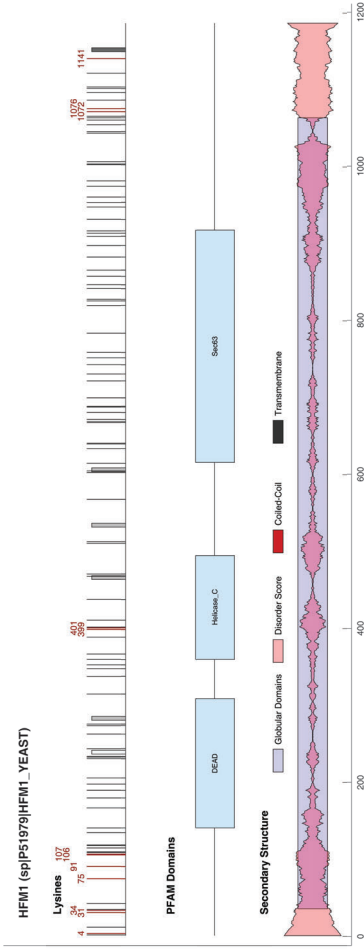
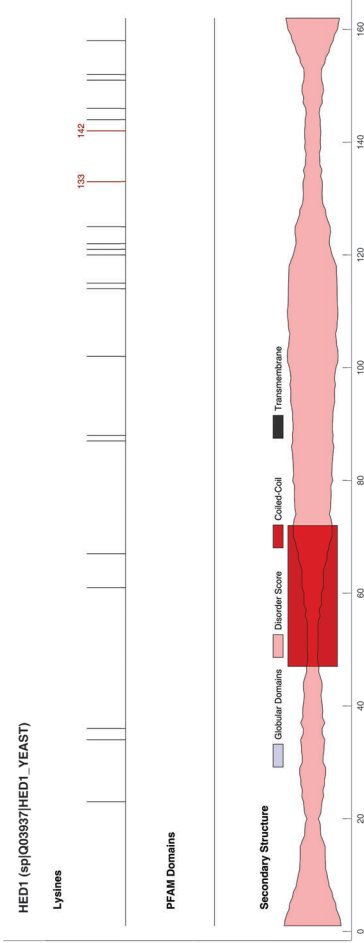


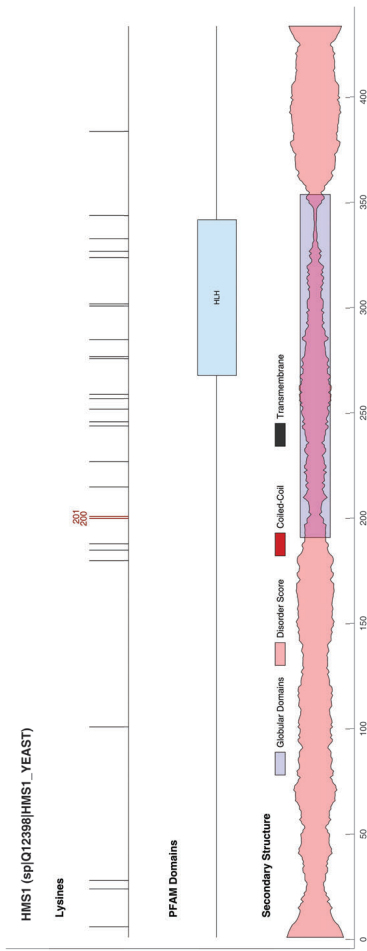
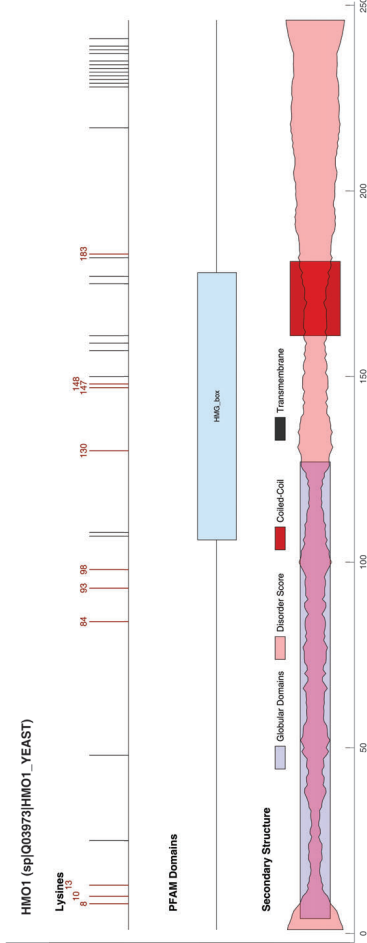
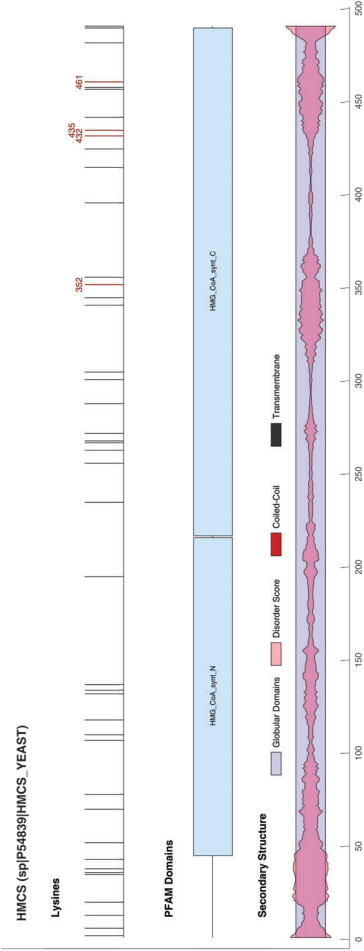
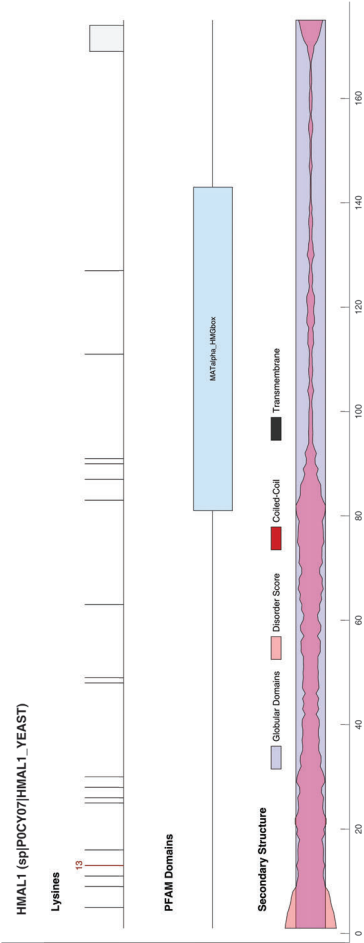


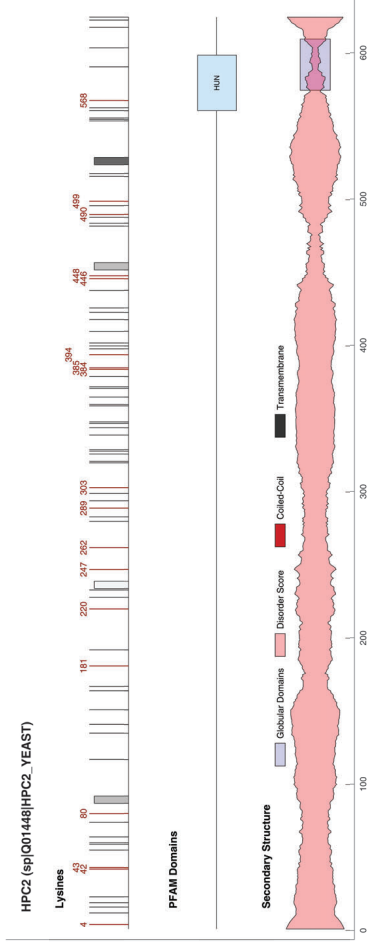
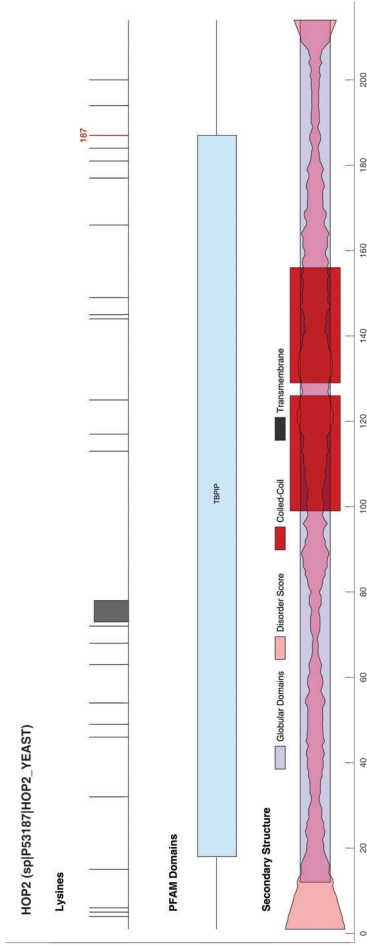
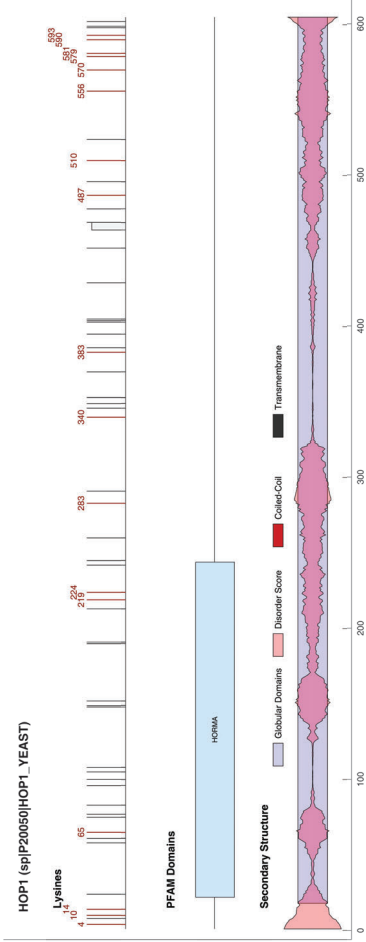
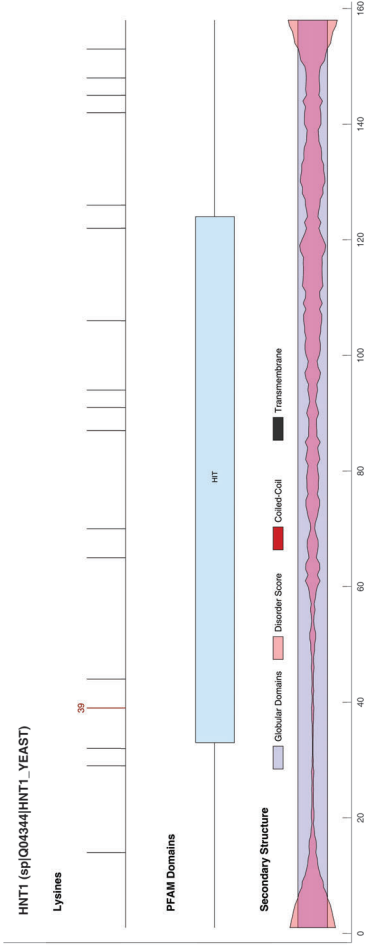


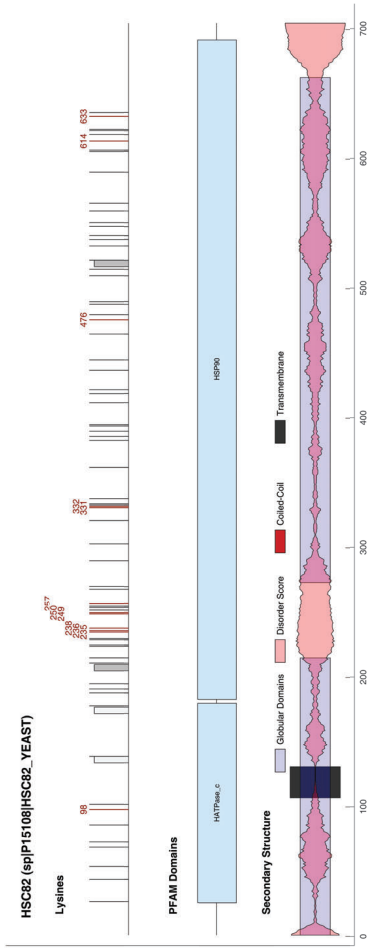
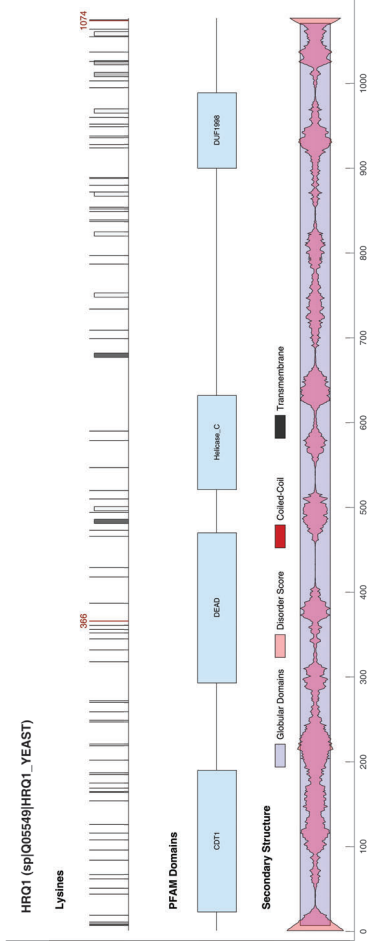
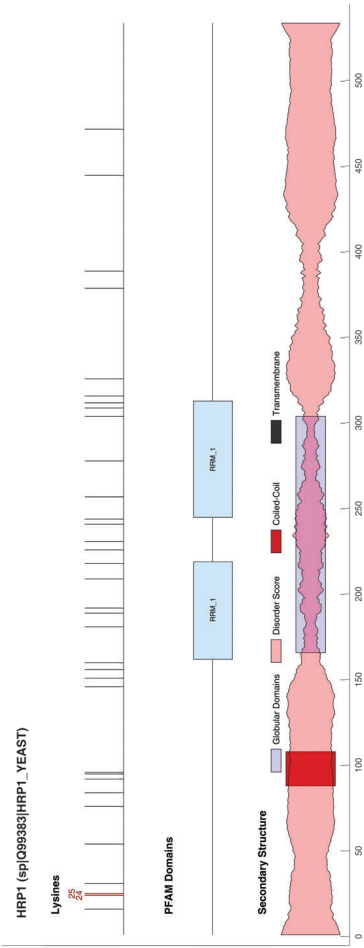
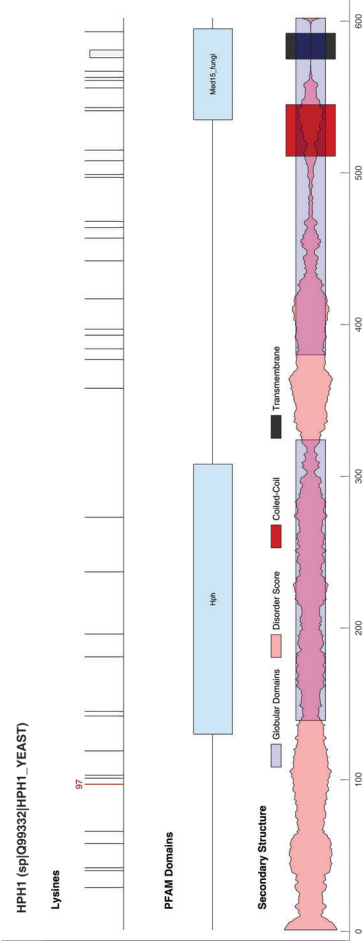




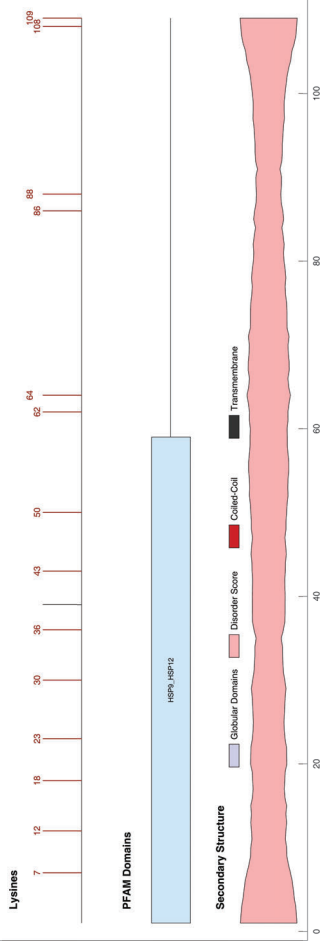




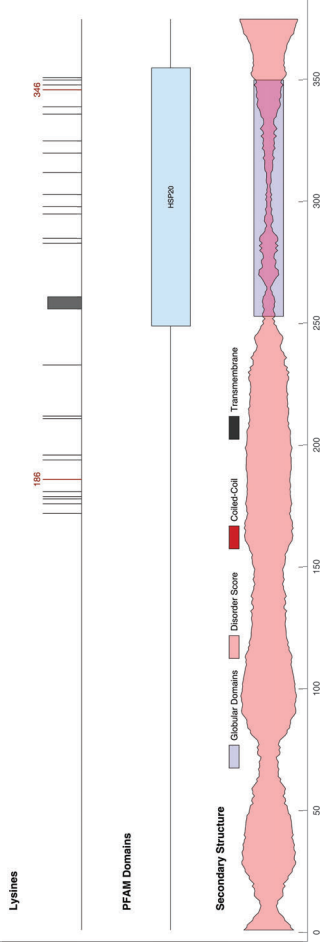




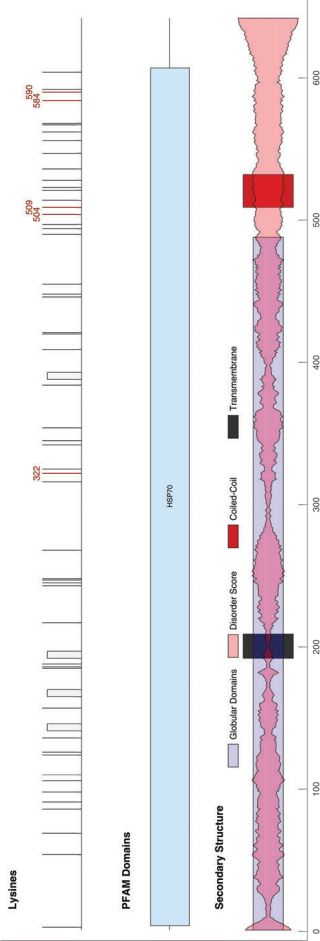
HSP12 (sp|P22943|HSP12_YEAST)



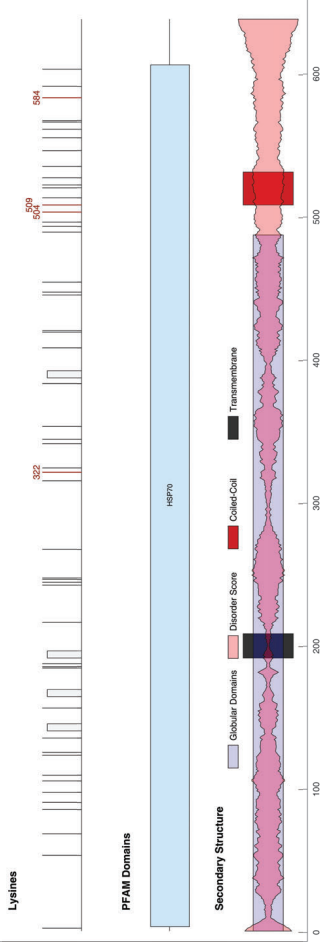
HSP42 (sp|Q12329|HSP42_YEAST)

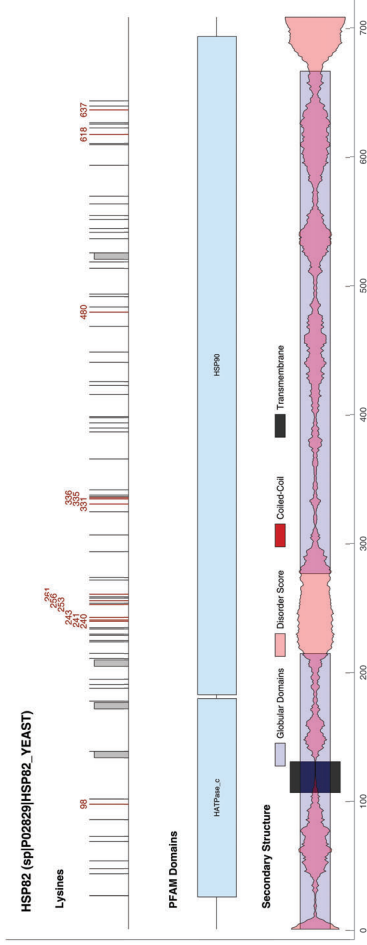
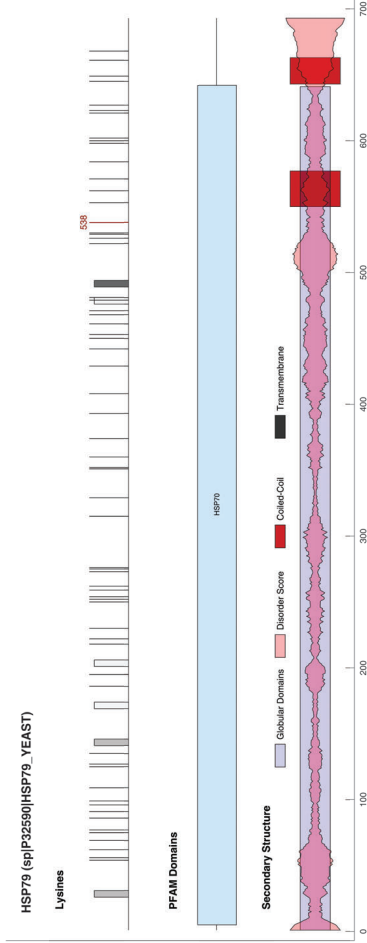
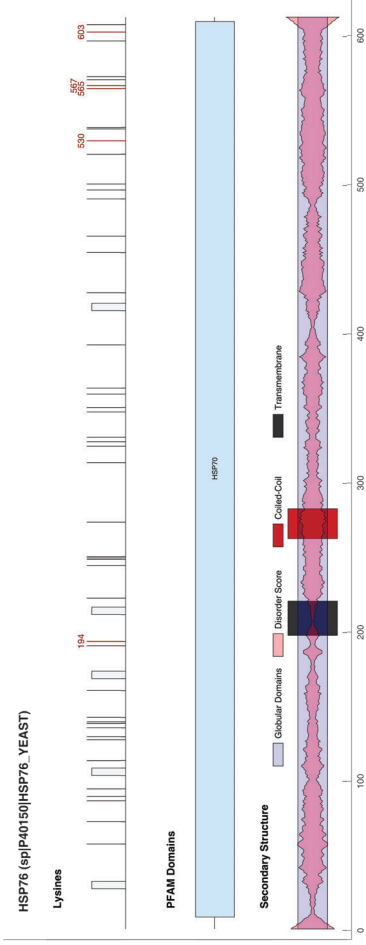
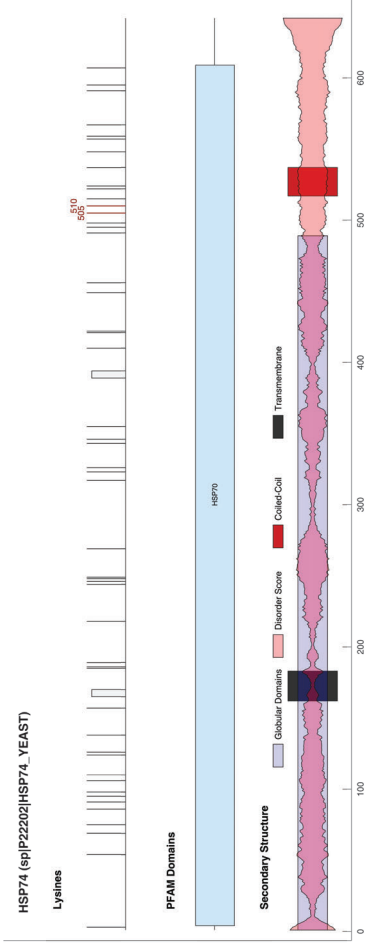


HSP71 (sp|P10591|HSP71_YEAST)

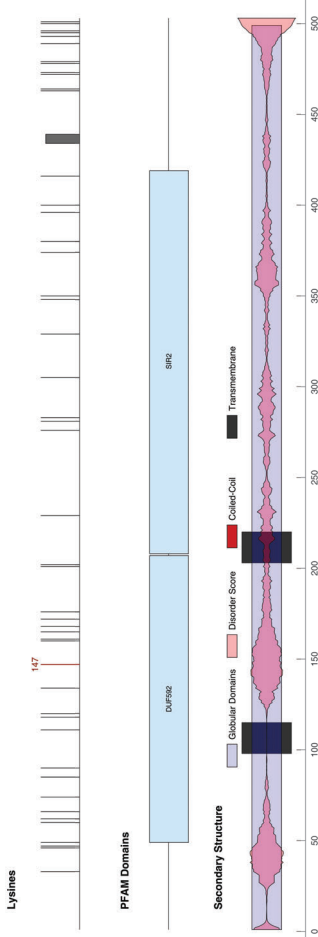


HSP72 (sp|P10592|HSP72_YEAST)

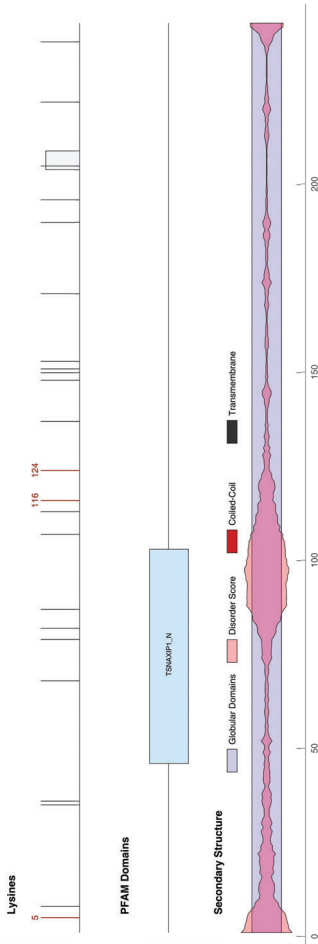




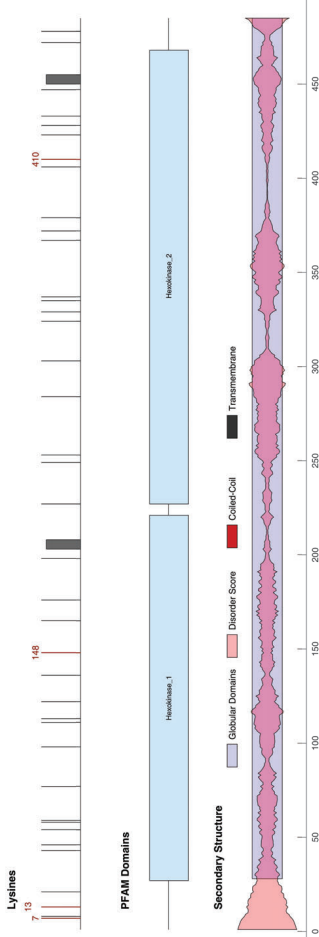
HST1 (sp|P53685|HST1_YEAST)



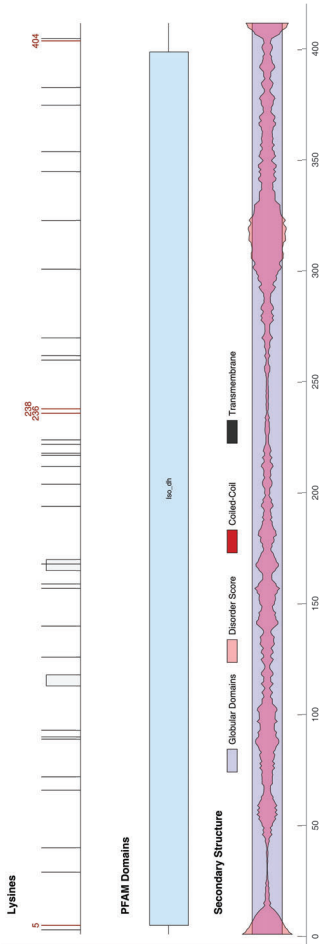
HUA2 (sp|Q12134|HUA2_YEAST)



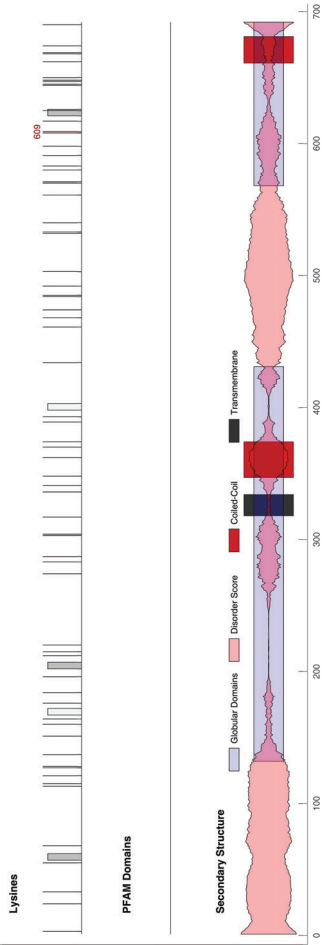
HXKA (sp|P04806|HXKA_YEAST)



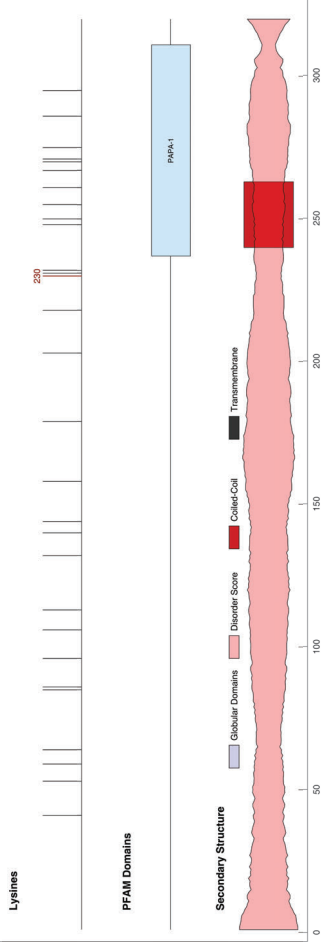
IDHC (sp|P41939|IDHC_YEAST)



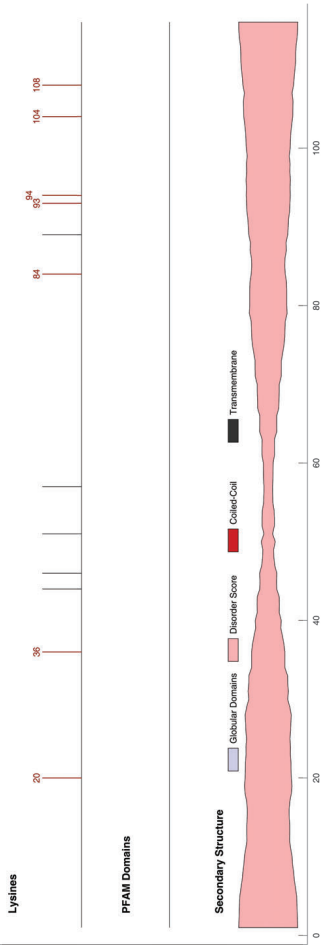
IES1 (sp|P43579|IES1_YEAST)



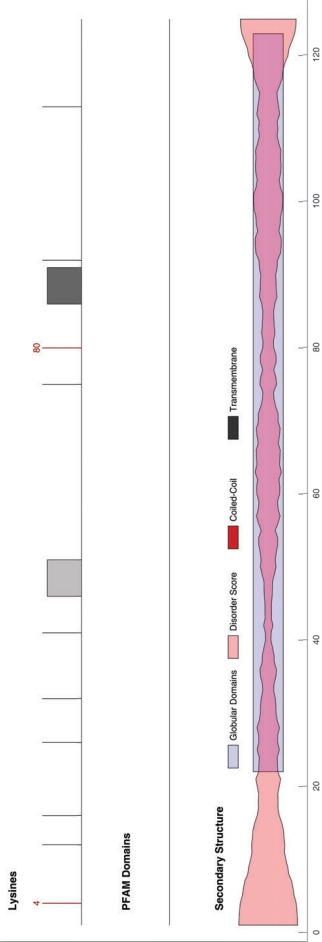
IES2 (sp|P40154|IES2_YEAST)



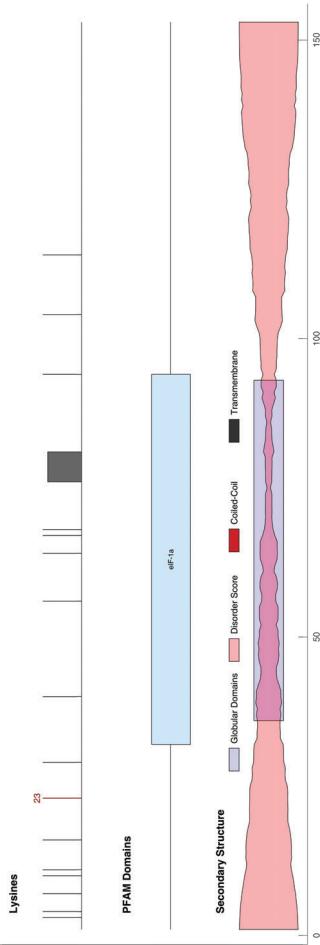
IES4 (sp|Q08561|IES4_YEAST)



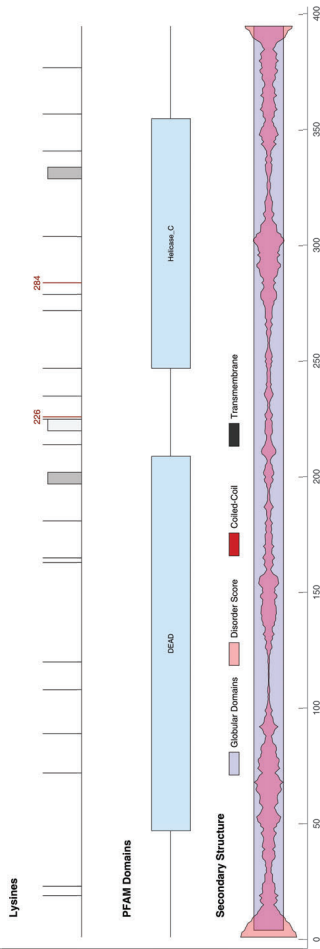
IES5 (sp|P40060|IES5_YEAST)



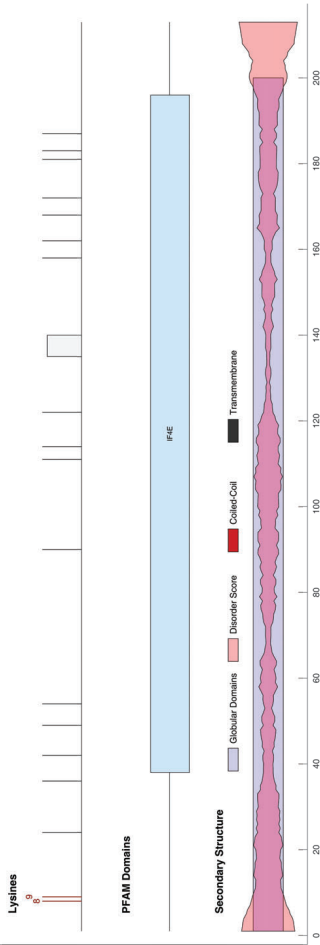
IF1A (sp|P38912|IF1A_YEAST)



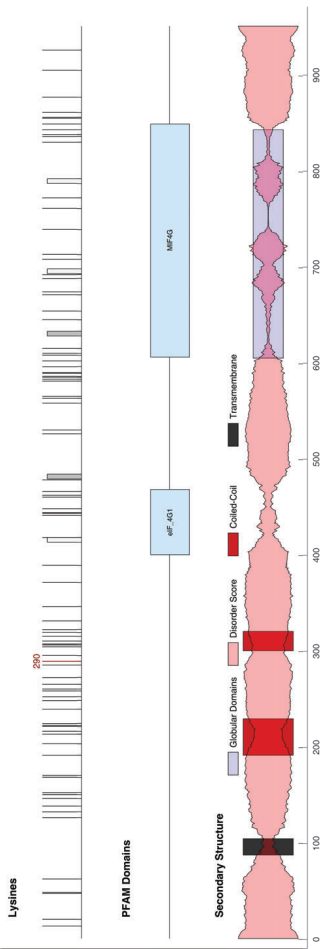
IF4A (sp|P10081|IF4A_YEAST)

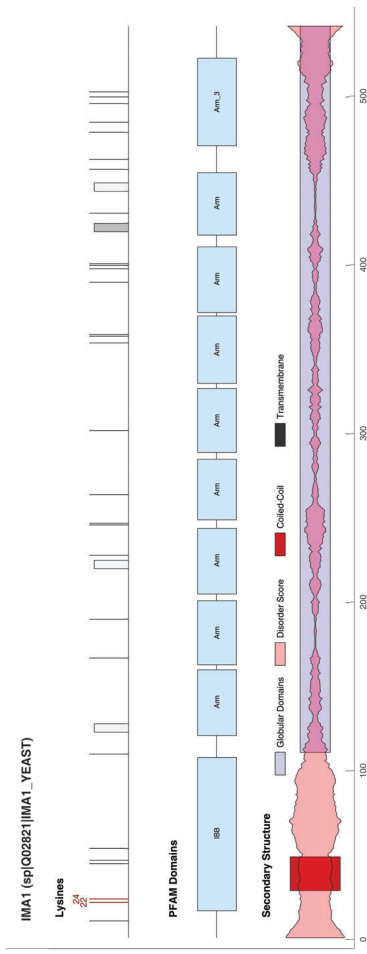
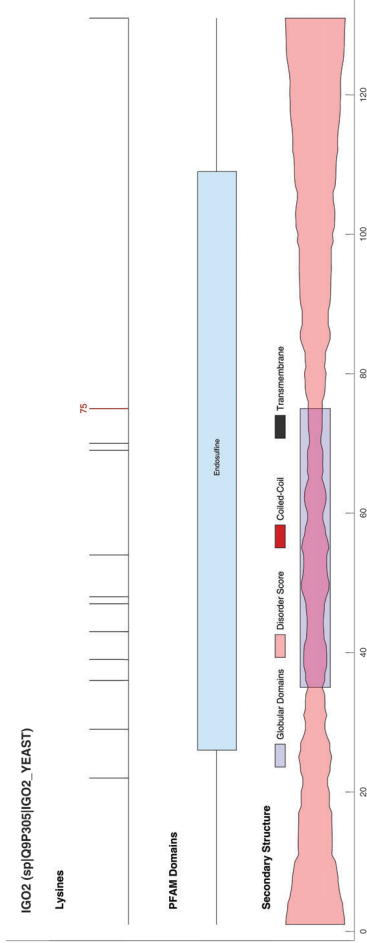
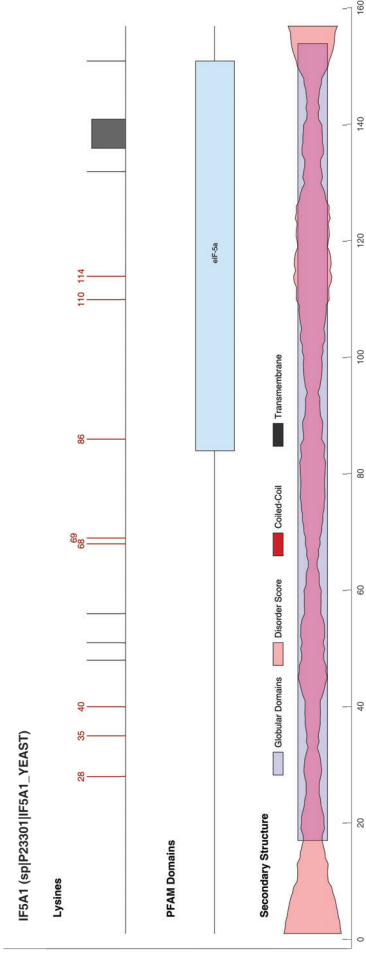
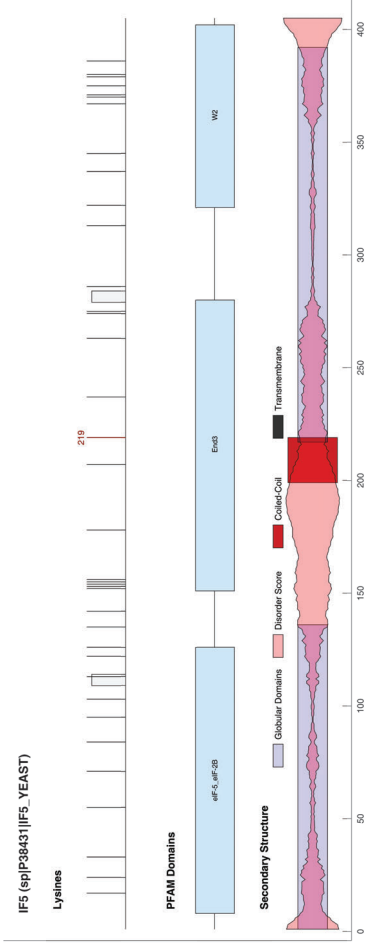


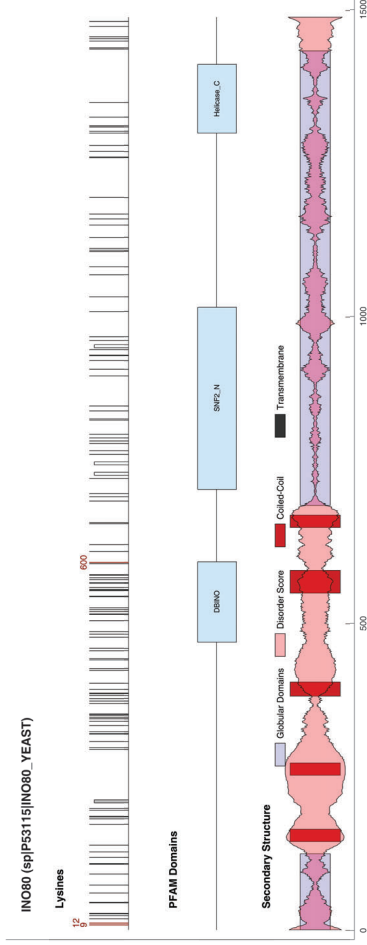
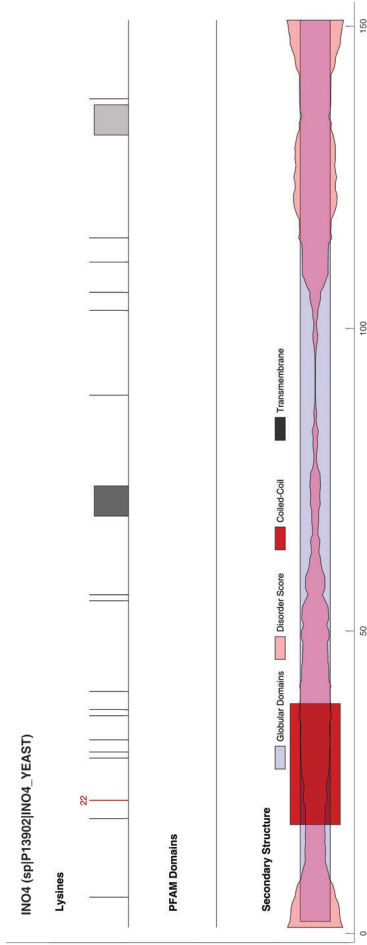
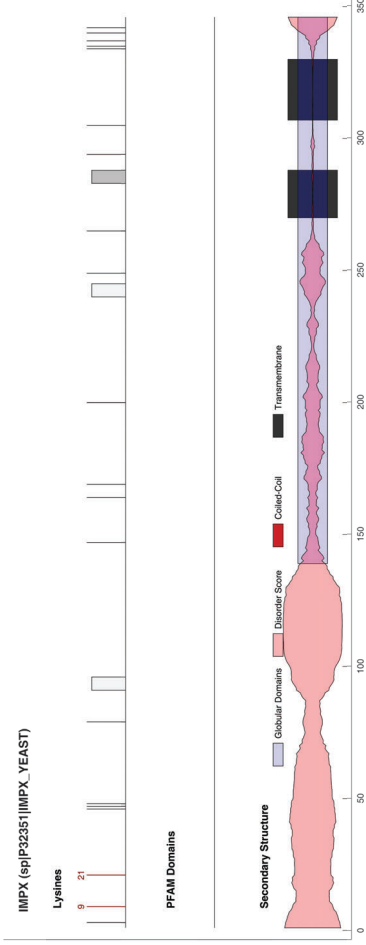
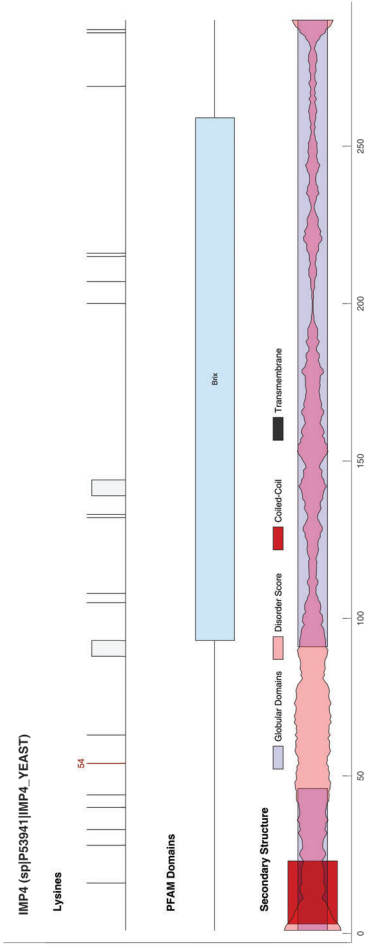
IF4E (sp|P07260|IF4E_YEAST)

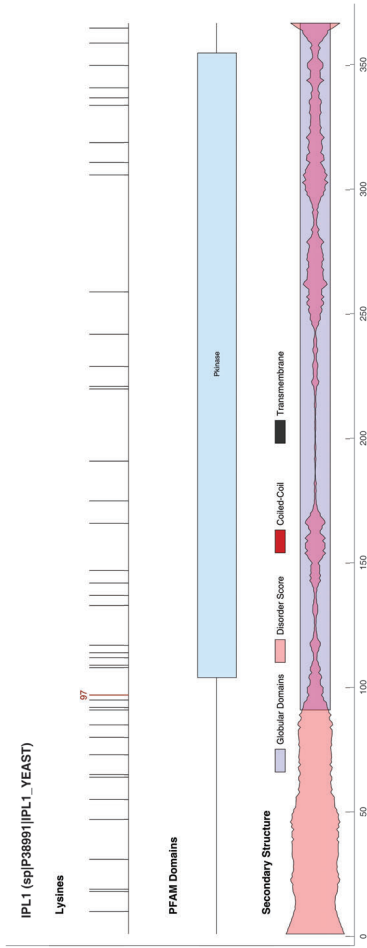
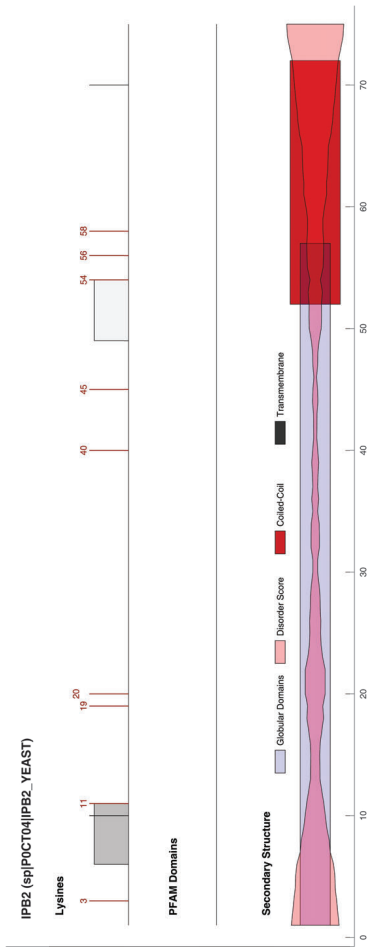
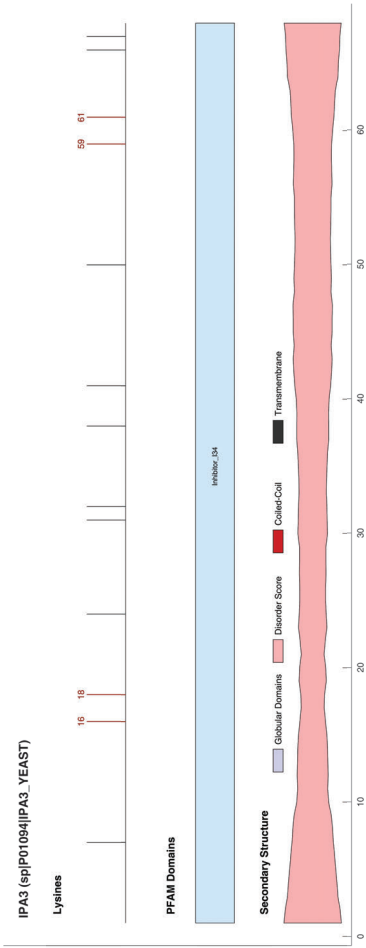
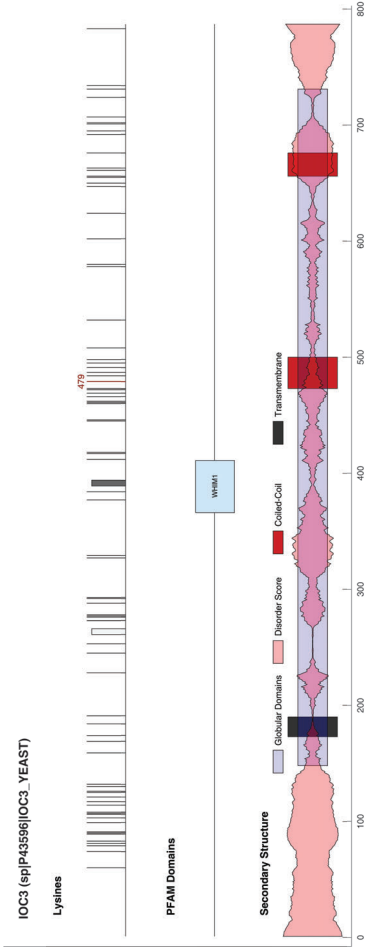


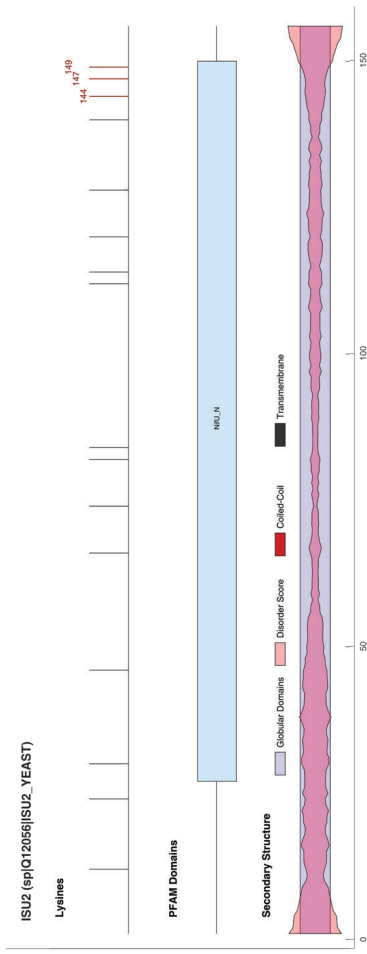
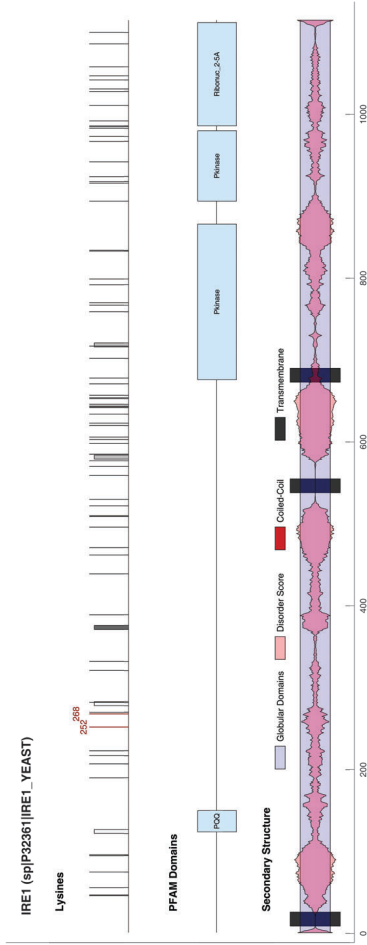
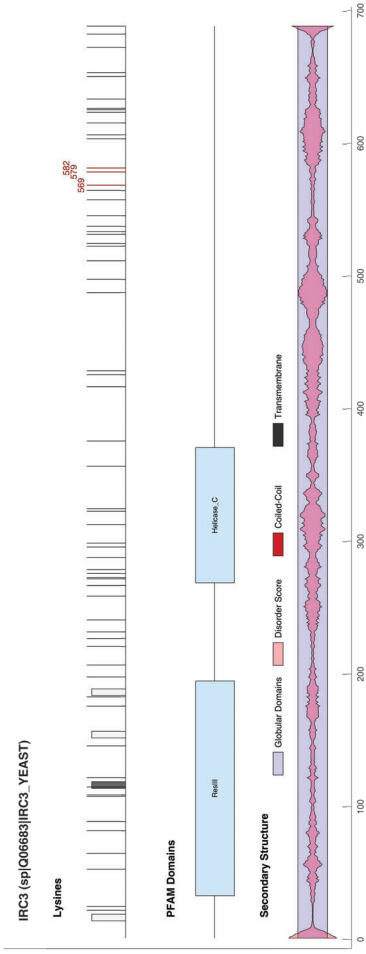
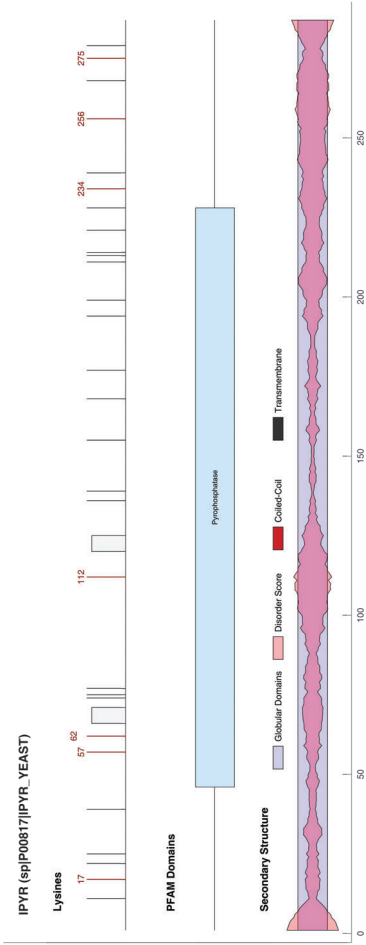
IF4F1 (sp|P39535|IF4F1_YEAST)

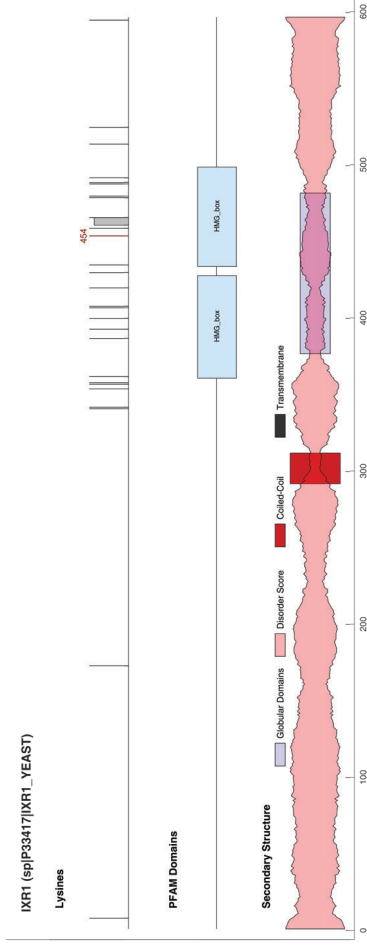
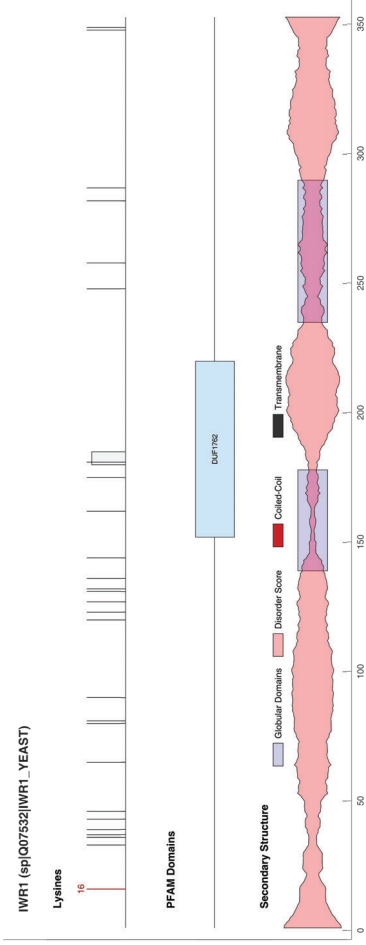
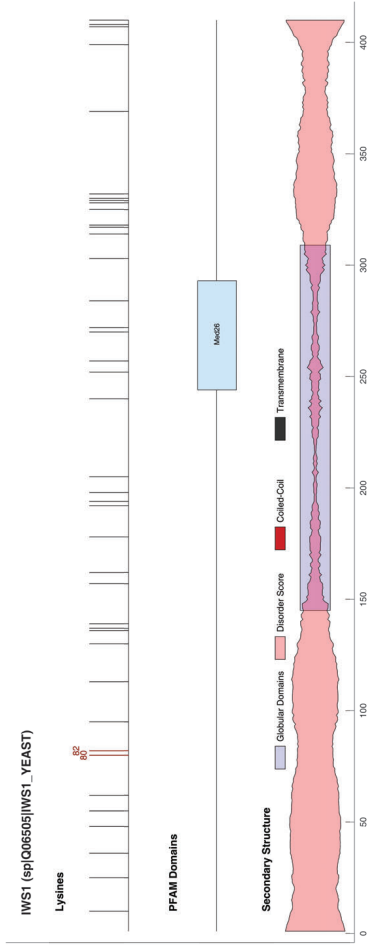
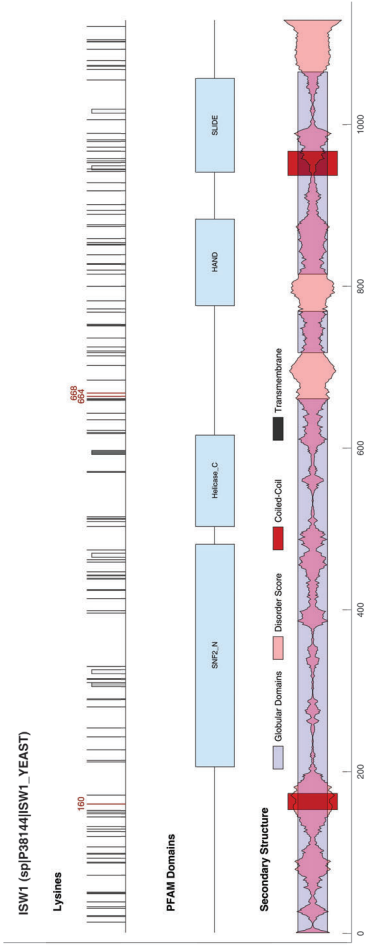


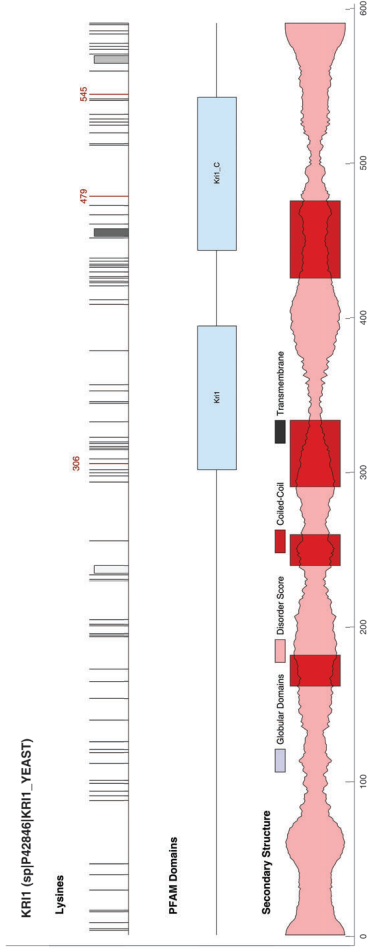
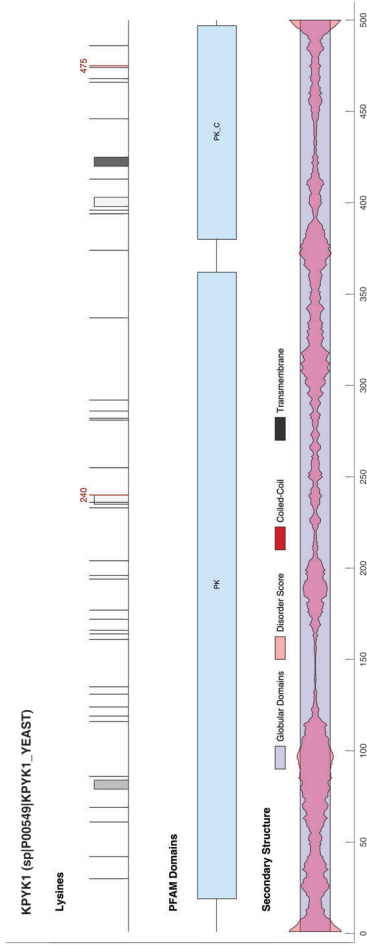
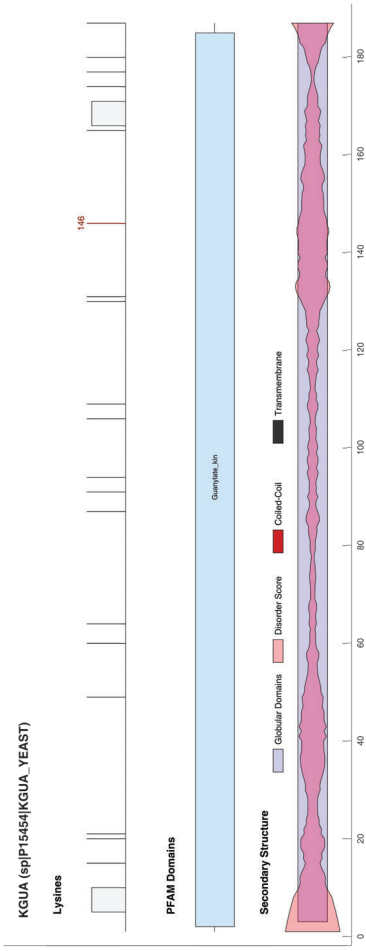
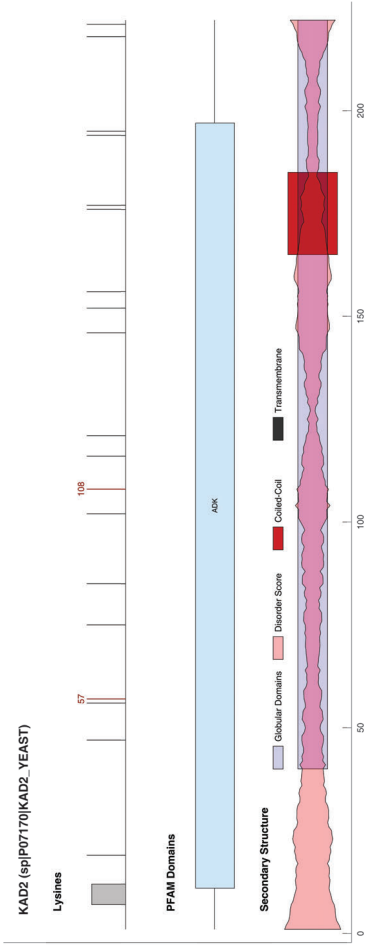


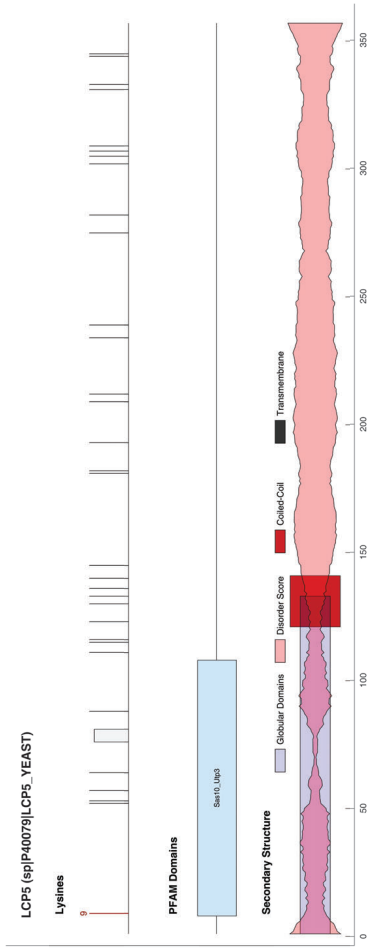
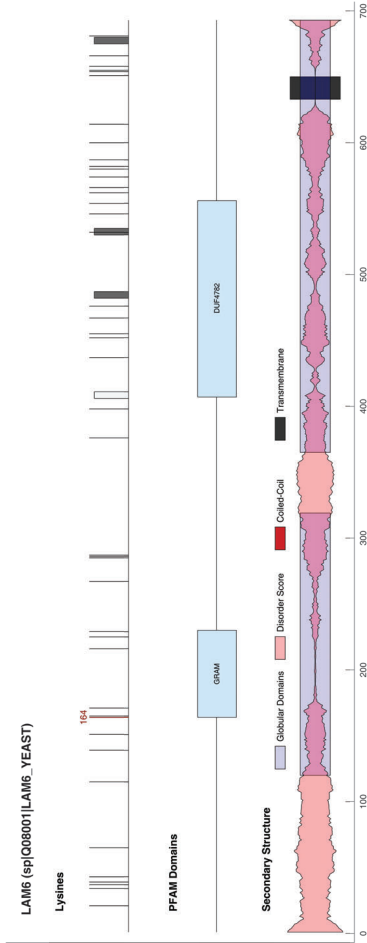
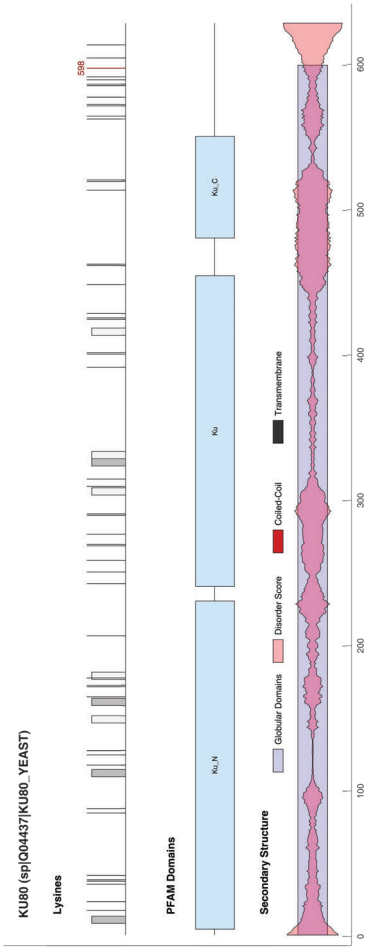
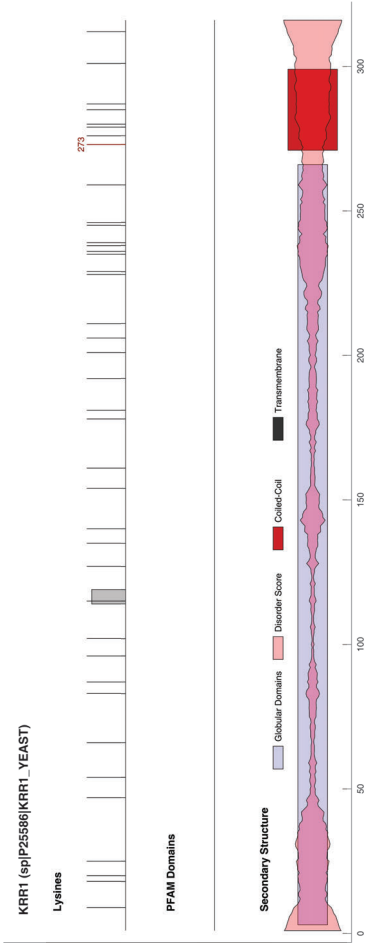


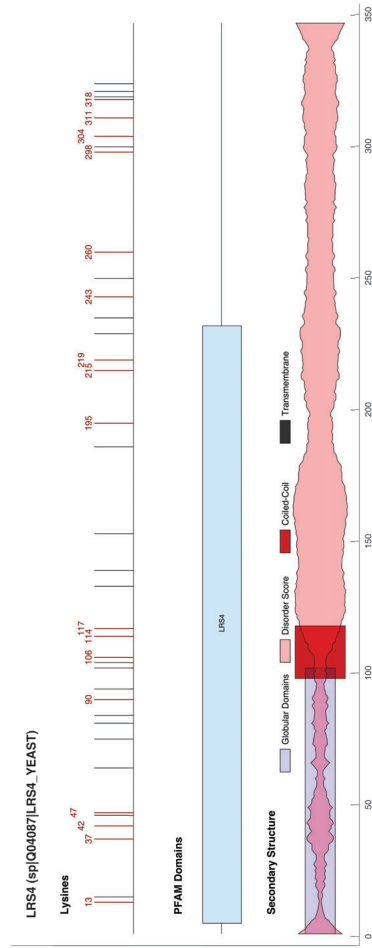
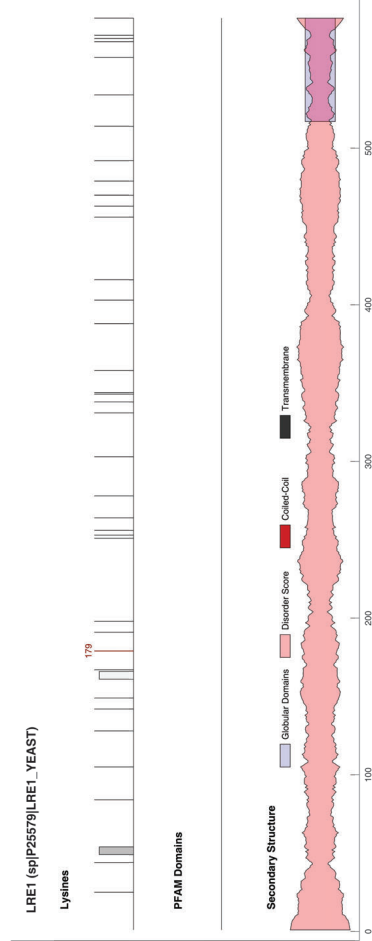
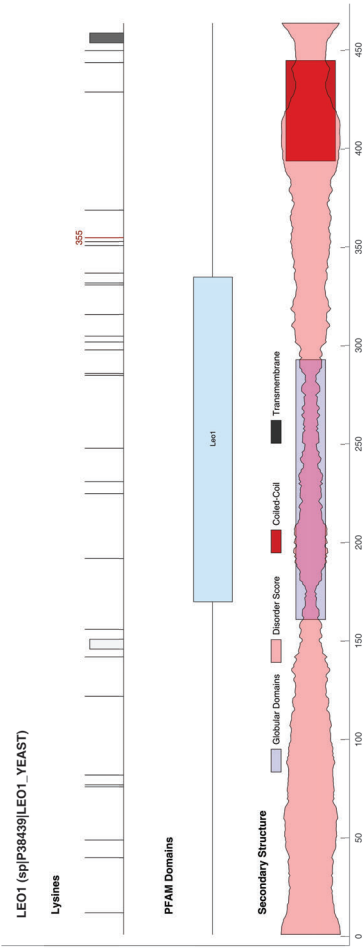
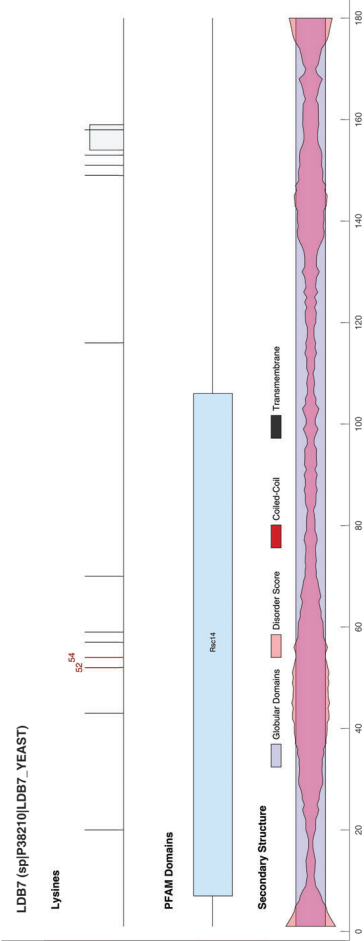


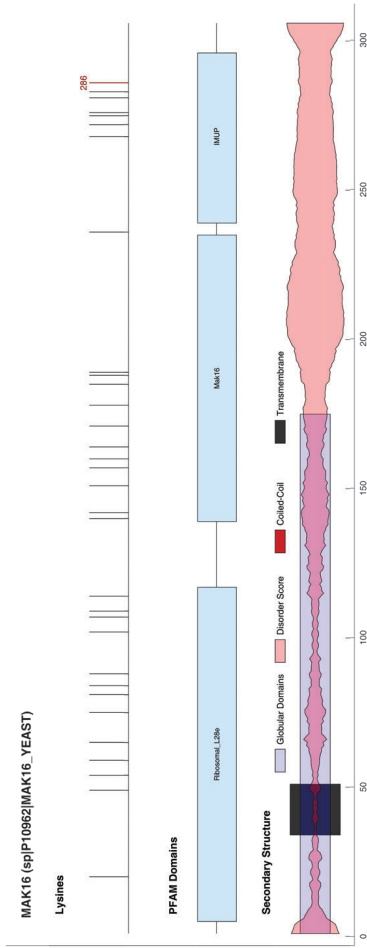
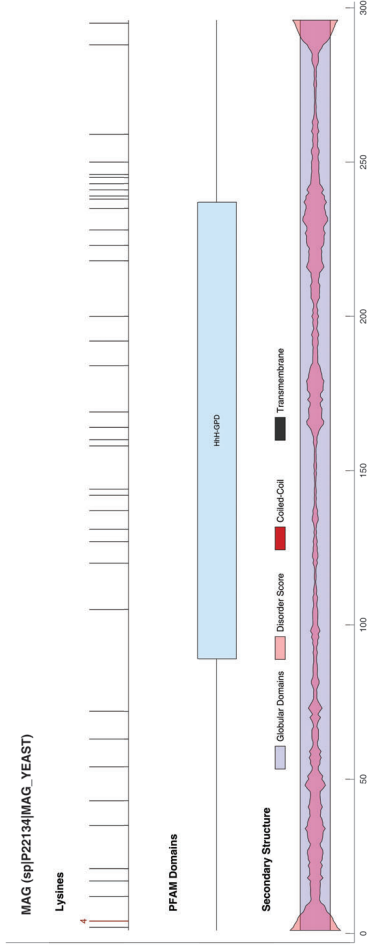
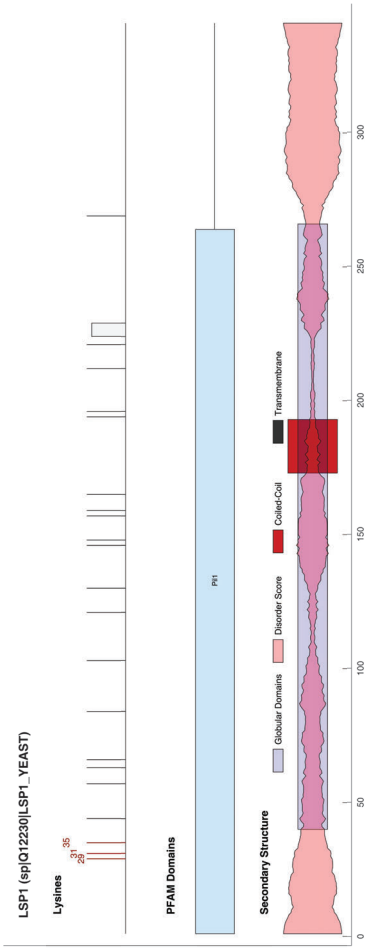
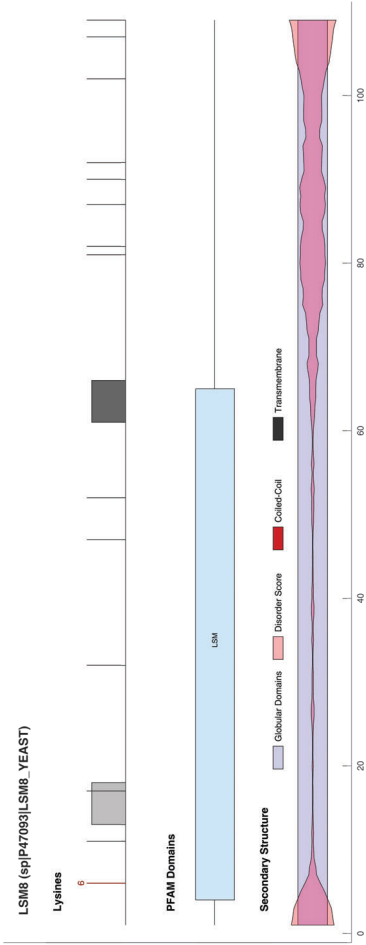












MAM1 (sp|P40065|MAM1_YEAST)

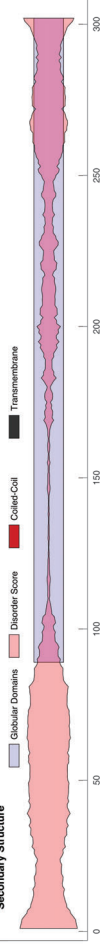
Lysines



PFAM Domains



Secondary Structure



MASY (sp|P30952|MASY_YEAST)

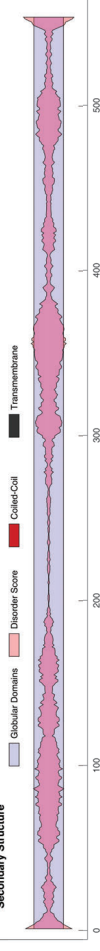
Lysines



PFAM Domains



Secondary Structure



MBF1 (sp|O14467|MBF1_YEAST)

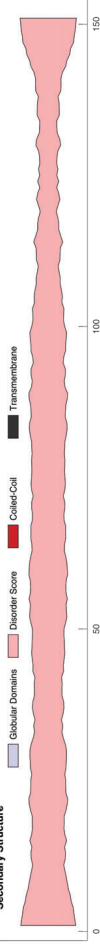
Lysines



PFAM Domains



Secondary Structure



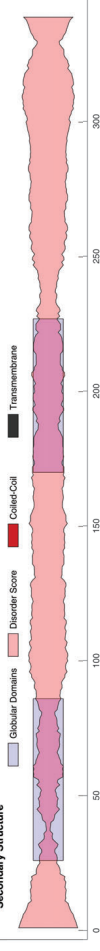
MBR1 (sp|P23493|MBR1_YEAST)

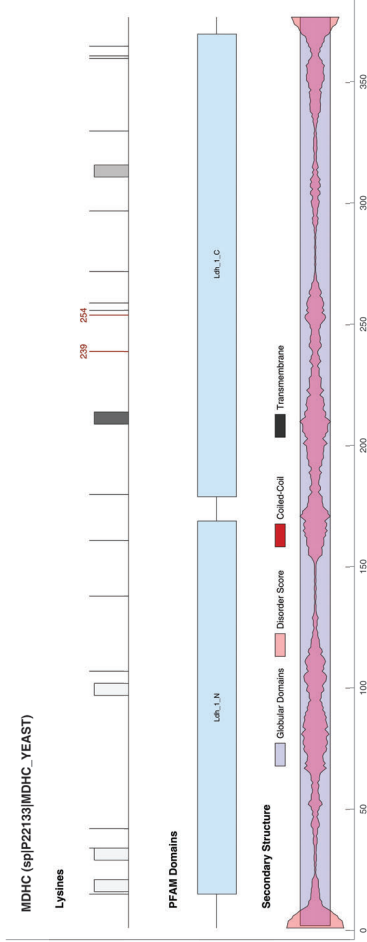
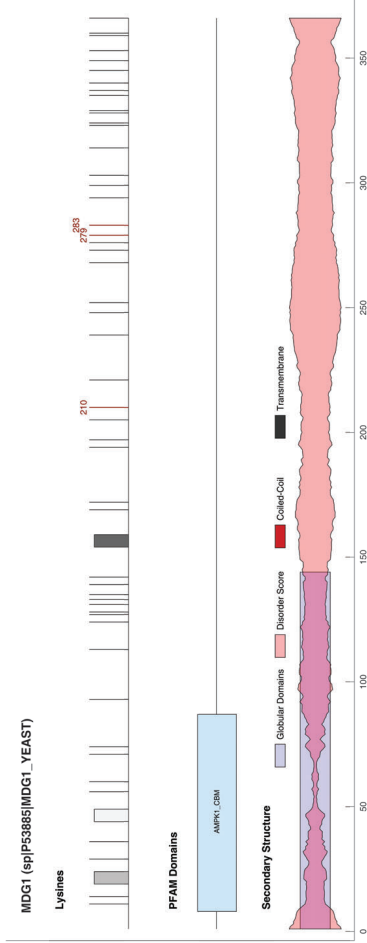
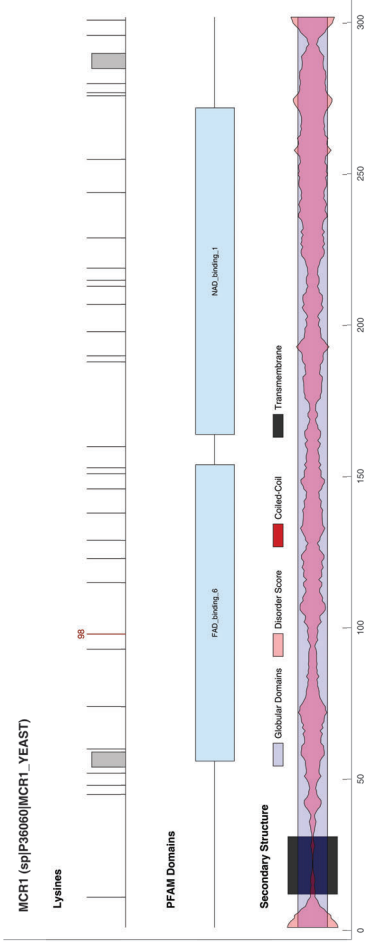
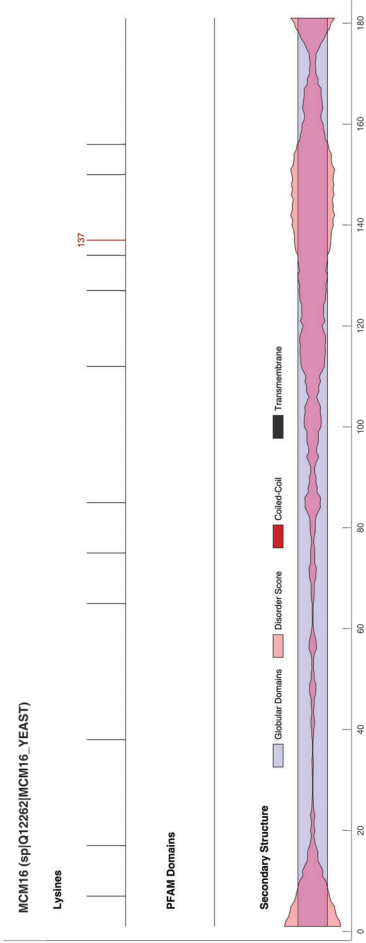
Lysines

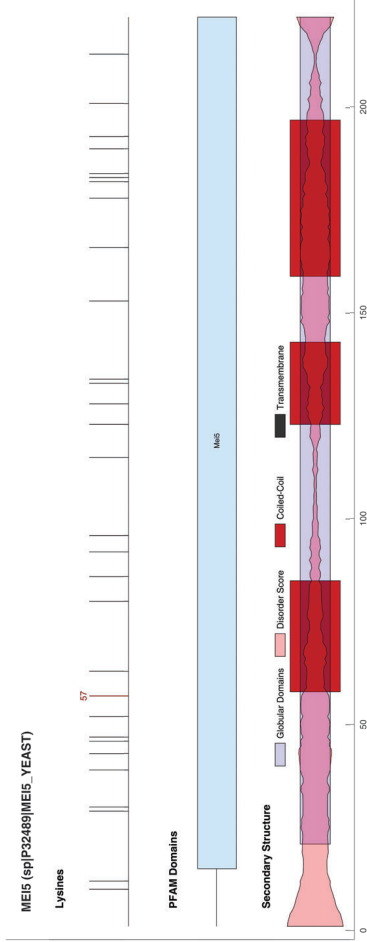
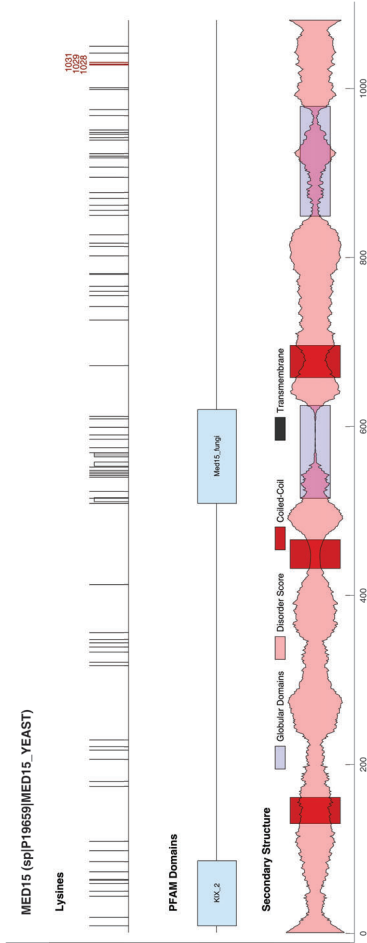
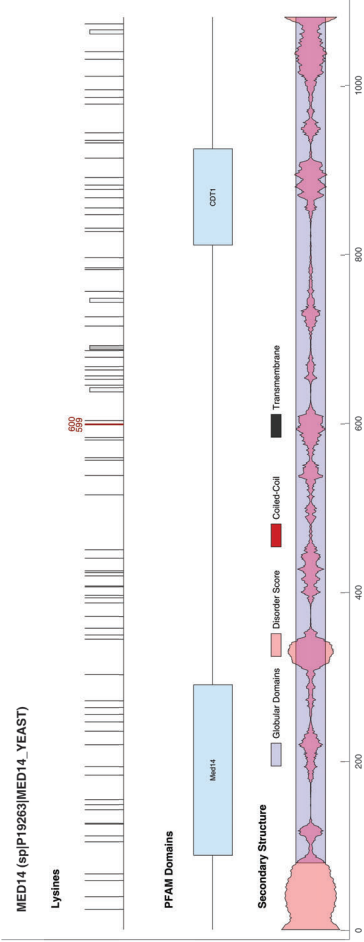
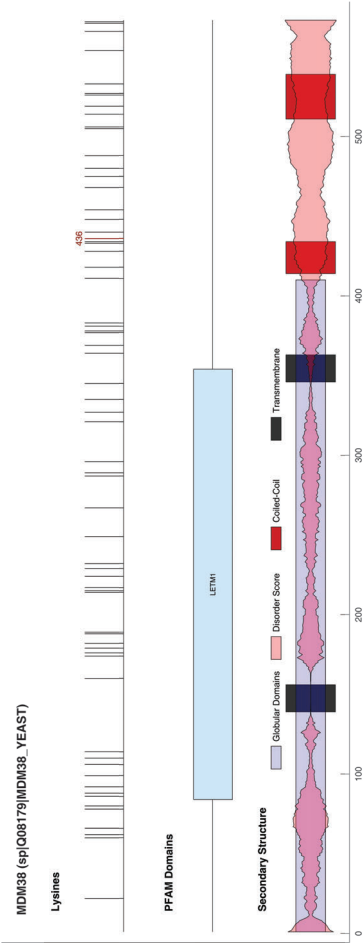


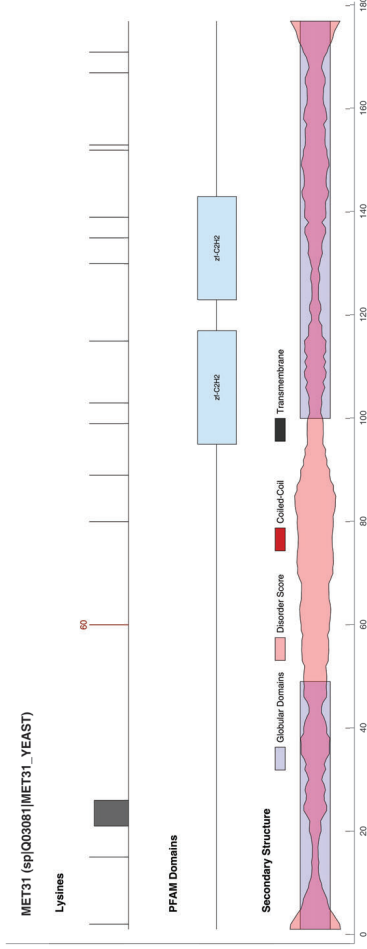
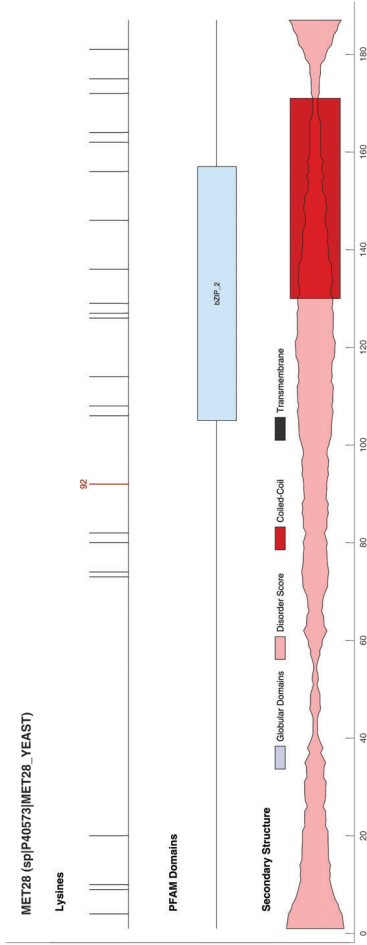
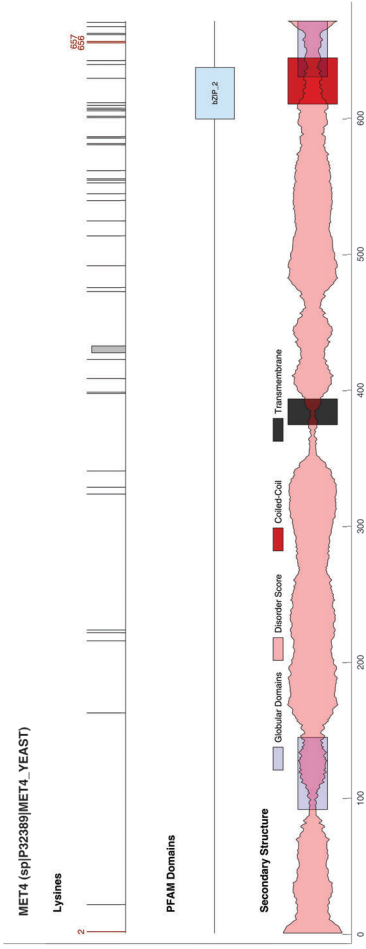
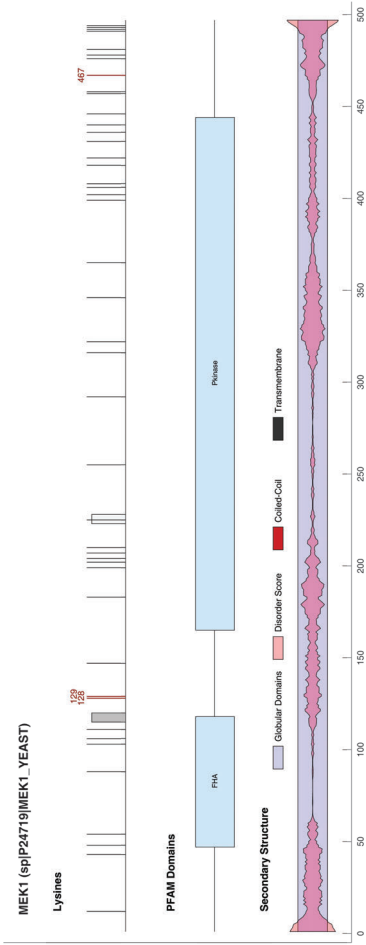
PFAM Domains

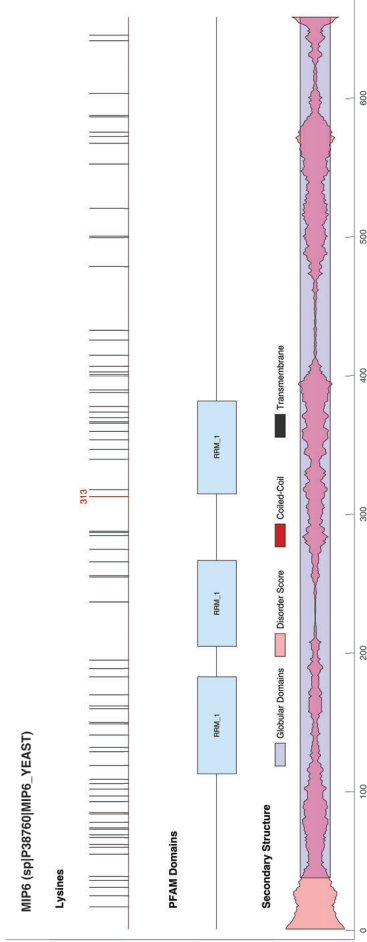
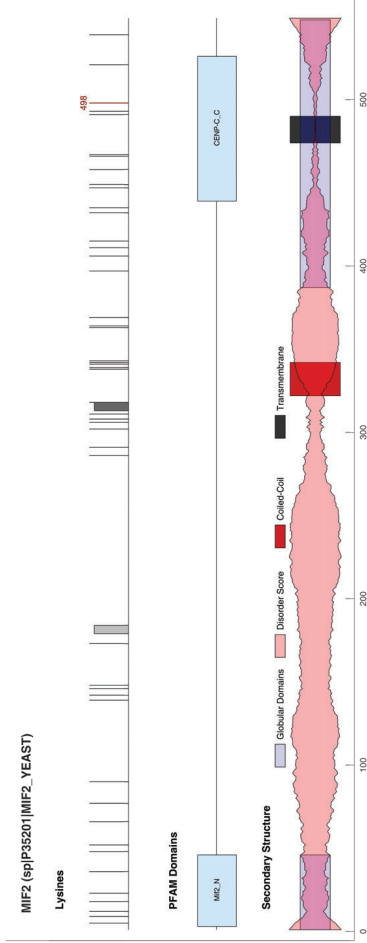
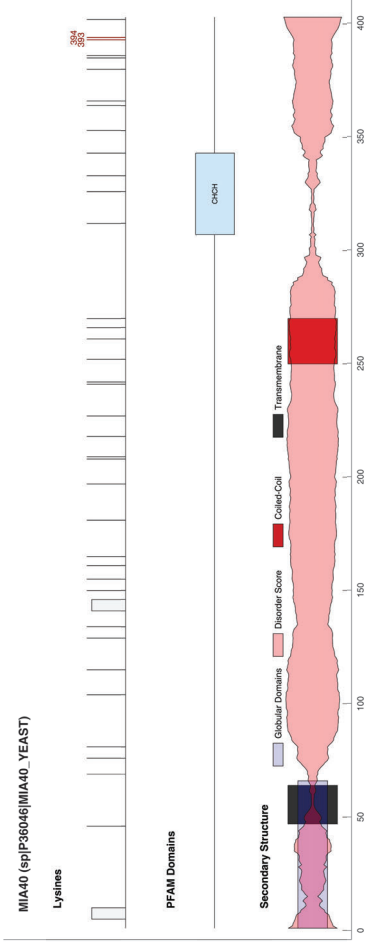
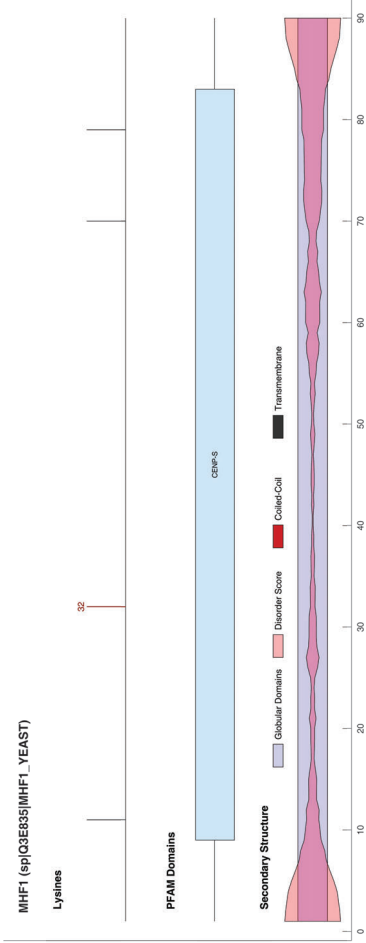
Secondary Structure

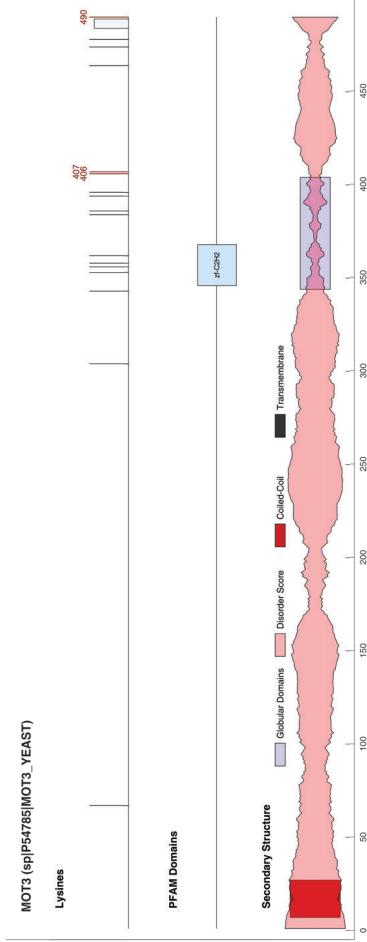
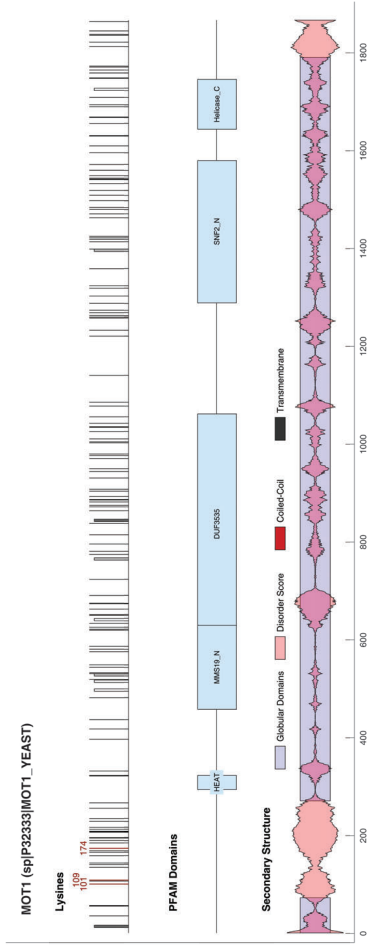
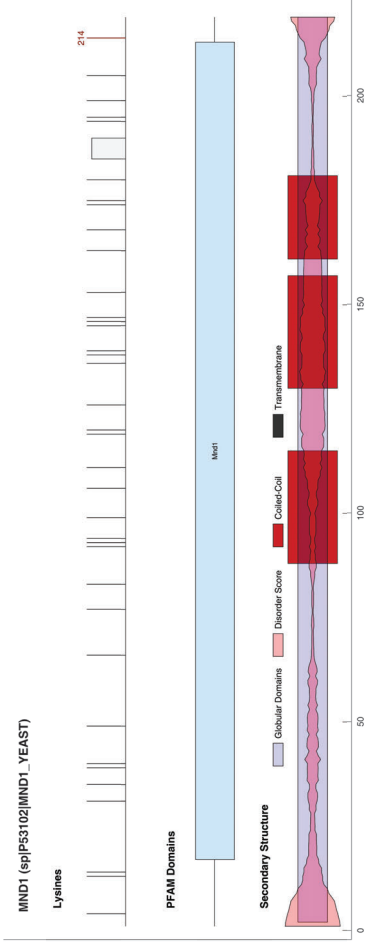
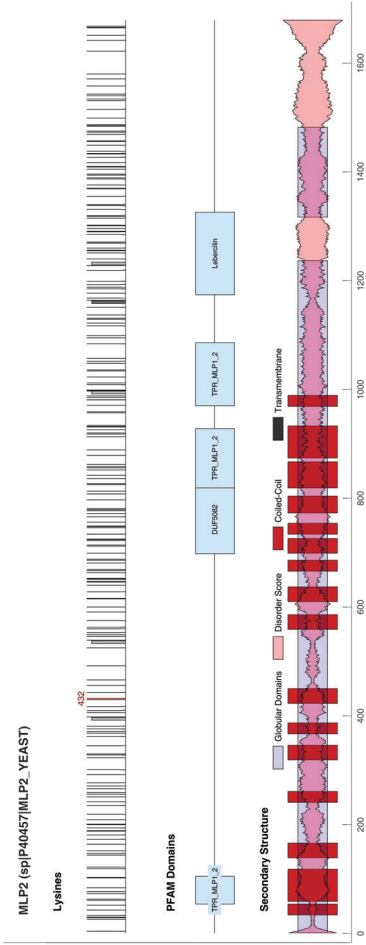












MPG1 (sp|P41940|MPG1_YEAST)

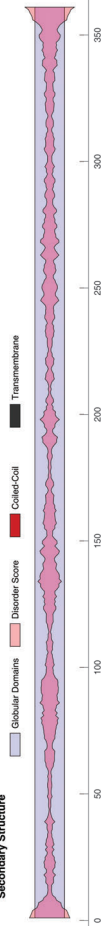
Lysines



PFAM Domains

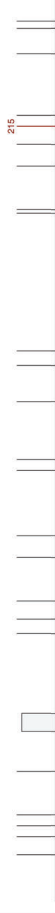


Secondary Structure

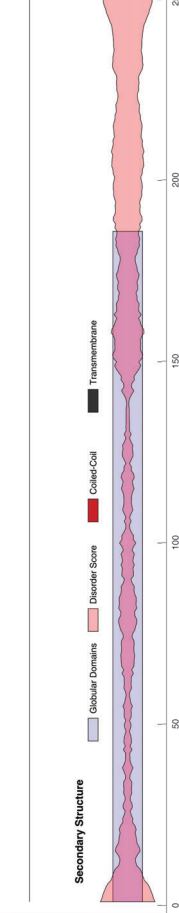


MPM1 (sp|P40364|MPM1_YEAST)

Lysines



PFAM Domains



MRC1 (sp|P25588|MRC1_YEAST)

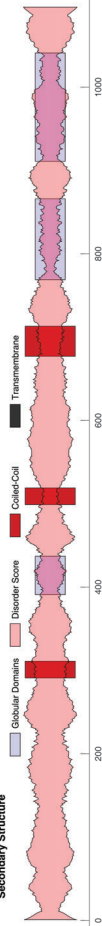
Lysines



PFAM Domains



Secondary Structure

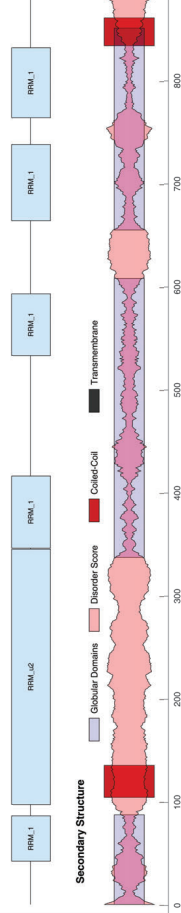


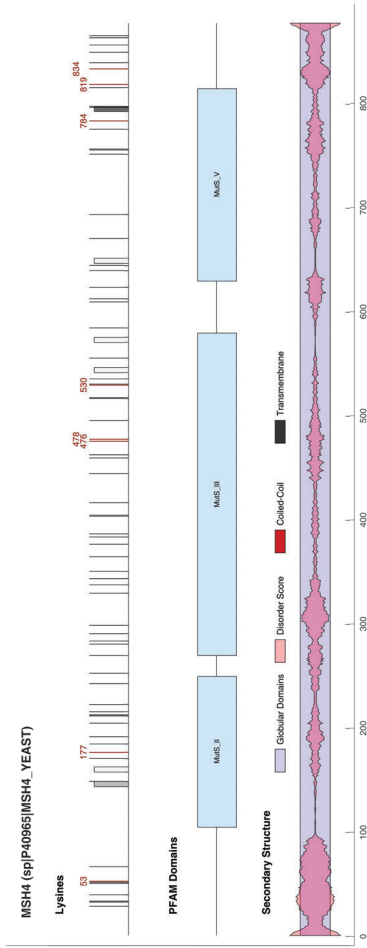
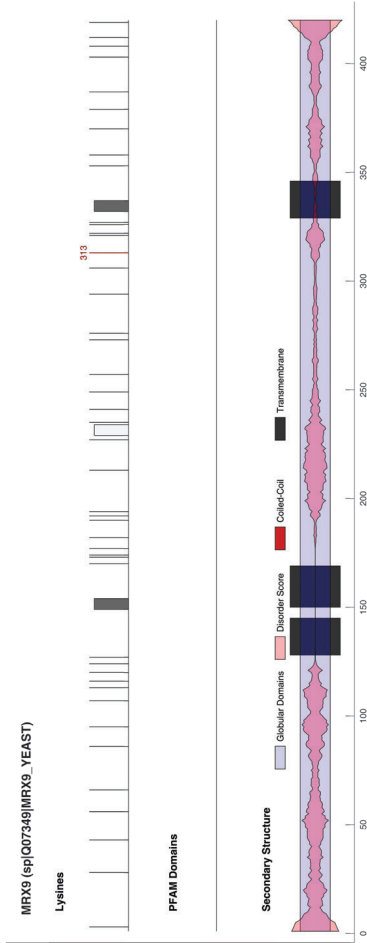
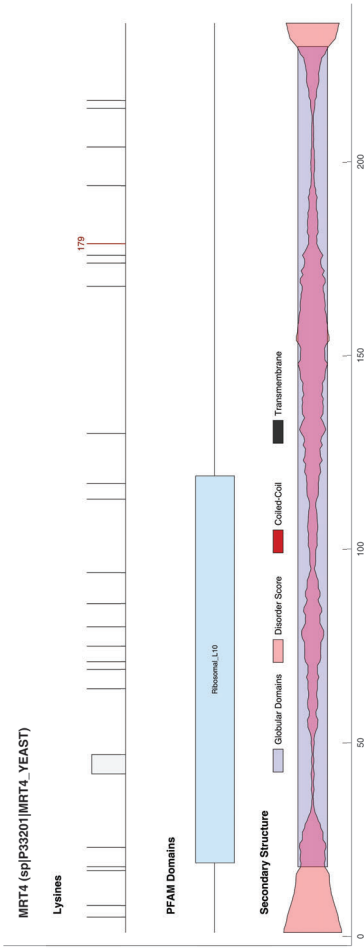
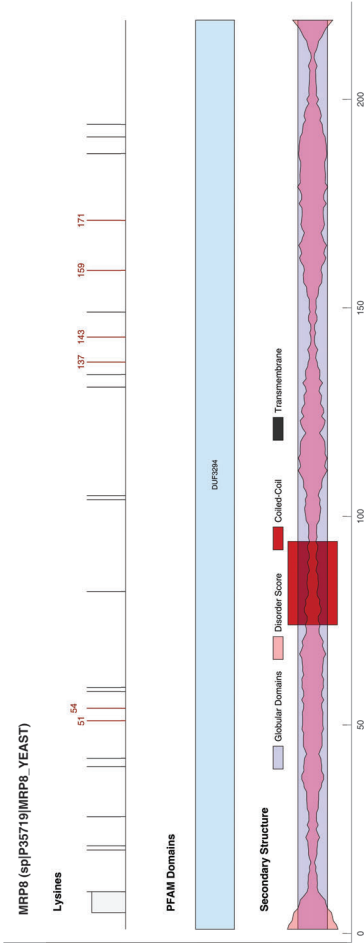
MRD1 (sp|Q06106|MRD1_YEAST)

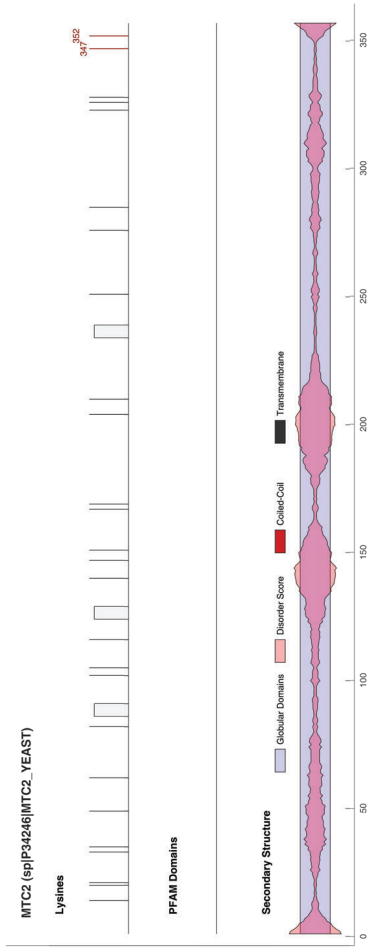
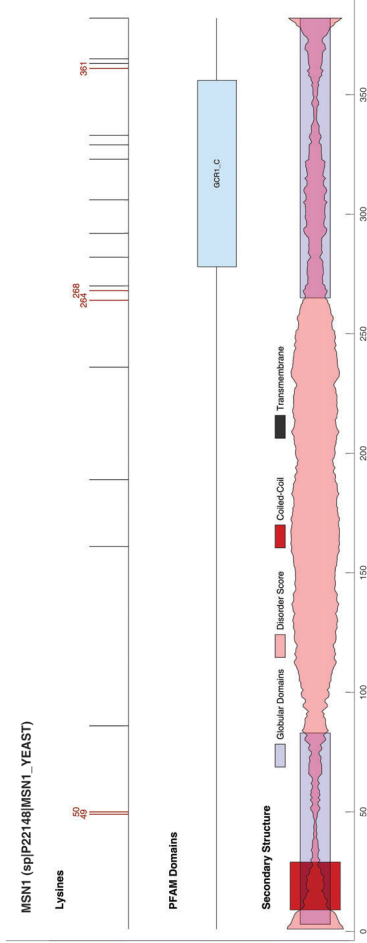
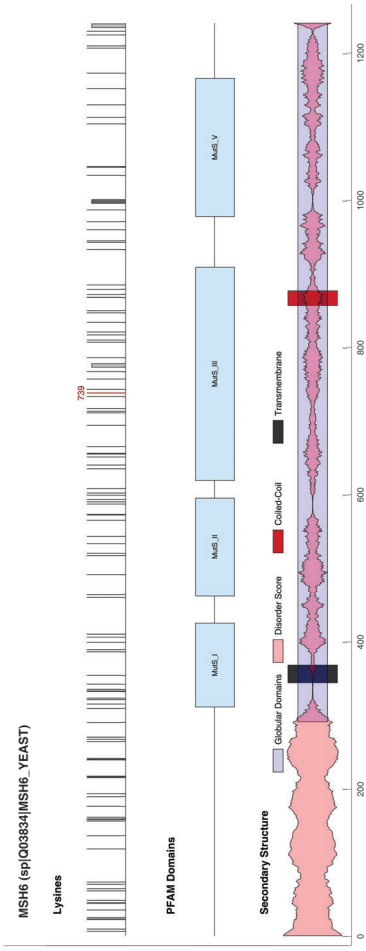
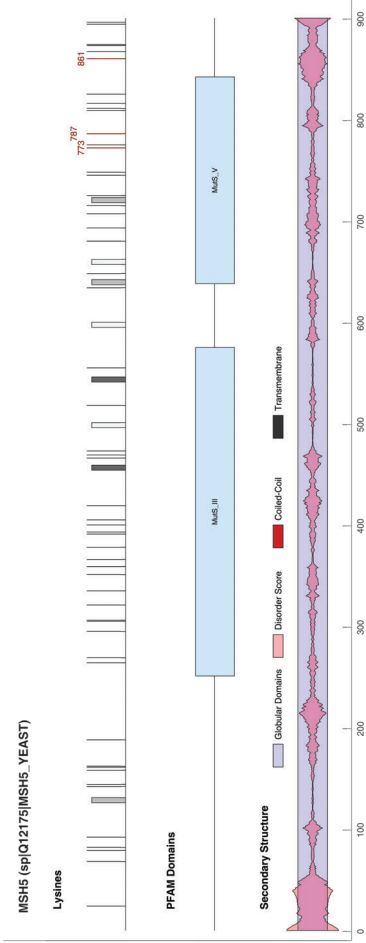
Lysines



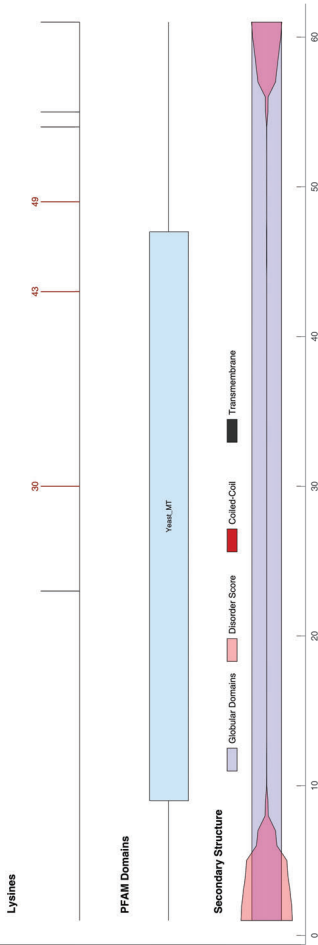
PFAM Domains



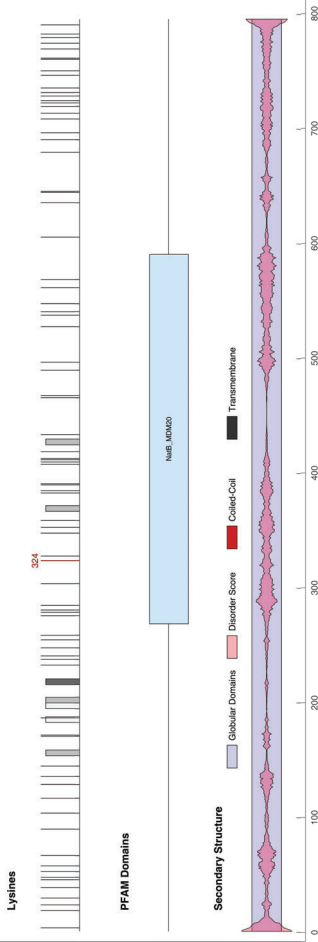




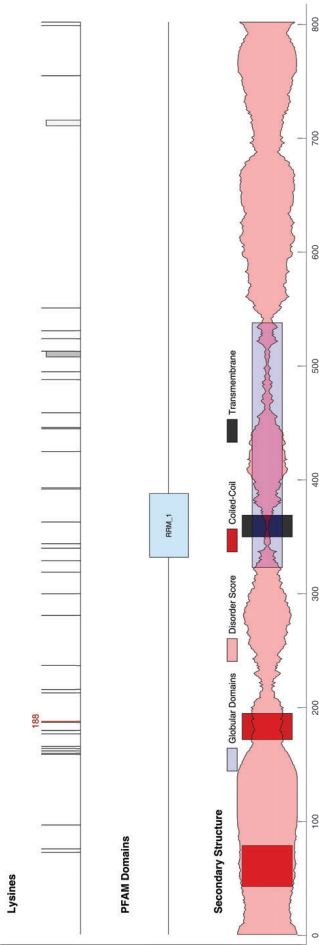
MTCU2 (sp|POCX81|MTCU2_YEAST)



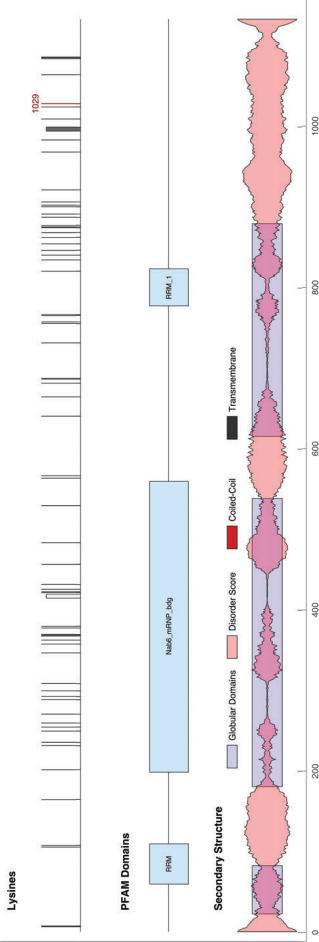
NAA25 (sp|Q12387|NAA25_YEAST)



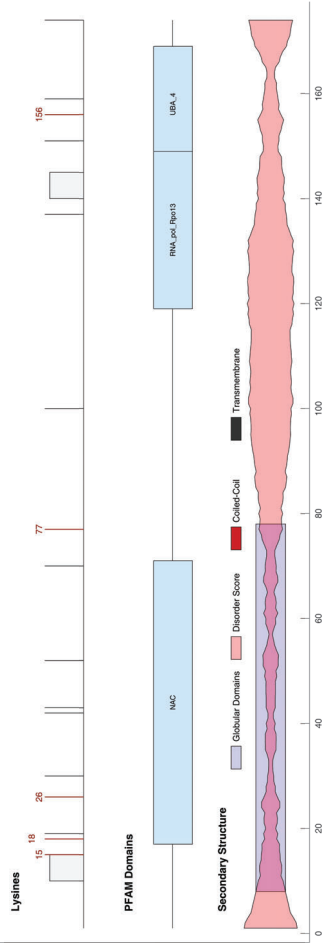
NAB3 (sp|P38996|NAB3_YEAST)



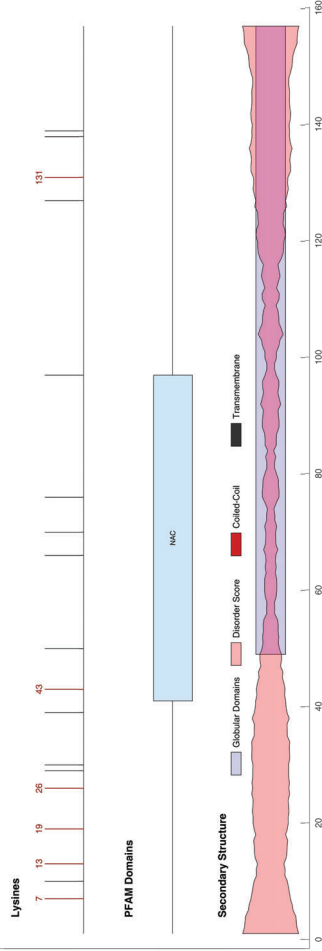
NAB6 (sp|Q03735|NAB6_YEAST)



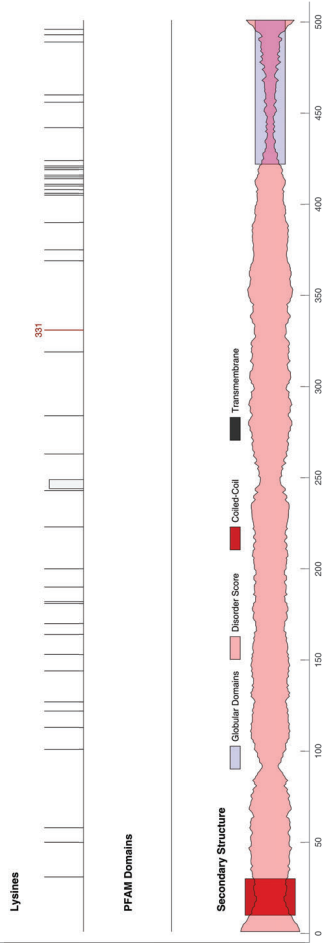
NACA (sp|P38879|NACA_YEAST)



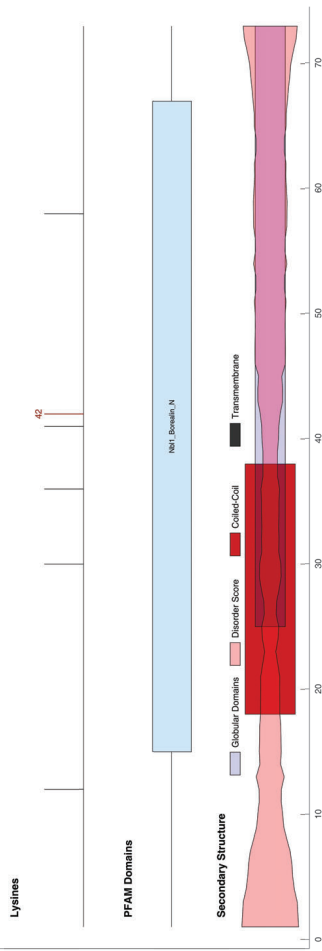
NACB1 (sp|Q02642|NACB1_YEAST)

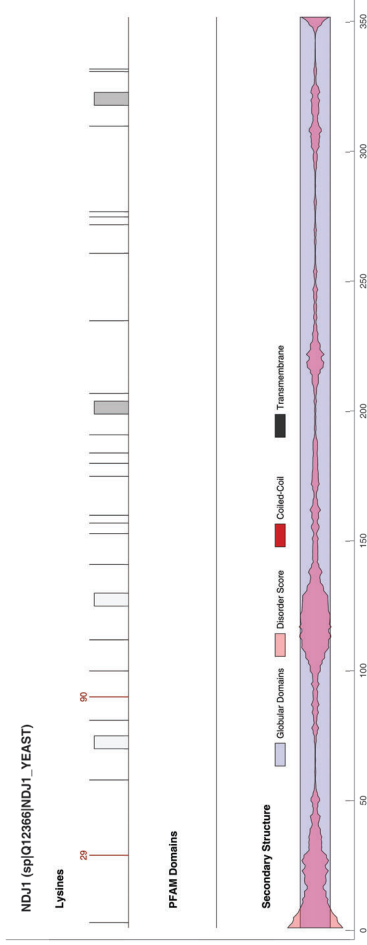
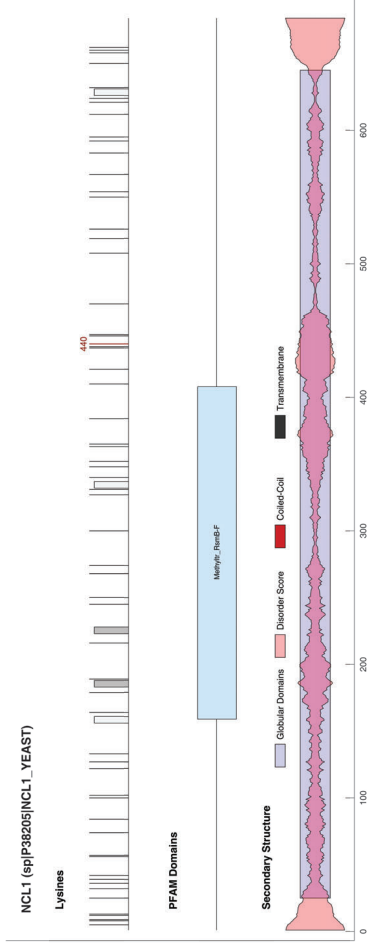
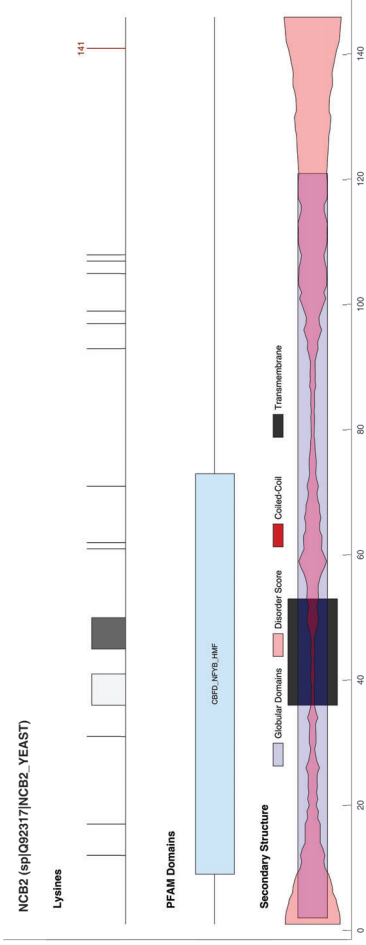
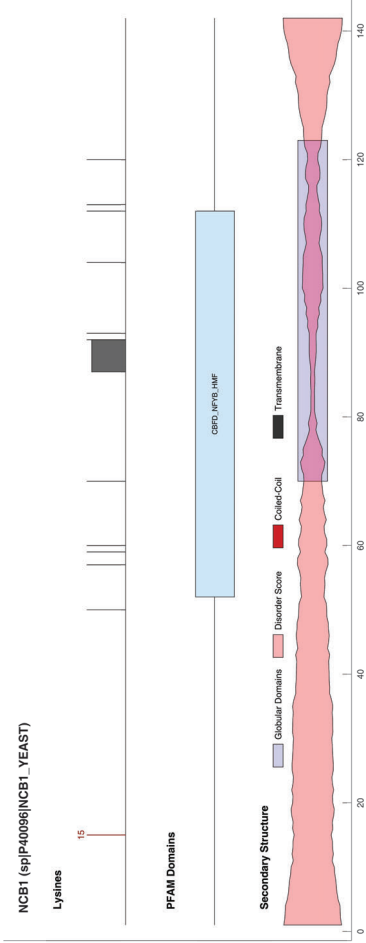


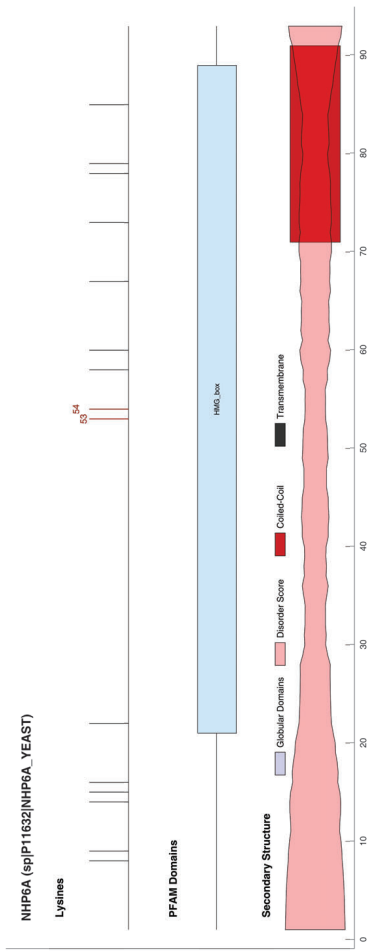
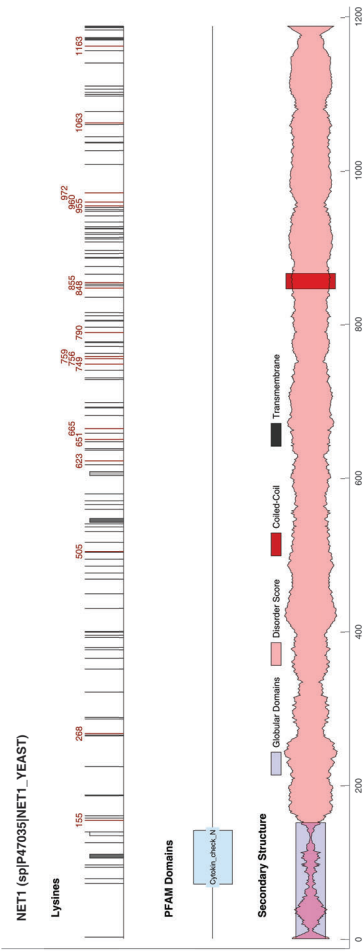
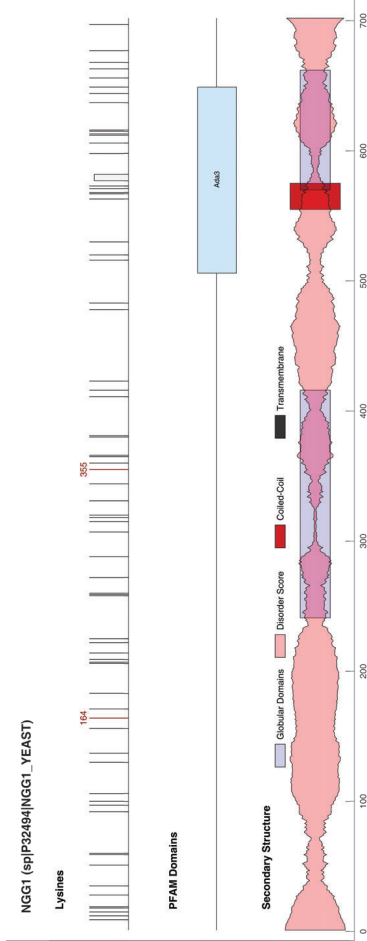
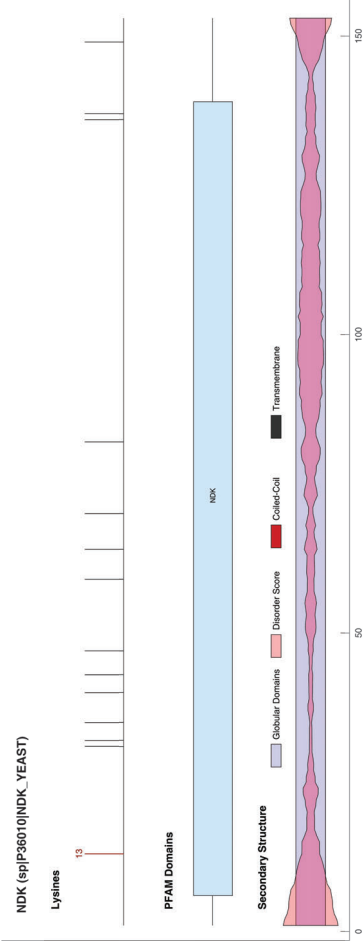
NBA1 (sp|Q08229|NBA1_YEAST)

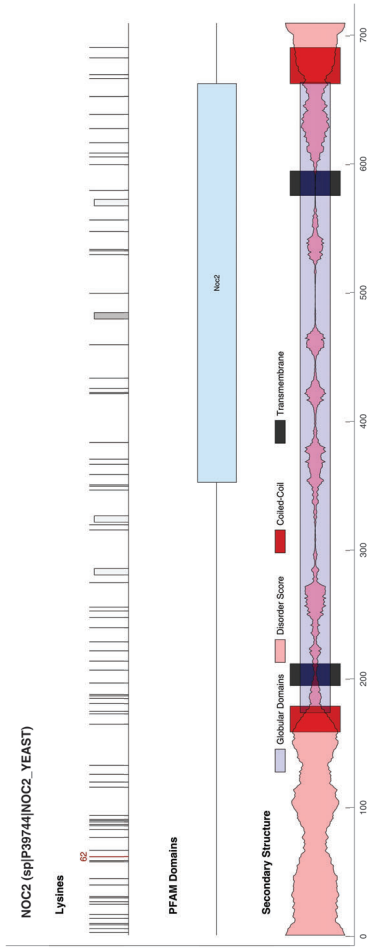
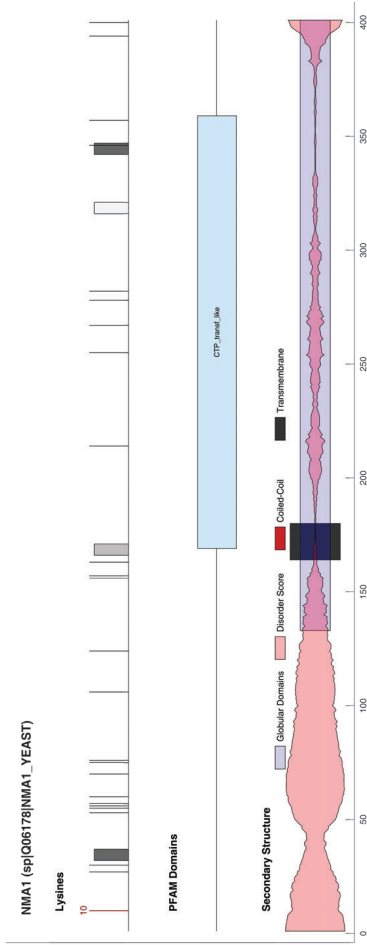
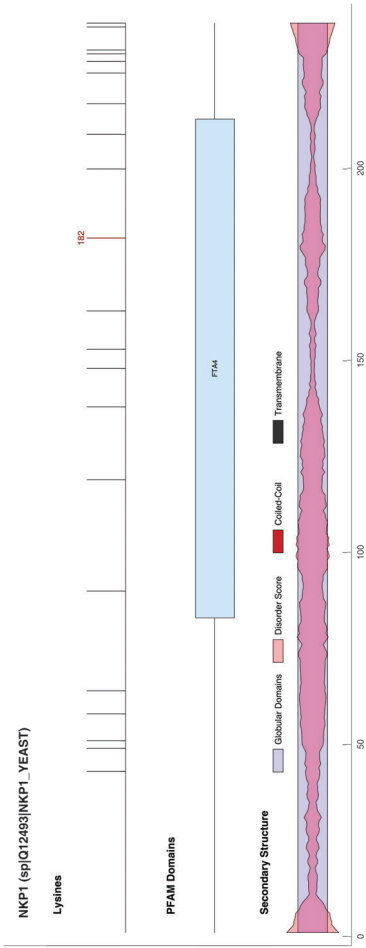
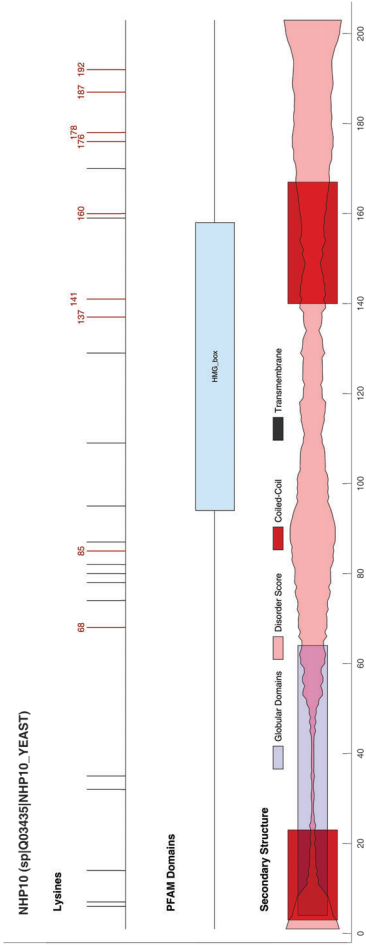


NBL1 (sp|Q3E7Y6|NBL1_YEAST)

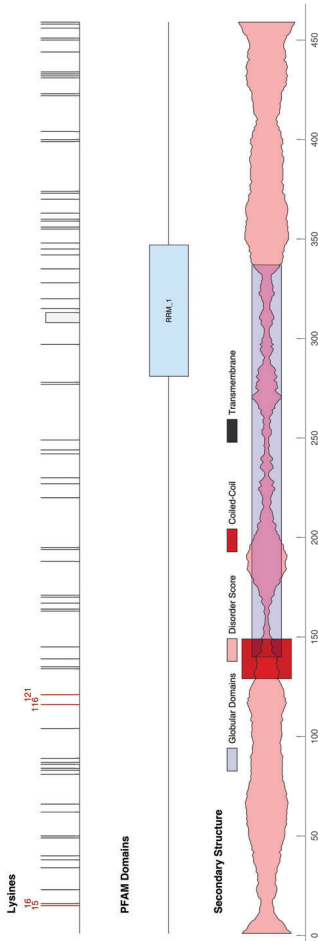




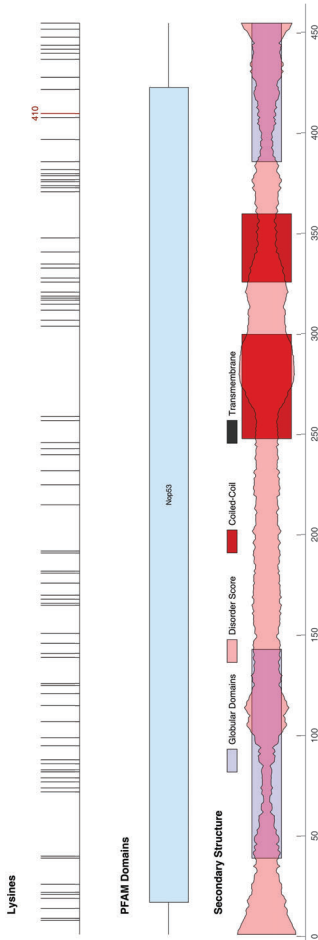




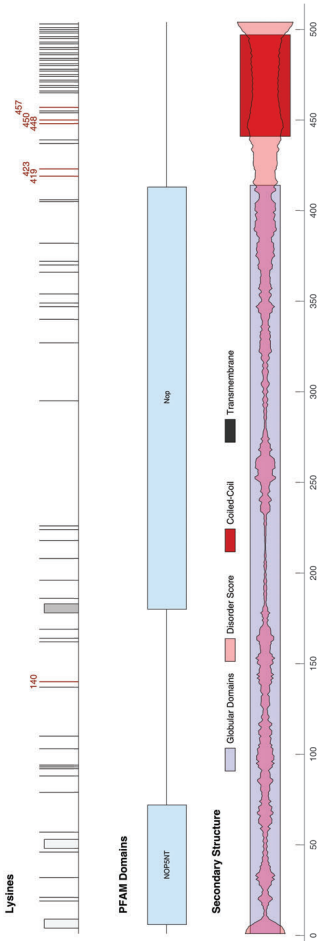
NOP12 (ep|Q08208|NOP12_YEAST)



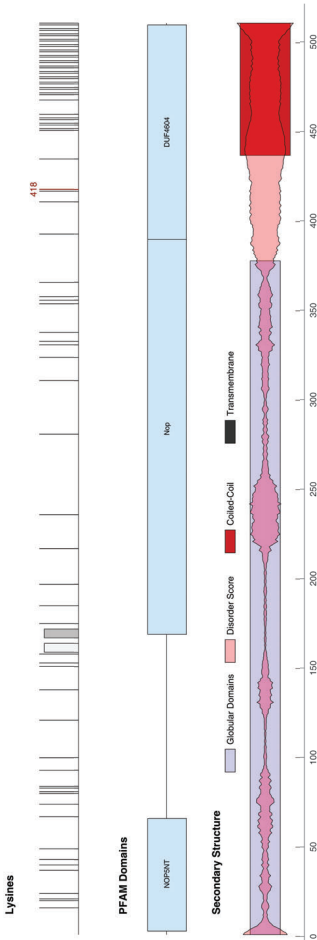
NOP53 (ep|Q12080|NOP53_YEAST)

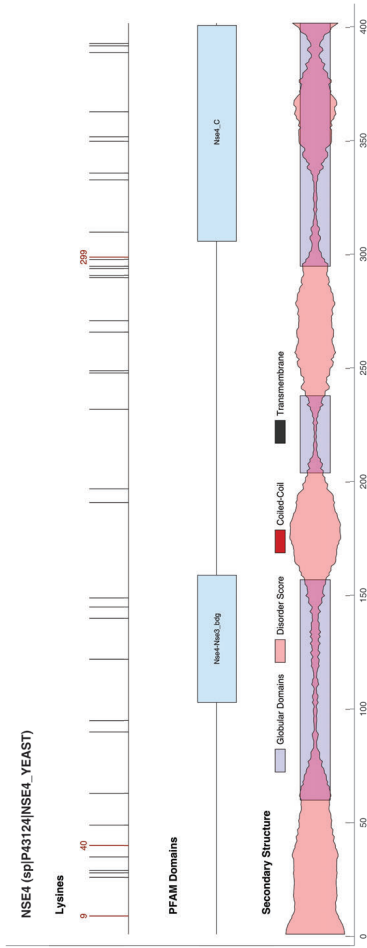
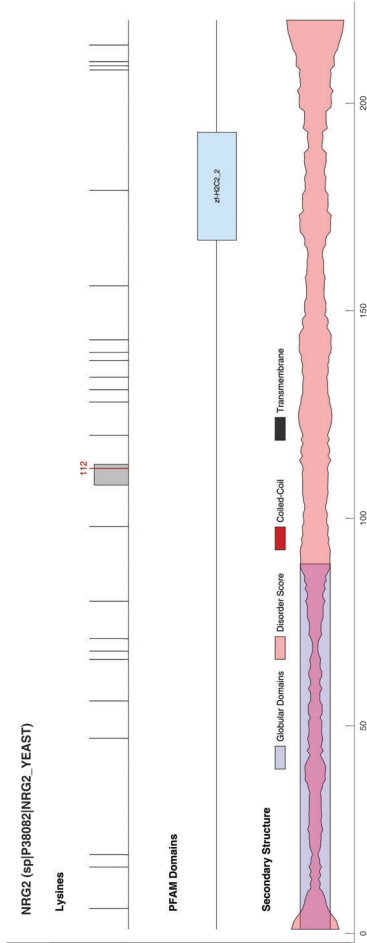
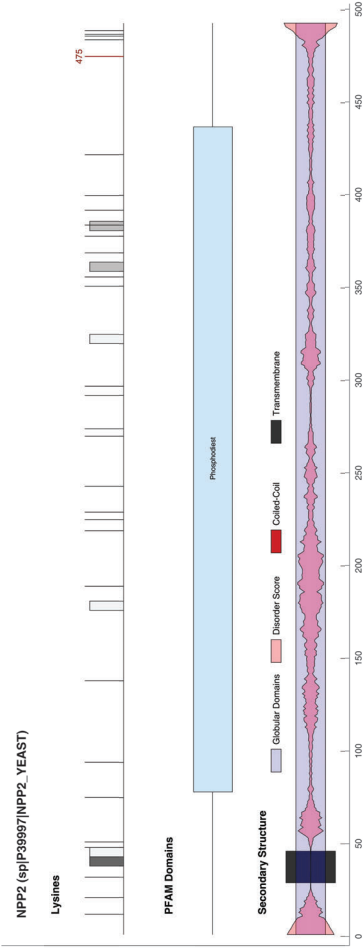
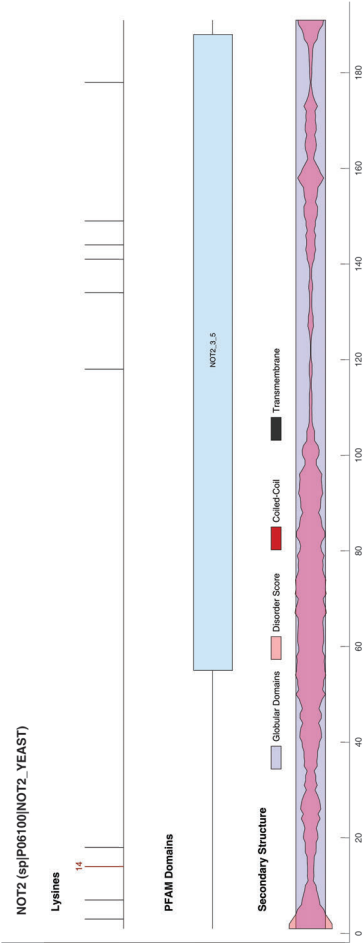


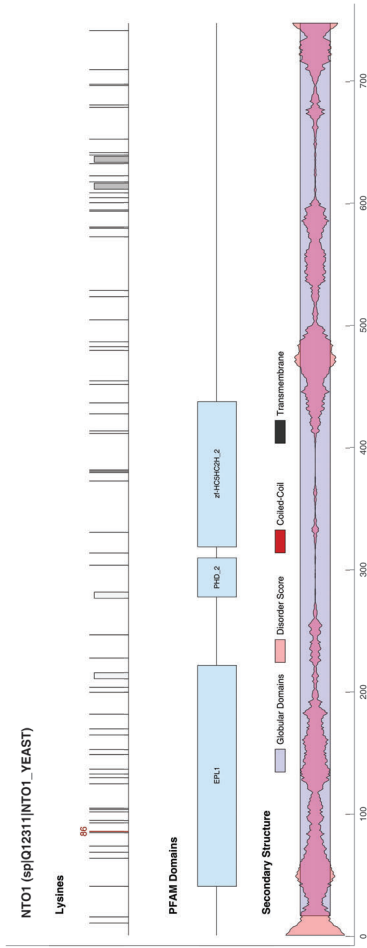
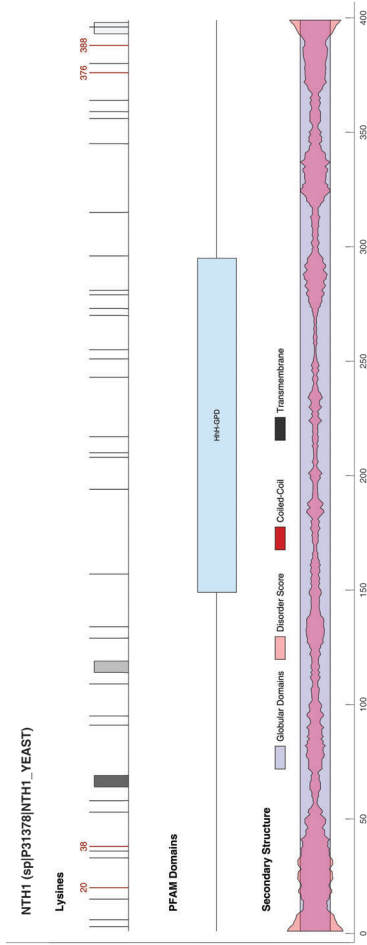
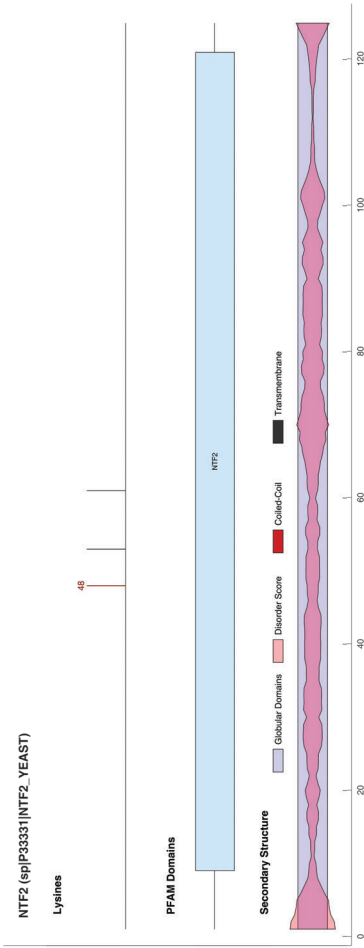
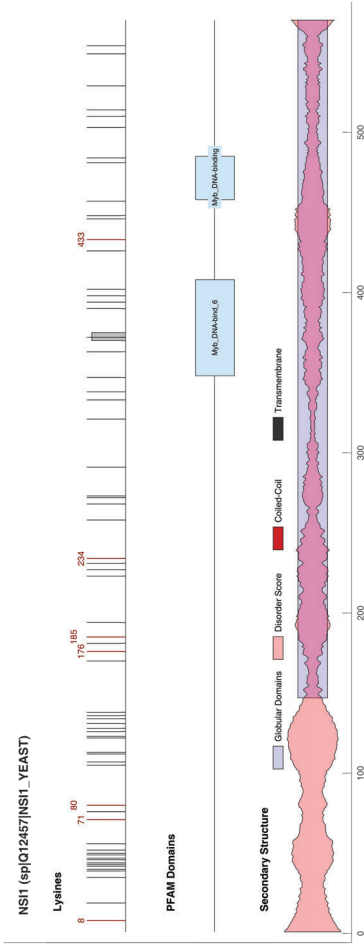
NOP56 (ep|Q12460|NOP56_YEAST)

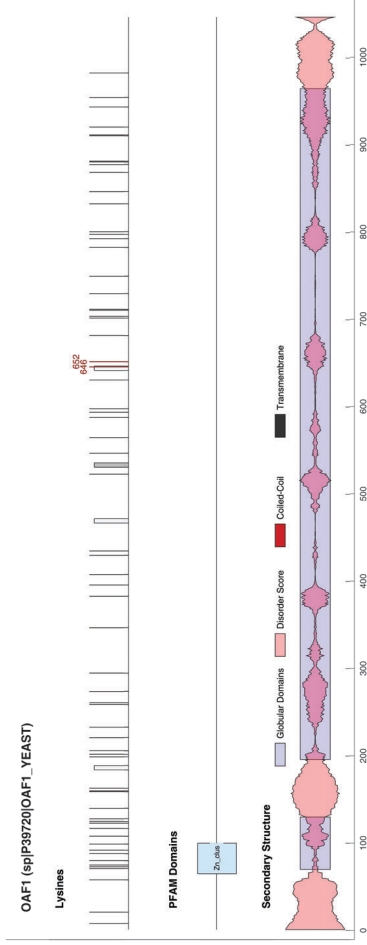
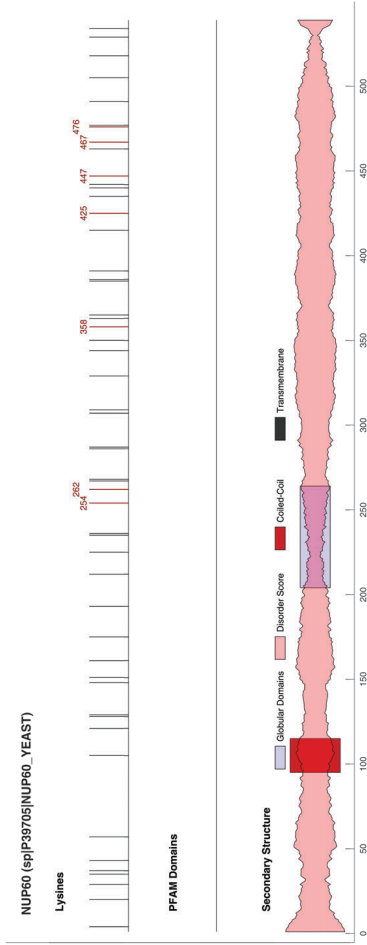
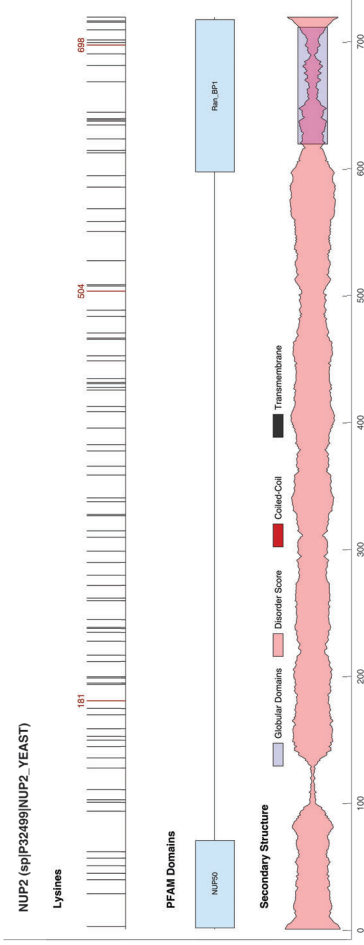
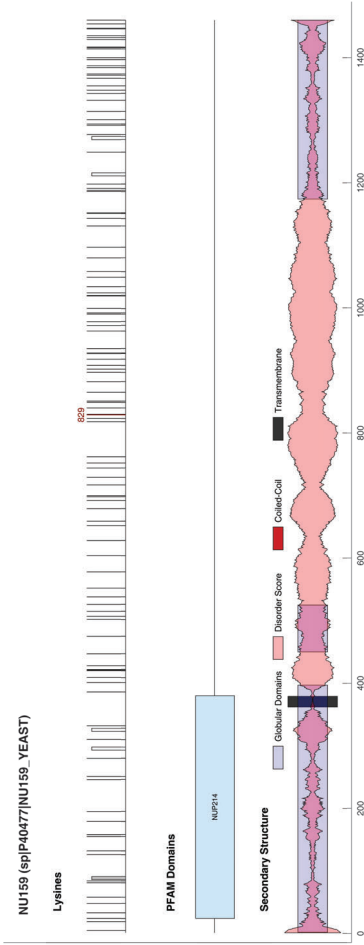


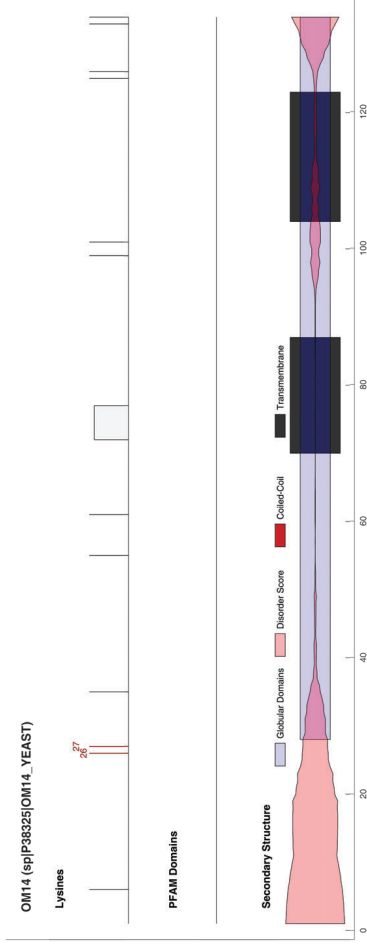
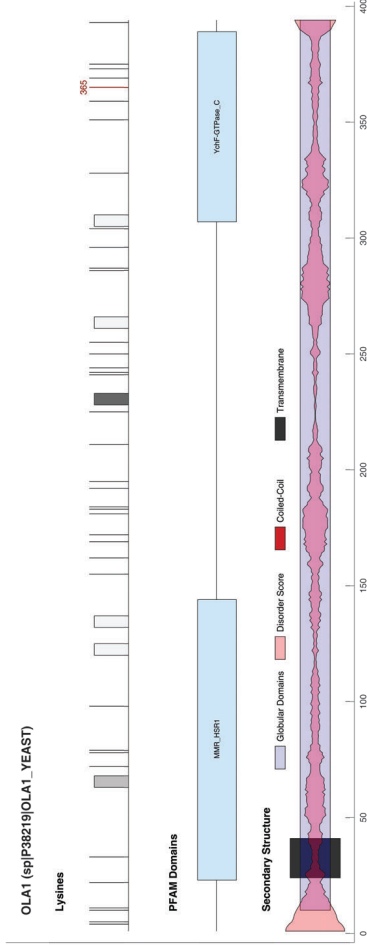
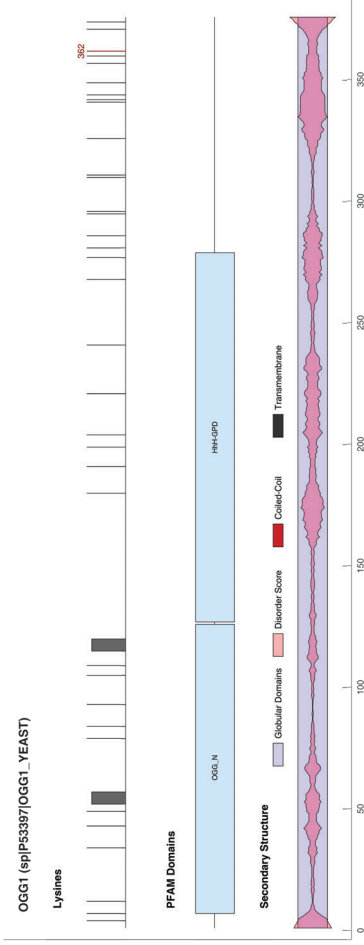
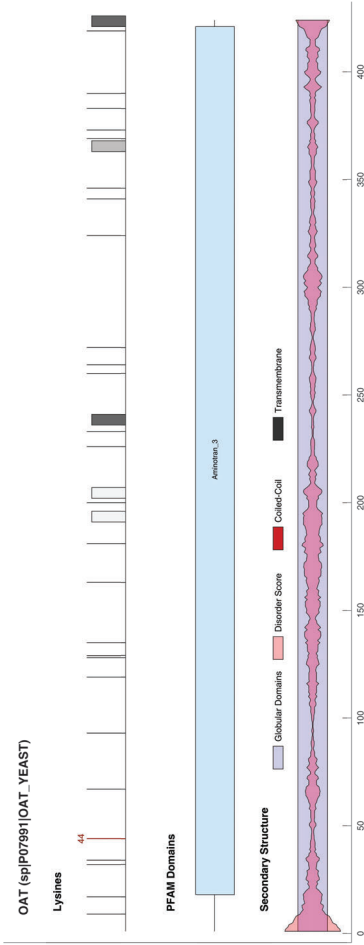
NOP58 (ep|Q12459|NOP58_YEAST)

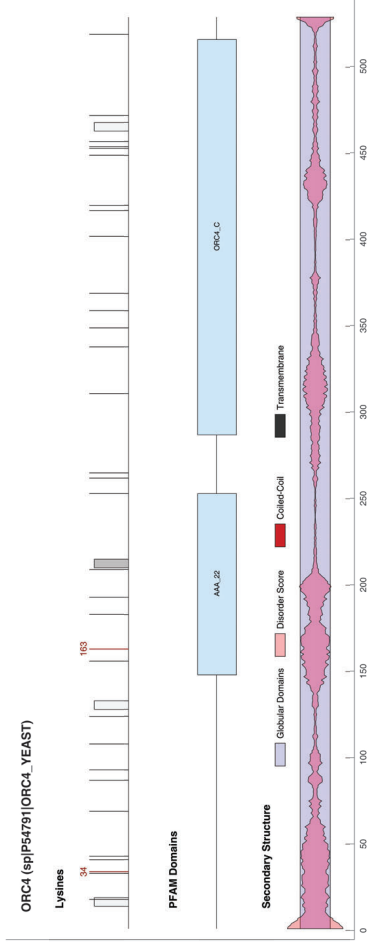
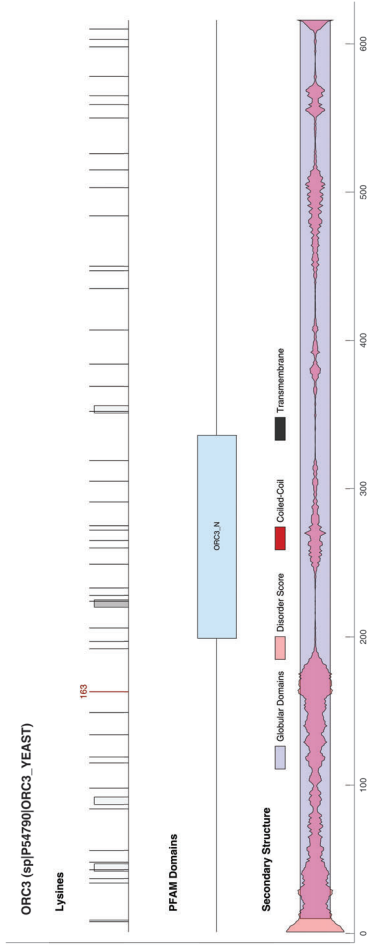
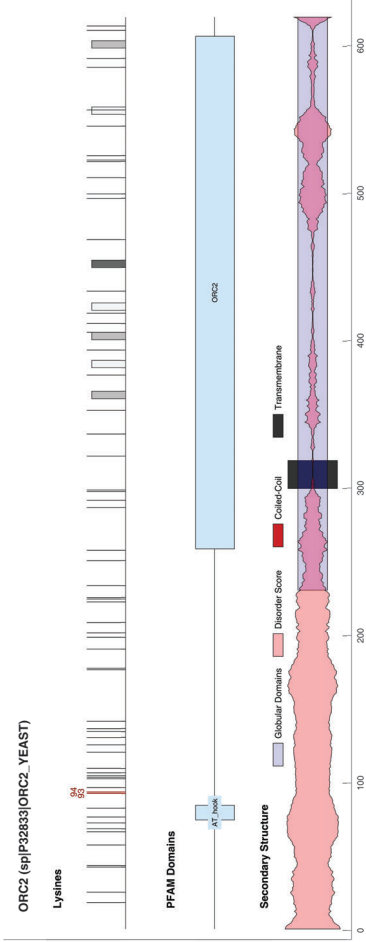
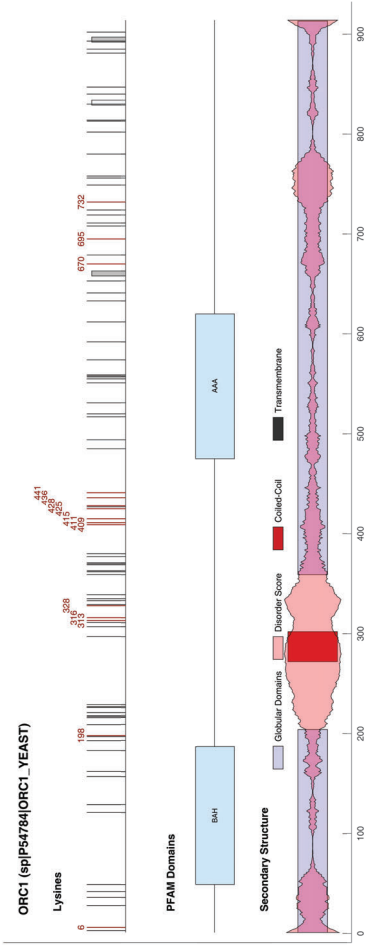


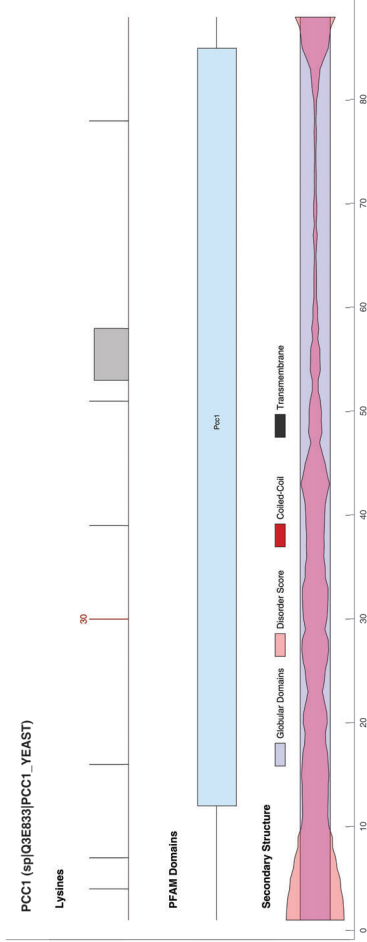
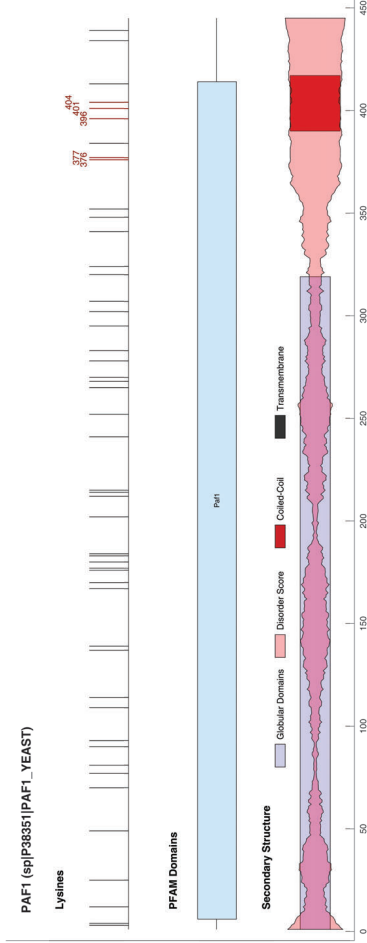
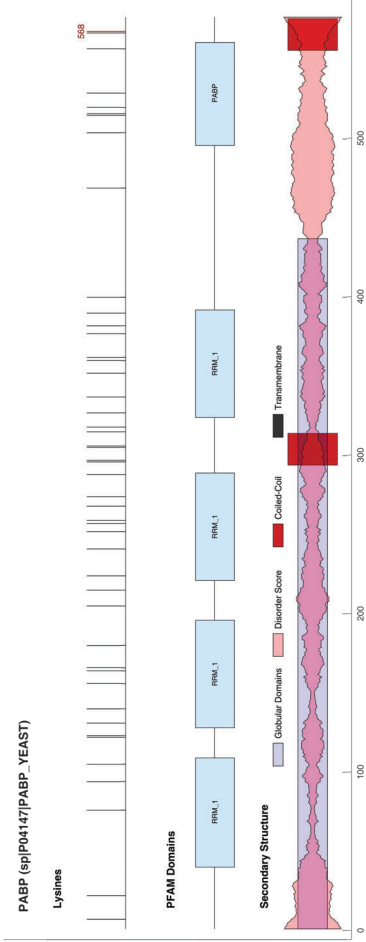
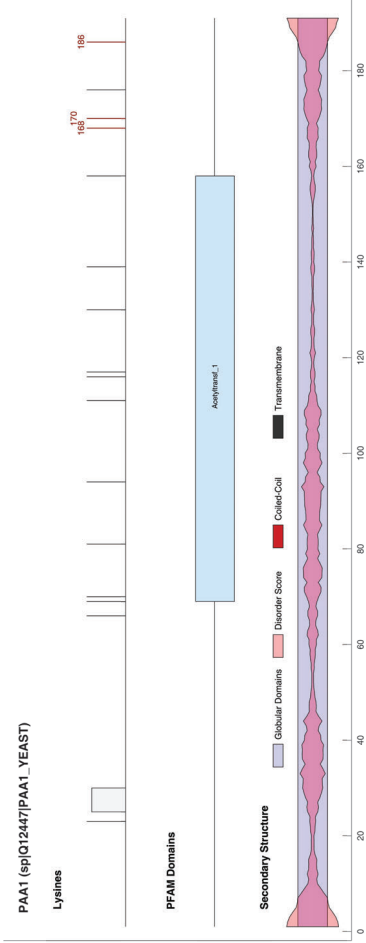


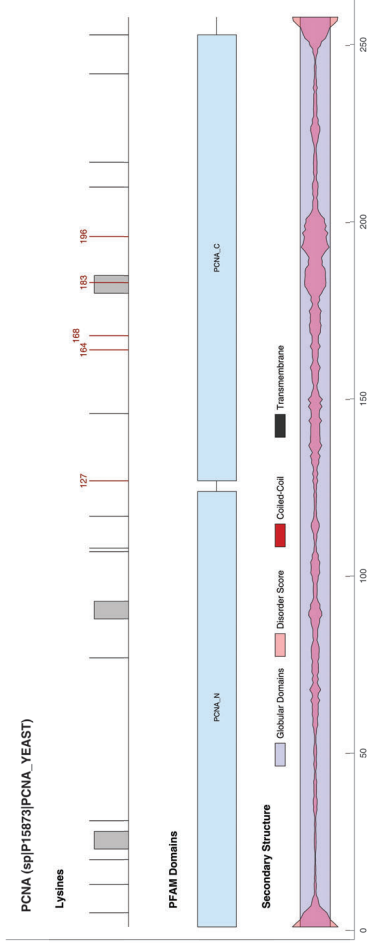
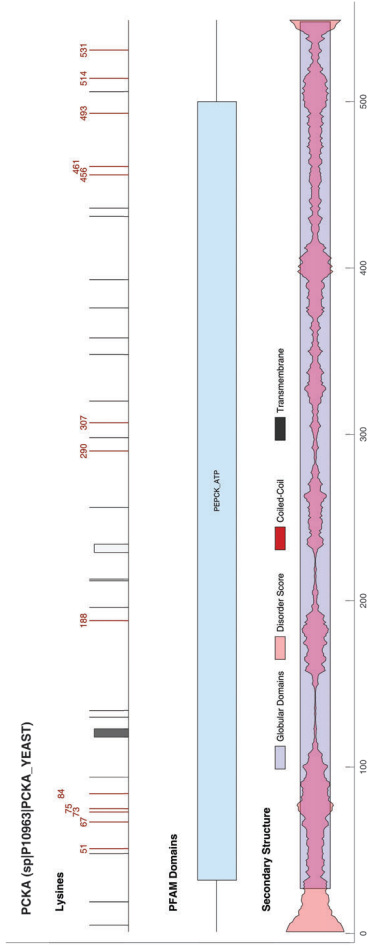
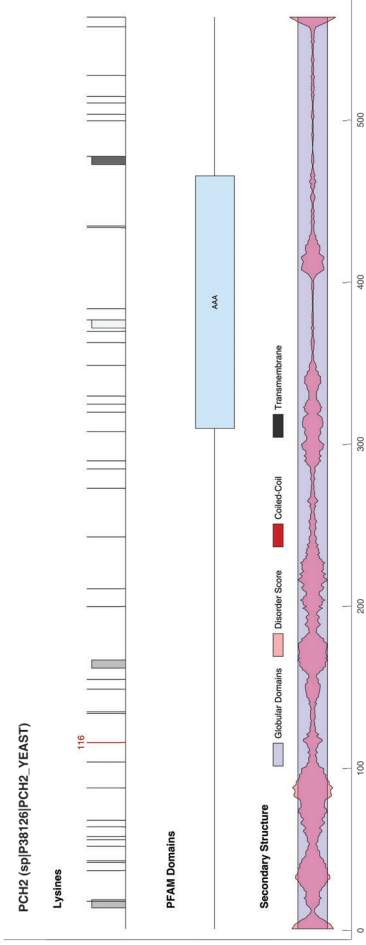
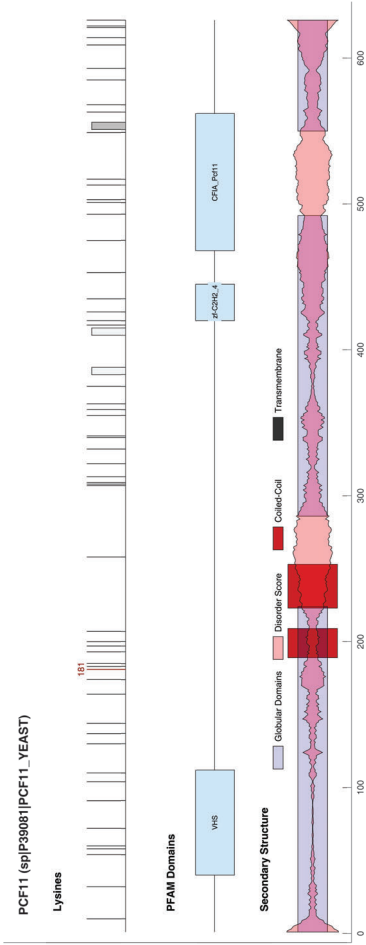


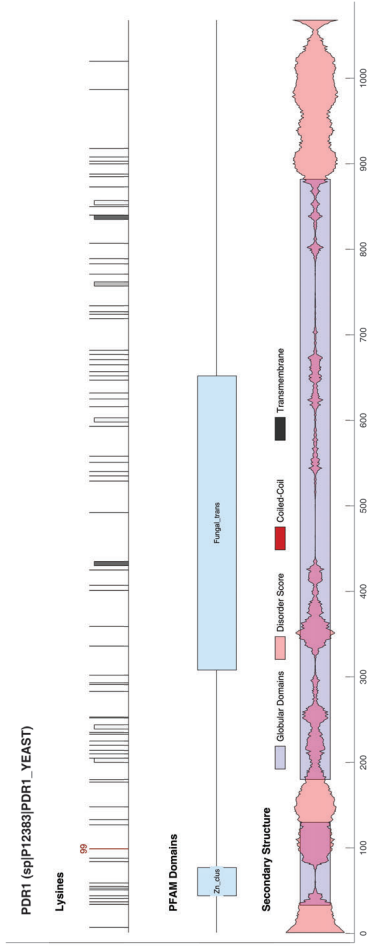
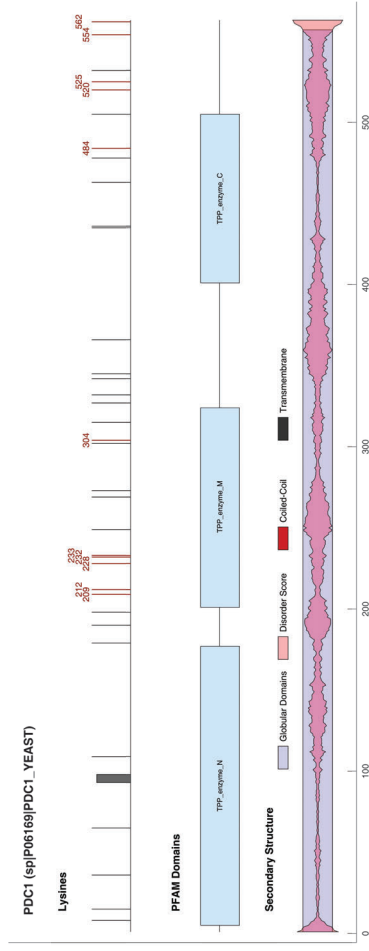


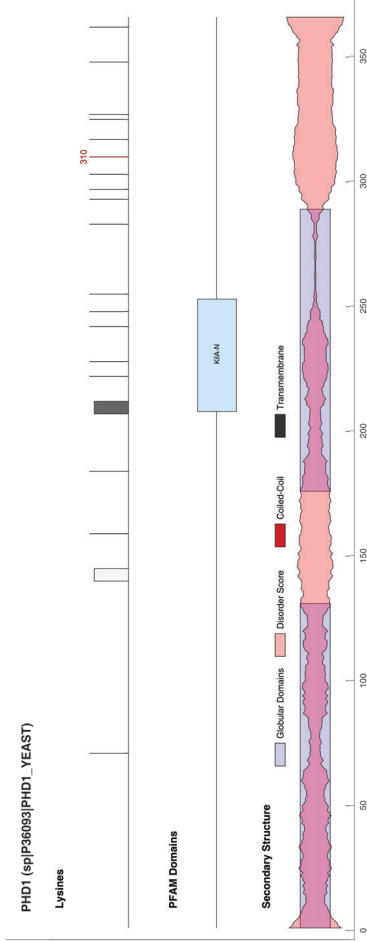
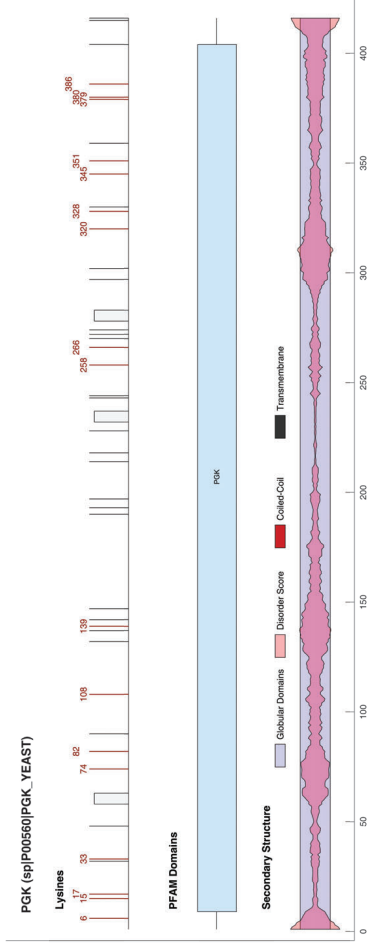
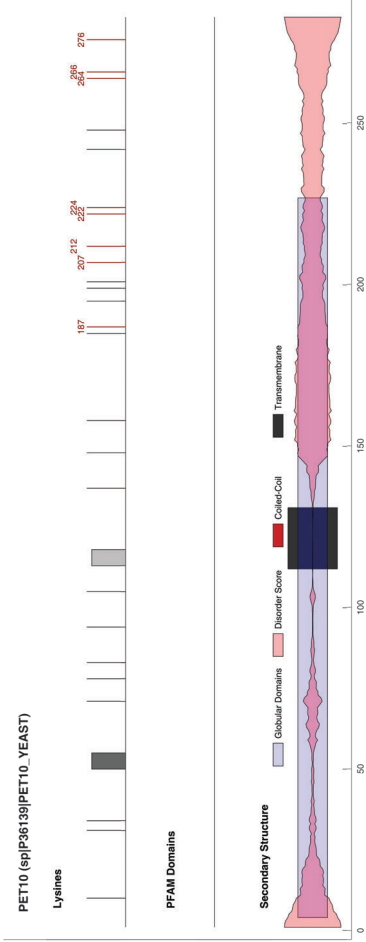
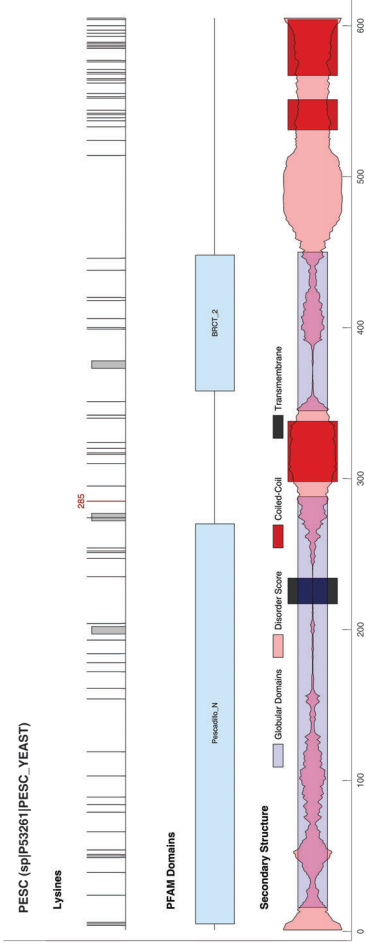












PHO23 (sp|P50947|PHO23_YEAST)

Lysines



PFAM Domains



Secondary Structure

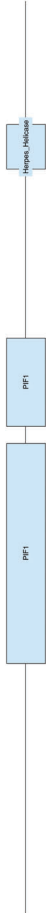


PIF1 (sp|P07271|PIF1_YEAST)

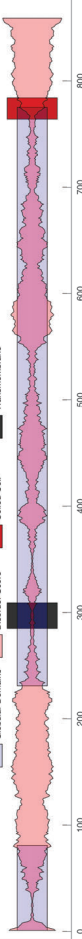
Lysines



PFAM Domains



Secondary Structure



PHSG (sp|P06738|PHSG_YEAST)

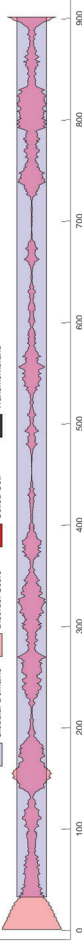
Lysines



PFAM Domains



Secondary Structure



PIG1 (sp|Q06216|PIG1_YEAST)

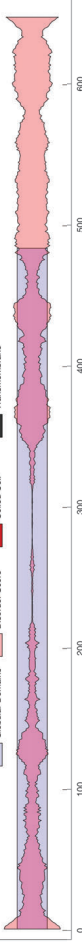
Lysines

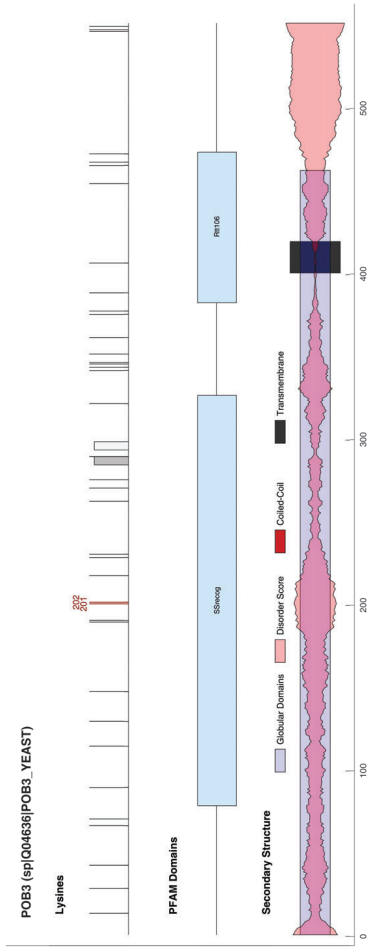
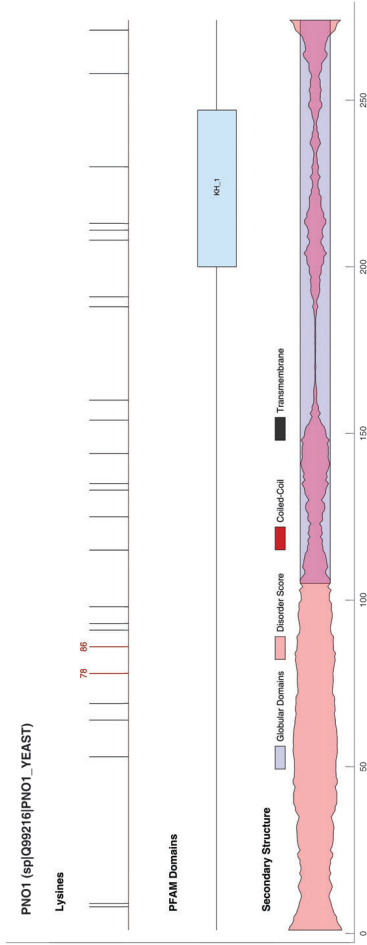
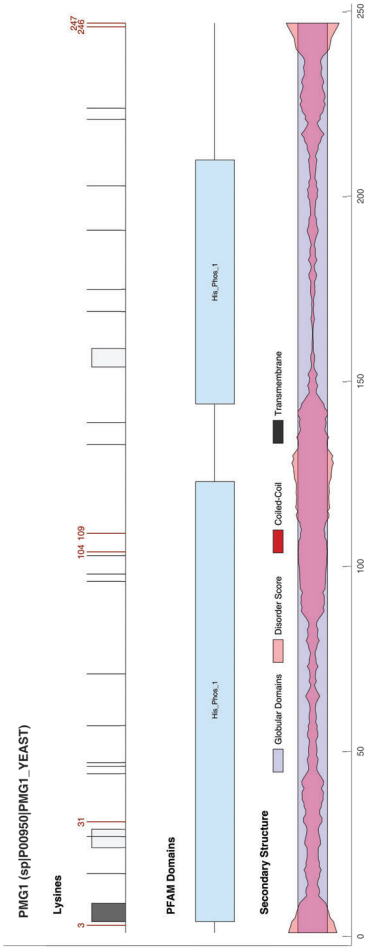
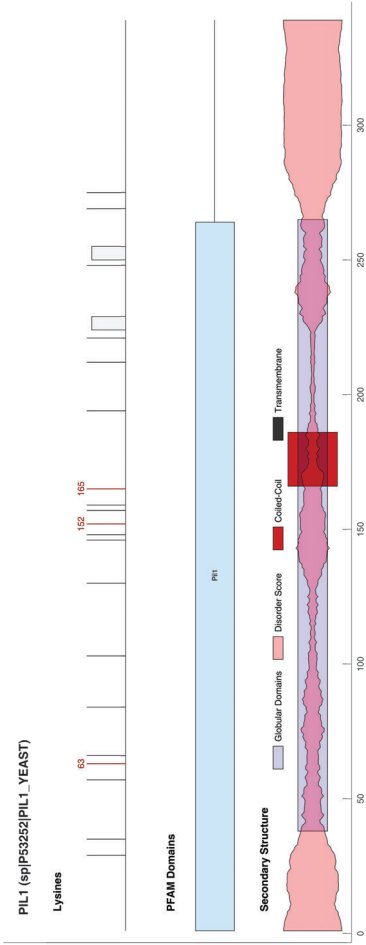


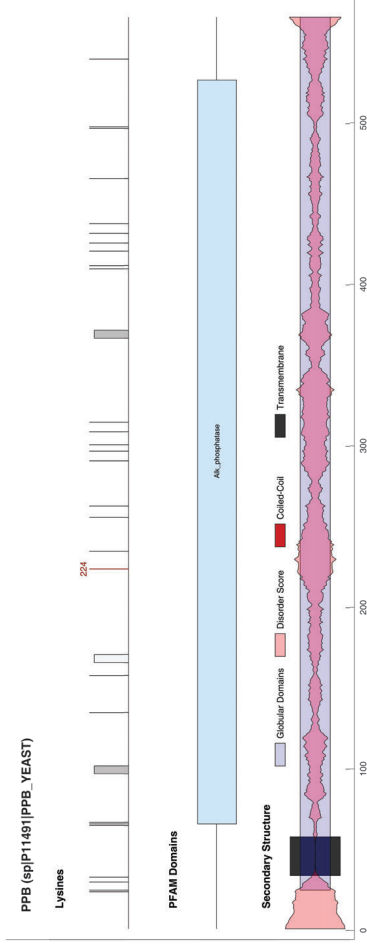
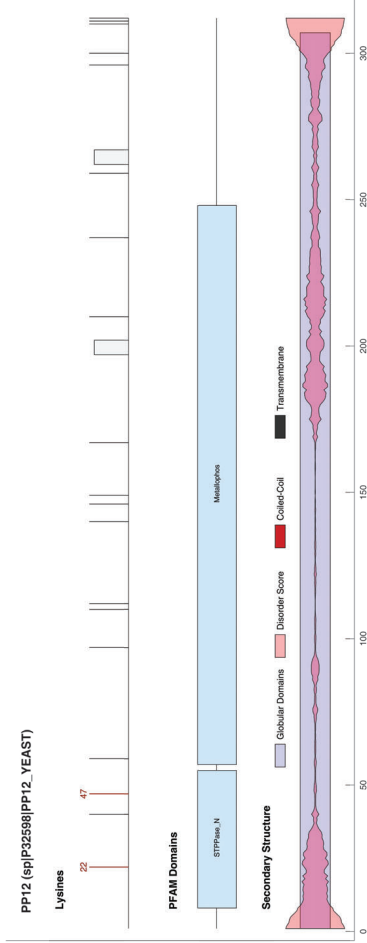
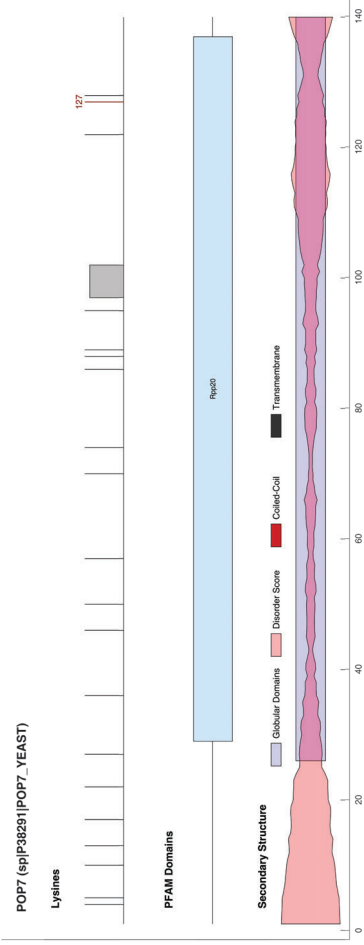
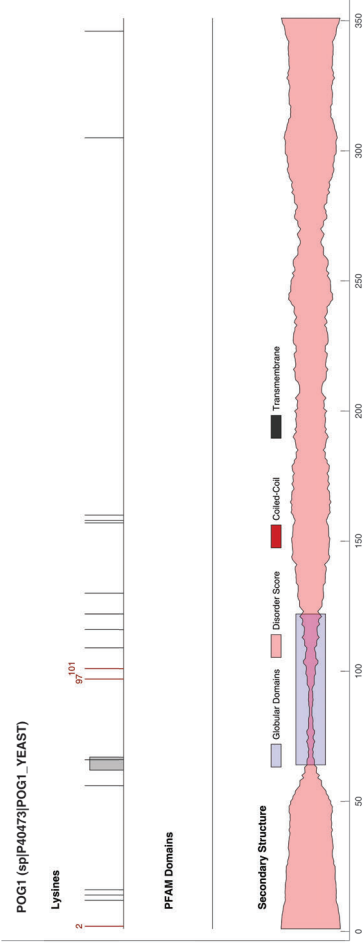
PFAM Domains

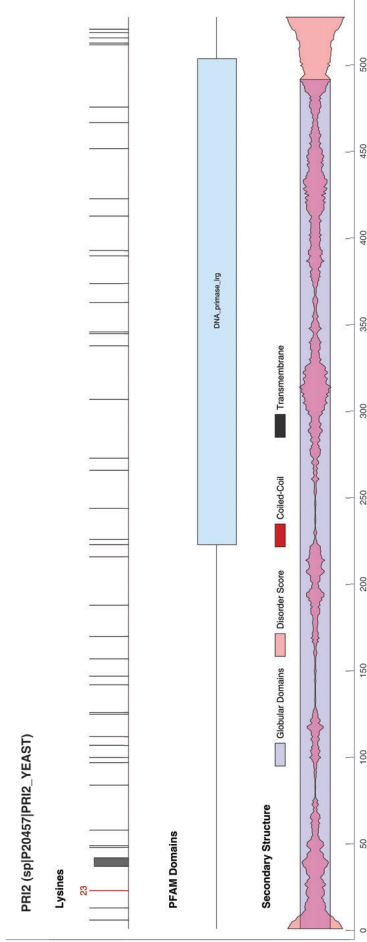
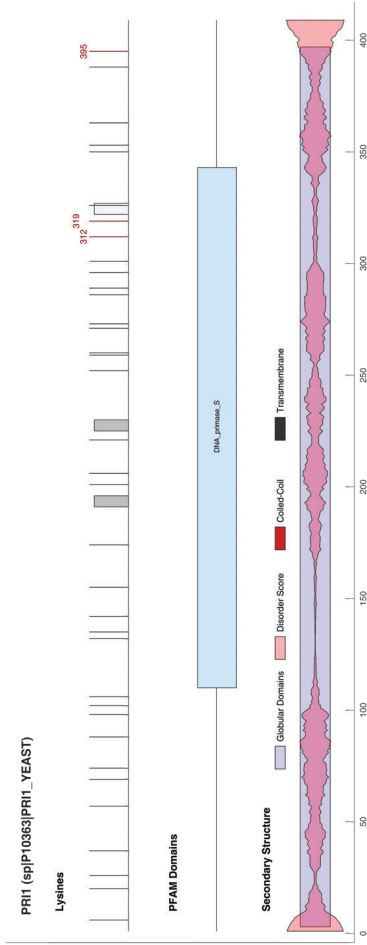
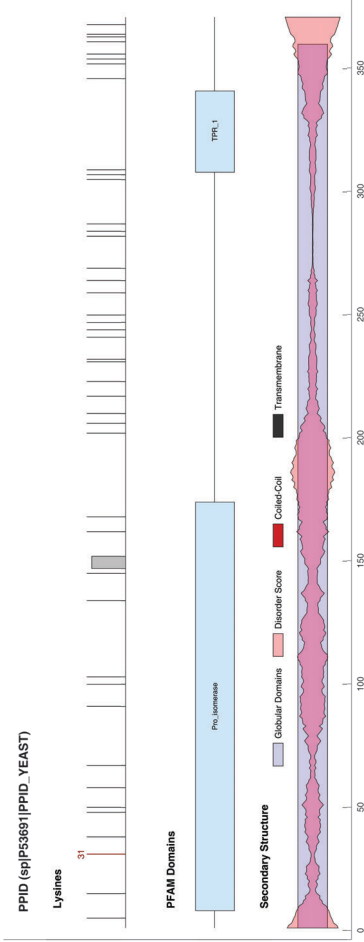
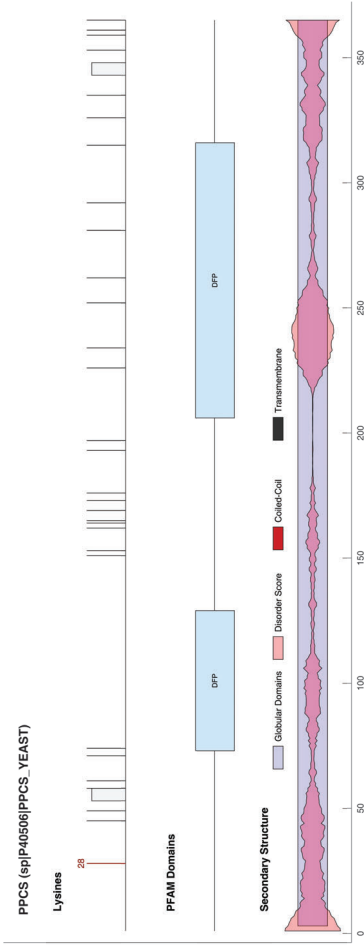


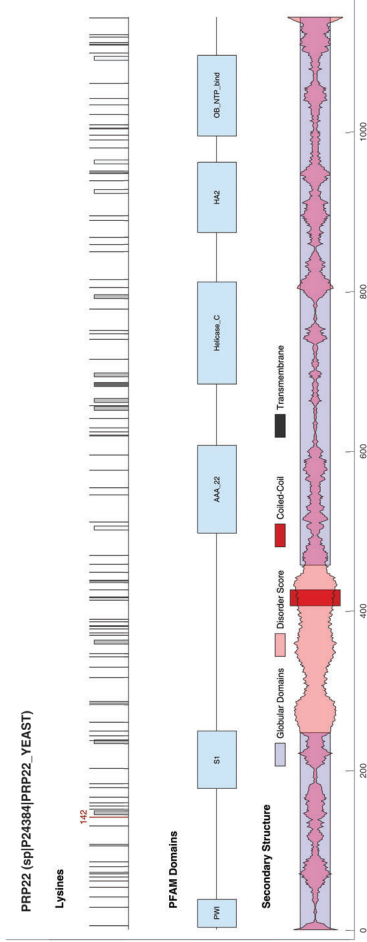
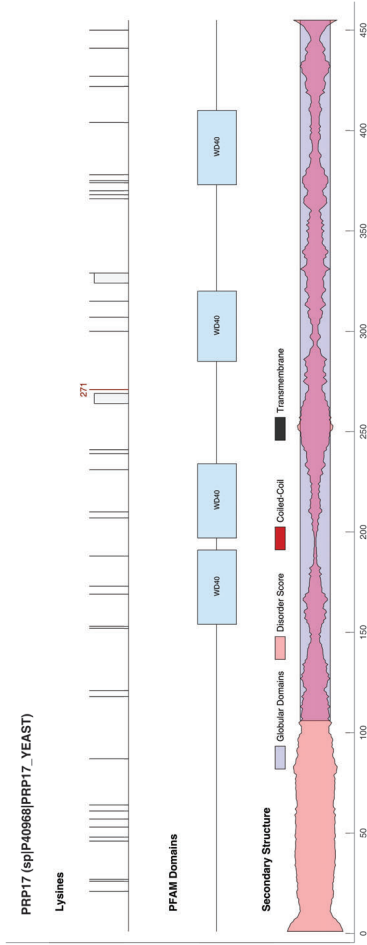
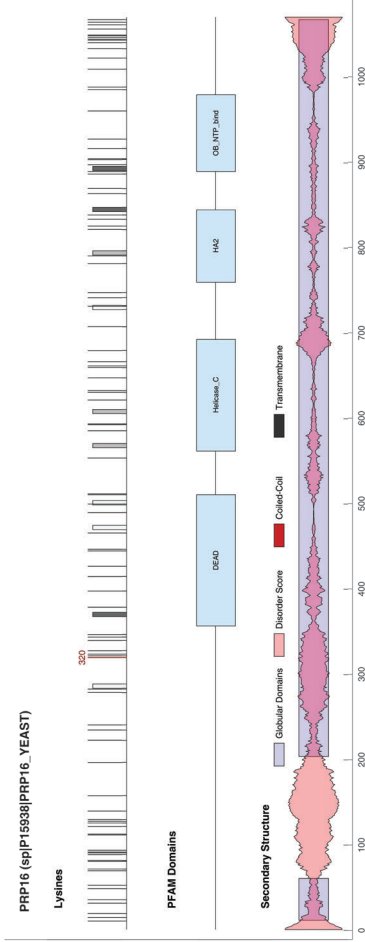
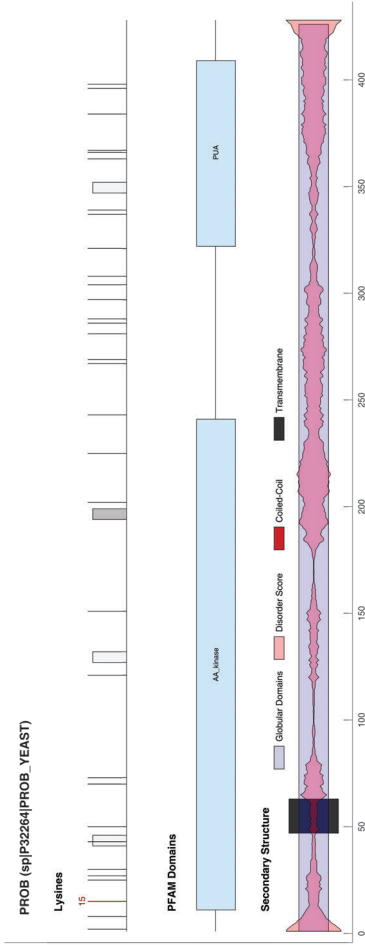
Secondary Structure



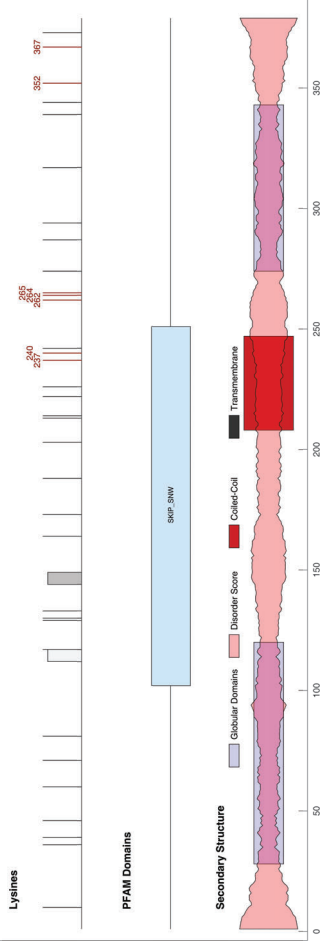




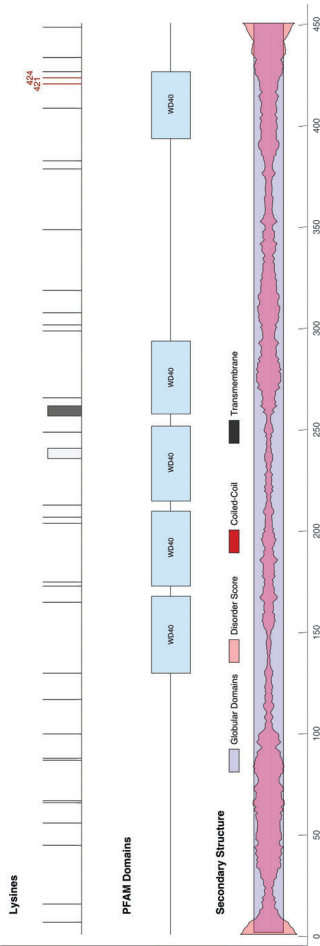




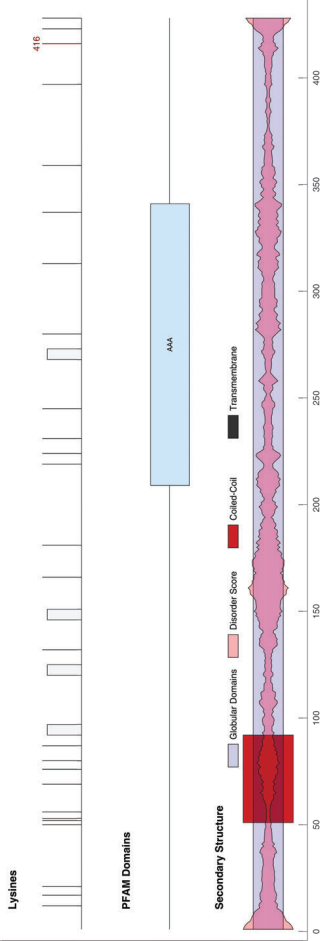
PRP45 (sp|P28004|PRP45_YEAST)



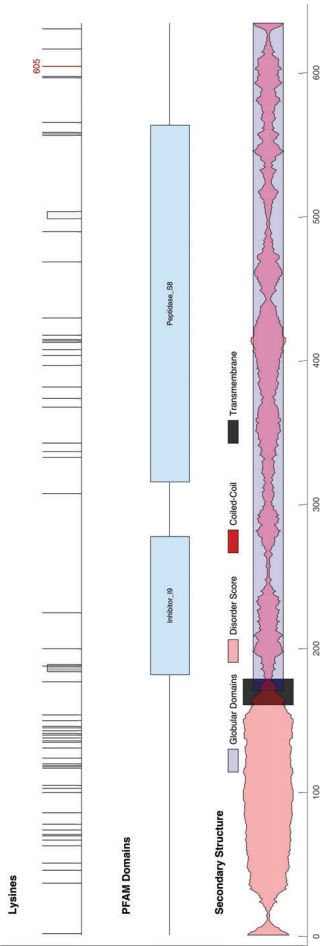
PRP46 (sp|Q12417|PRP46_YEAST)



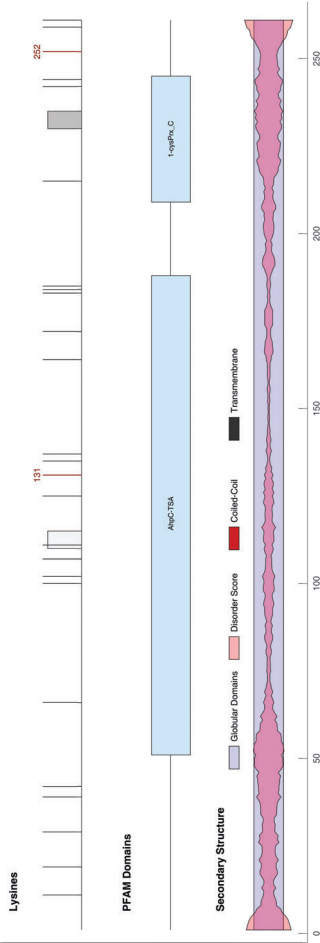
PRS6B (sp|P33258|PRS6B_YEAST)



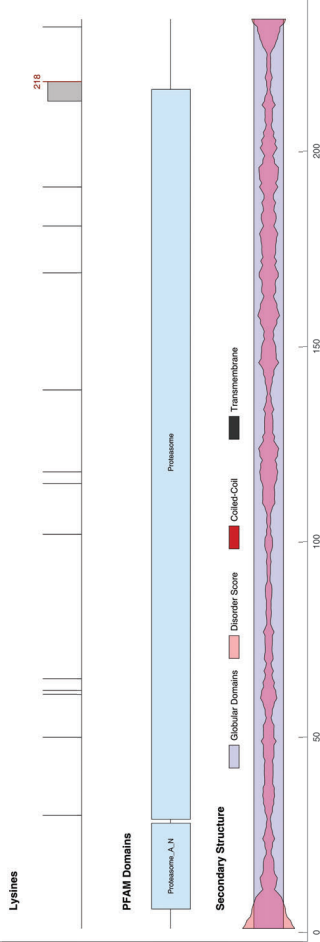
PRTB (sp|P09232|PRTB_YEAST)



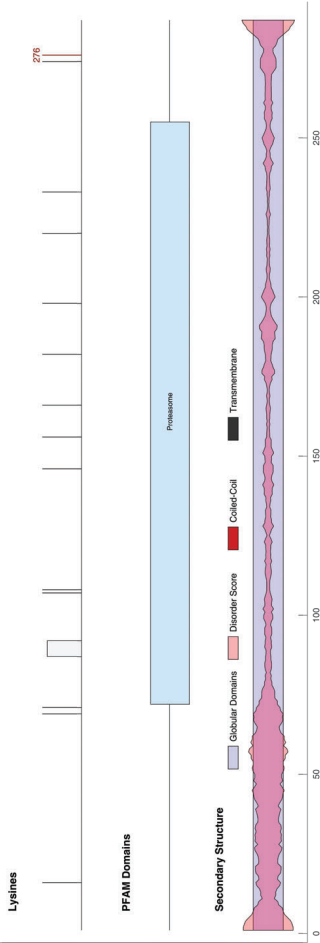
PRX1 (sp|P34227|PRX1_YEAST)



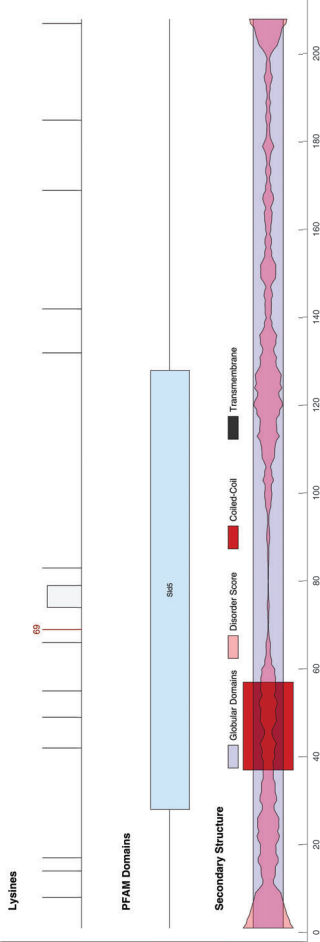
PSA6 (sp|P40302|PSA6_YEAST)

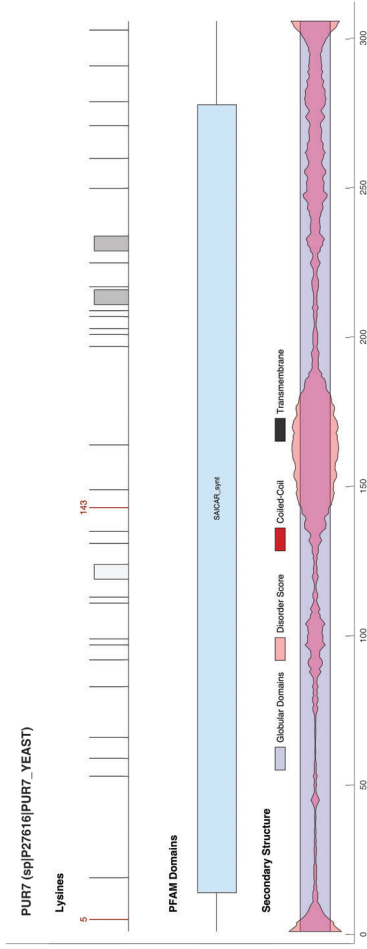
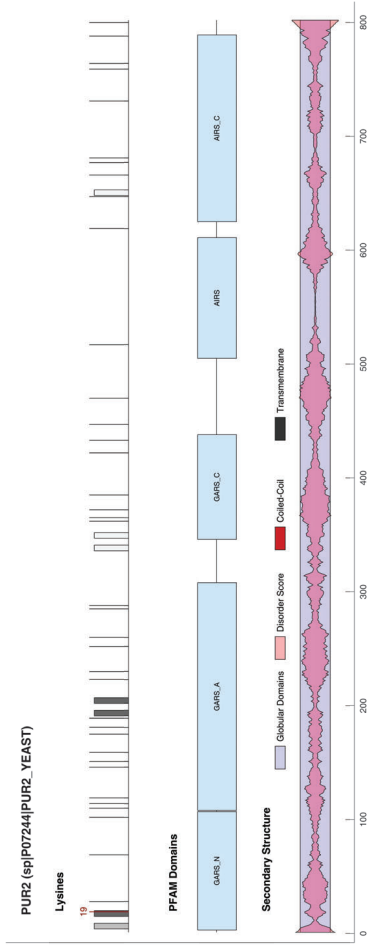
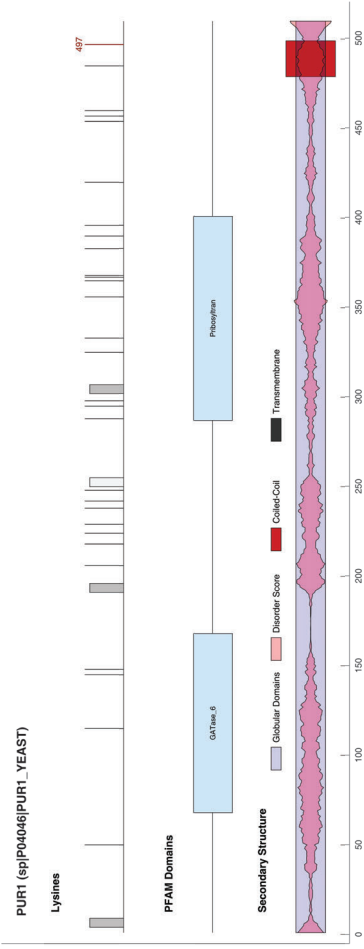
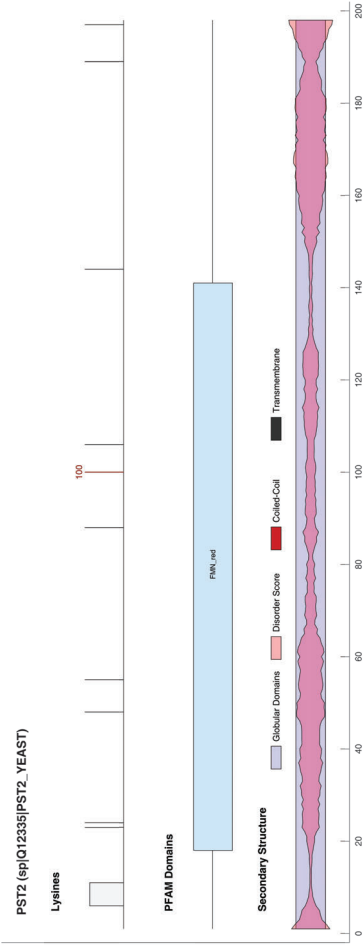


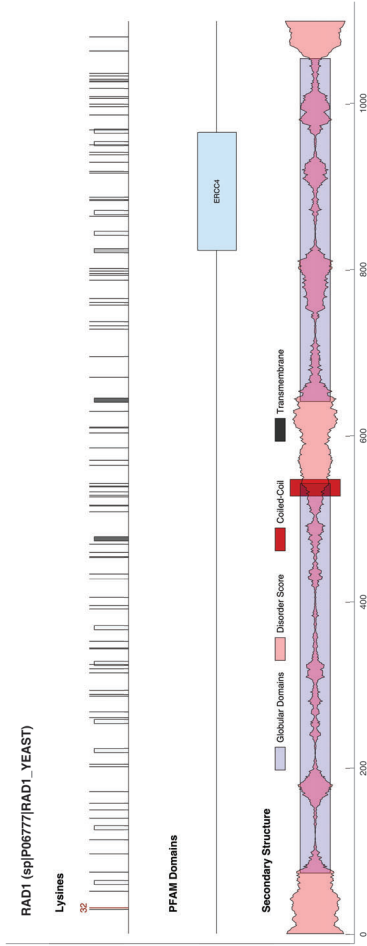
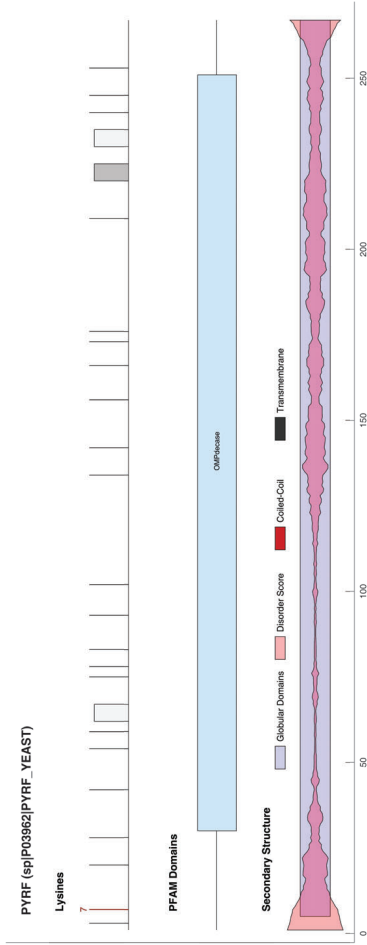
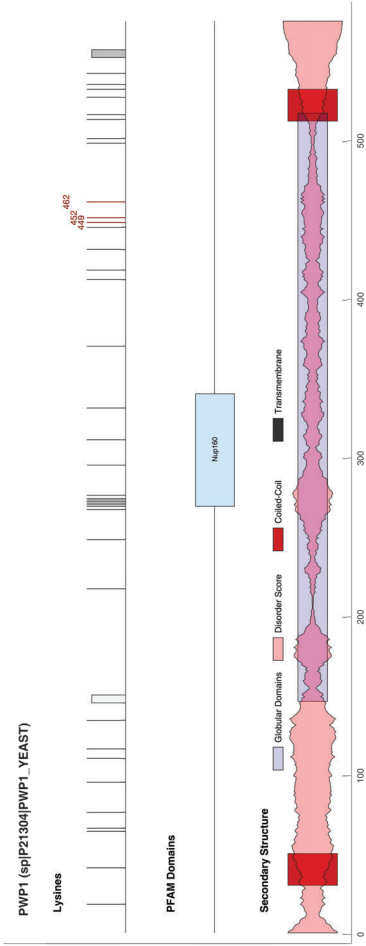
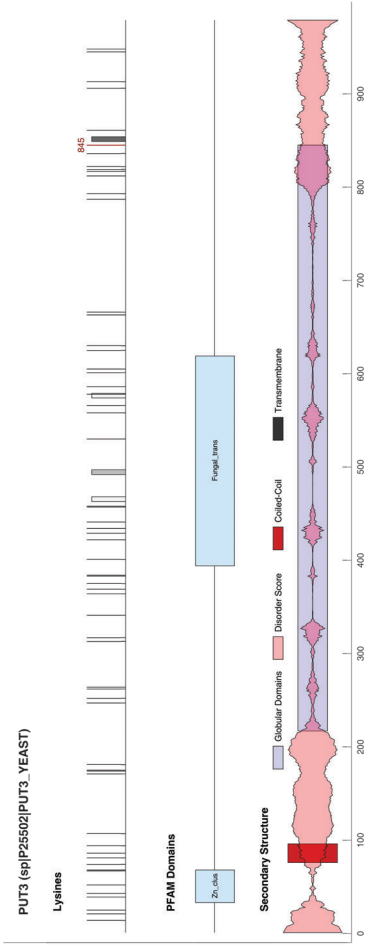
PSB5 (sp|P30656|PSB5_YEAST)

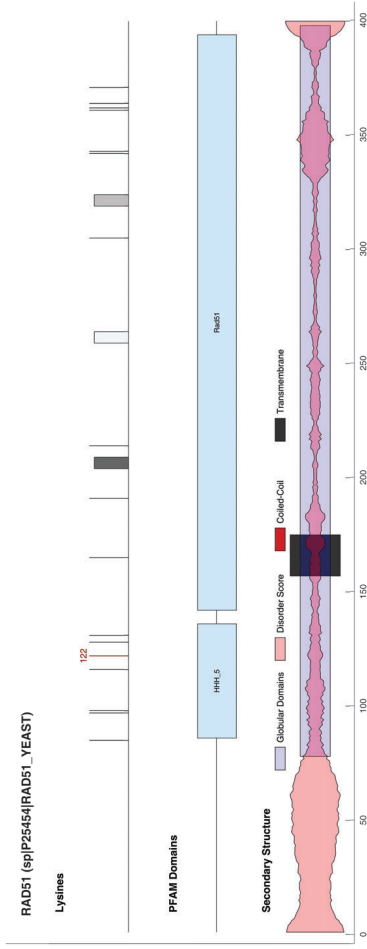
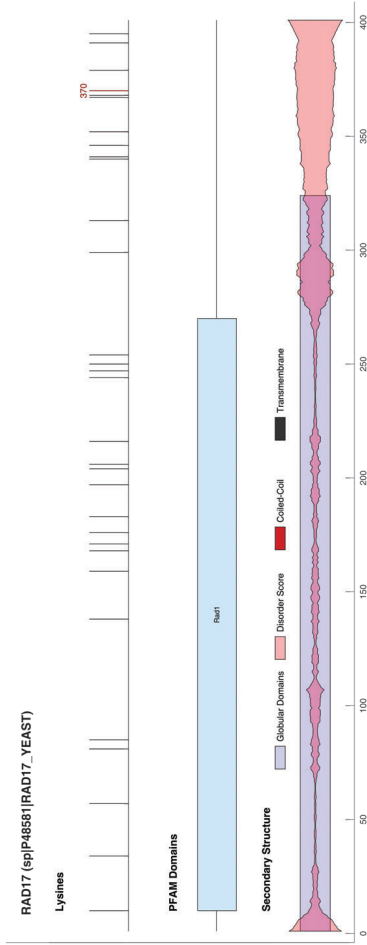
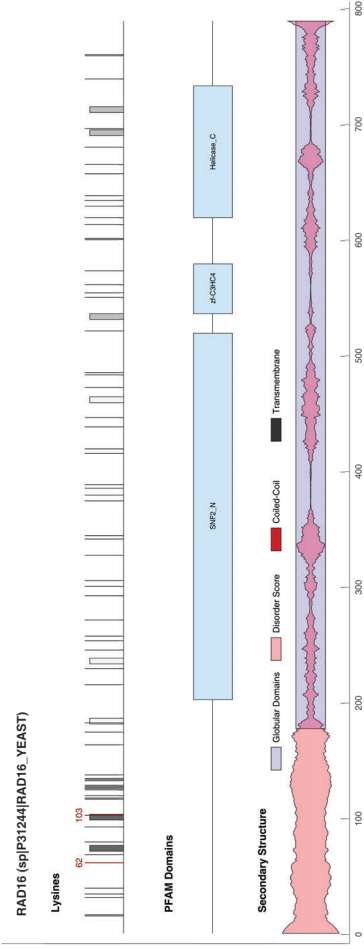
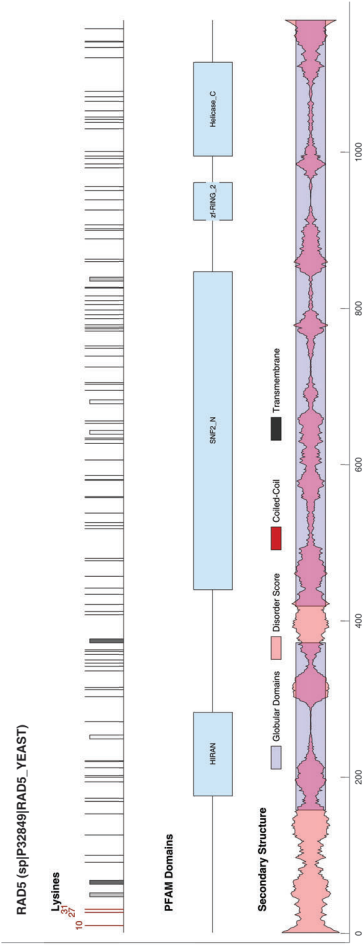


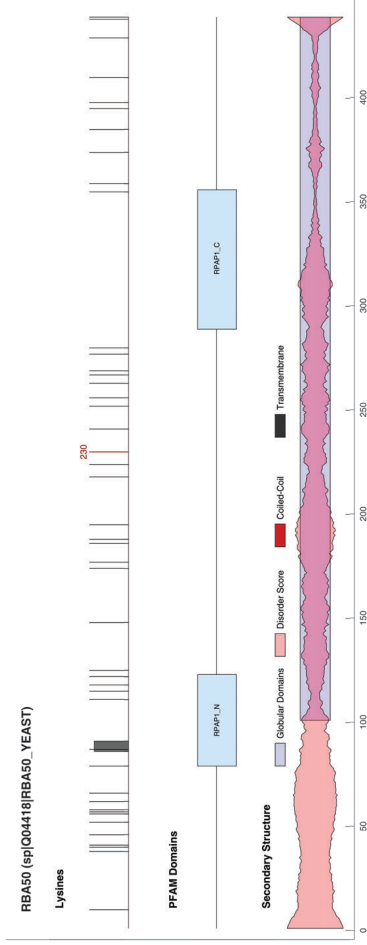
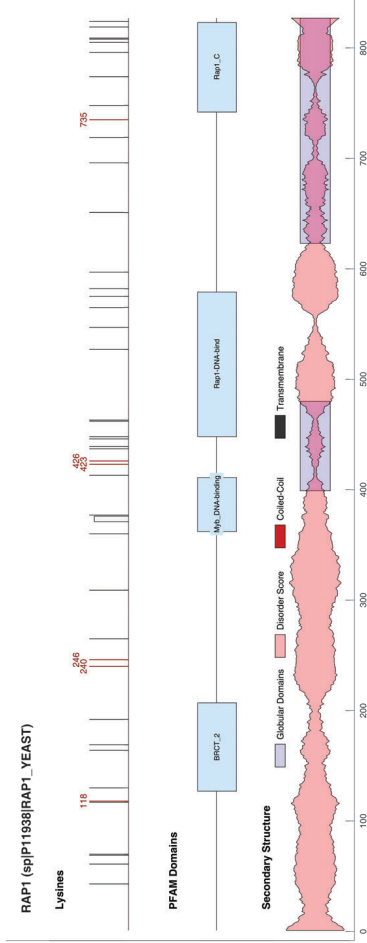
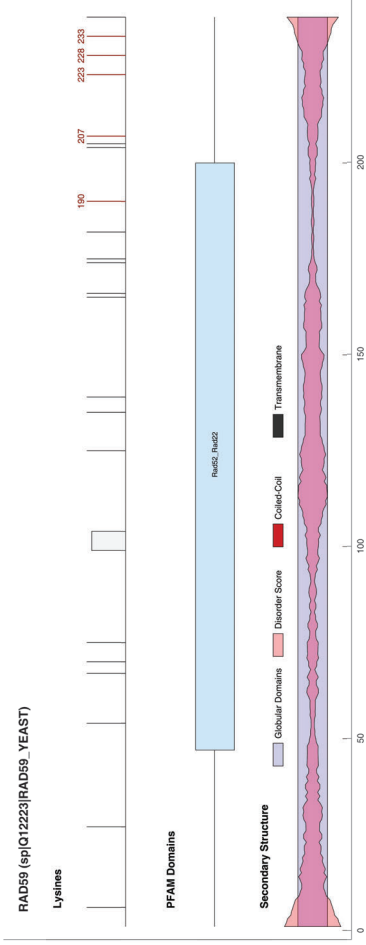
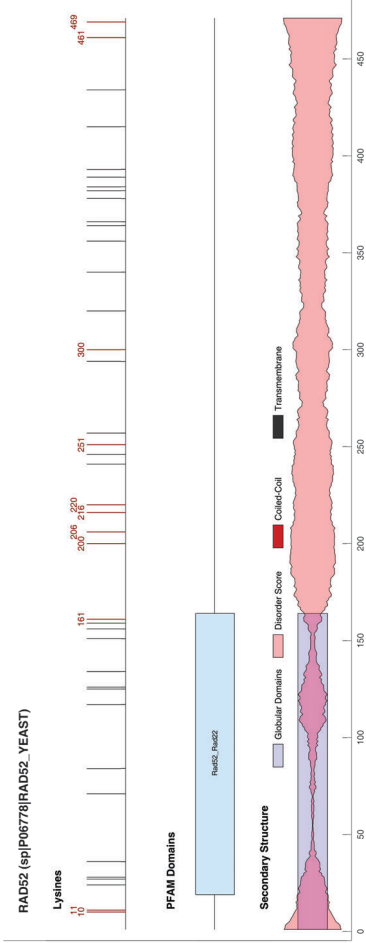
PSF1 (sp|Q12488|PSF1_YEAST)

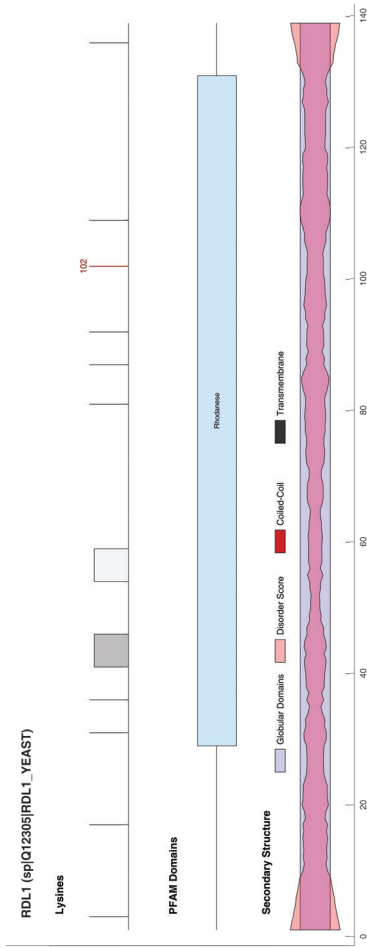
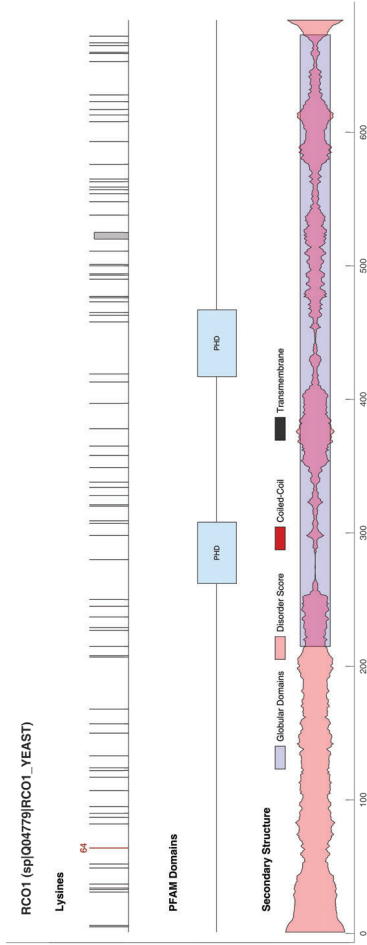
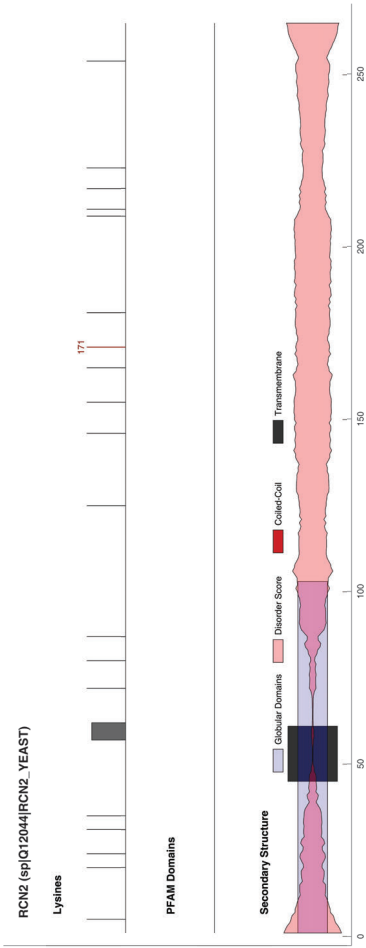
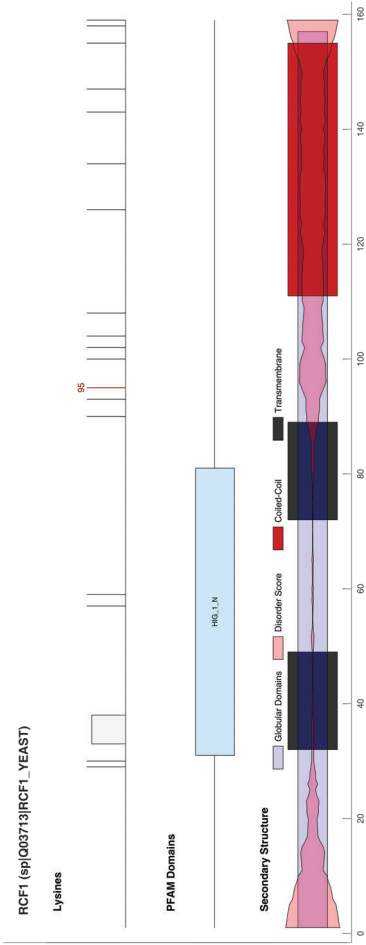


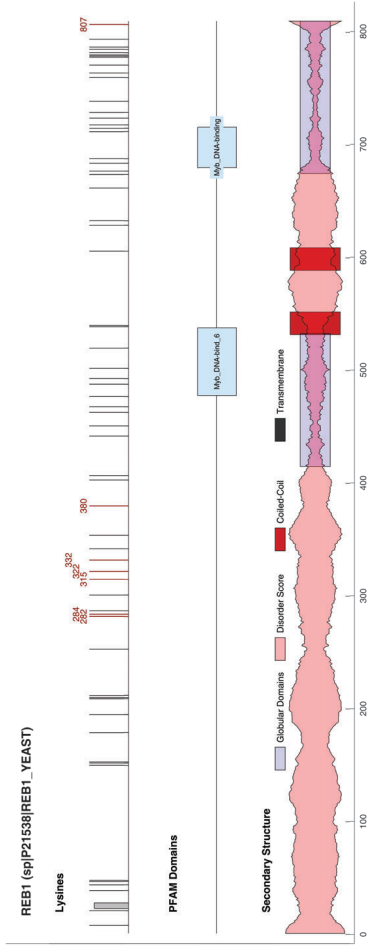
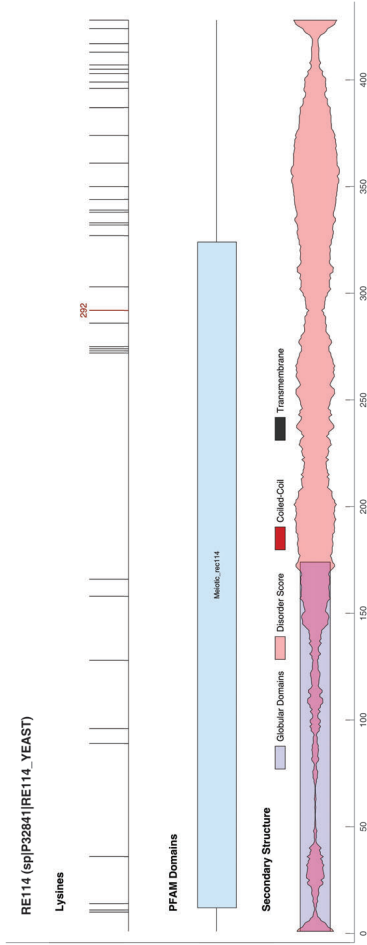
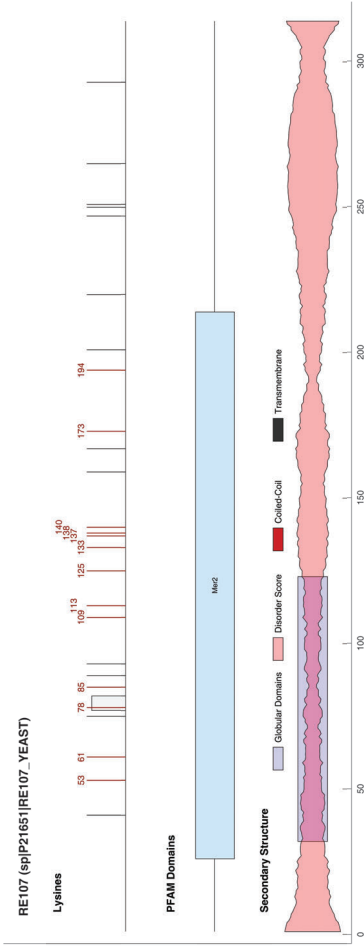
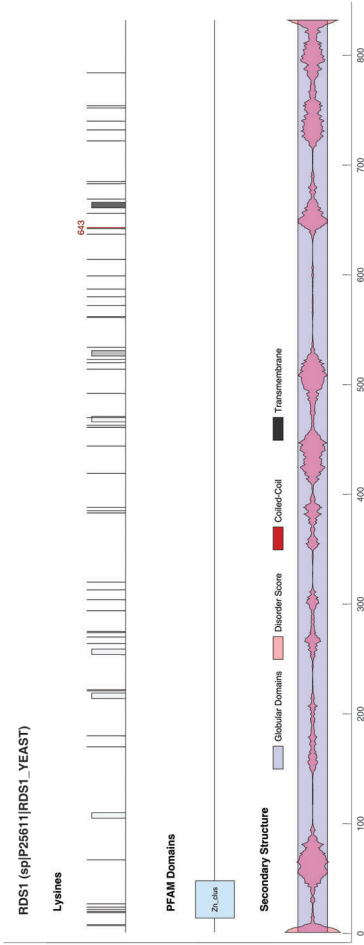


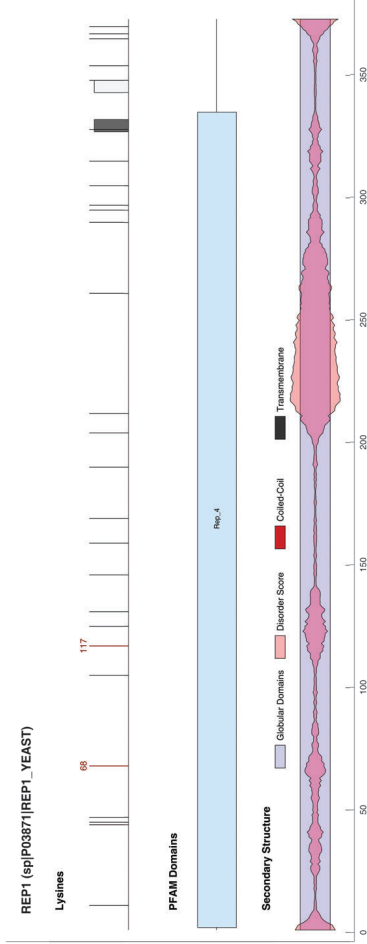
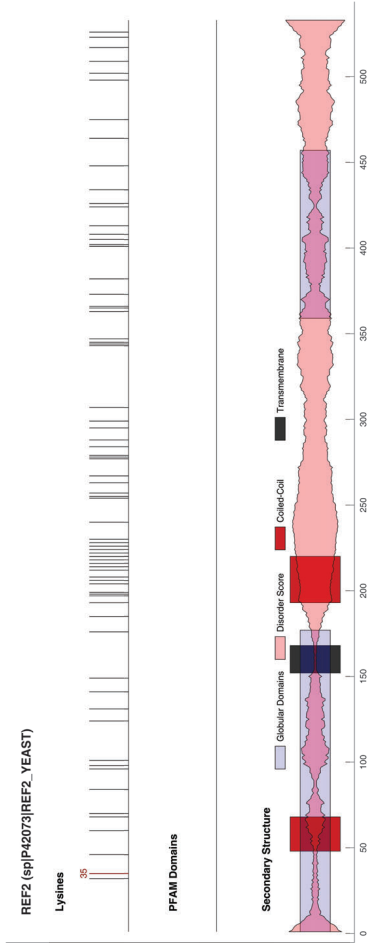
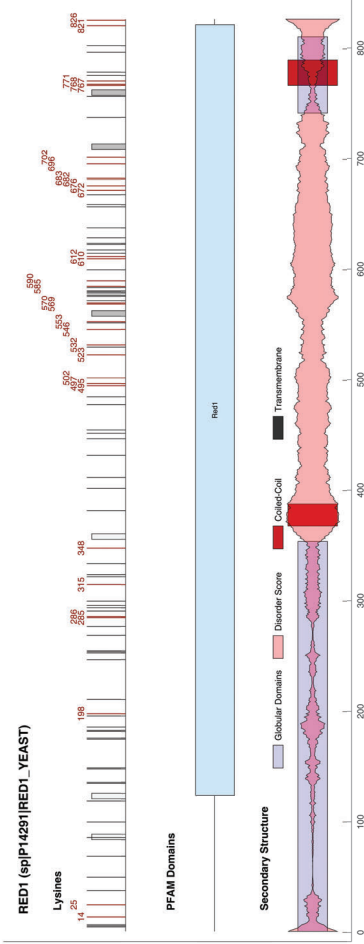
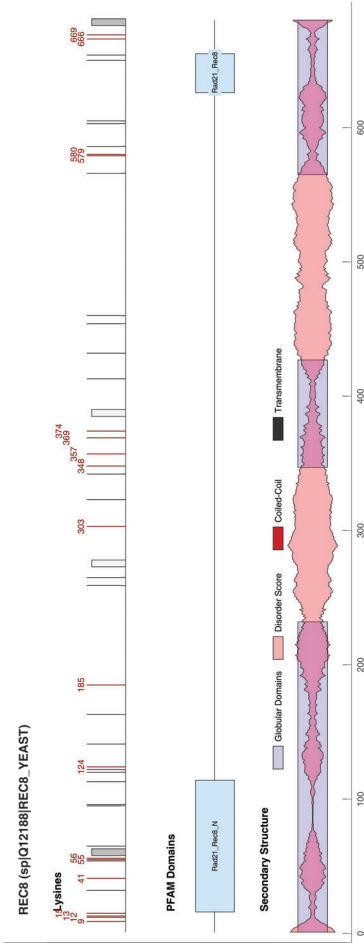




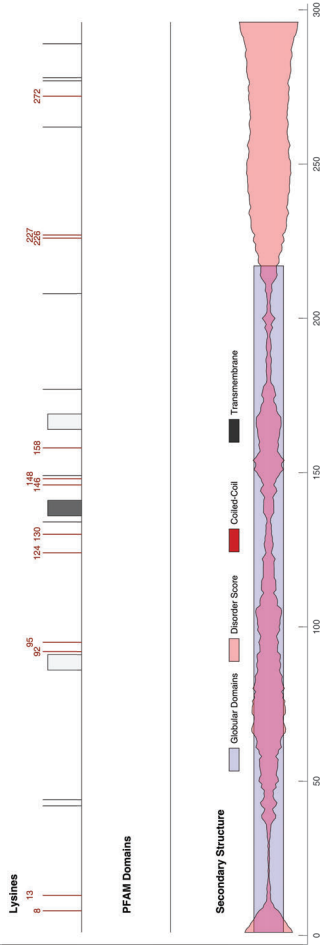




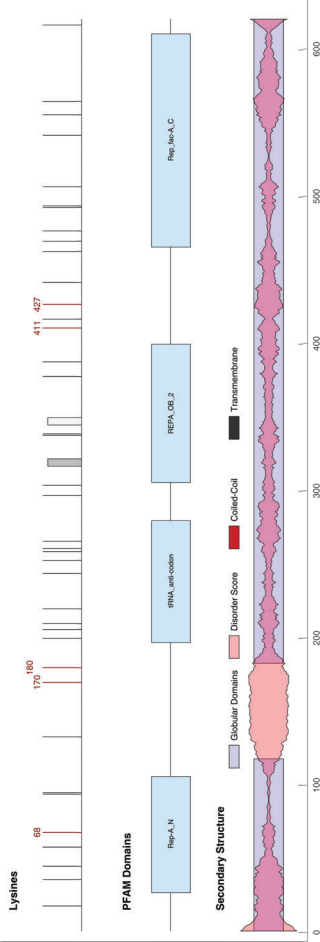




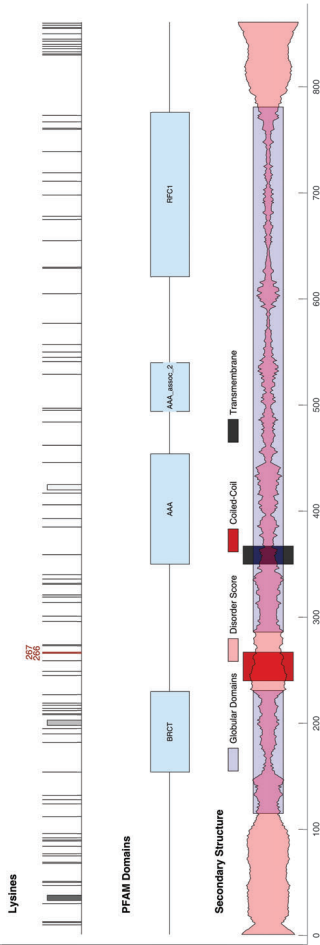
REP2 (sp|P03872|REP2_YEAST)



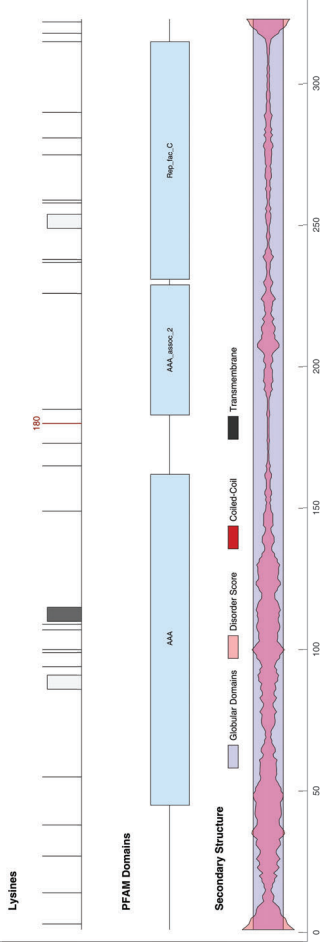
RFA1 (sp|P22336|RFA1_YEAST)

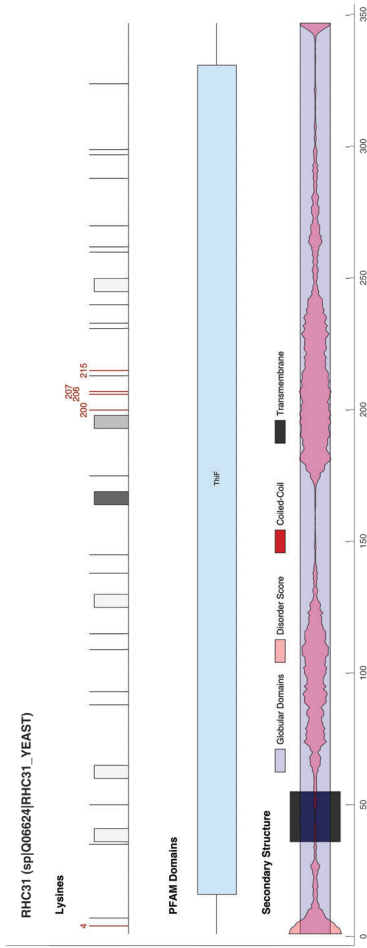
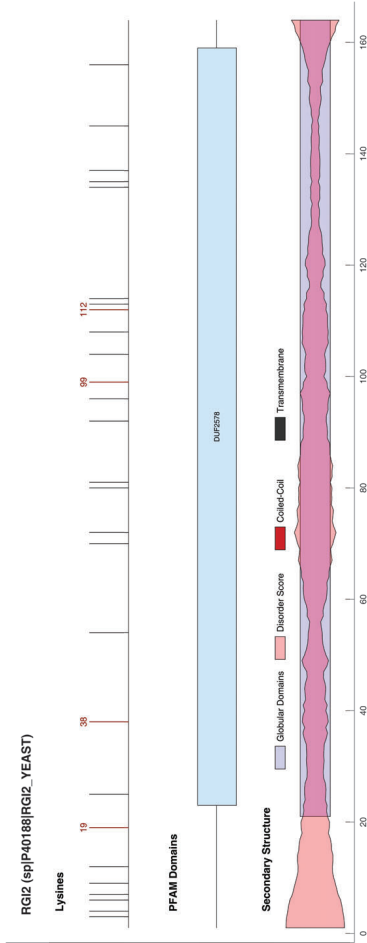
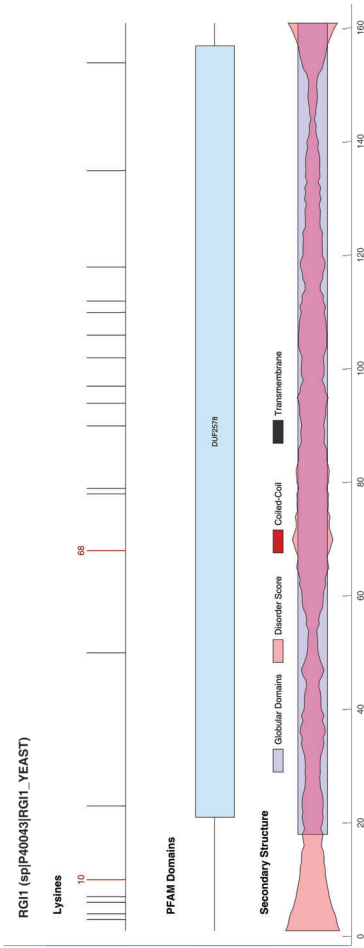
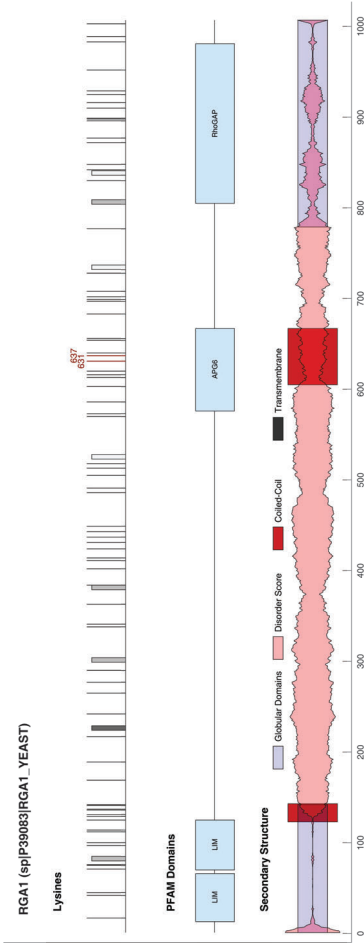


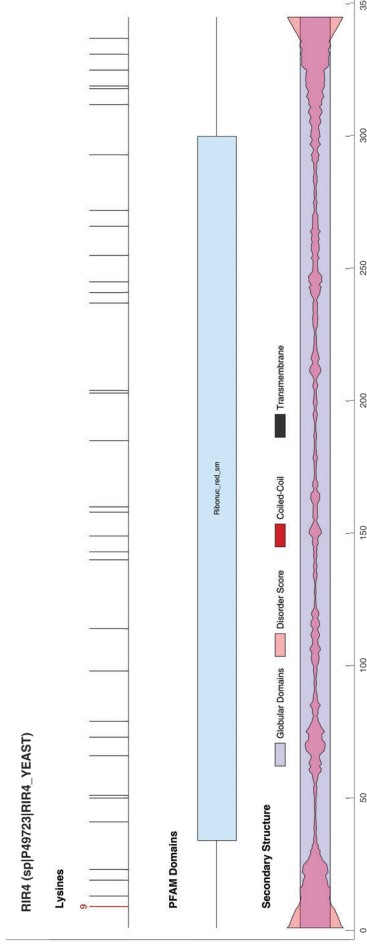
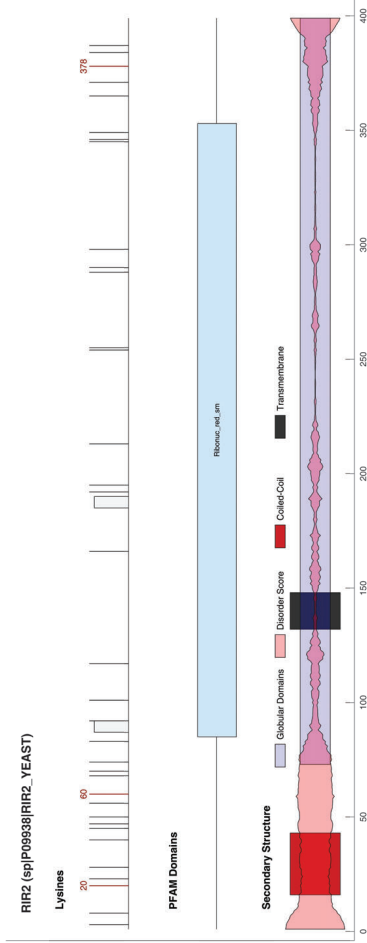
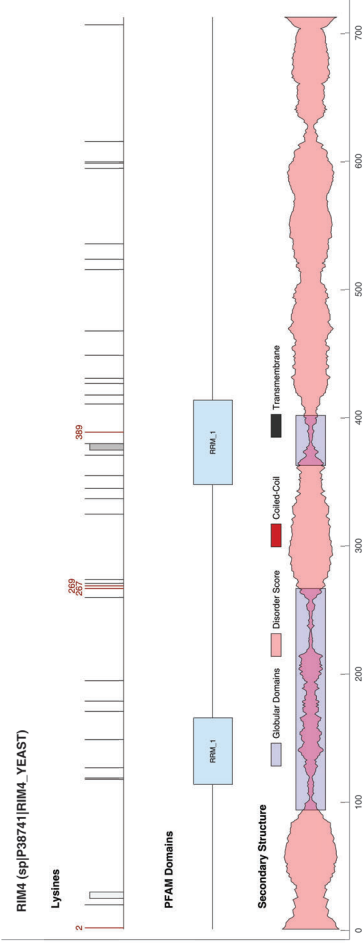
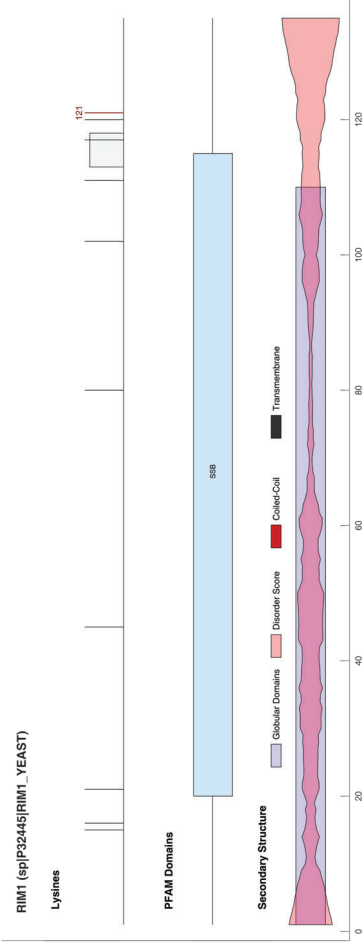
RFC1 (sp|P38630|RFC1_YEAST)



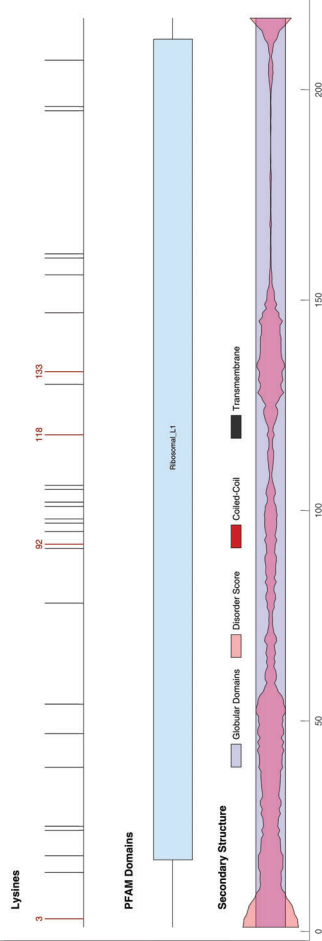
RFC4 (sp|P40339|RFC4_YEAST)



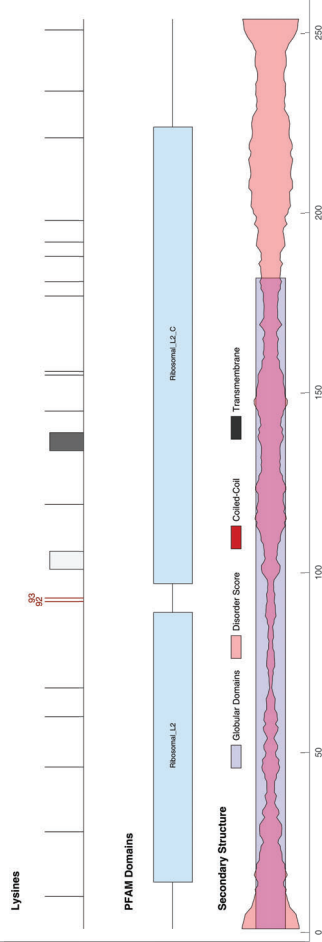




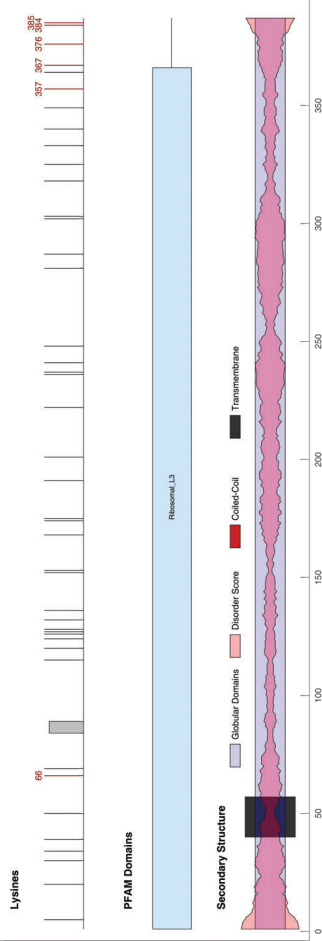
RL1B (sp|P0CX44|RL1B_YEAST)



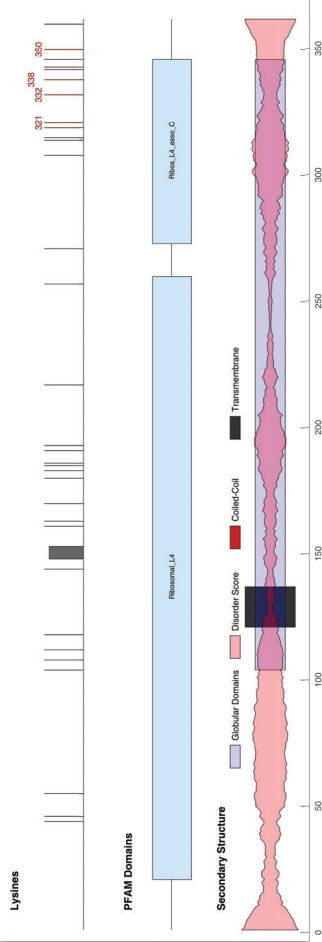
RL2B (sp|P0CX46|RL2B_YEAST)



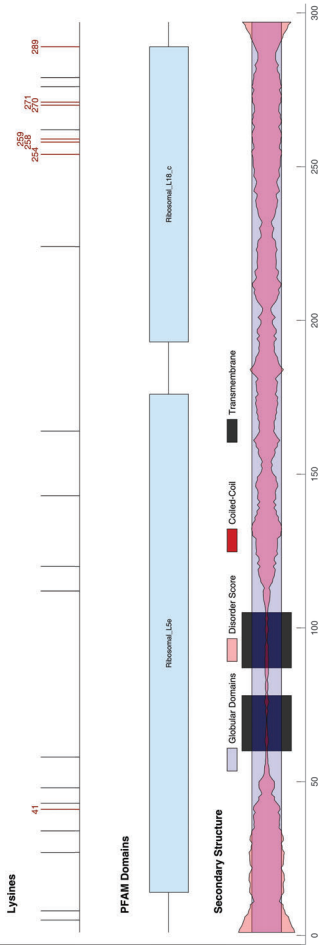
RL3 (sp|P14126|RL3_YEAST)



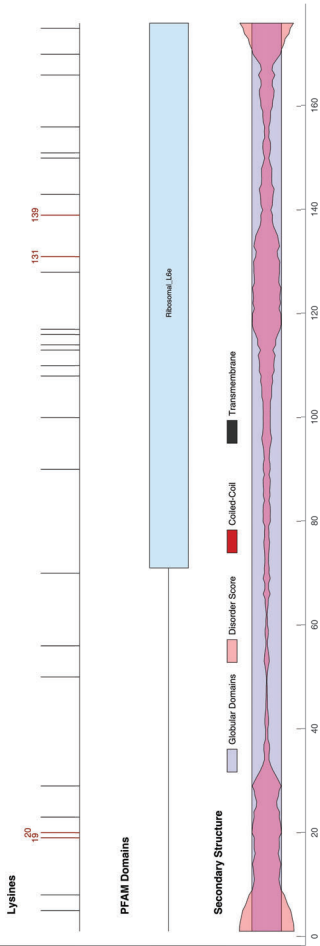
RL4A (sp|P10664|RL4A_YEAST)



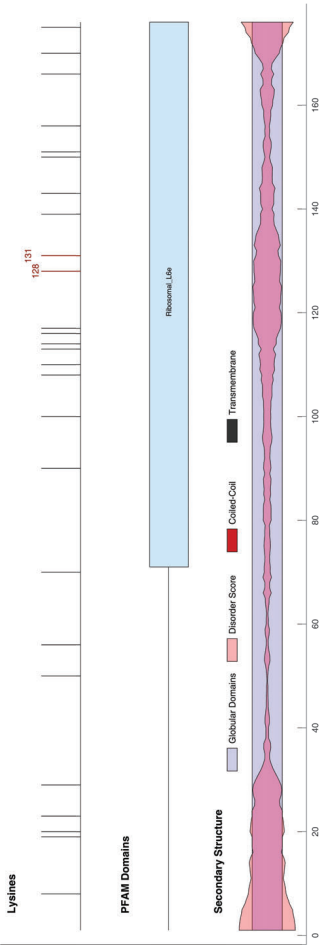
RL5 (sp|P26321|RL5_YEAST)



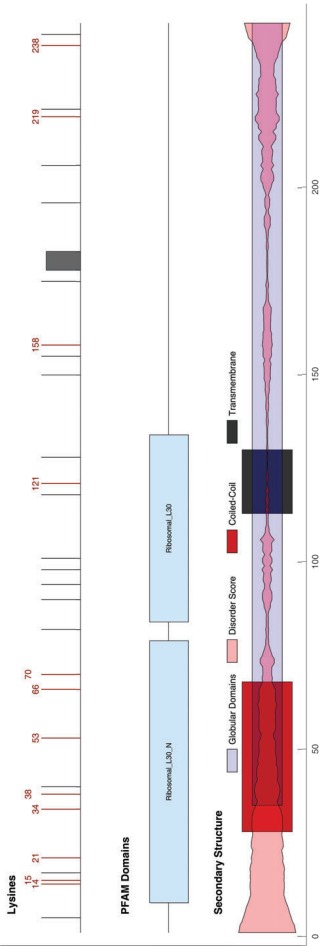
RL6A (sp|Q02326|RL6A_YEAST)

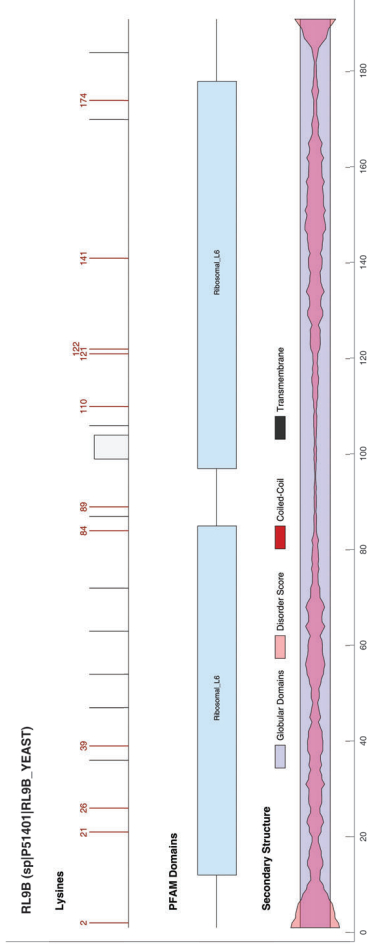
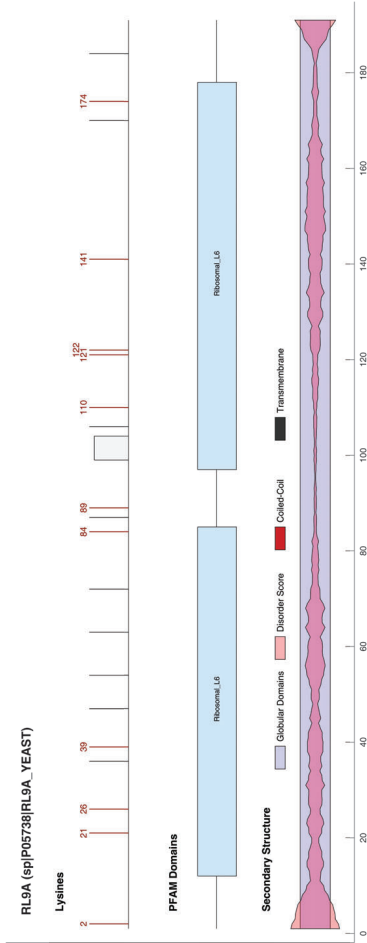
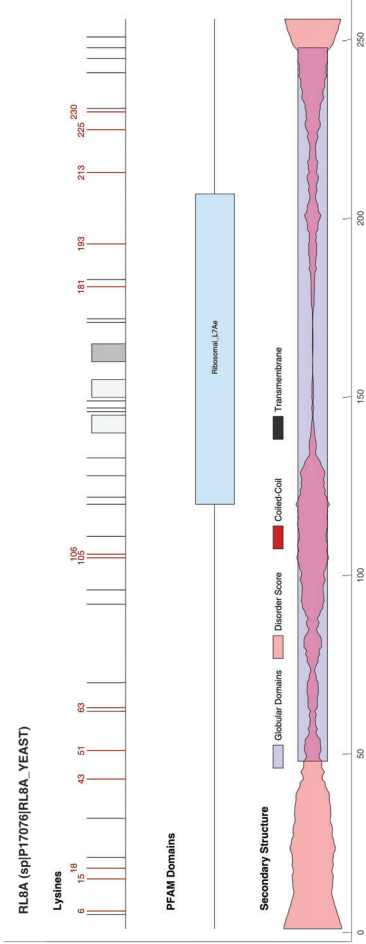
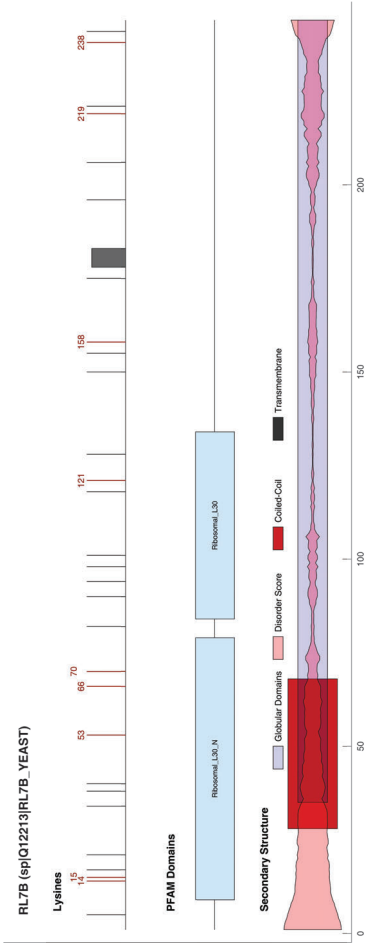


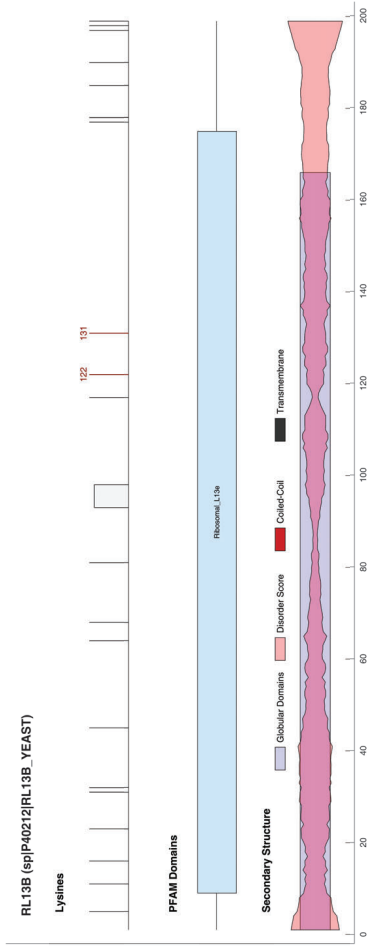
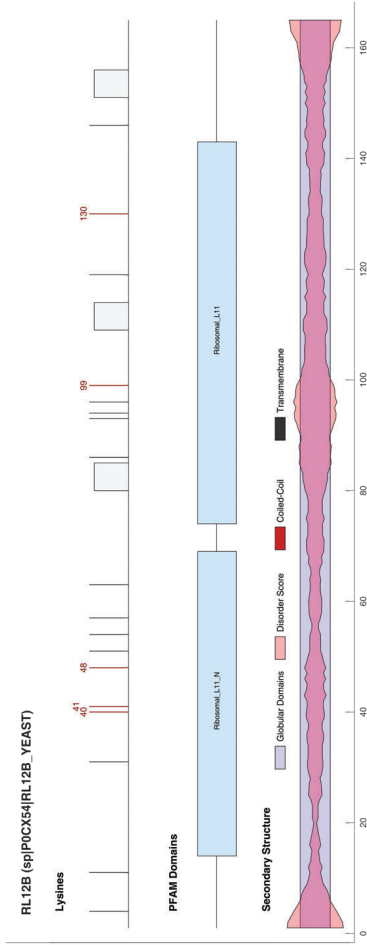
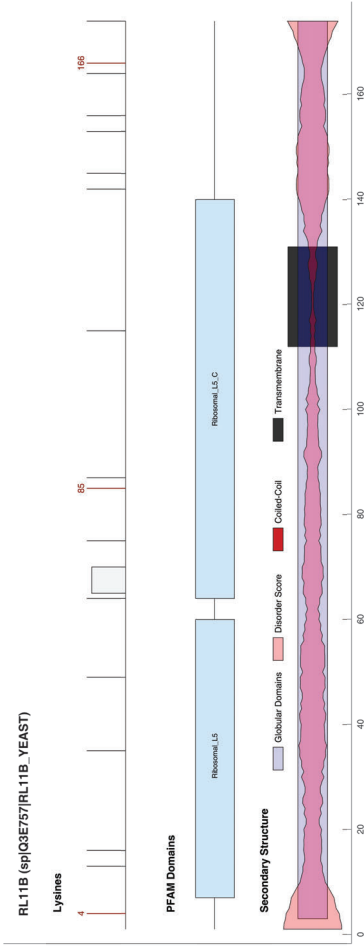
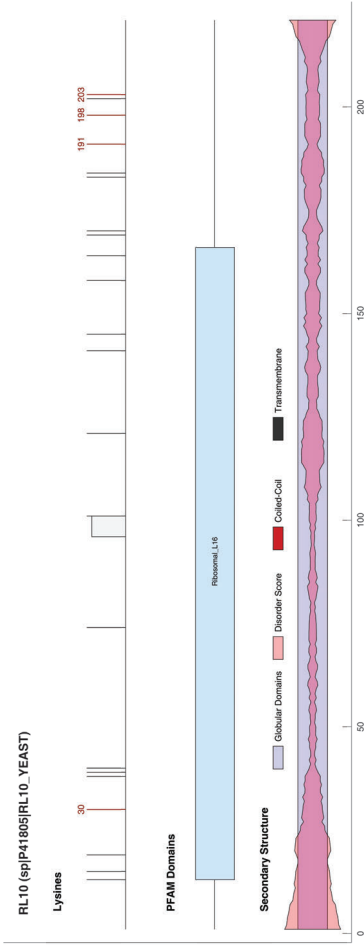
RL6B (sp|P05739|RL6B_YEAST)

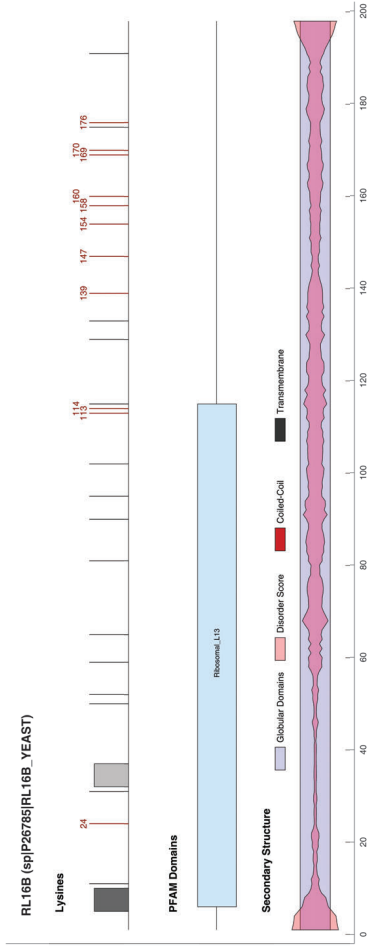
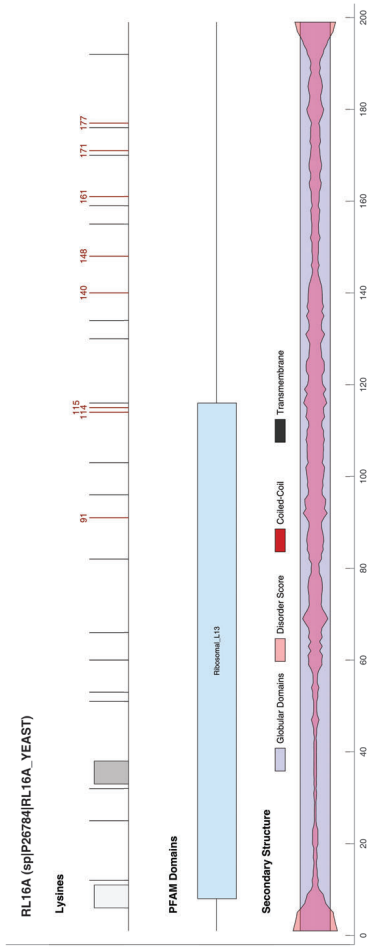
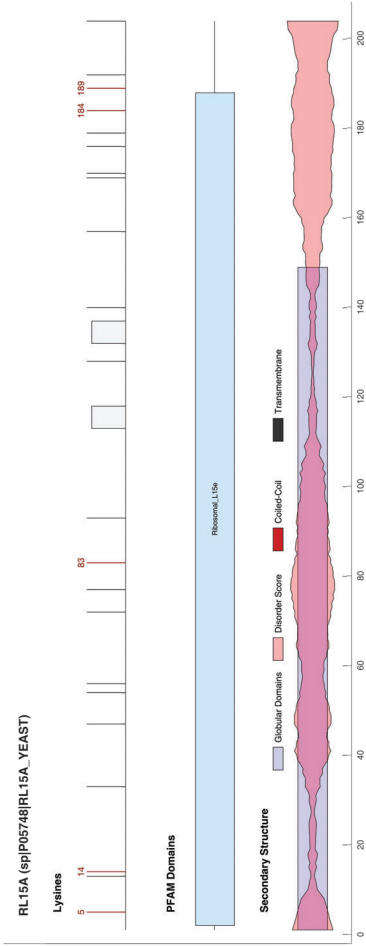
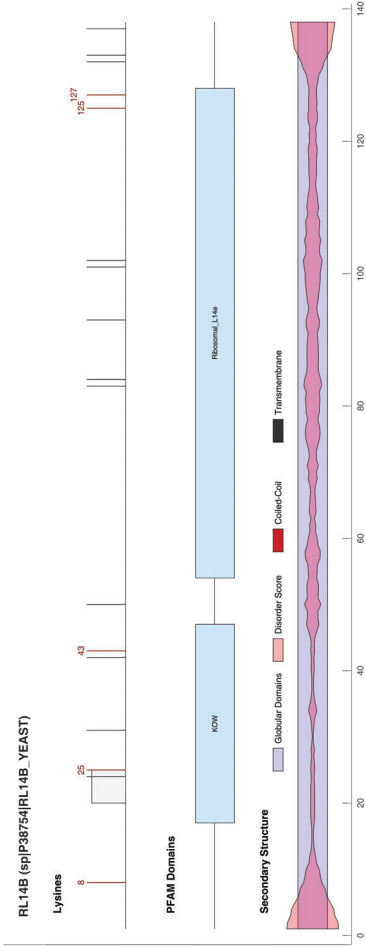


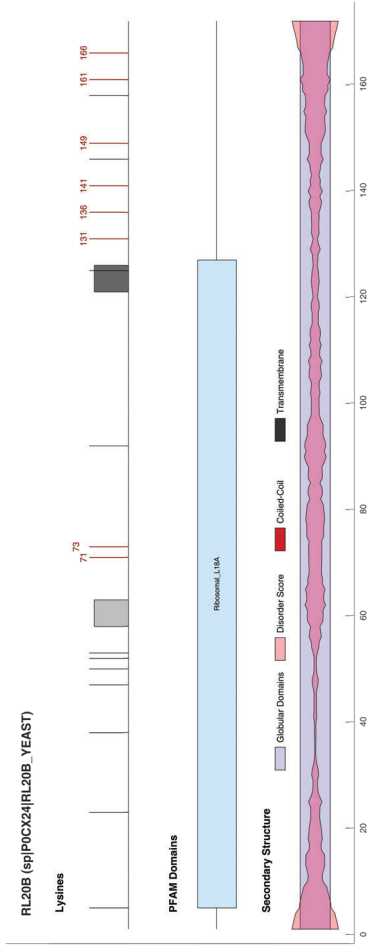
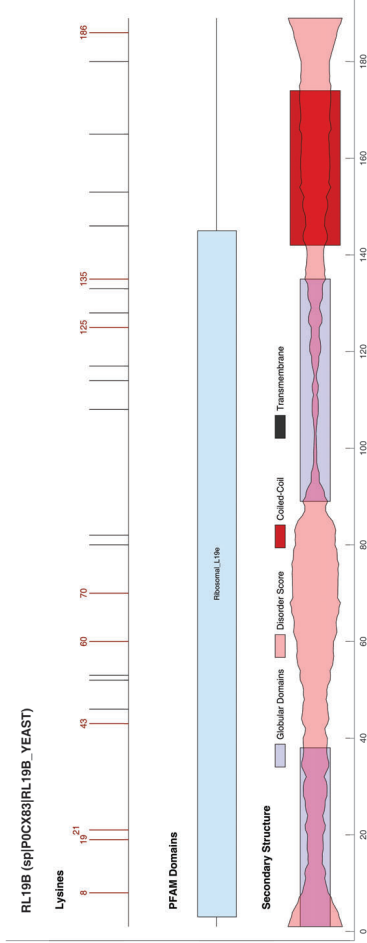
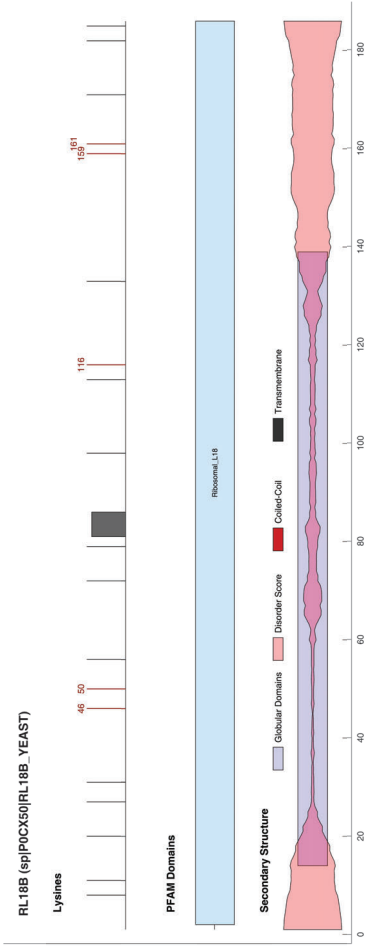
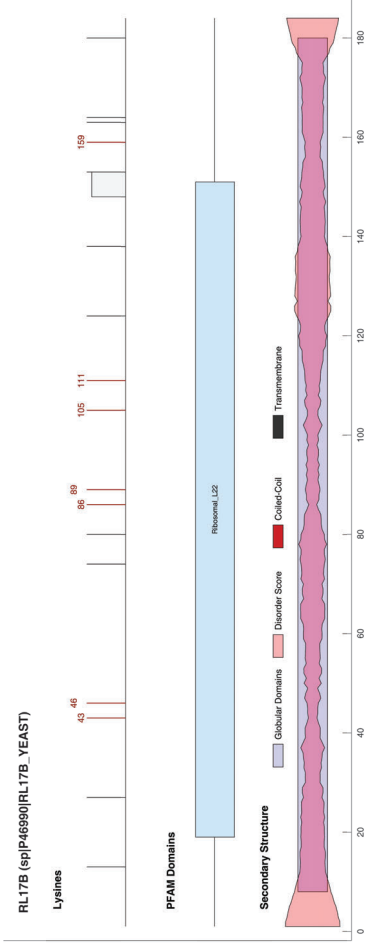
RL7A (sp|P05737|RL7A_YEAST)

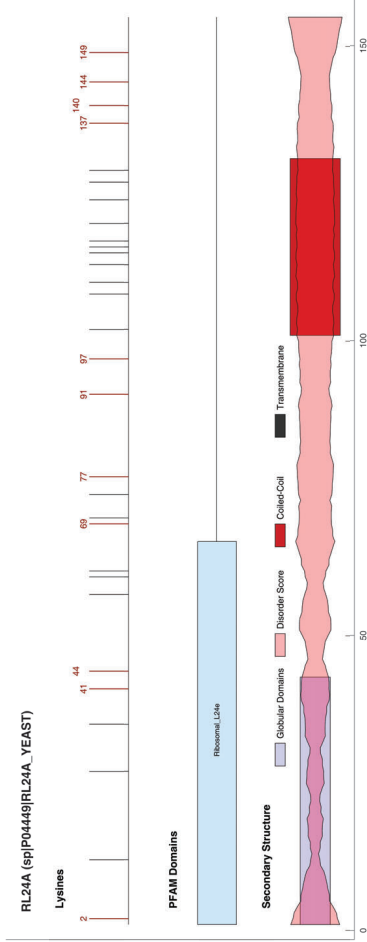
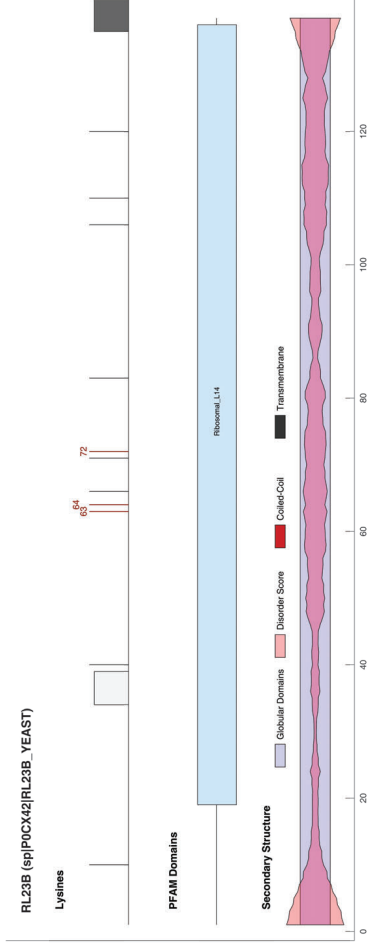
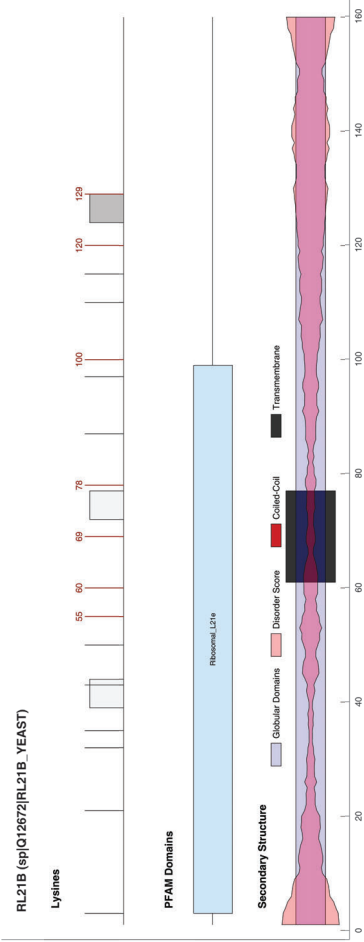
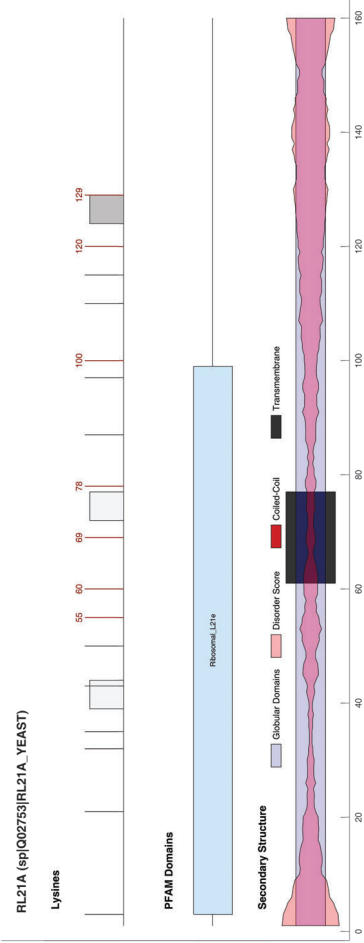


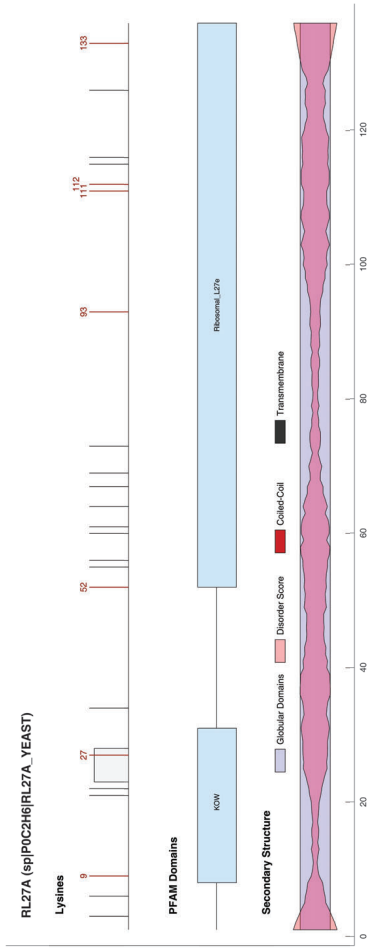
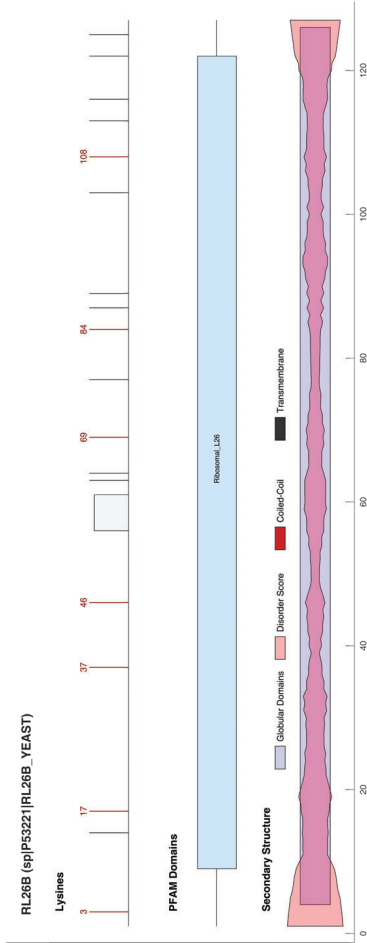
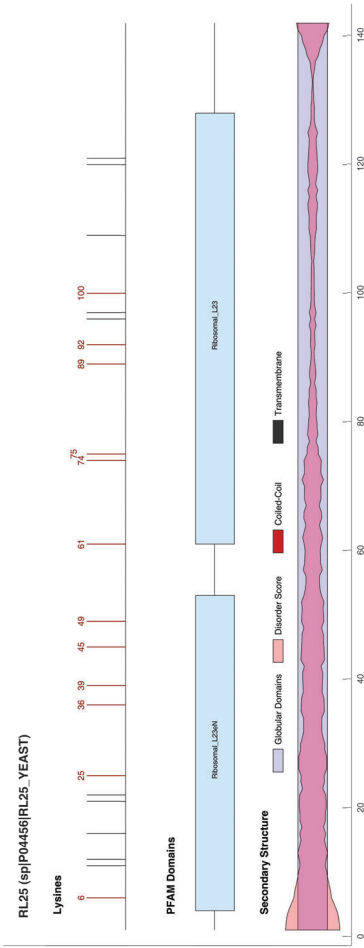
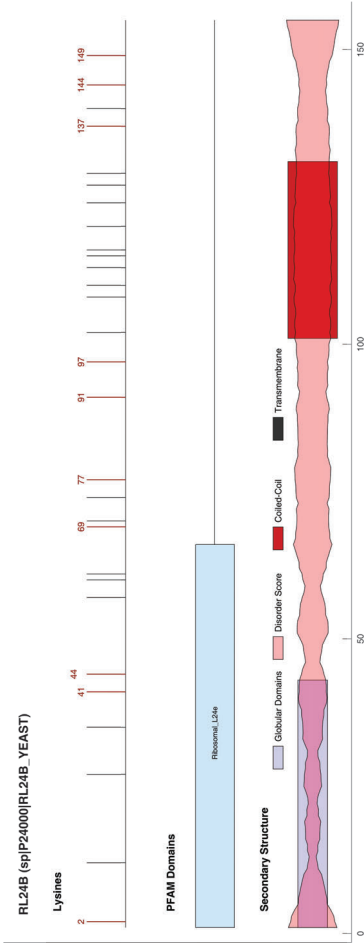




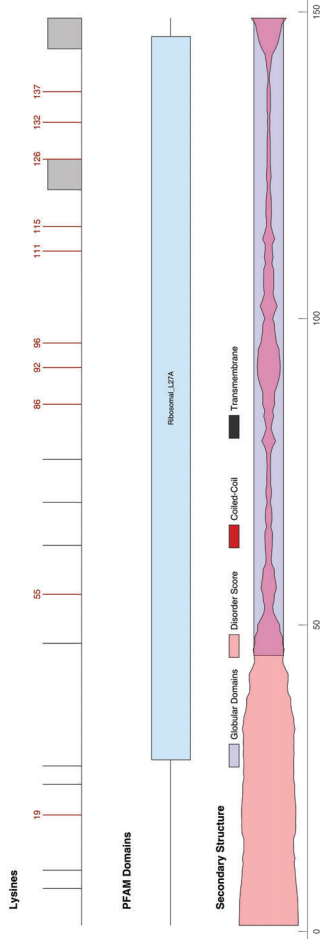




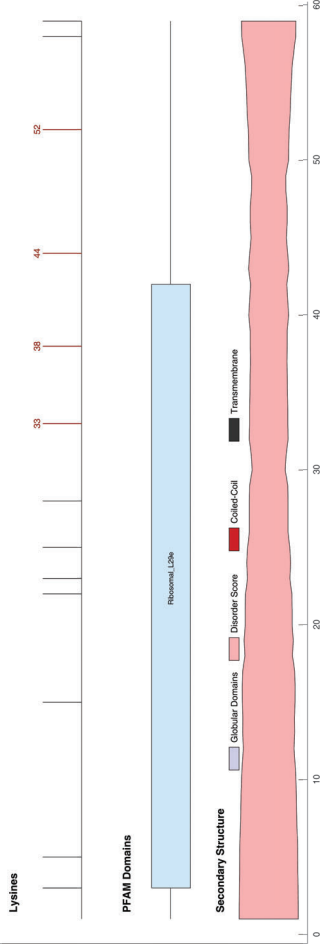




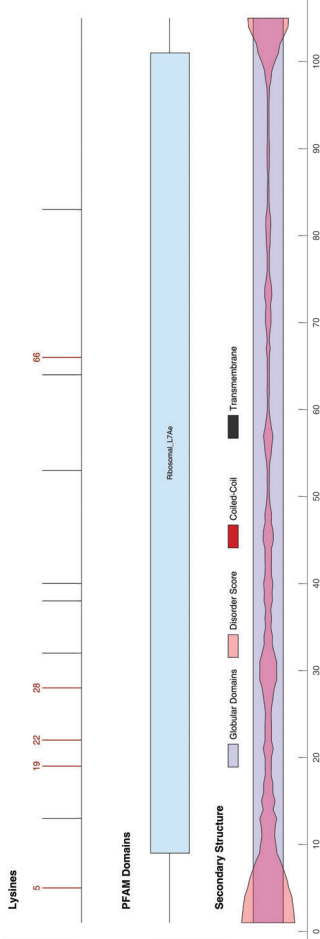
RL28 (sp|P02406|RL28_YEAST)



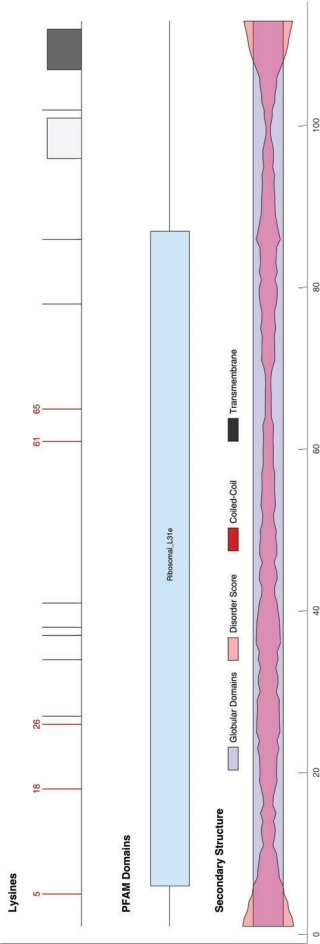
RL29 (sp|P05747|RL29_YEAST)



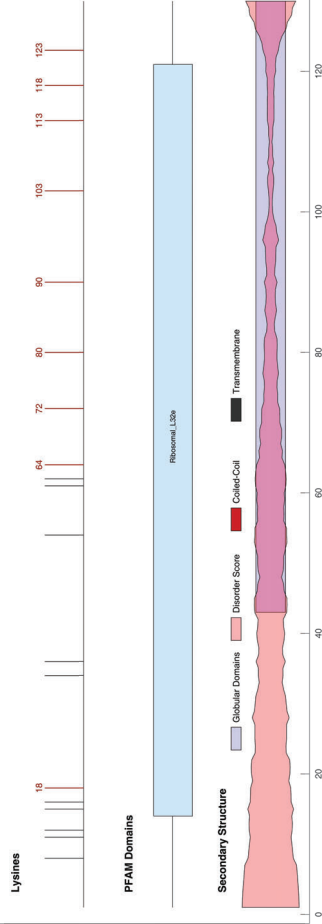
RL30 (sp|P14120|RL30_YEAST)



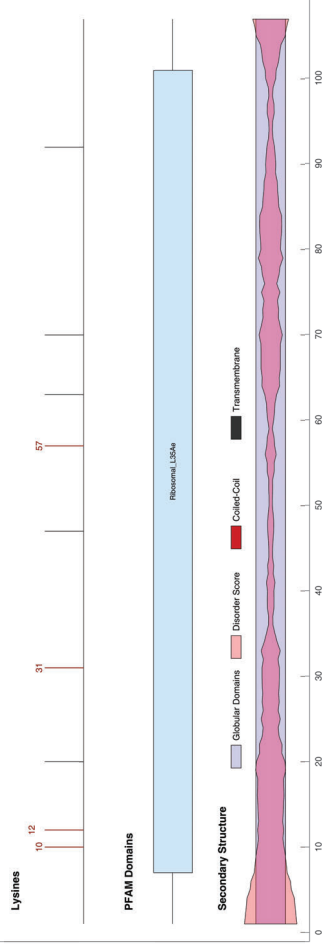
RL31A (sp|P0C2H8|RL31A_YEAST)



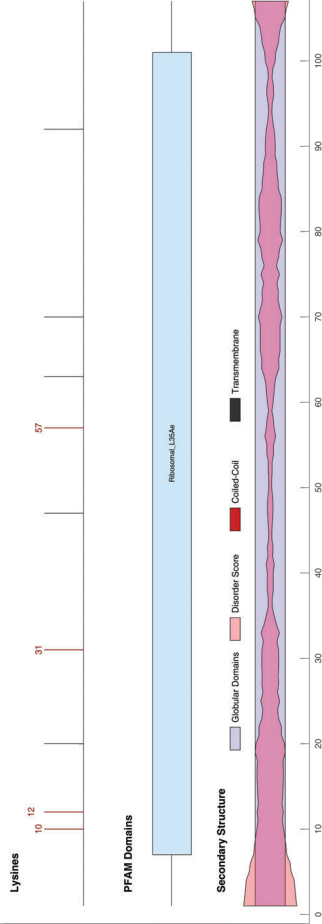
RL32 (sp|P38061|RL32_YEAST)



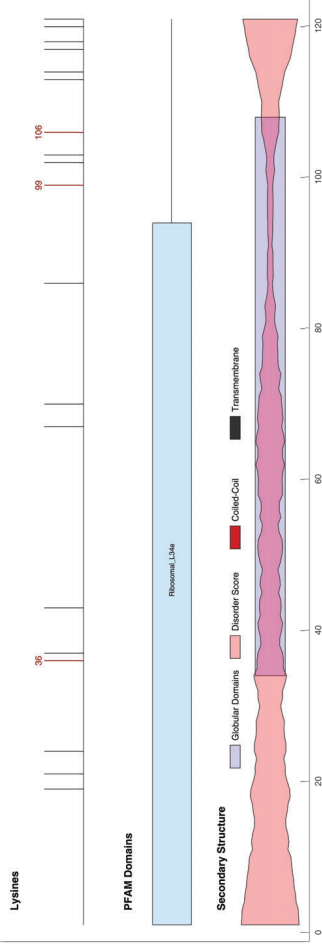
RL33A (sp|P05744|RL33A_YEAST)

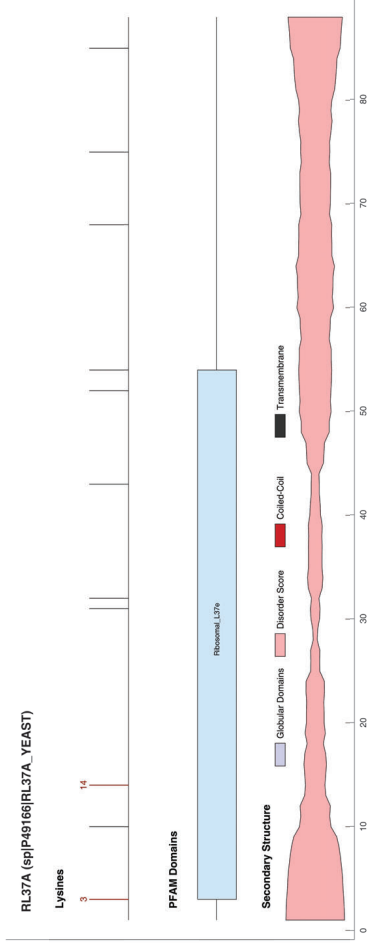
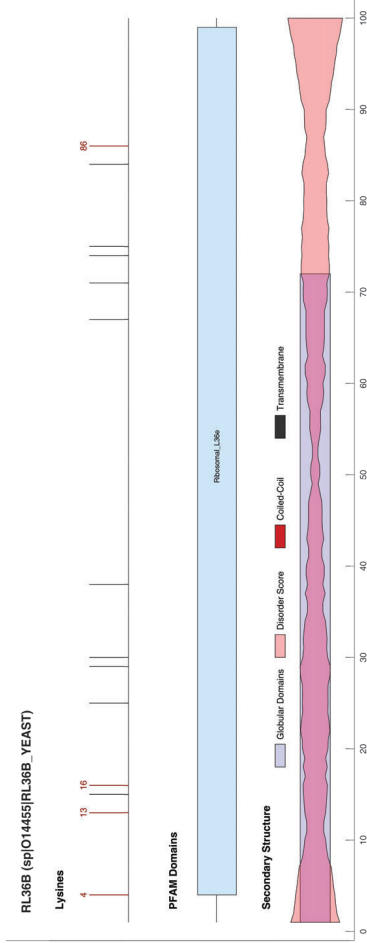
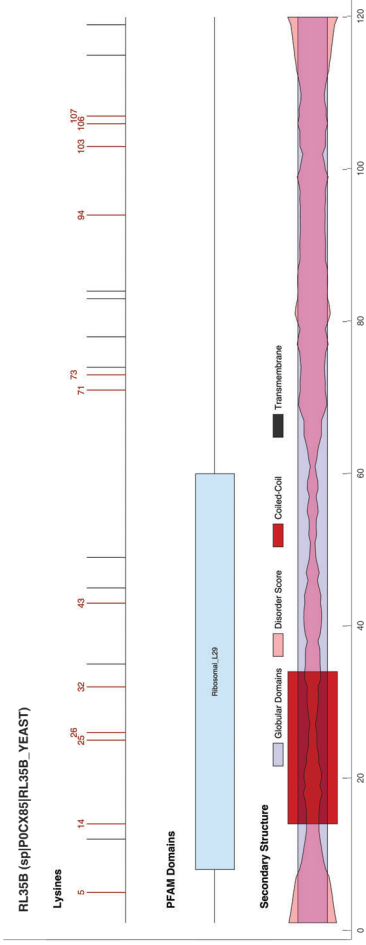
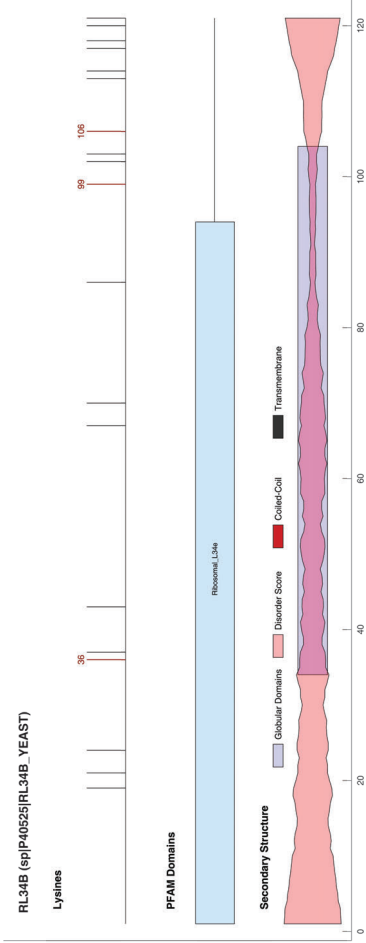


RL33E (sp|P41056|RL33E_YEAST)

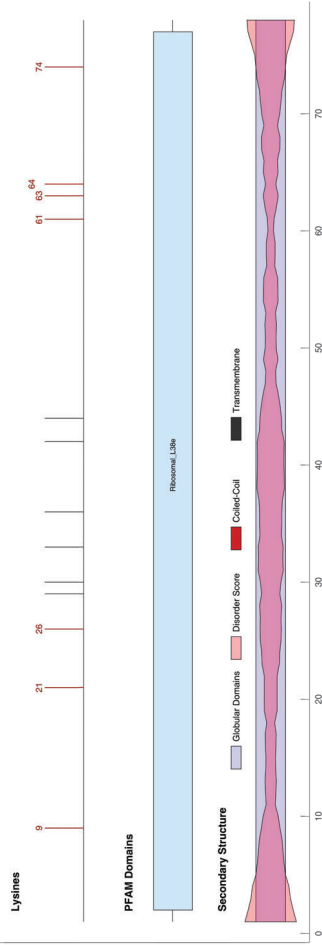


RL34A (sp|P87262|RL34A_YEAST)

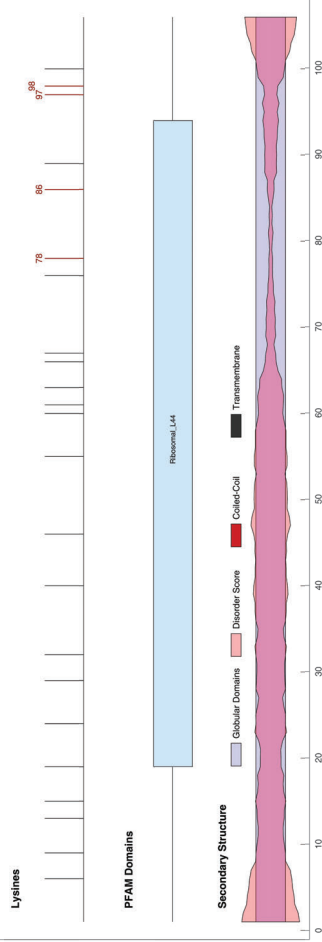




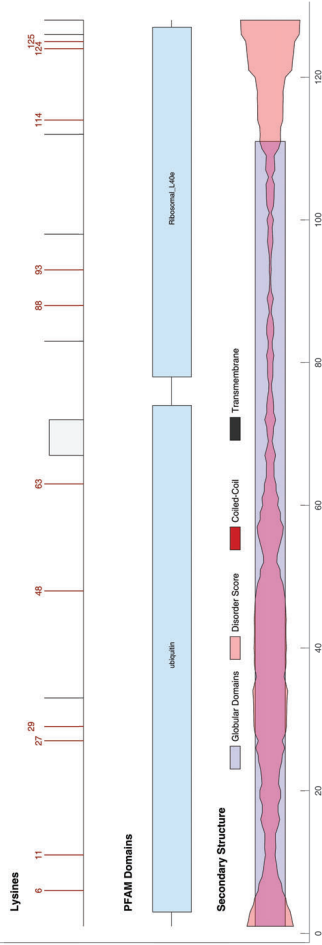
RL38 (sp|P49167|RL38_YEAST)



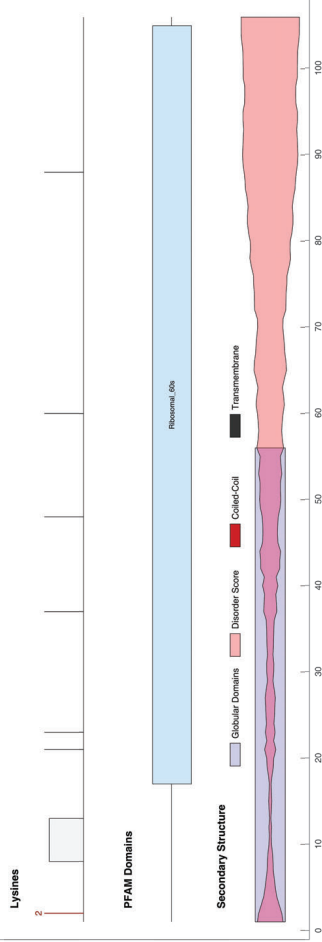
RL44B (sp|P0CX28|RL44B_YEAST)

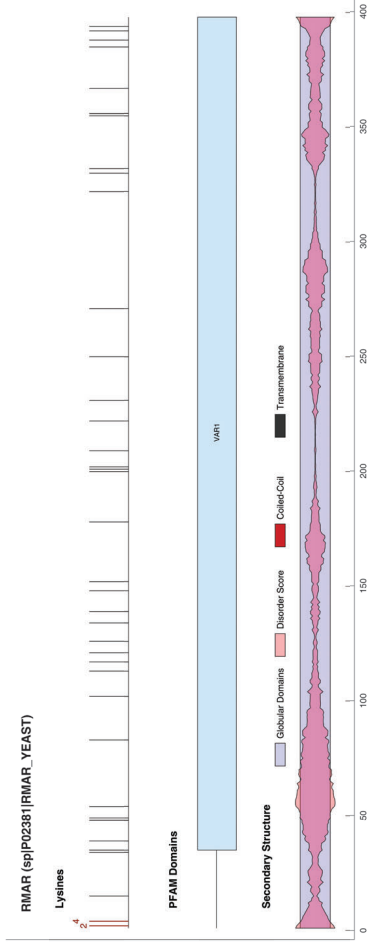
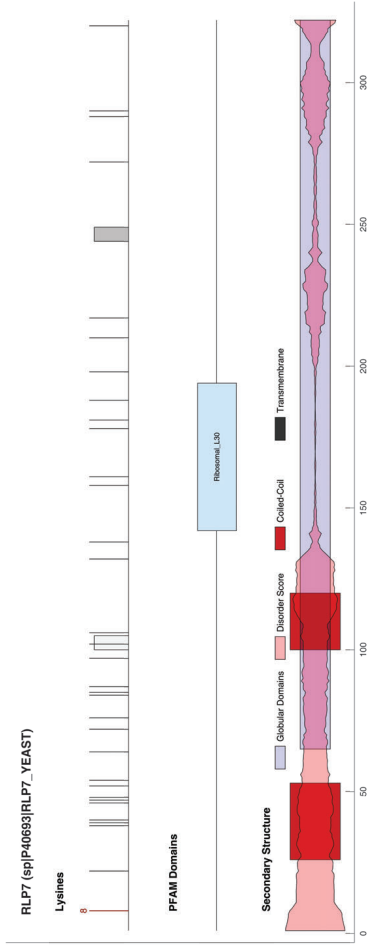
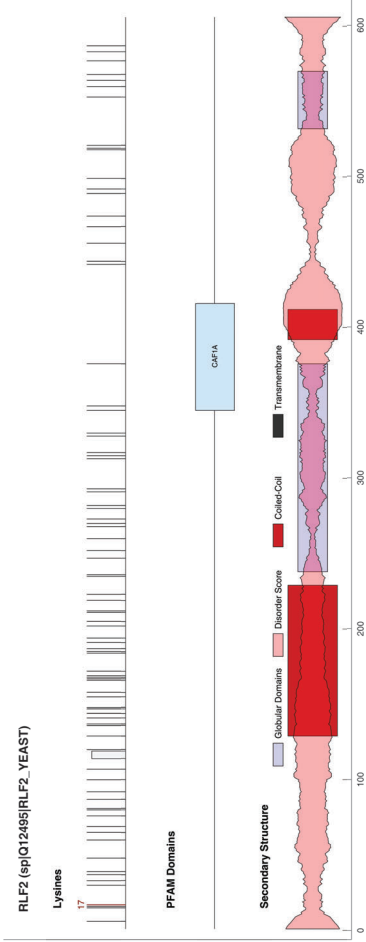
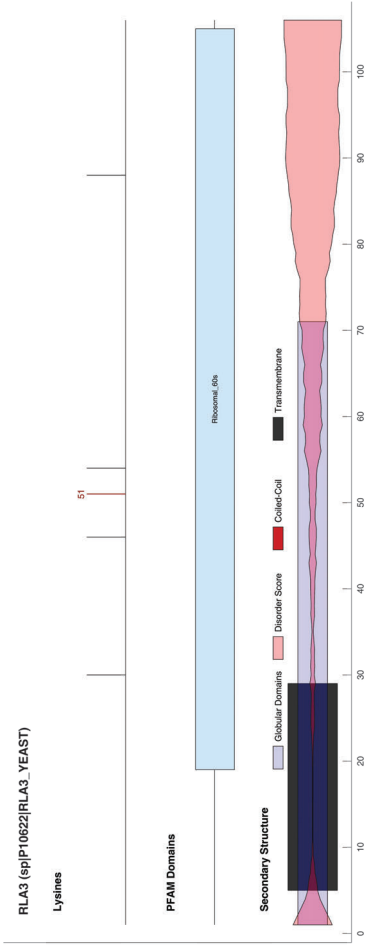


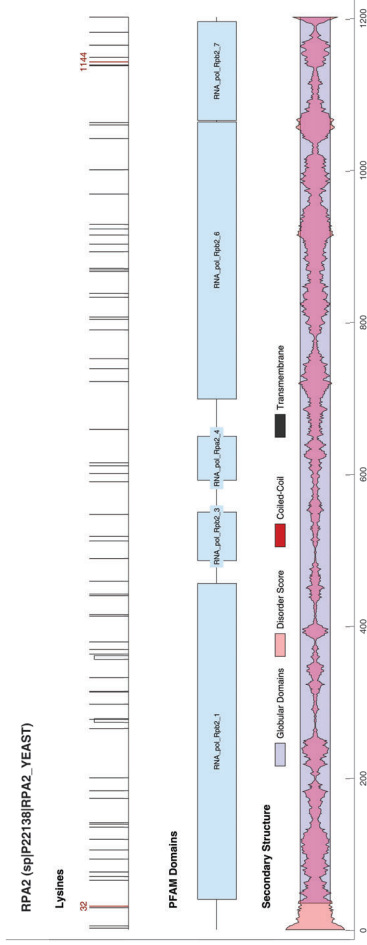
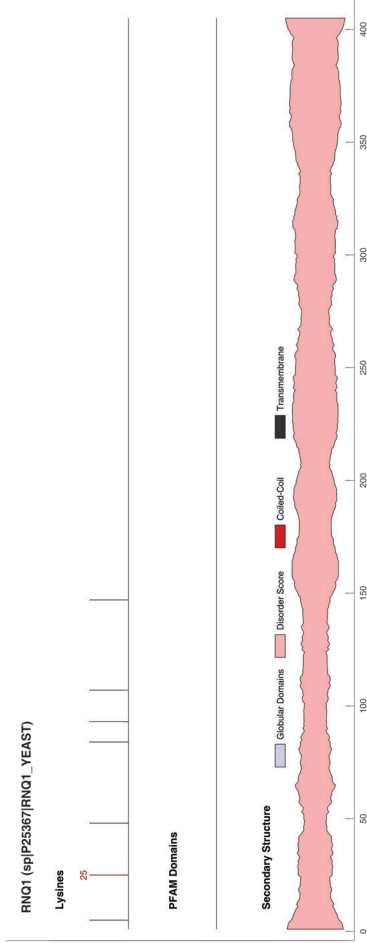
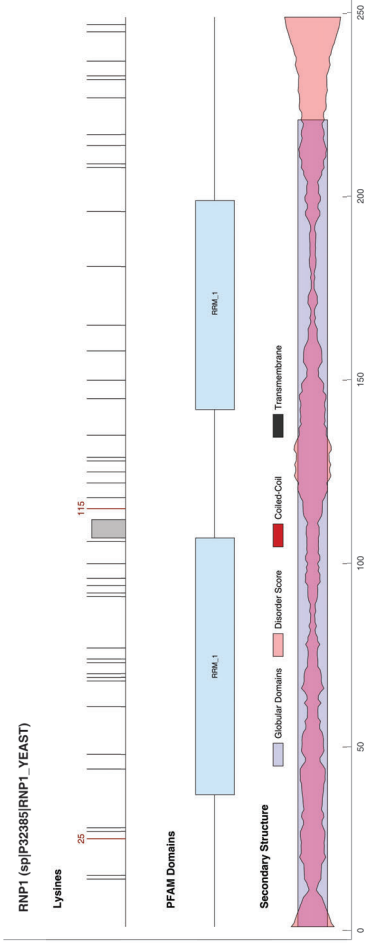
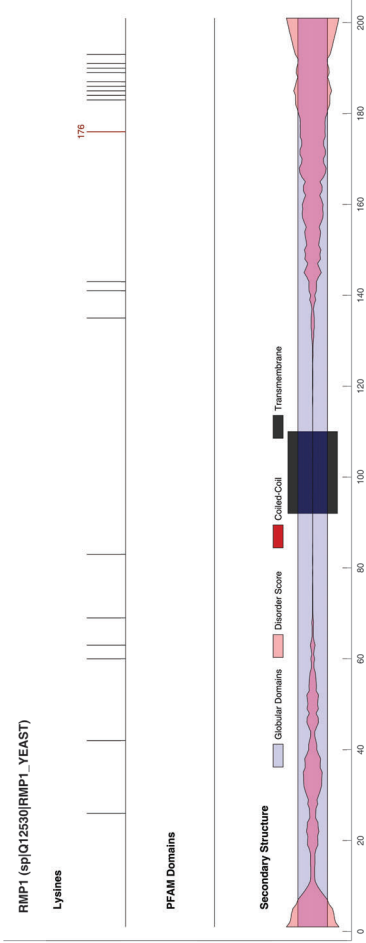
RL402 (sp|P0CH09|RL402_YEAST)

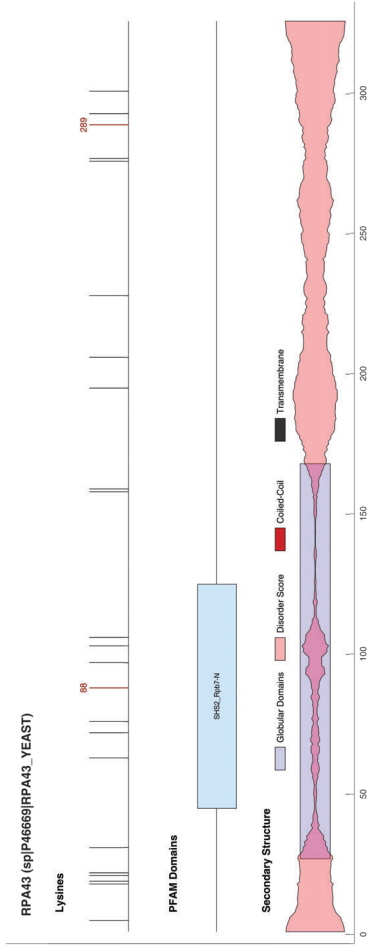
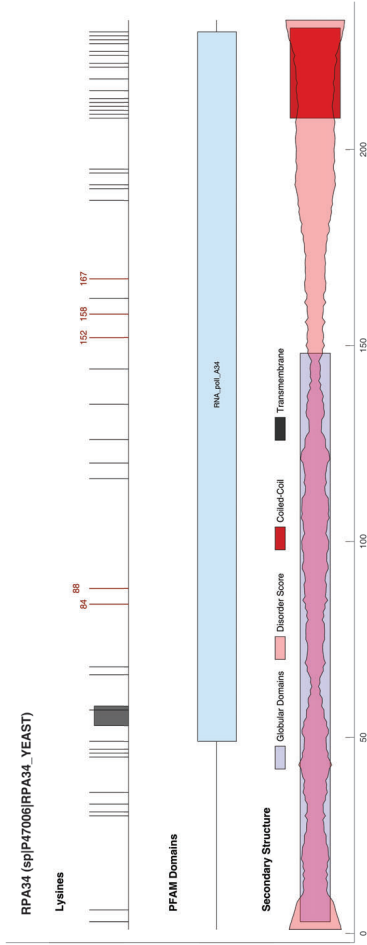
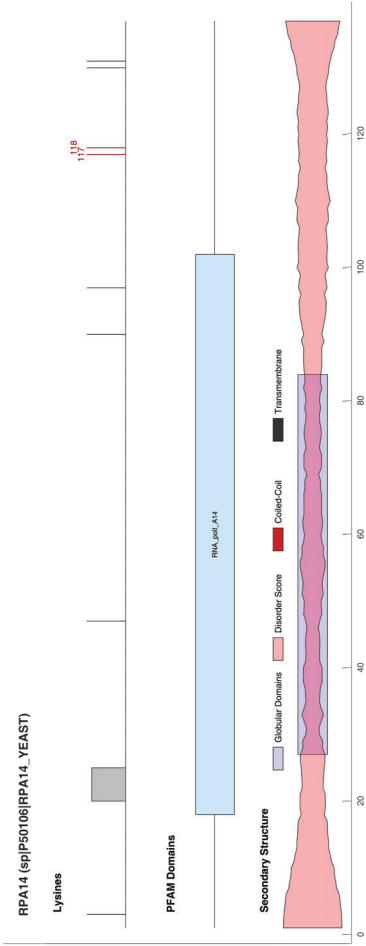
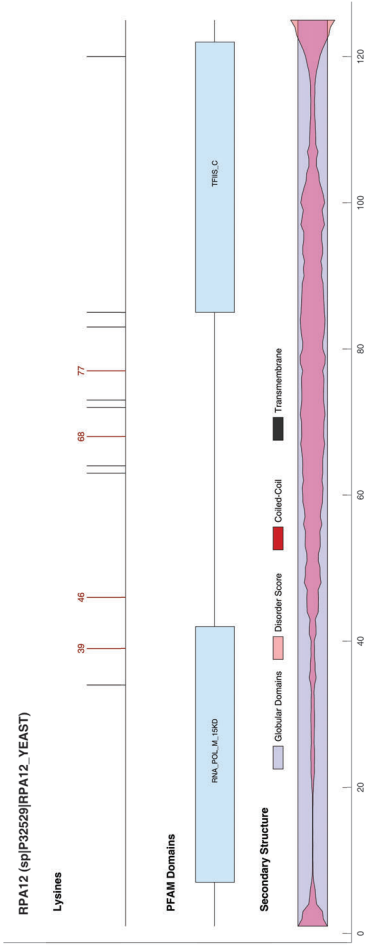


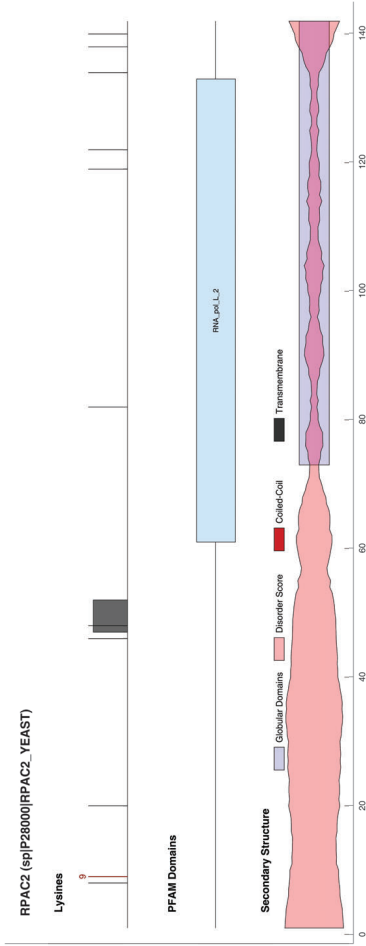
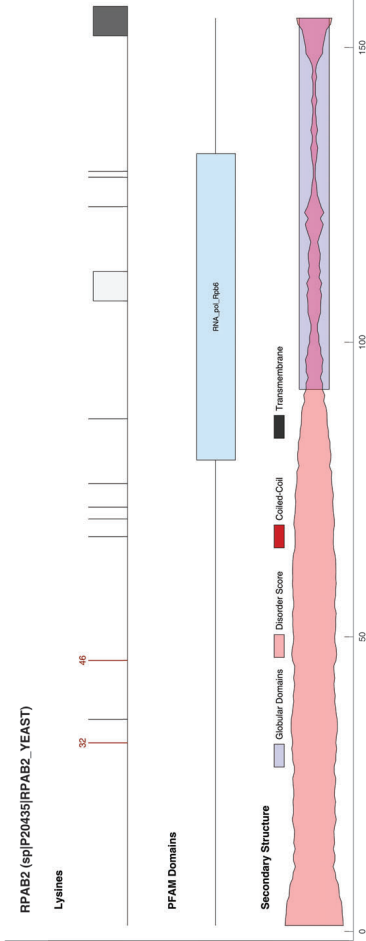
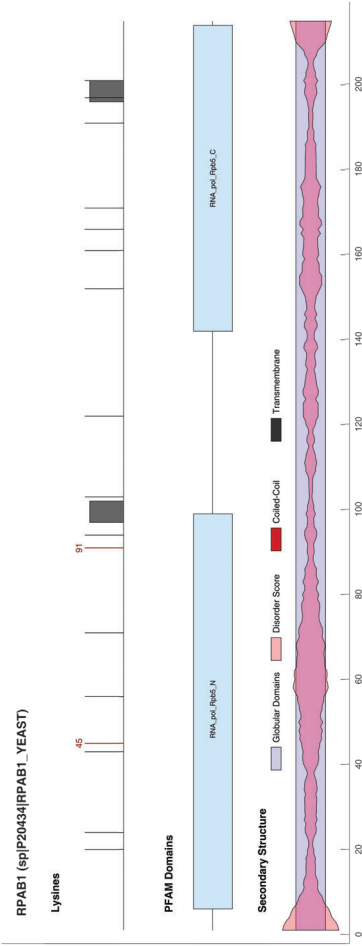
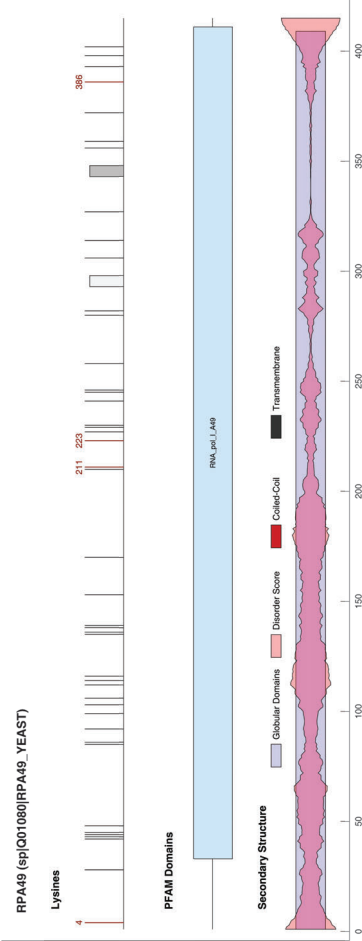
RLA2 (sp|P05319|RLA2_YEAST)



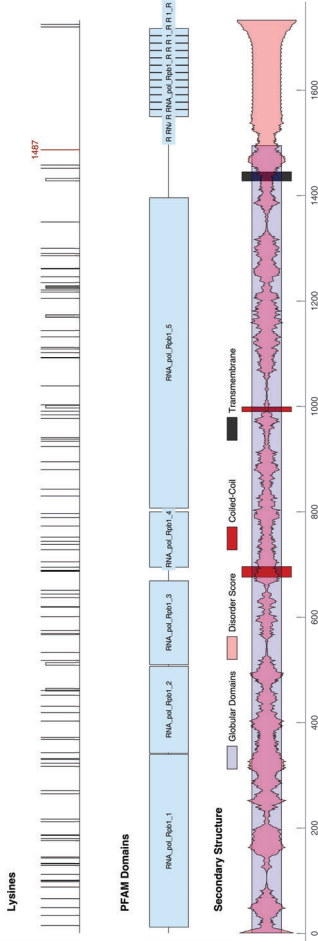




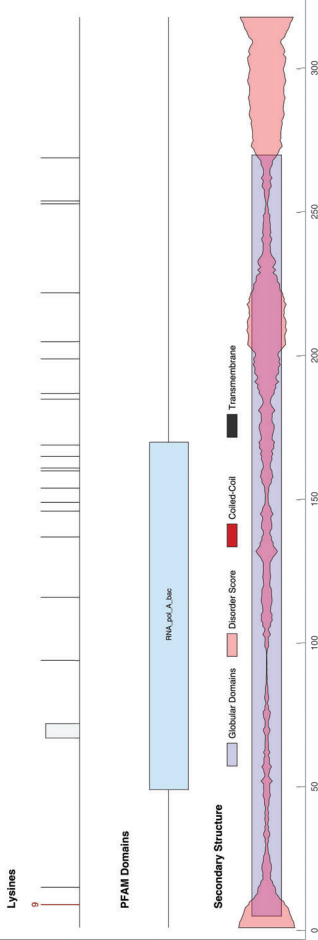




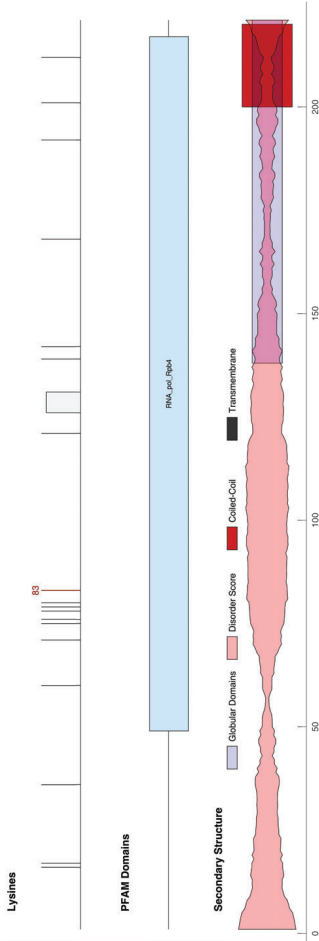
RPB1 (sp|P04050|RPB1_YEAST)



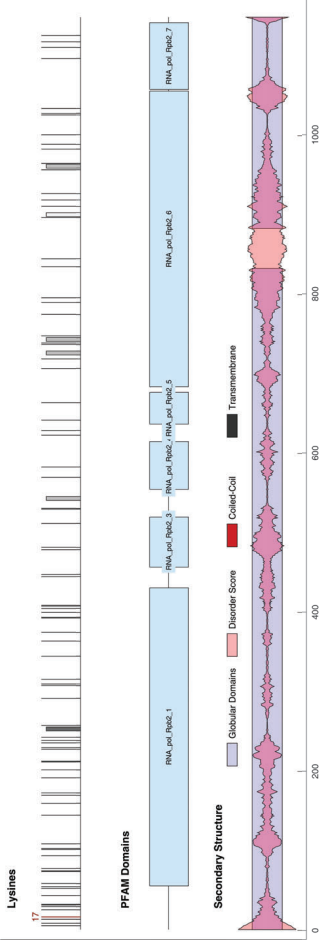
RPB3 (sp|P16370|RPB3_YEAST)



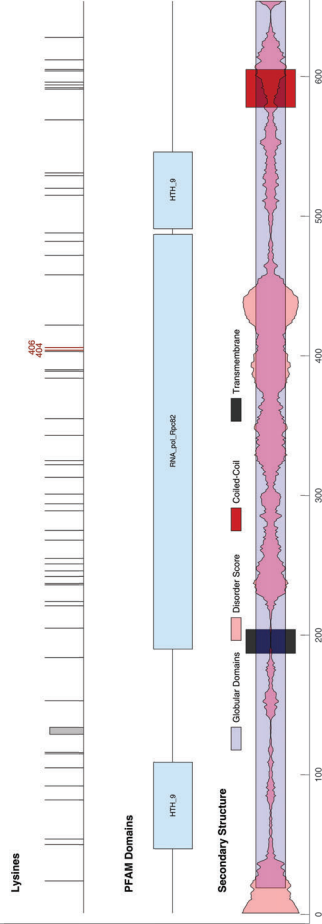
RPB4 (sp|P20433|RPB4_YEAST)



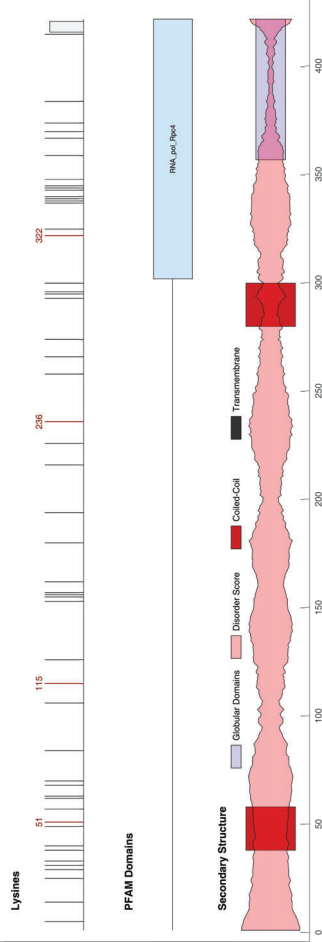
RPC2 (sp|P22276|RPC2_YEAST)



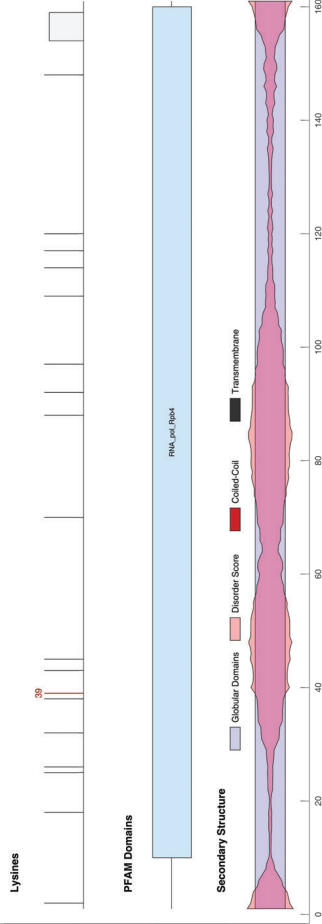
RPC3 (sp|P32349|RPC3_YEAST)



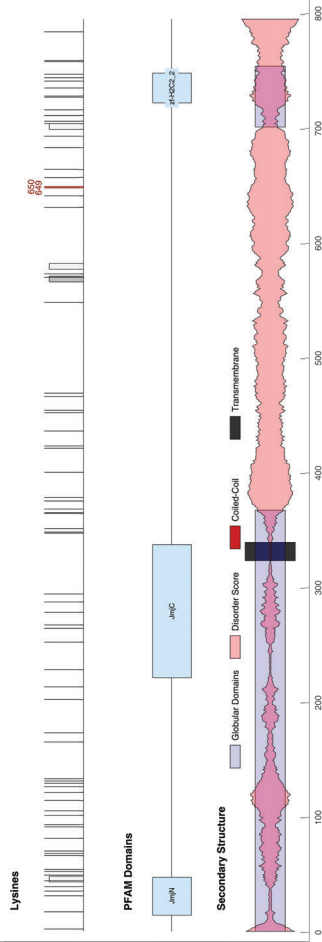
RPC4 (sp|P25441|RPC4_YEAST)



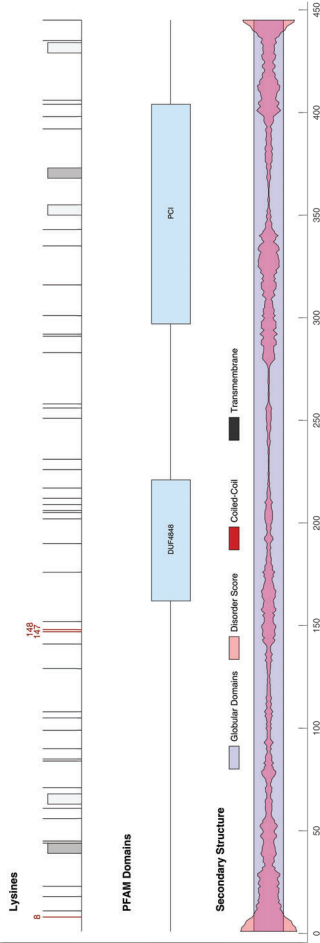
RPC9 (sp|P47076|RPC9_YEAST)



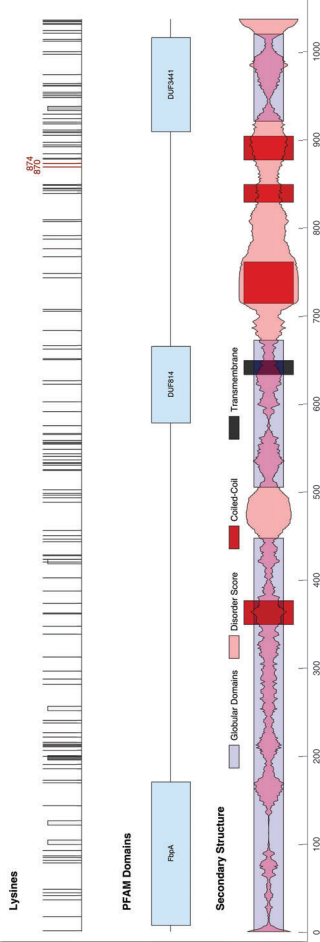
RPH1 (sp|P39556|RPH1_YEAST)



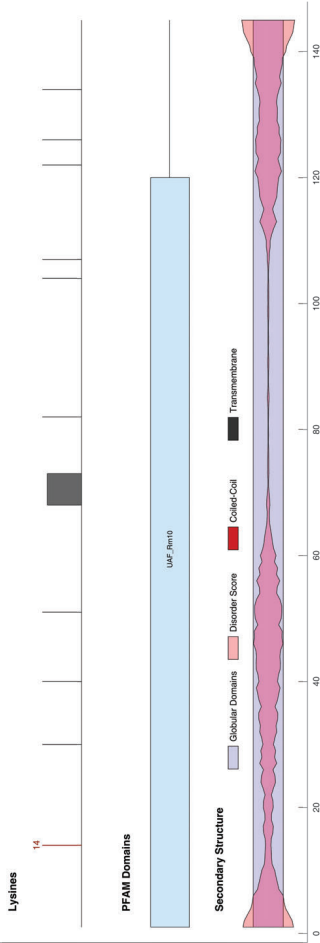
RPN5 (sp|Q1250|RPN5_YEAST)



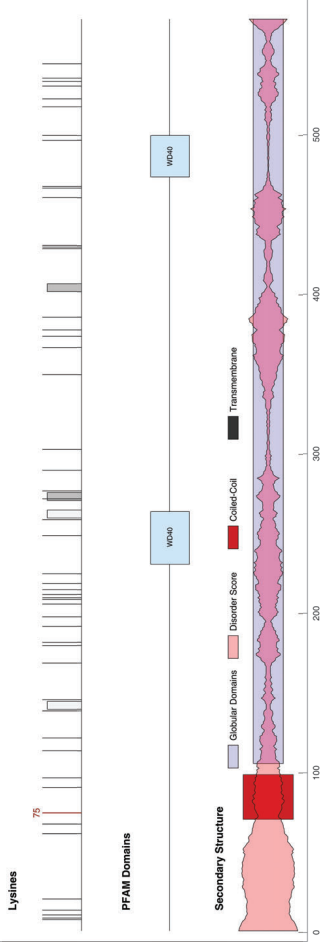
RQC2 (sp|Q12532|RQC2_YEAST)

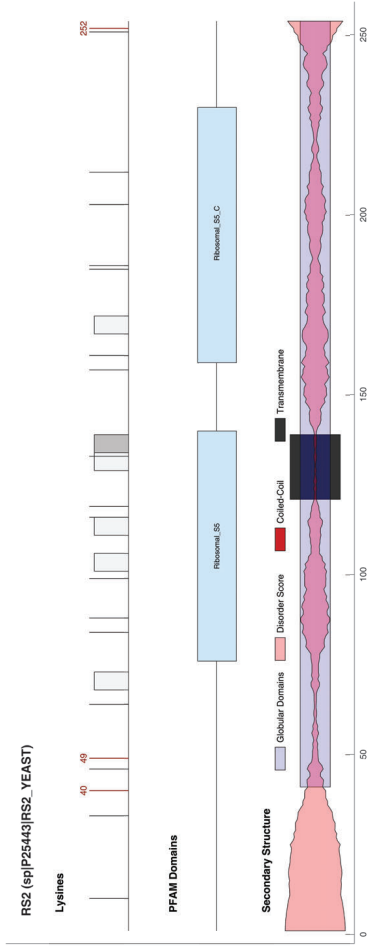
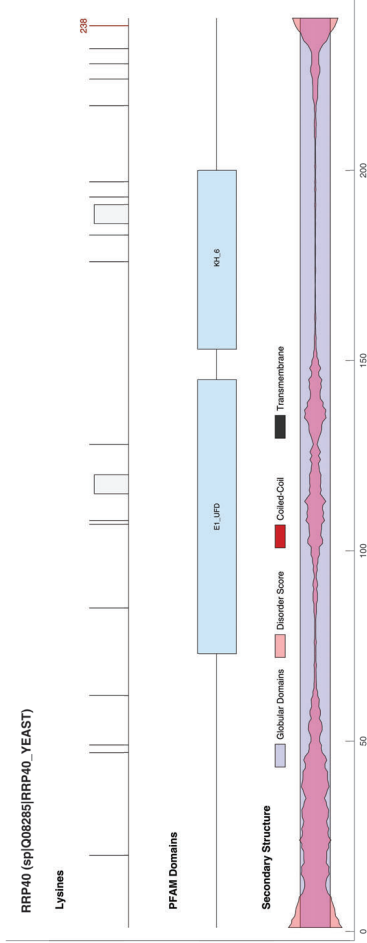
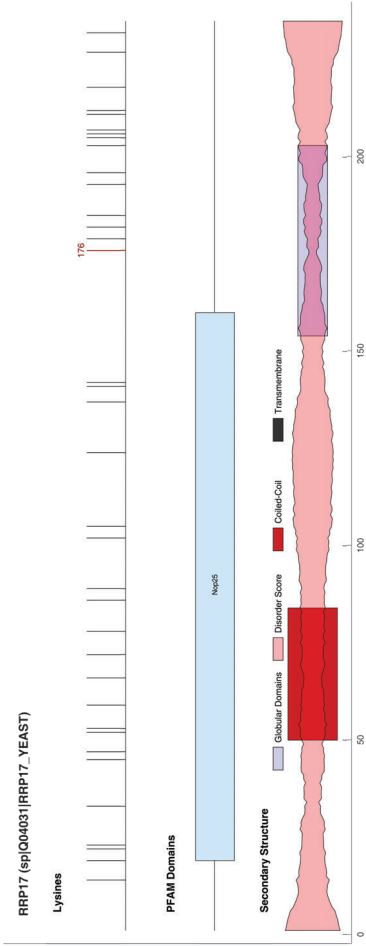
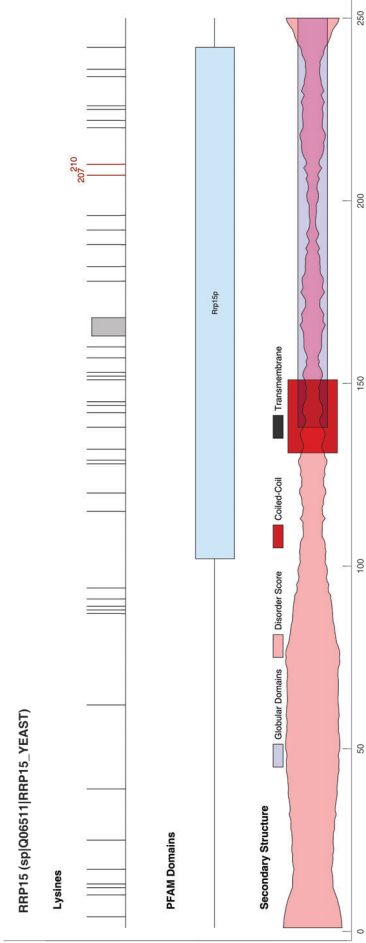


RRM10 (sp|P39204|RRM10_YEAST)

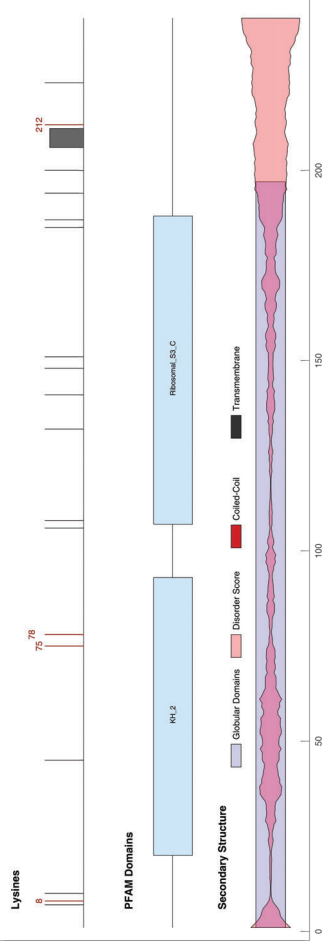


RRP9 (sp|Q06506|RRP9_YEAST)

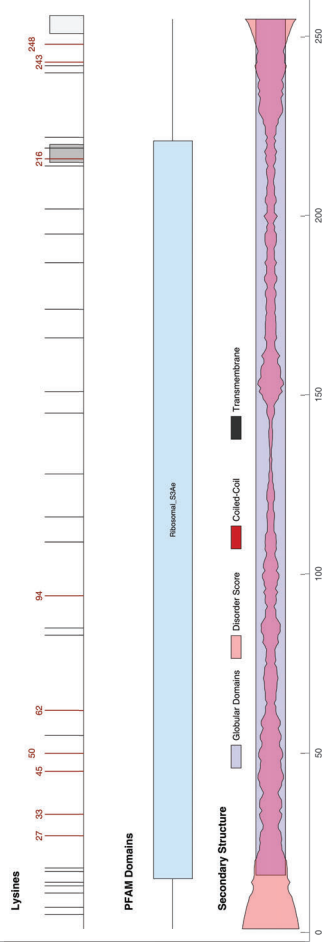




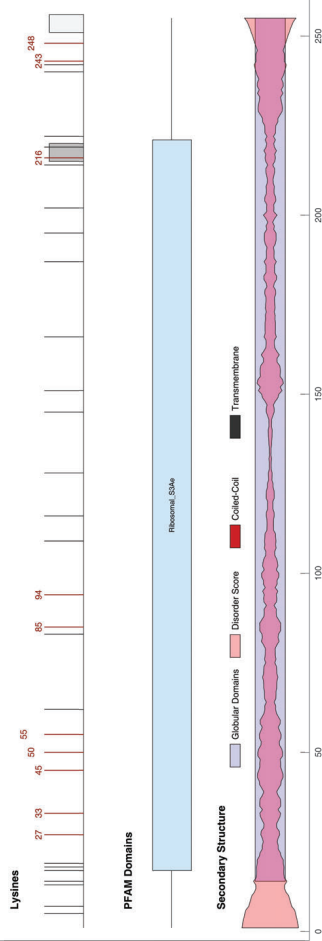
RS3 (sp|P05750|RS3_YEAST)



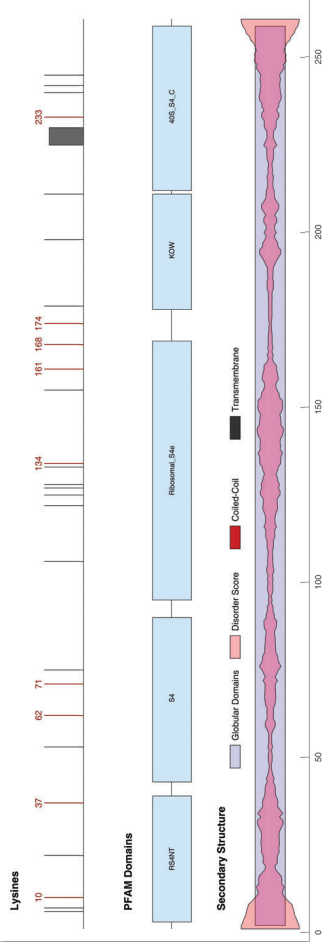
RS3A1 (sp|P33442|RS3A1_YEAST)

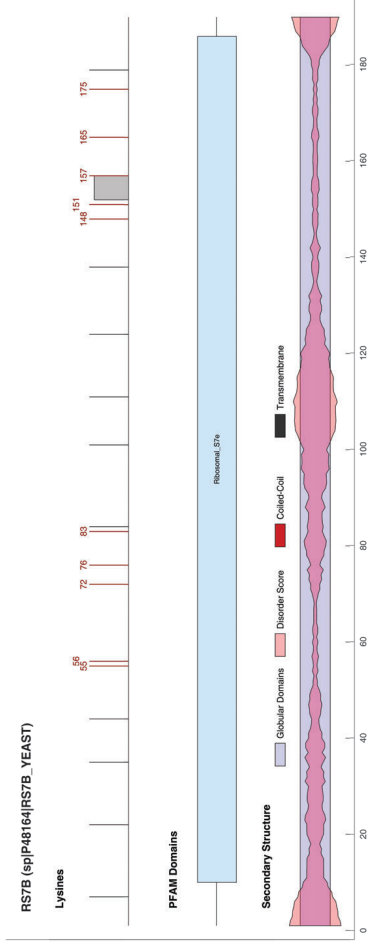
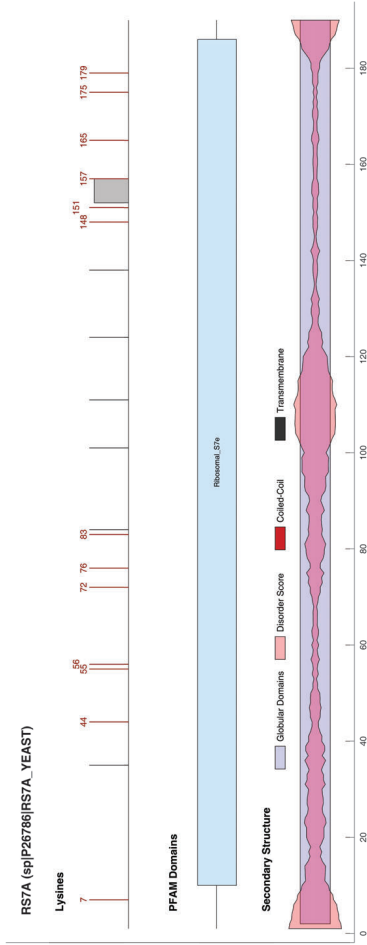
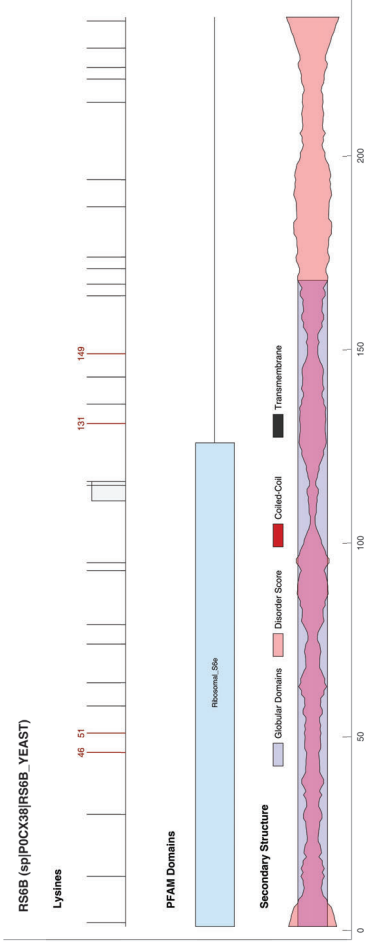
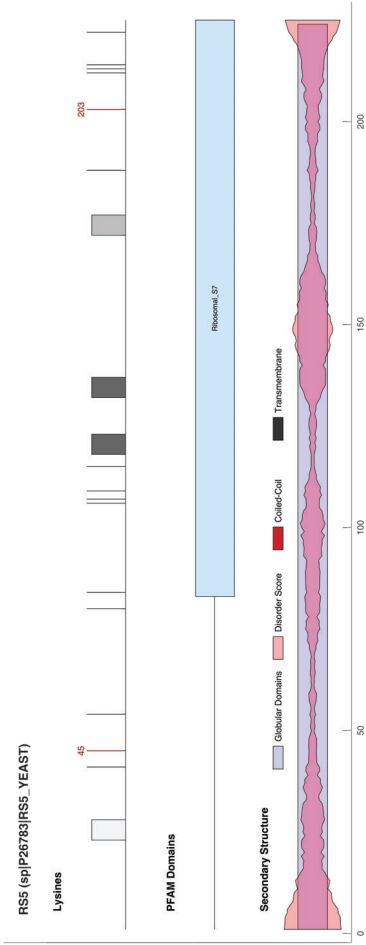


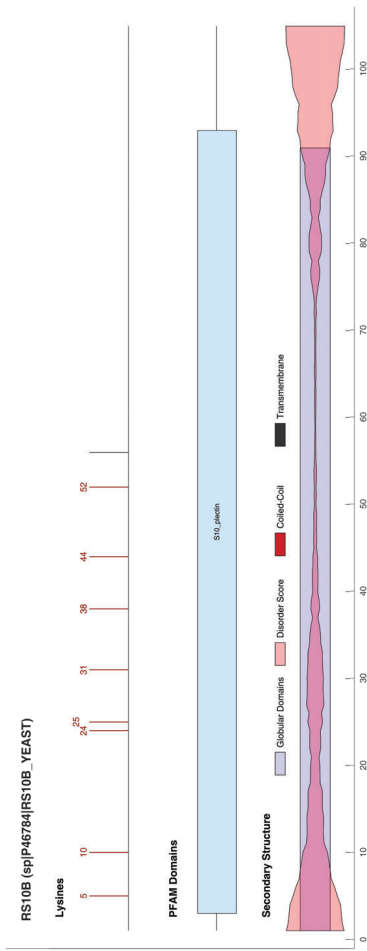
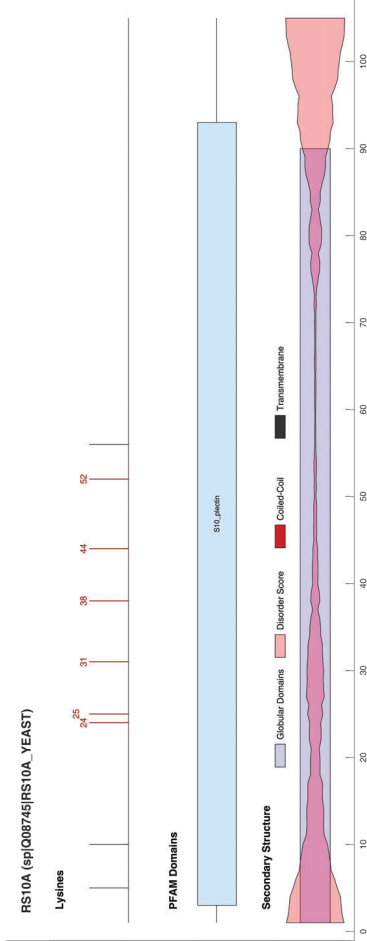
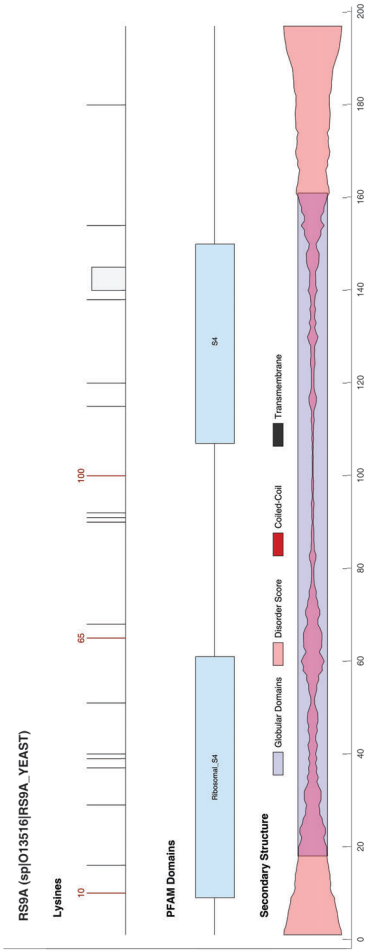
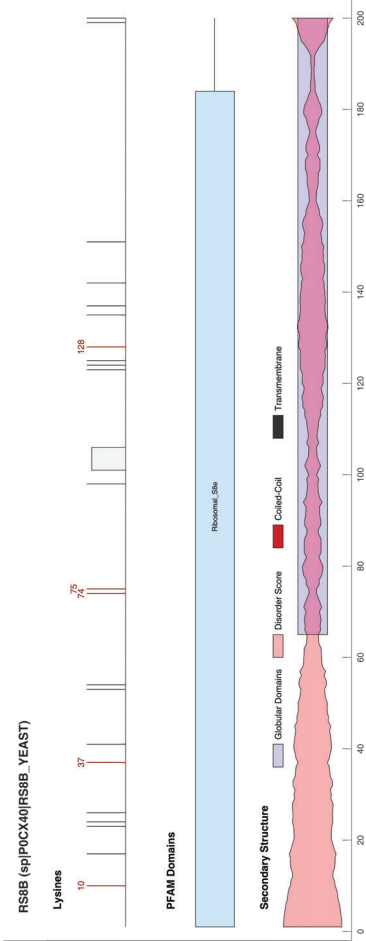
RS3A2 (sp|P23248|RS3A2_YEAST)

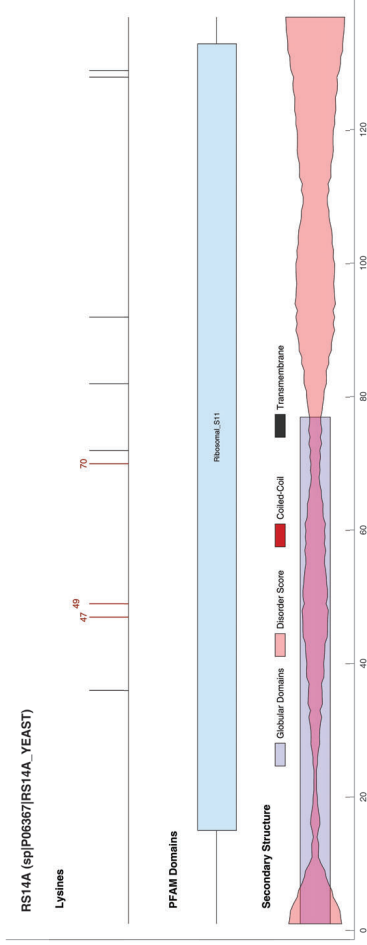
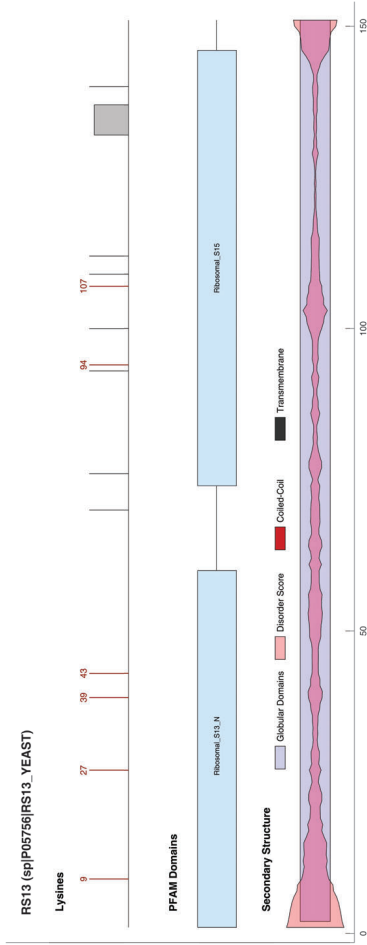
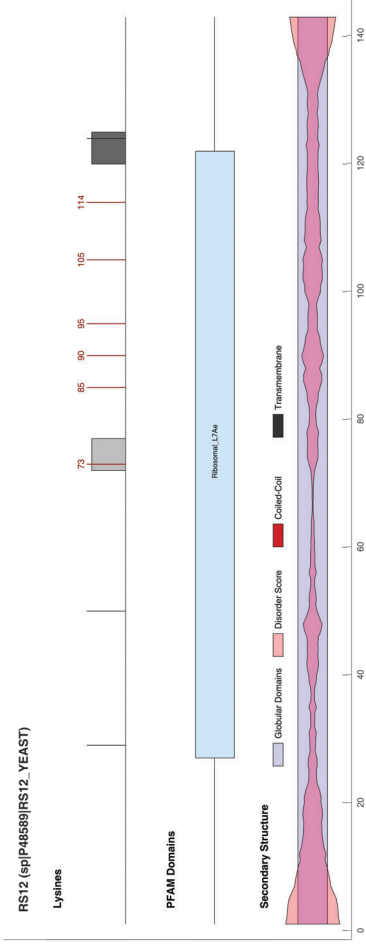
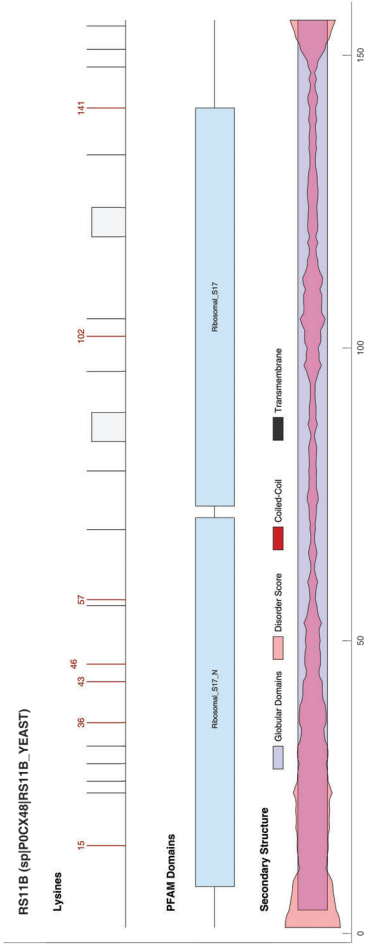


RS4B (sp|P0CX38|RS4B_YEAST)

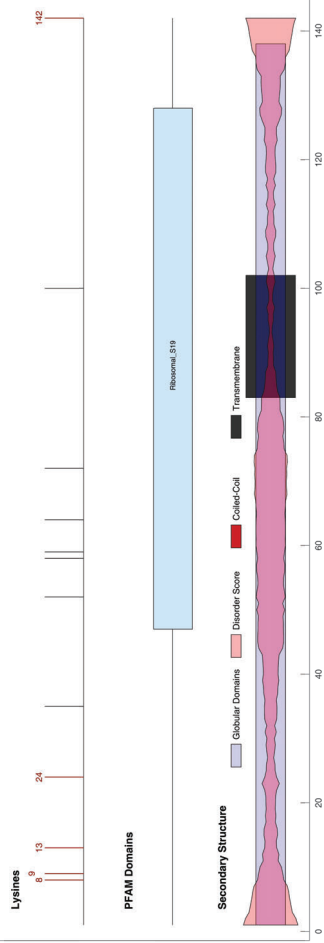




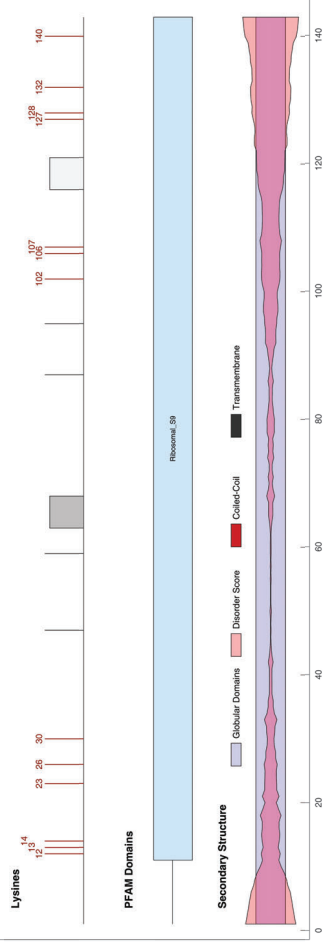




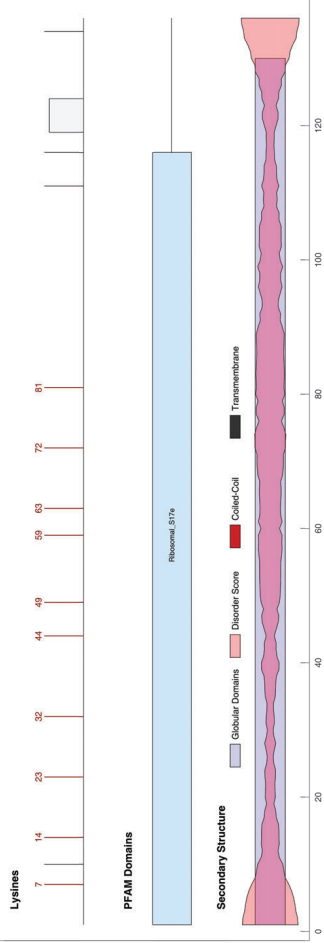
RS15 (sp|Q01855|RS15_YEAST)



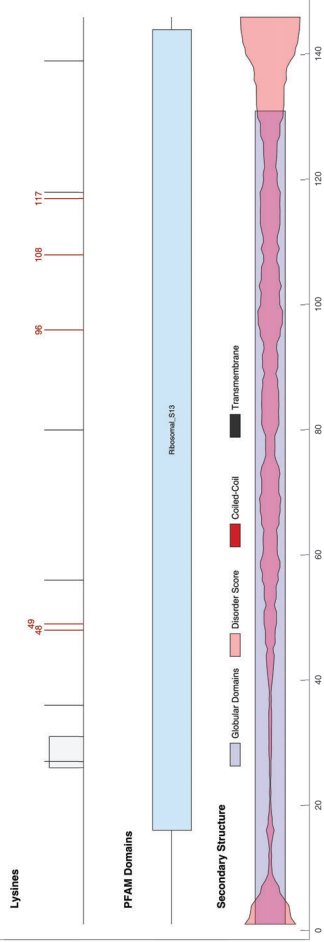
RS16E (sp|P0CX32|RS16E_YEAST)

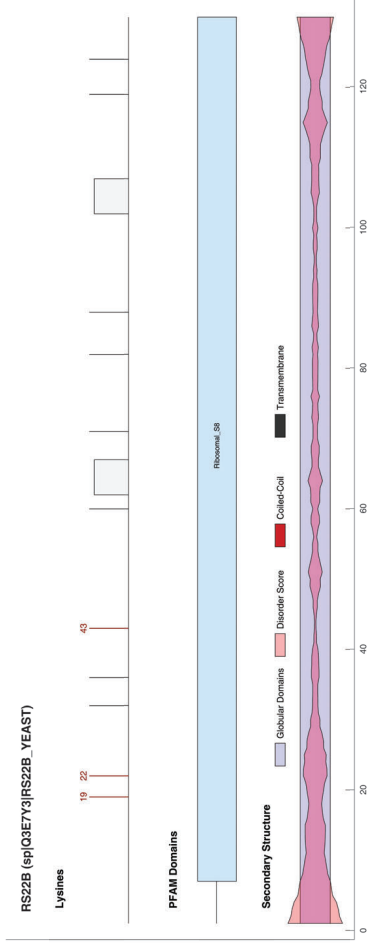
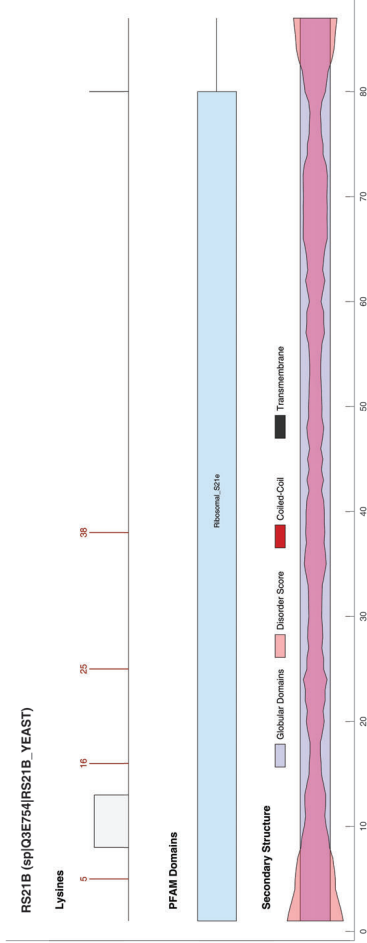
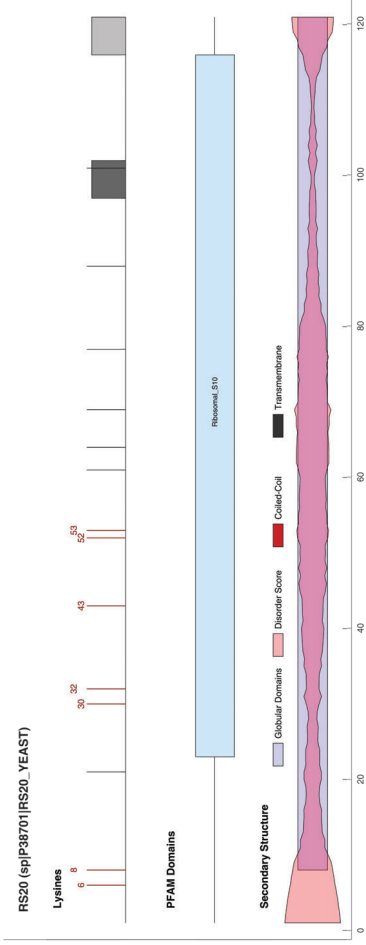
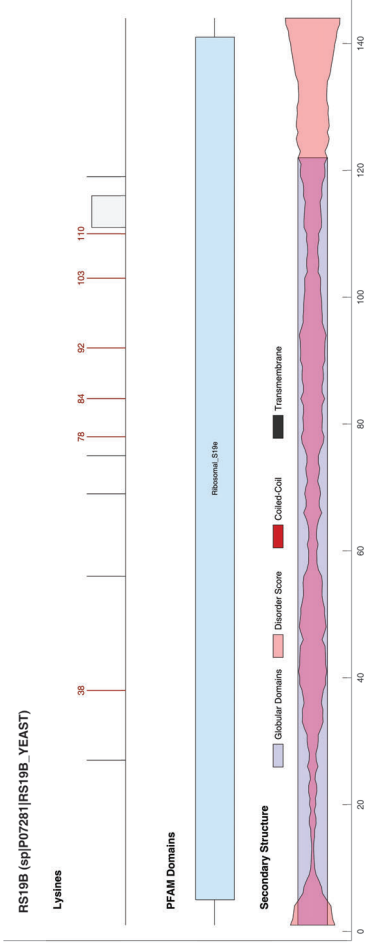


RS7B (sp|P14127|RS7B_YEAST)

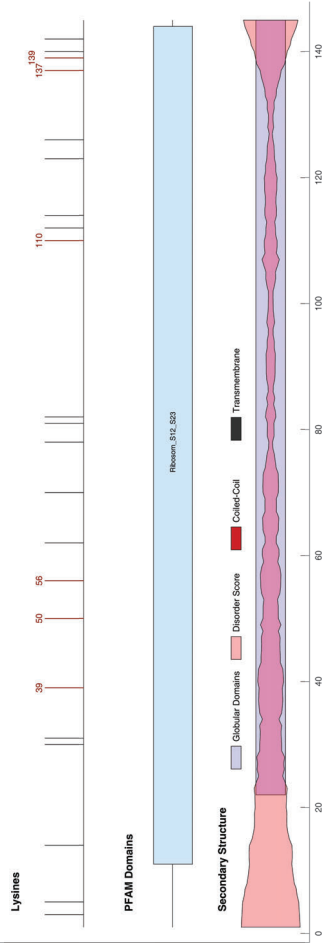


RS18E (sp|P0CX36|RS18E_YEAST)

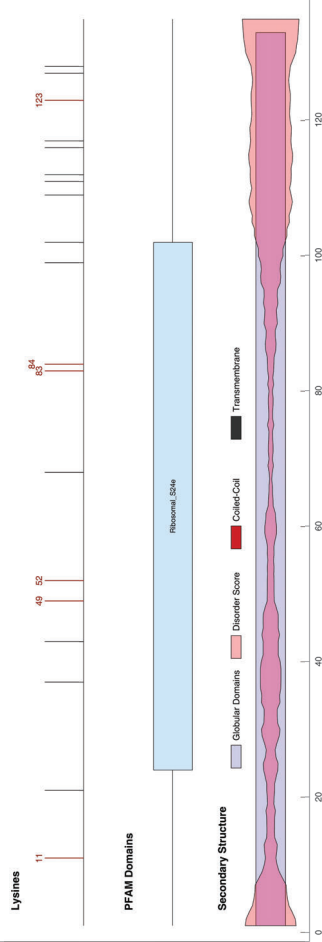




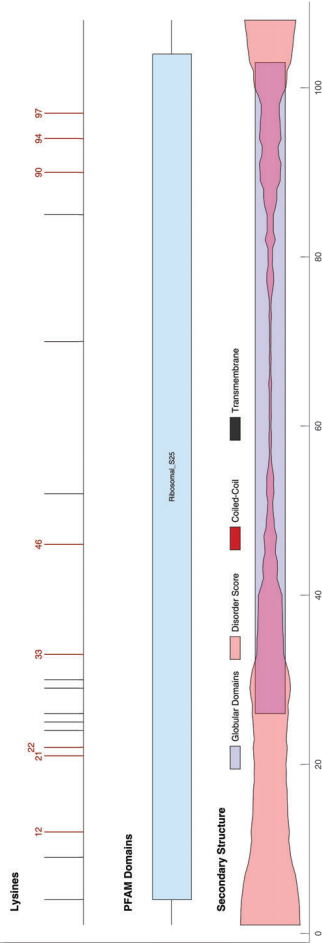
RS23B (sp|P0CX30|RS23B_YEAST)



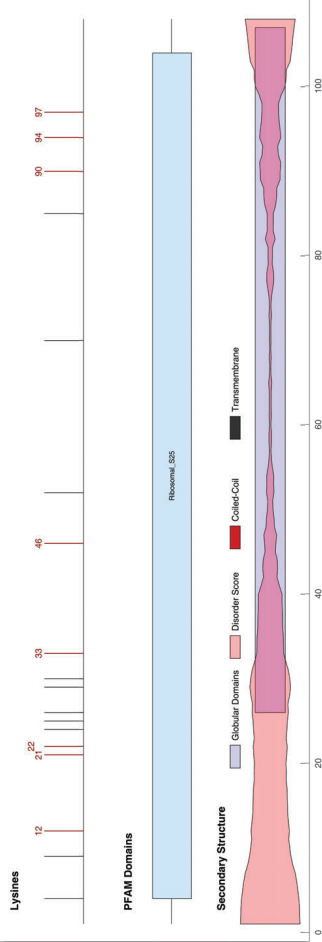
RS24B (sp|P0CX32|RS24B_YEAST)

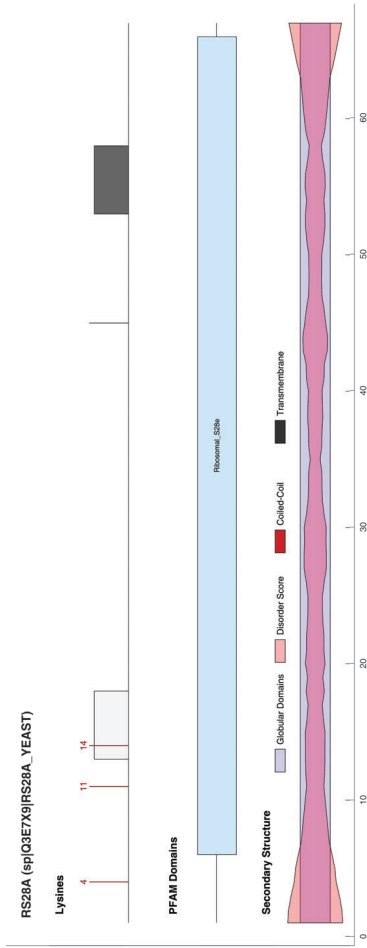
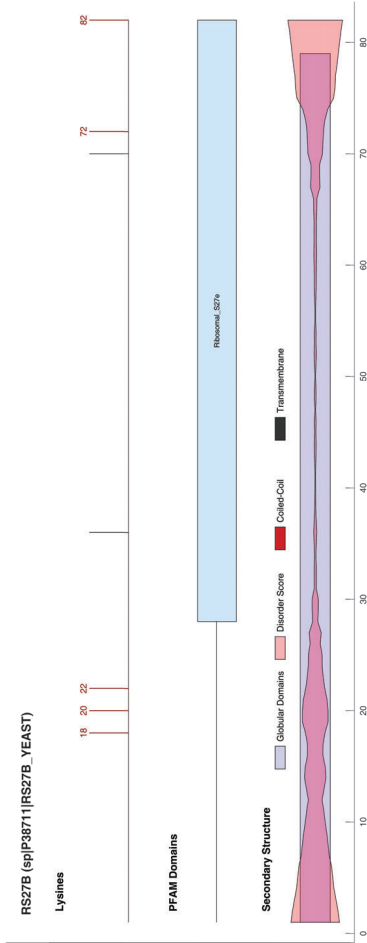
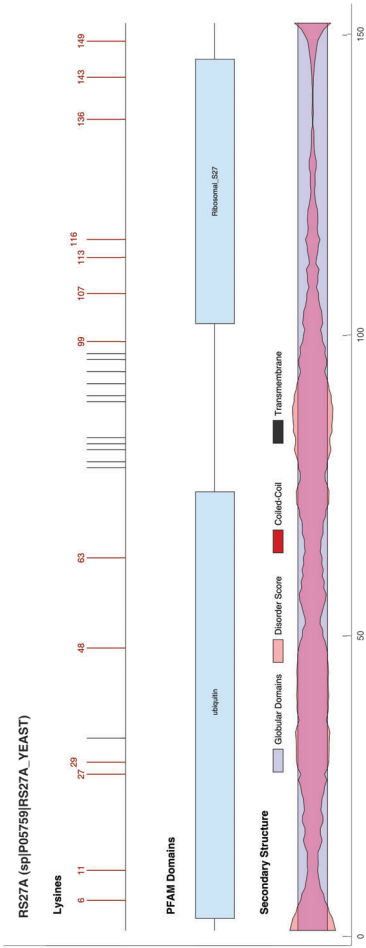
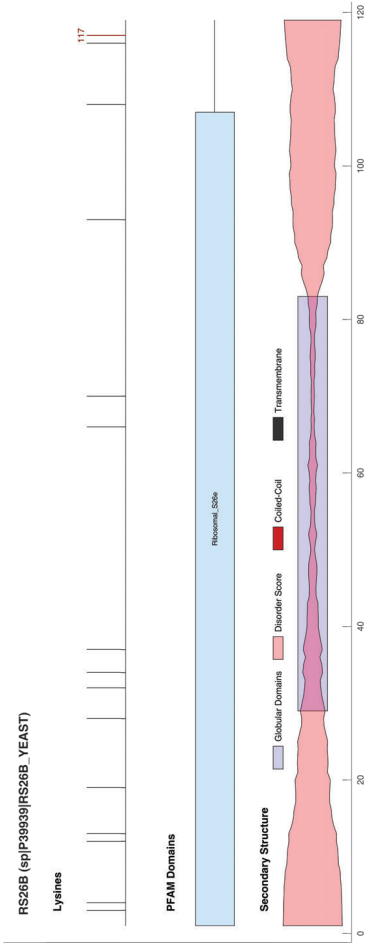


RS25A (sp|Q3E792|RS25A_YEAST)

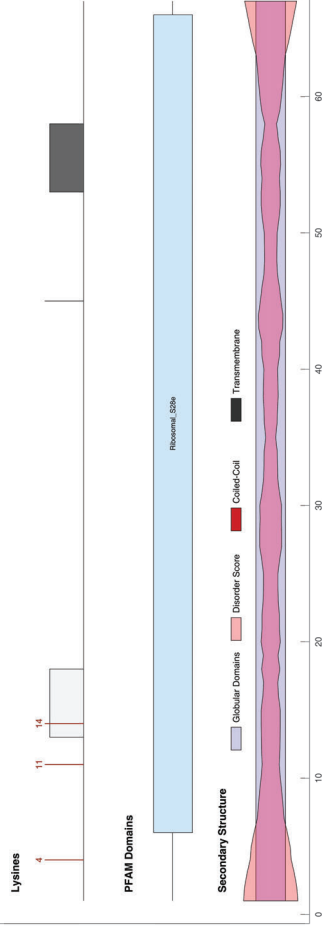


RS25B (sp|P0C074|RS25B_YEAST)

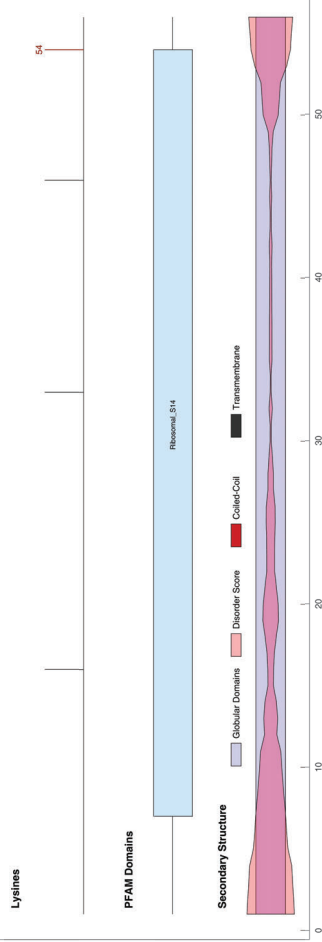




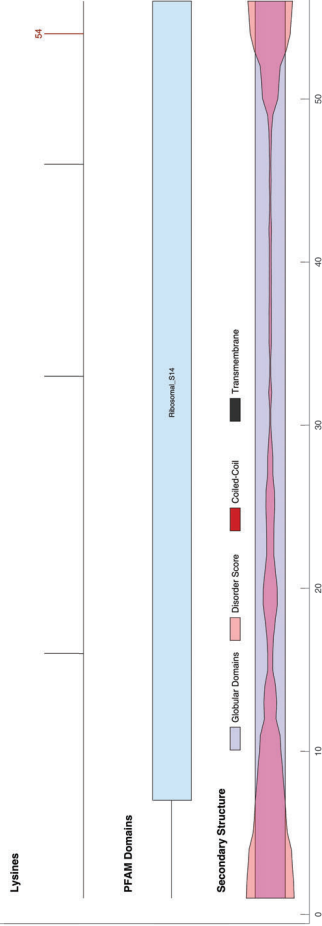
RS28E (sp|P0C0X0|RS28E_YEAST)



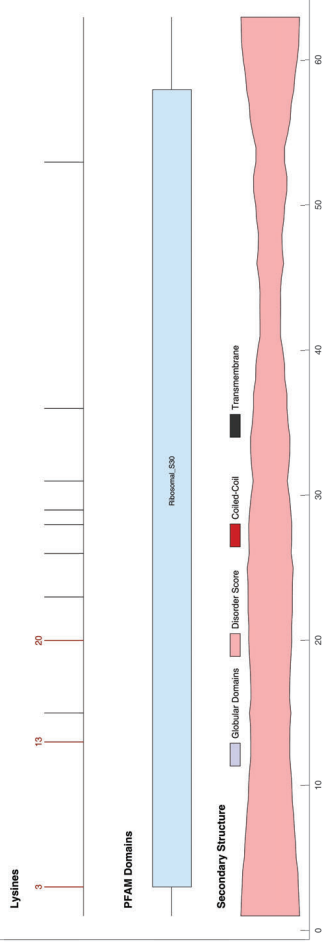
RS29A (sp|P41057|RS29A_YEAST)

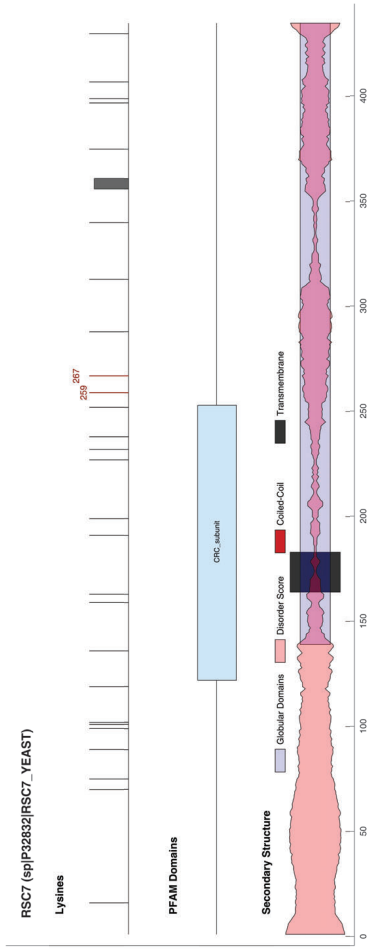
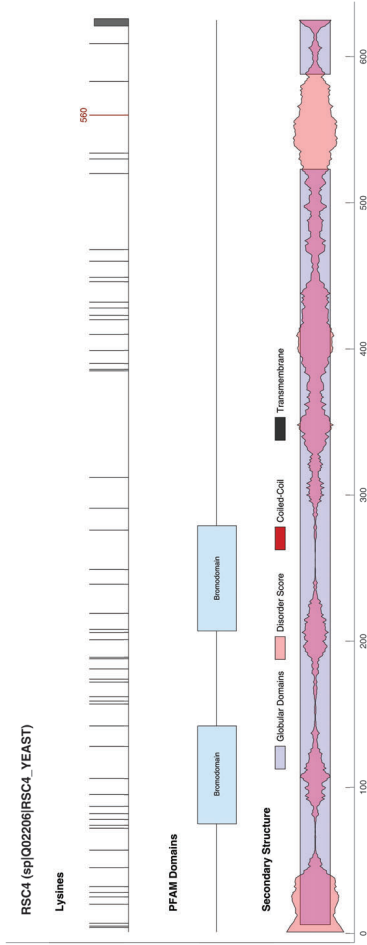
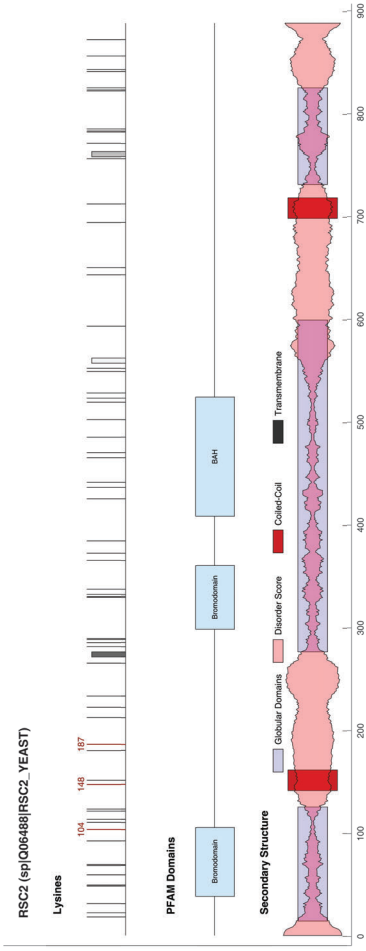
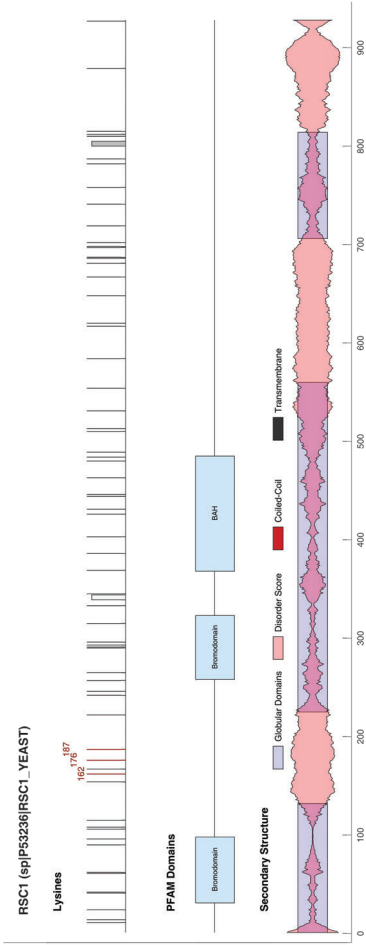


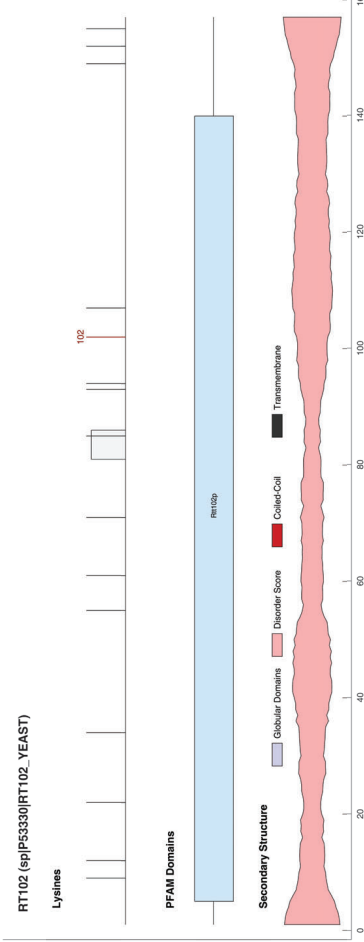
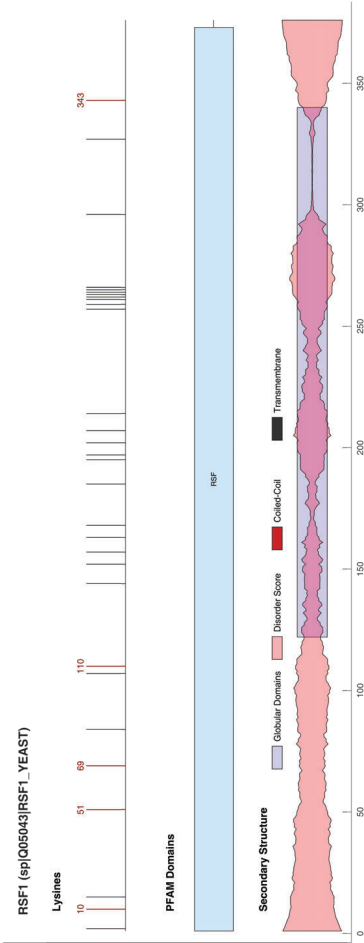
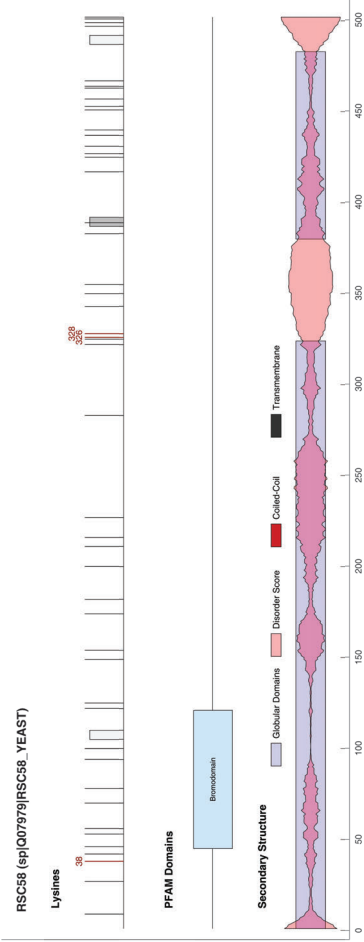
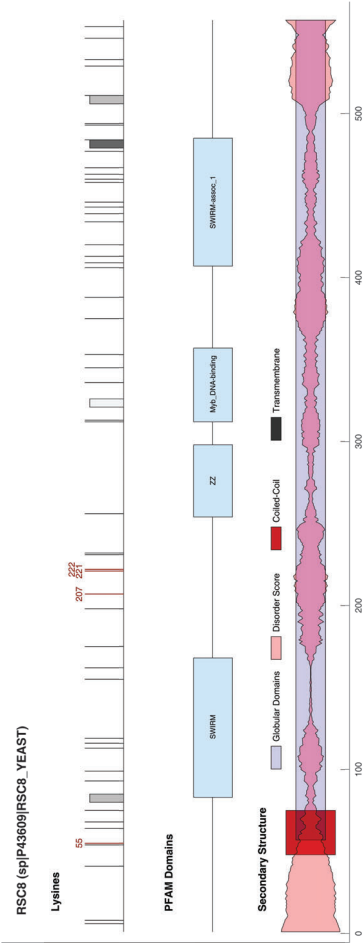
RS29E (sp|P41058|RS29E_YEAST)

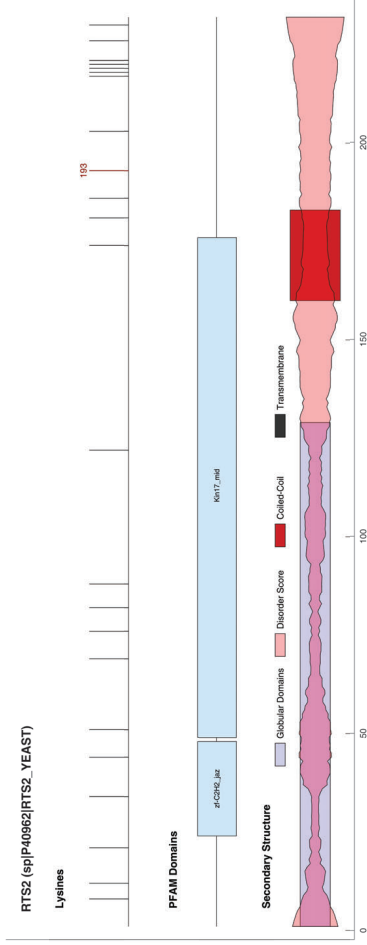
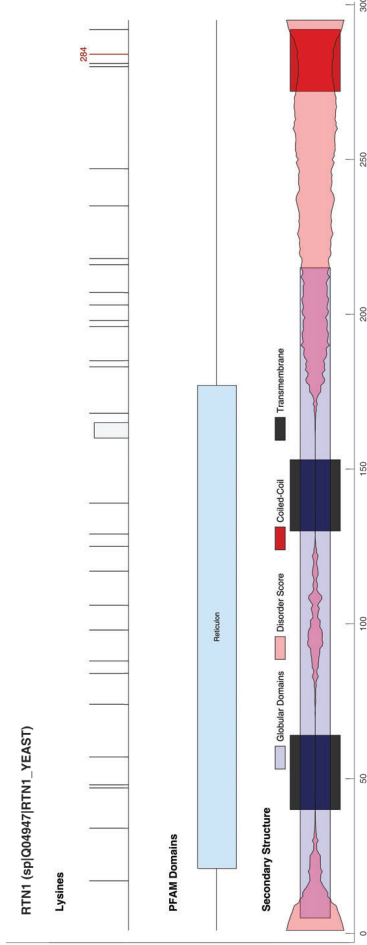
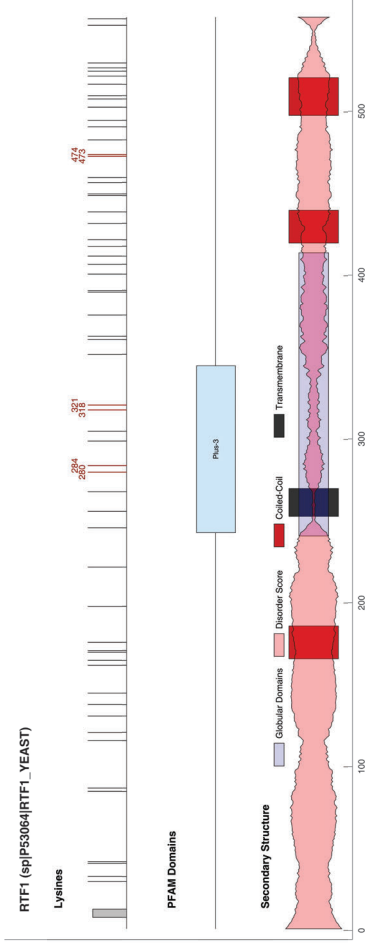
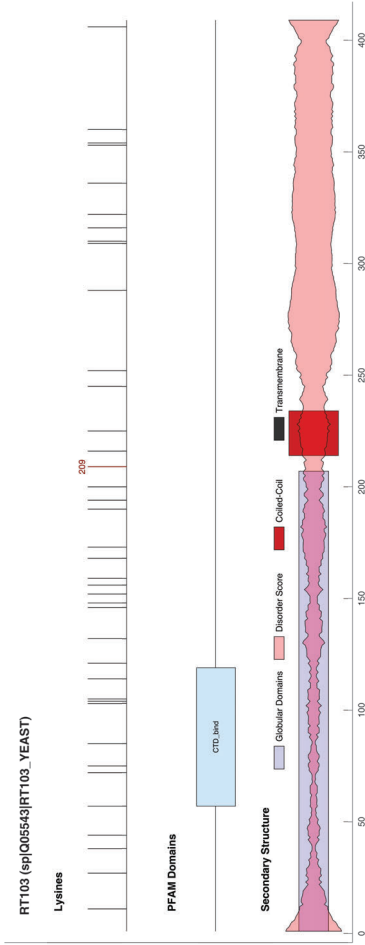


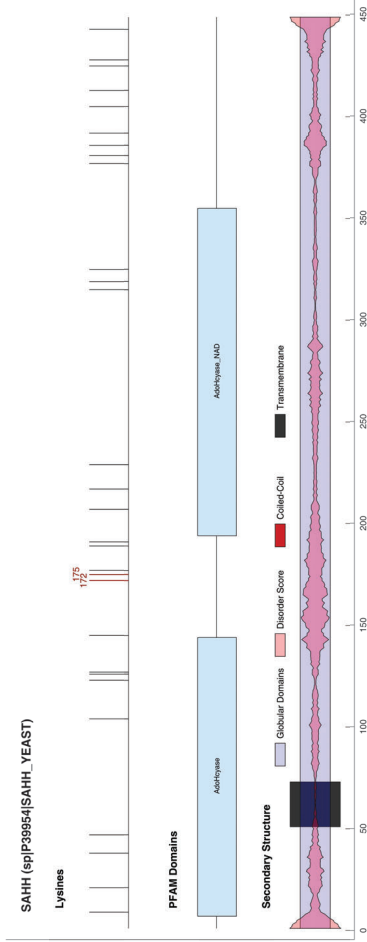
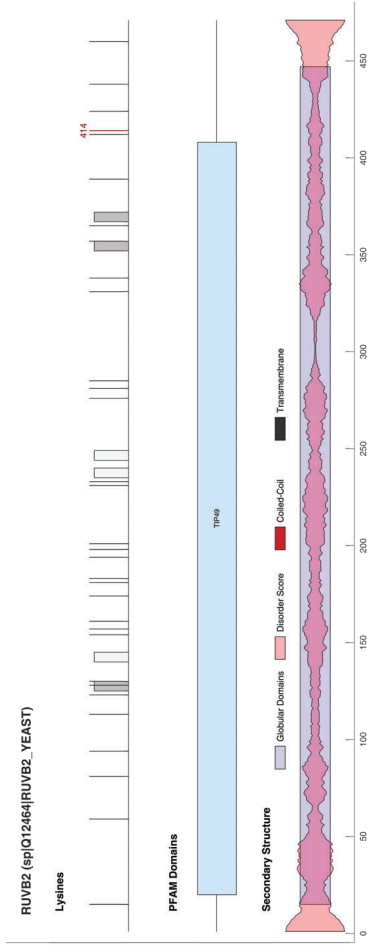
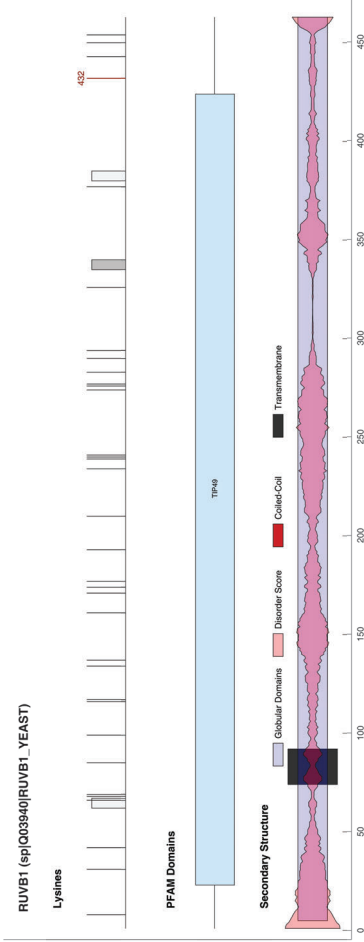
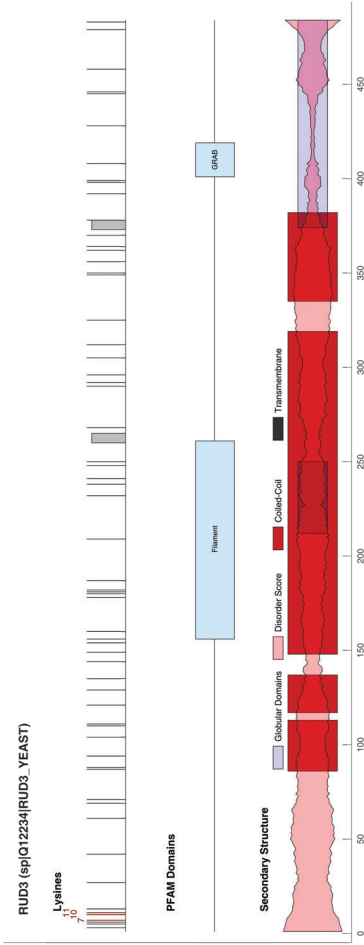
RS30E (sp|P0CX34|RS30E_YEAST)









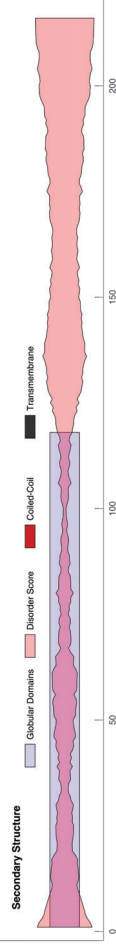


SBA1 (sp|P28707|SBA1_YEAST)

Lysines



PFAM Domains

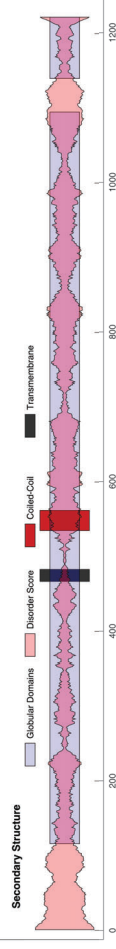


SC160 (sp|P06105|SC160_YEAST)

Lysines



PFAM Domains

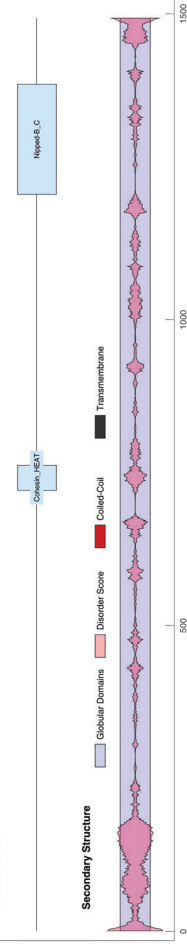


SCC2 (sp|Q04002|SCC2_YEAST)

Lysines



PFAM Domains

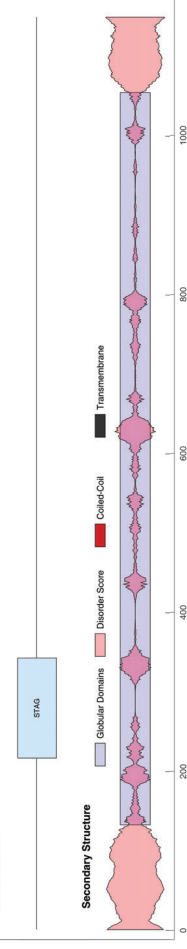


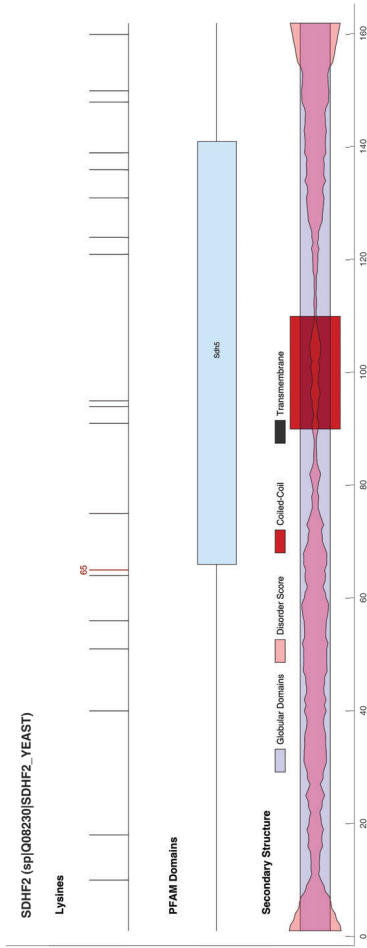
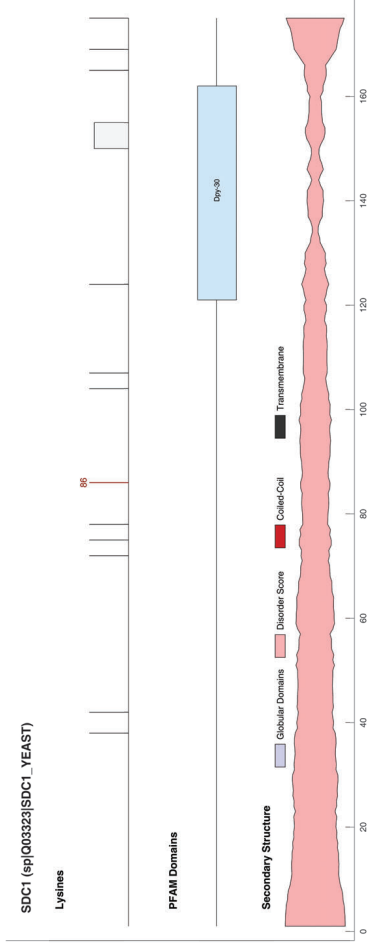
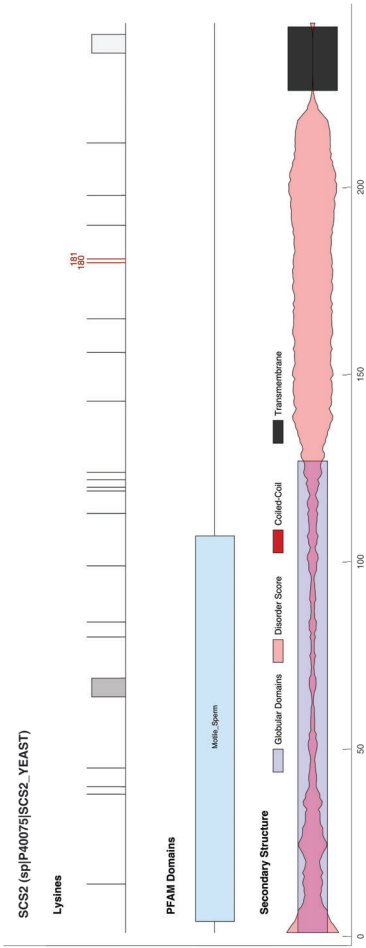
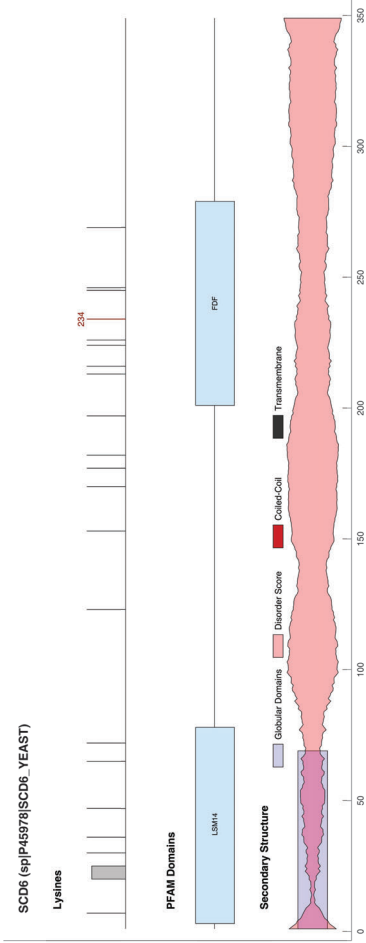
SCC3 (sp|P40541|SCC3_YEAST)

Lysines

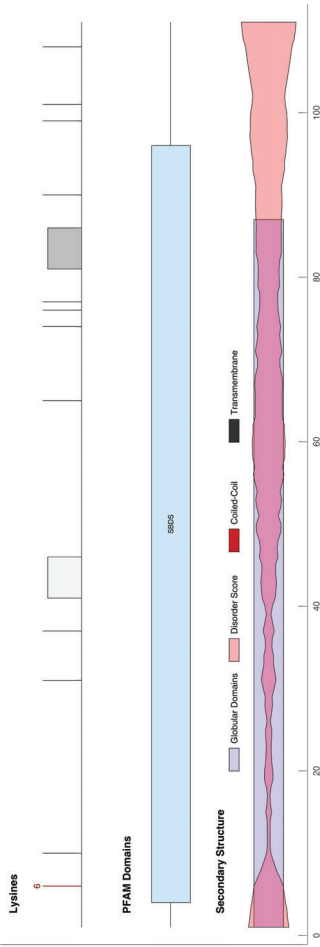


PFAM Domains

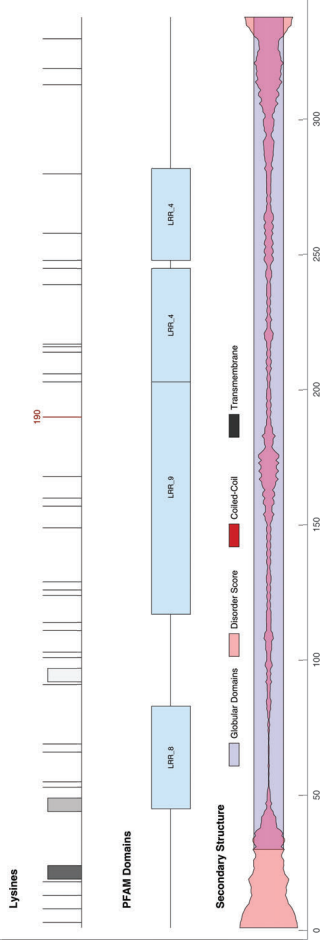




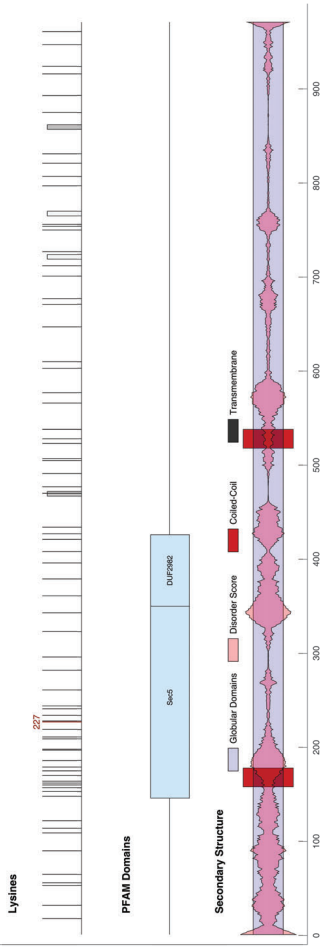
SDO1L (sp|P3804|SDO1L_YEAST)



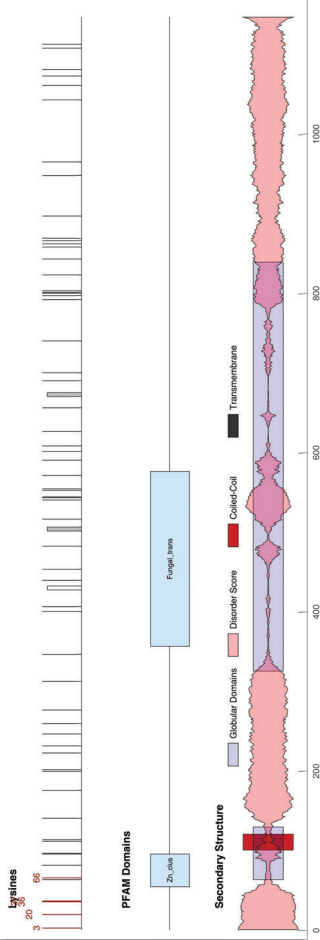
SDS22 (sp|P36047|SDS22_YEAST)

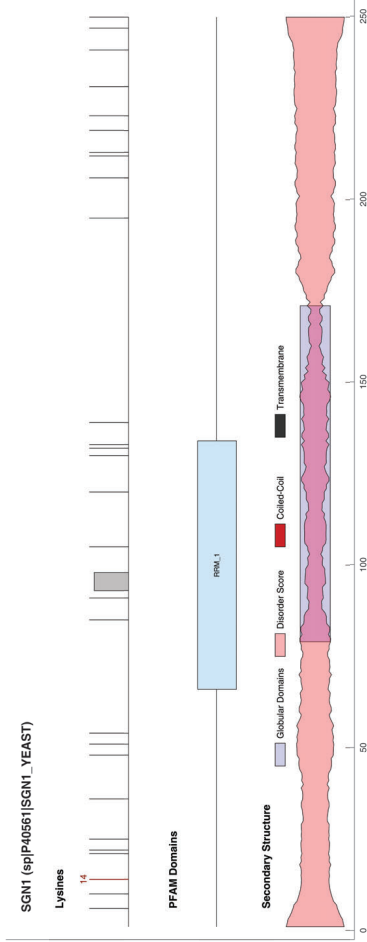
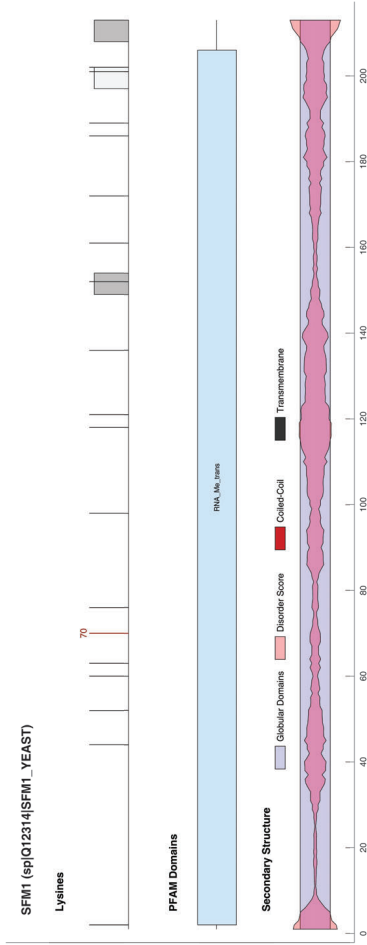
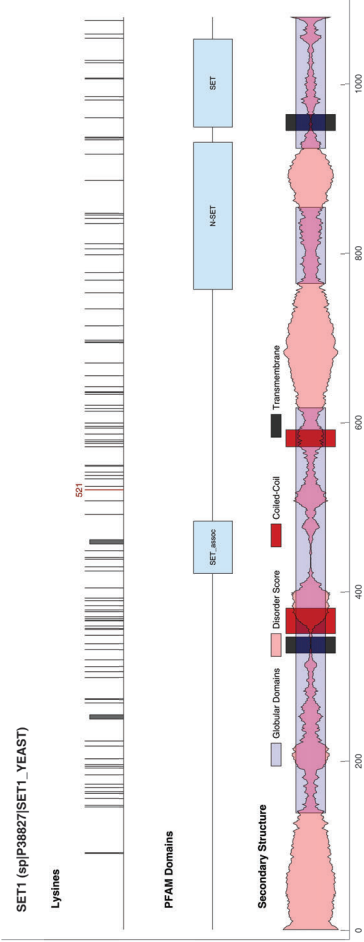
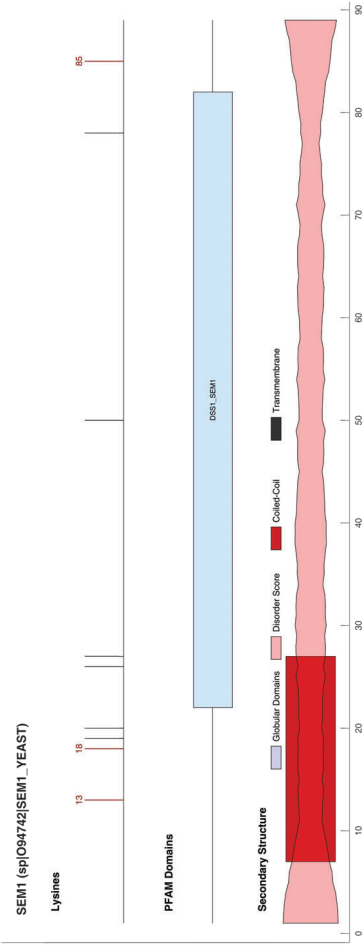


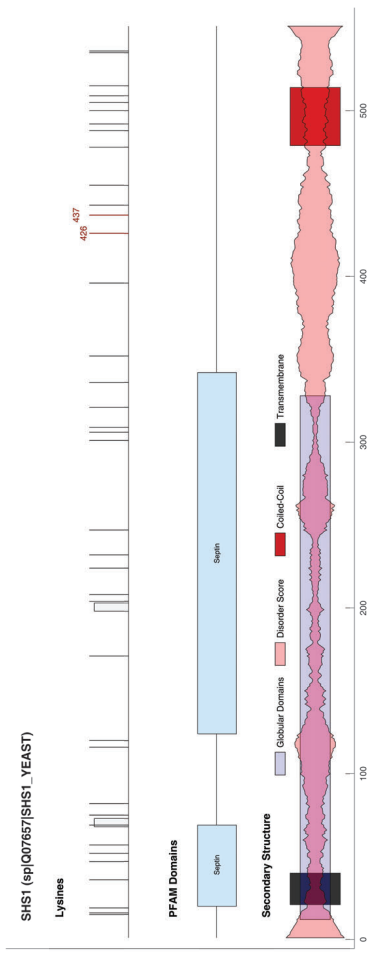
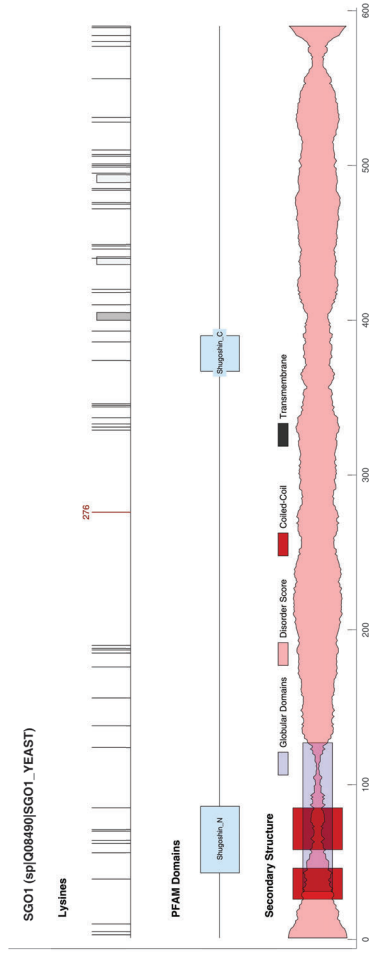
SEC5 (sp|P89102|SEC5_YEAST)

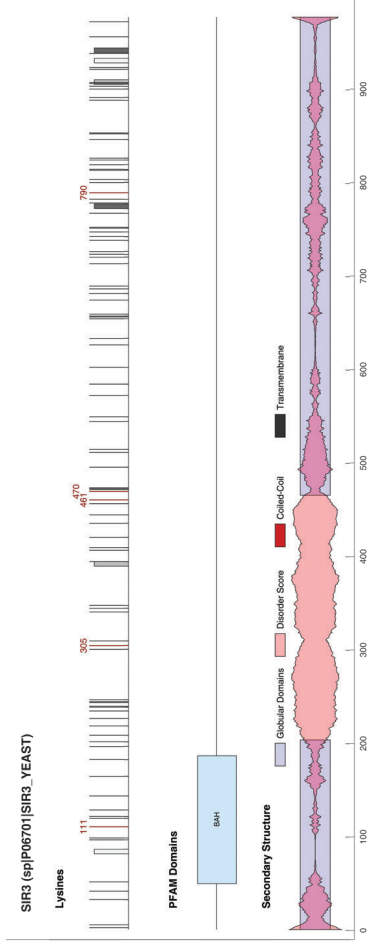
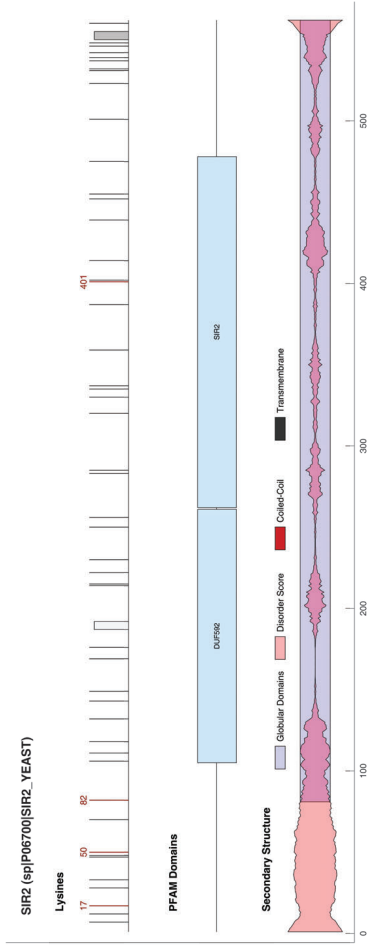
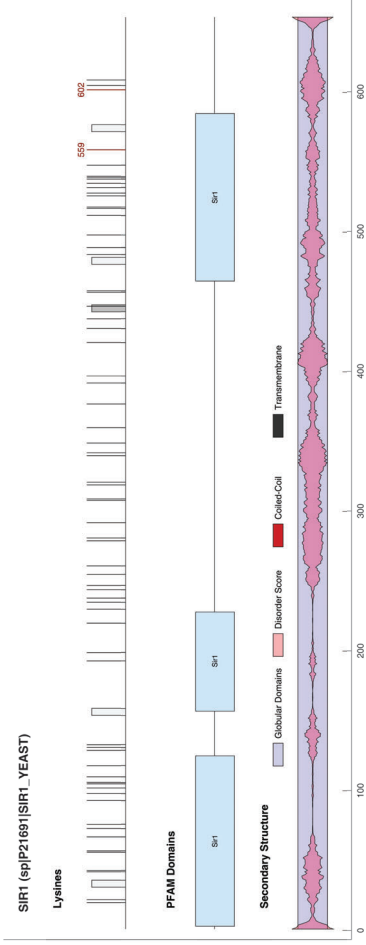
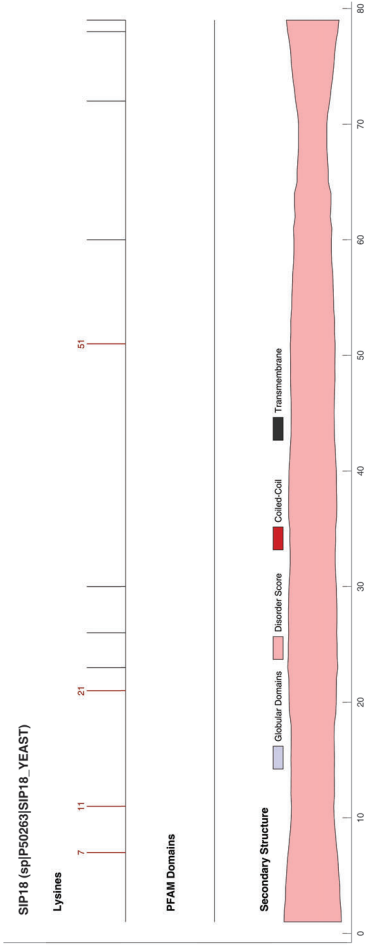


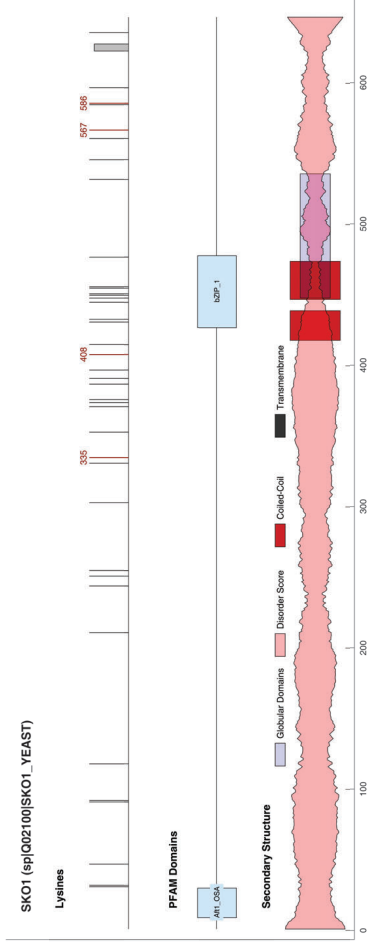
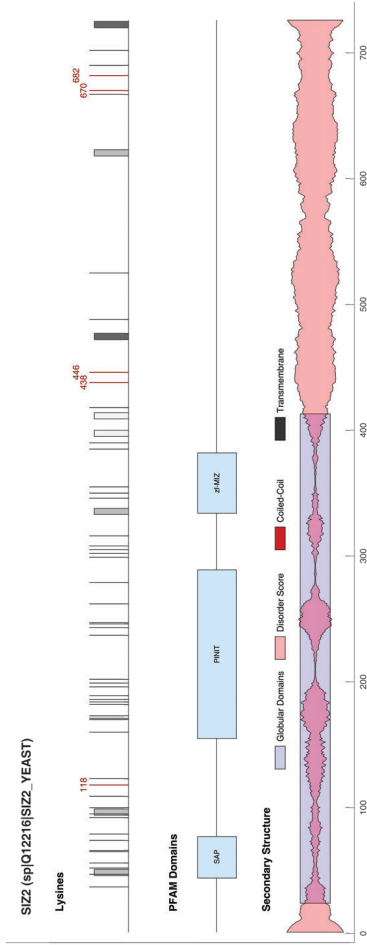
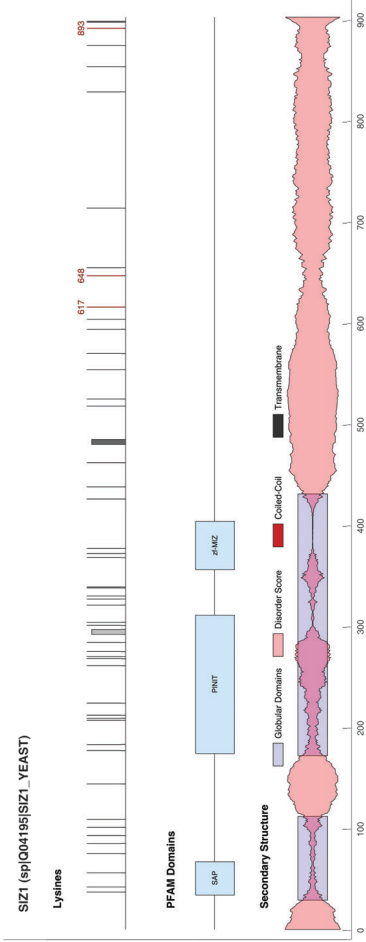
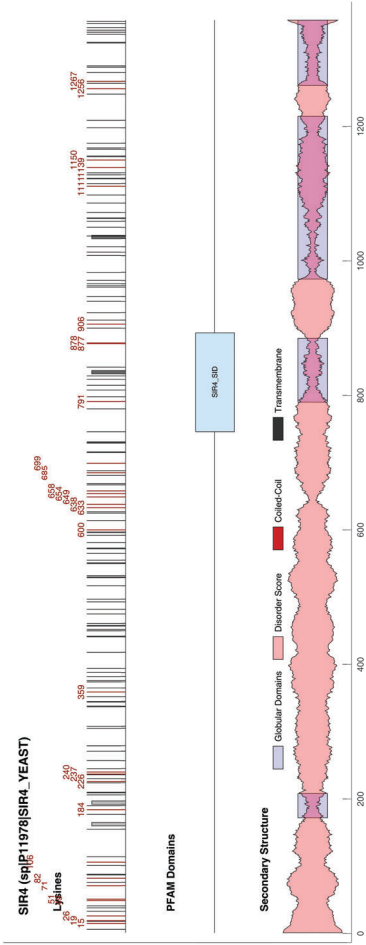
SEF1 (sp|P34228|SEF1_YEAST)

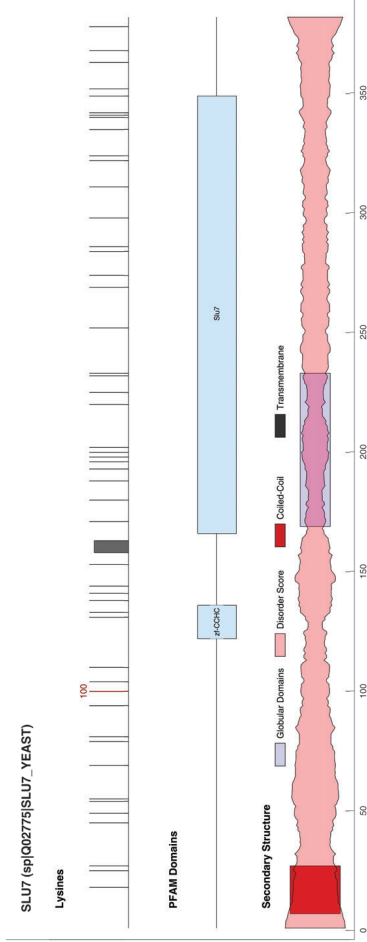
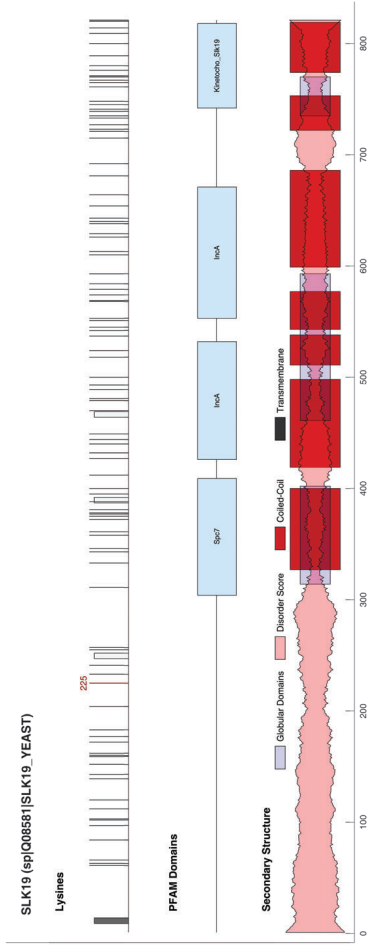
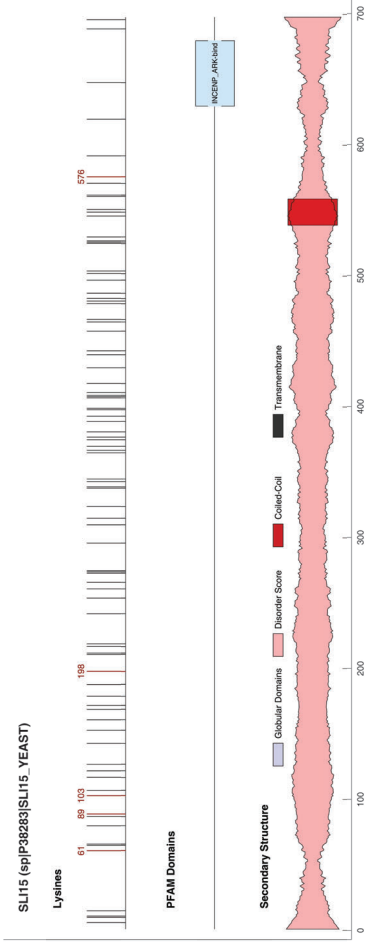
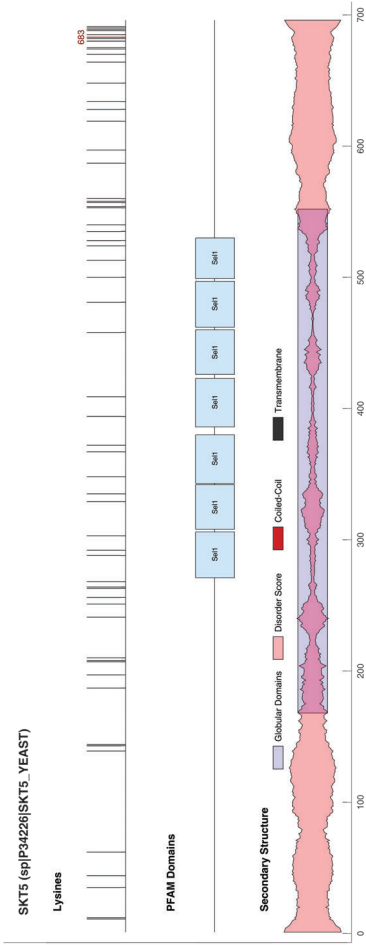


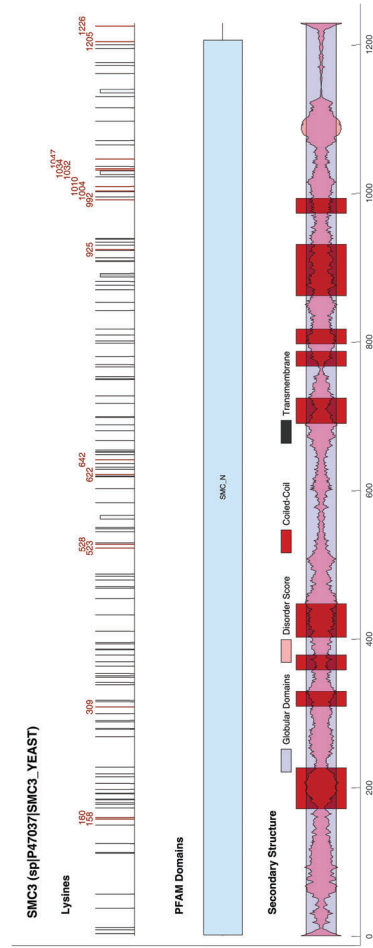
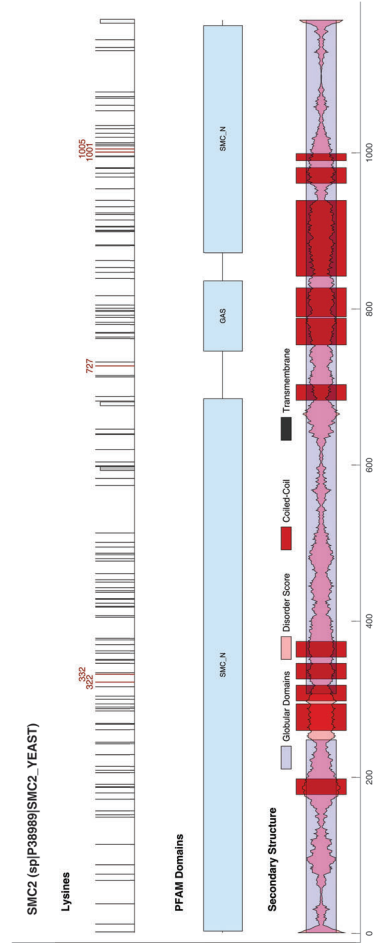
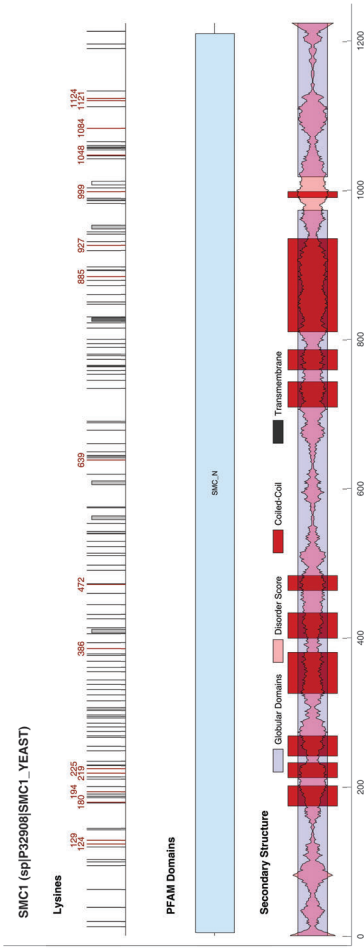
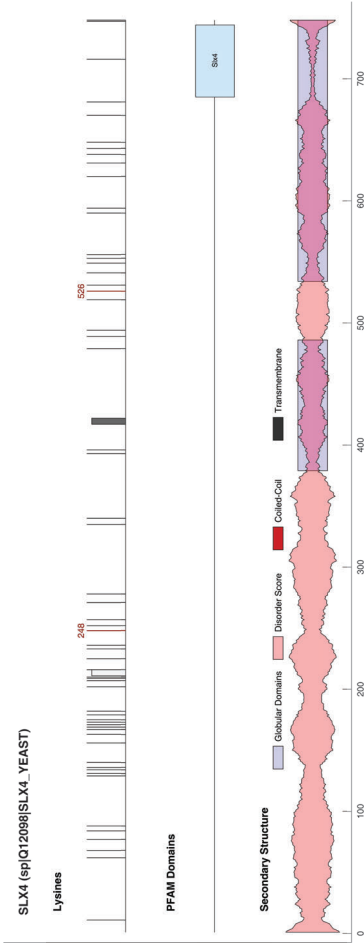


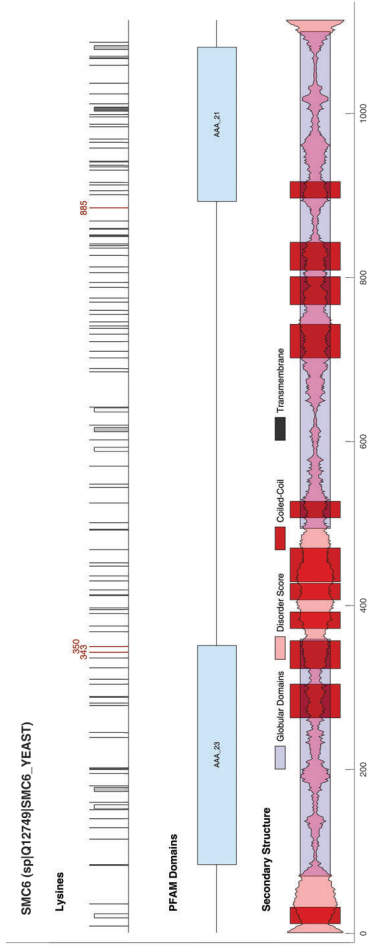
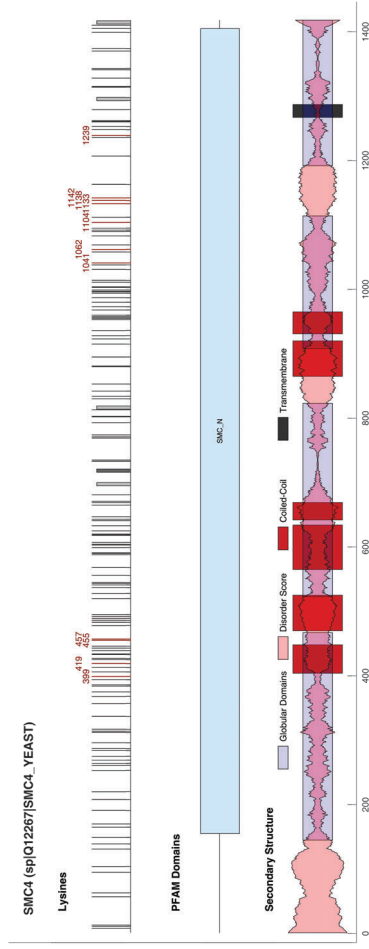




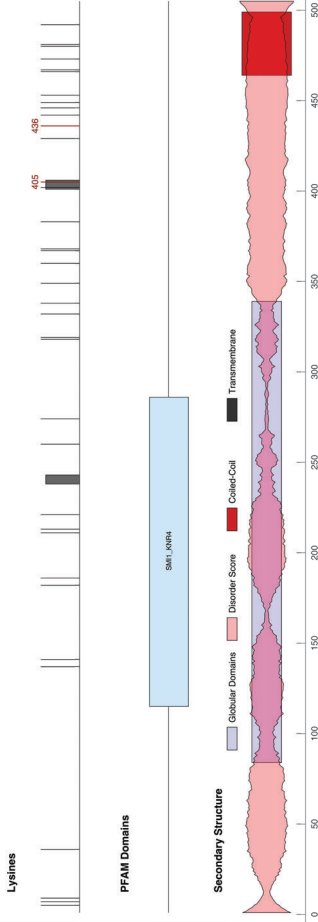




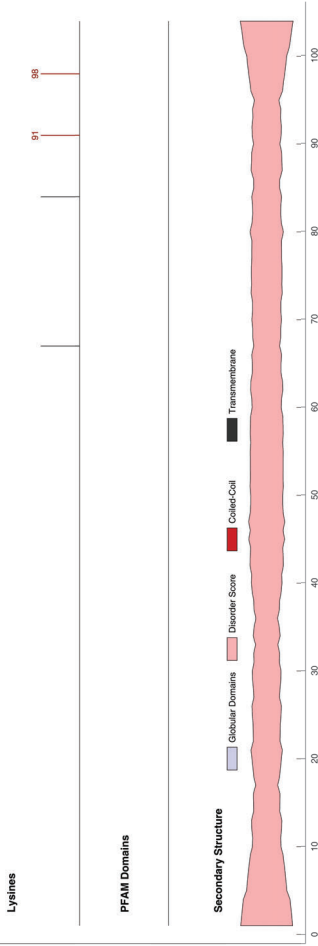




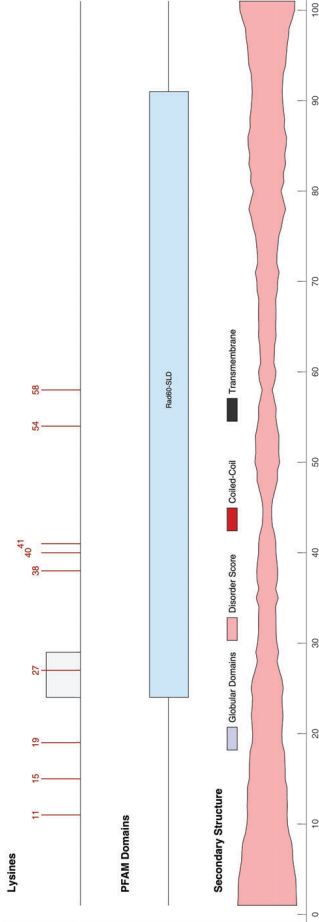
SMT1 (sp|P32566|SMT1_YEAST)



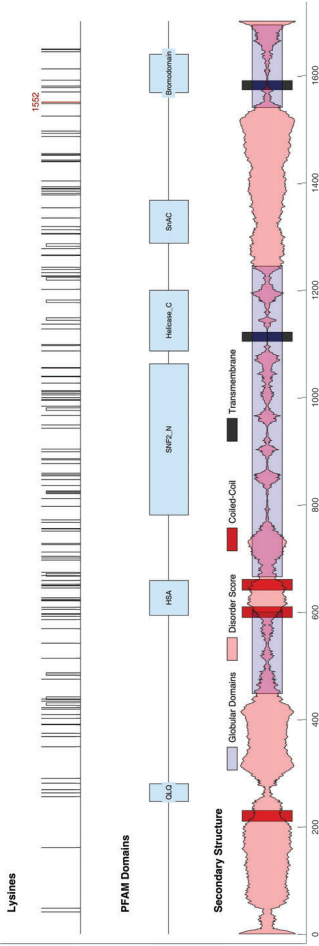
SML1 (sp|Q04964|SML1_YEAST)

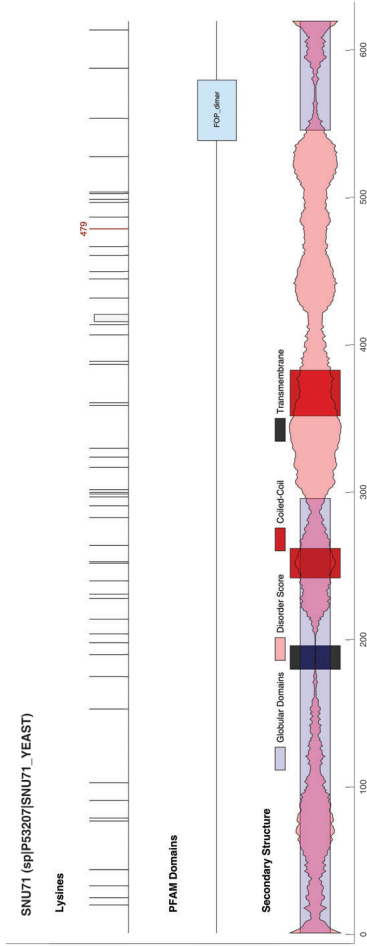
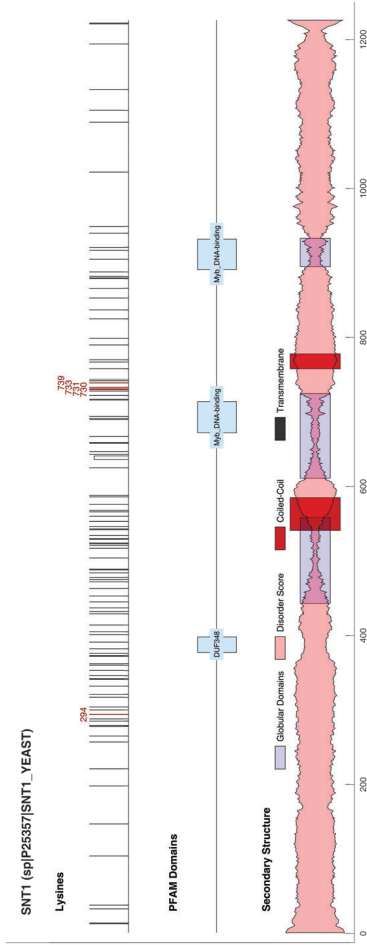
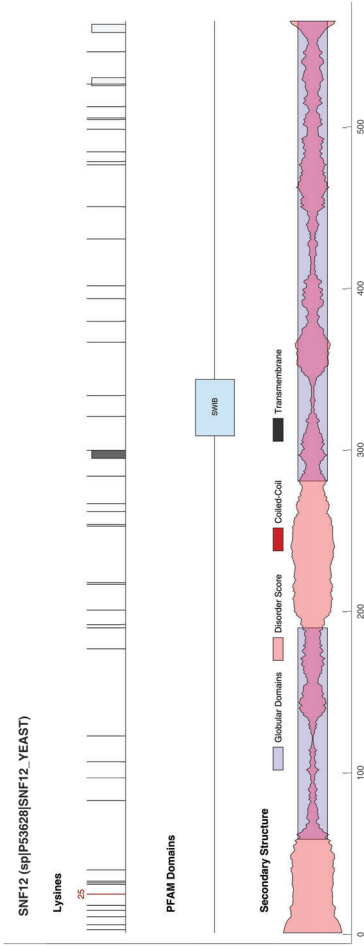
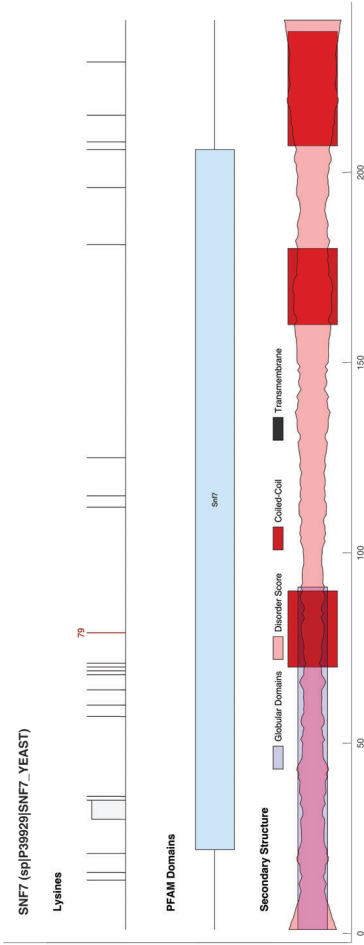


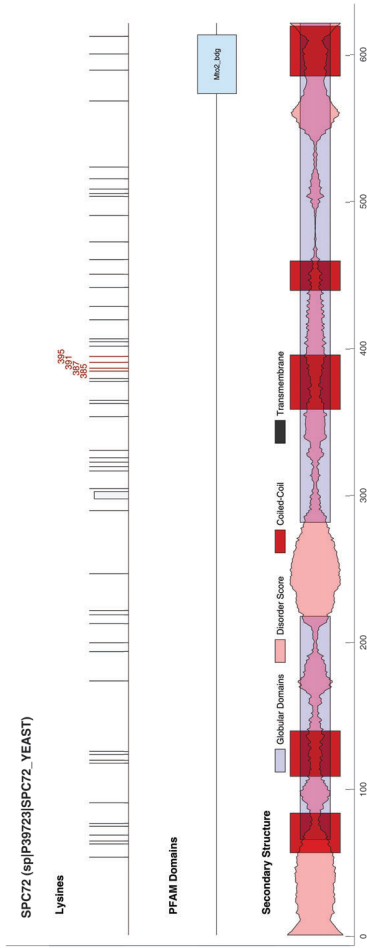
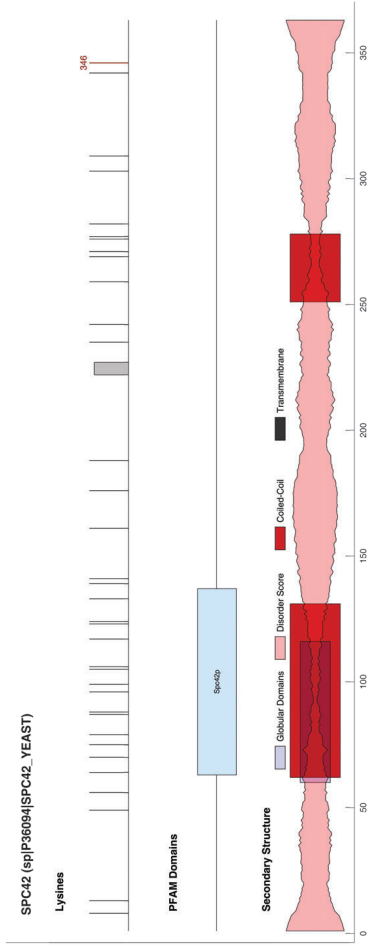
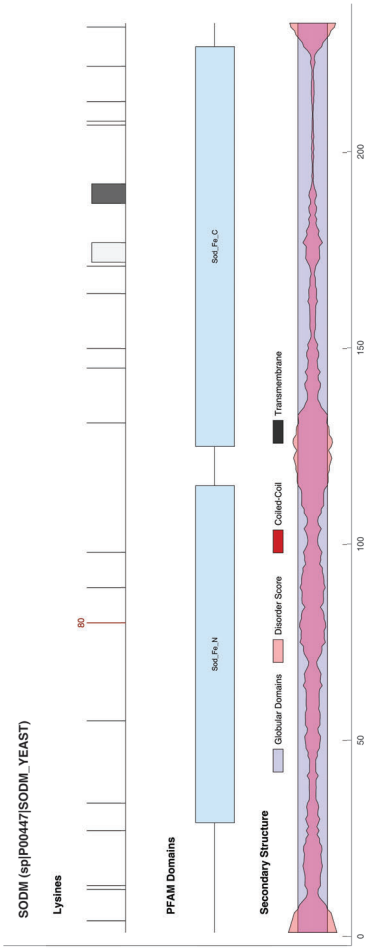
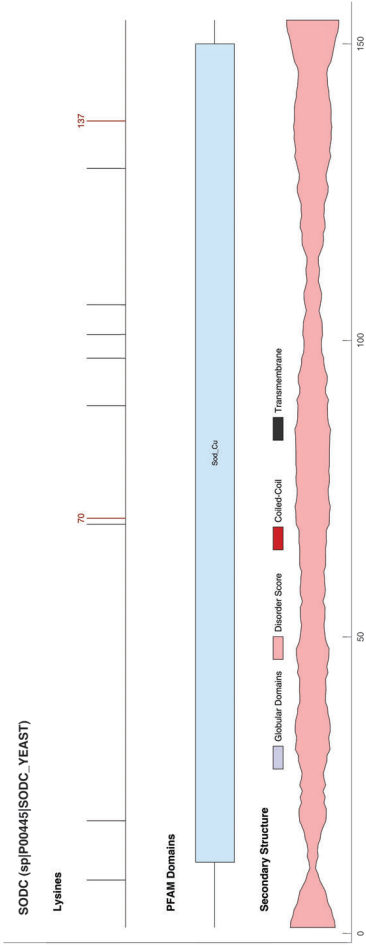
SMT3 (sp|Q12306|SMT3_YEAST)

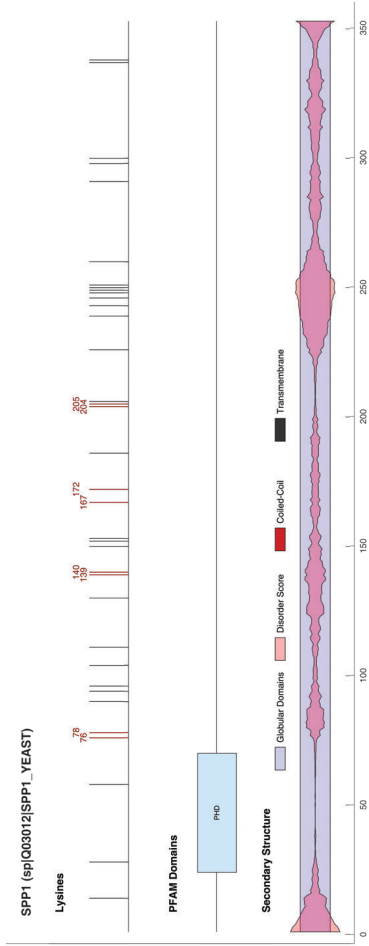
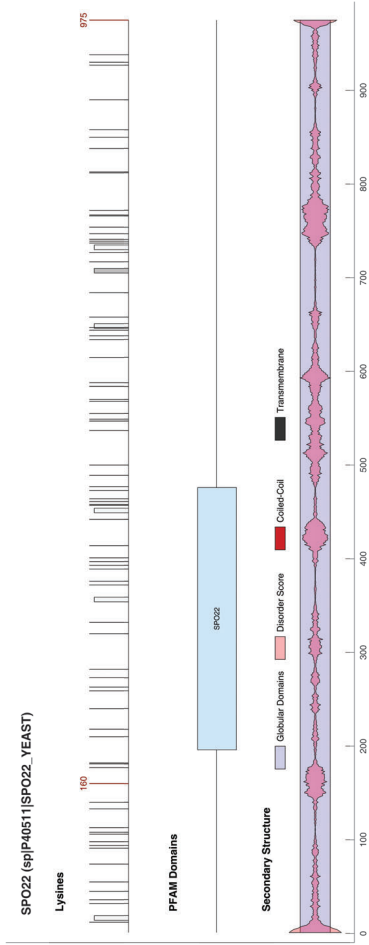
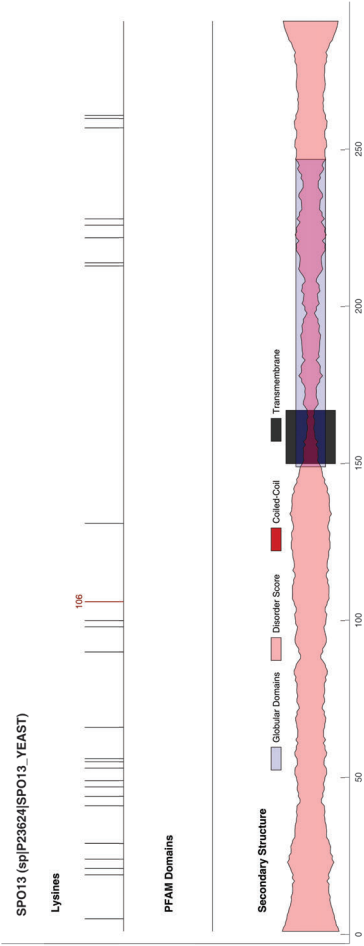
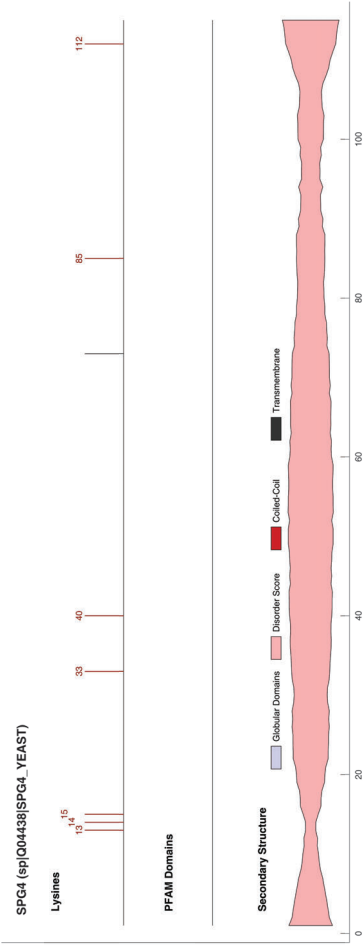


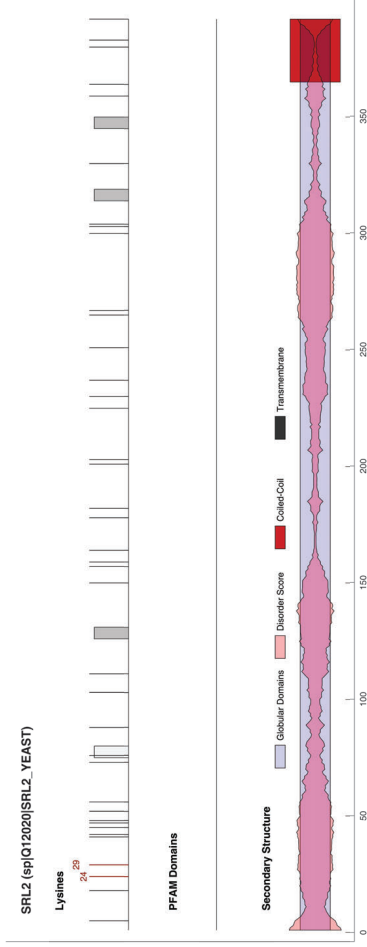
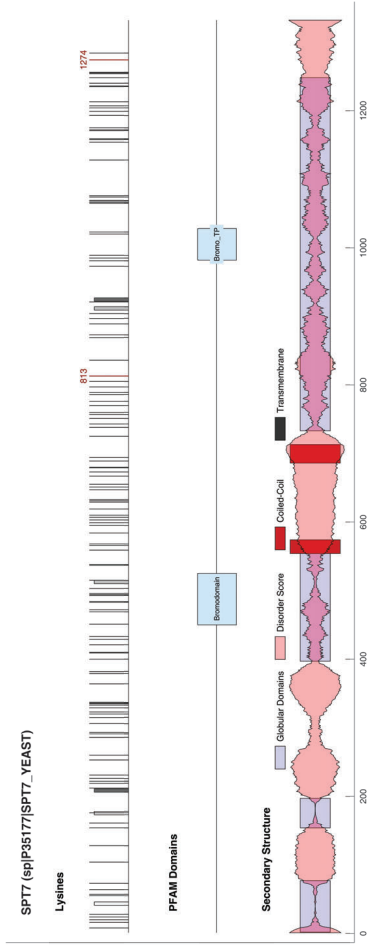
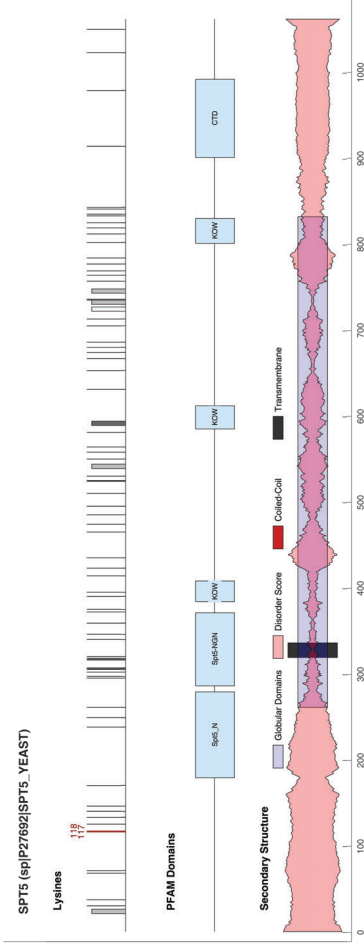
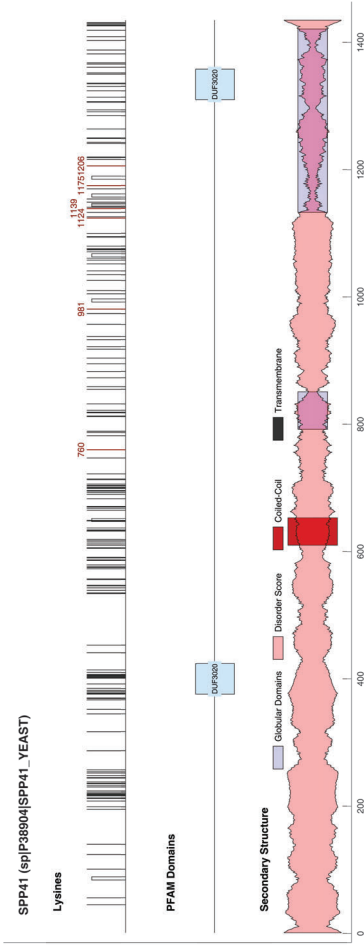
SNF2 (sp|P22082|SNF2_YEAST)

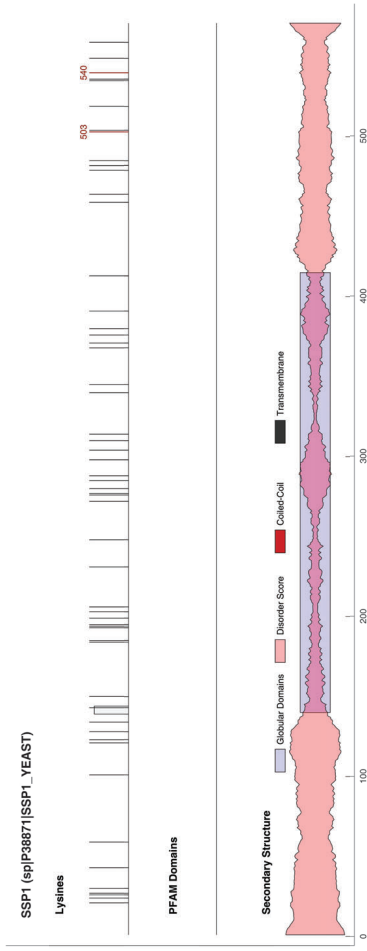
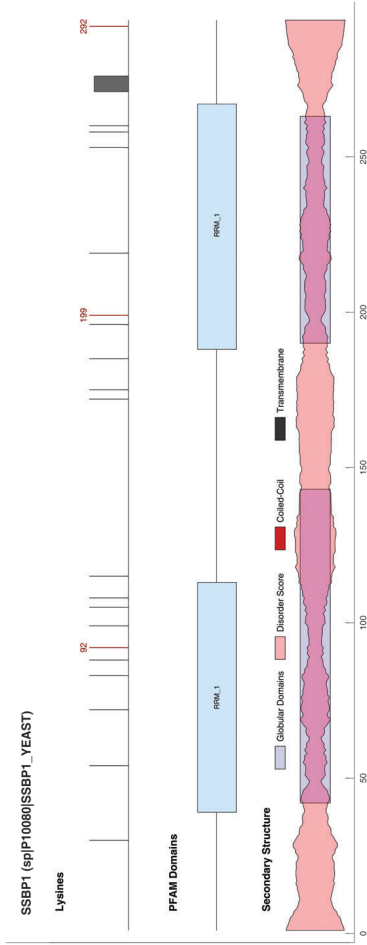
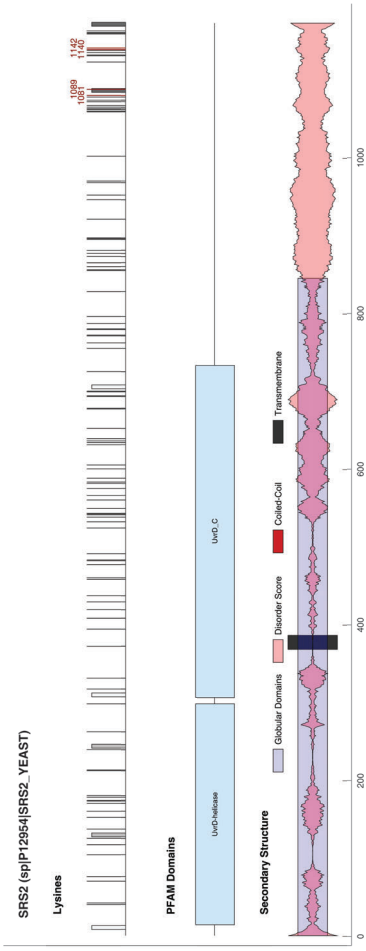
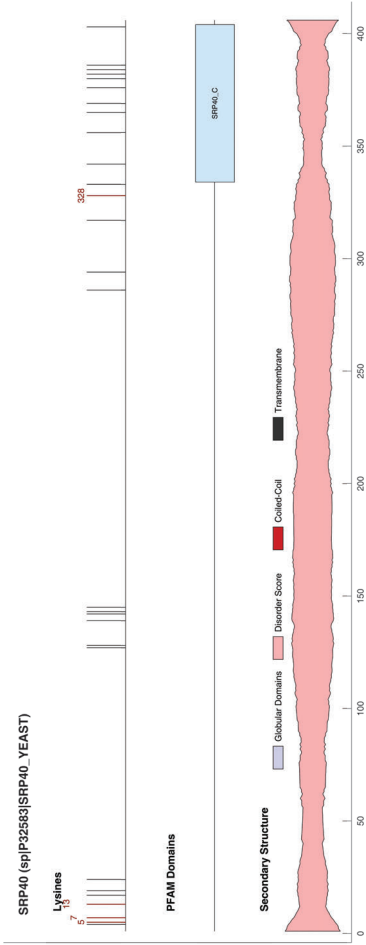


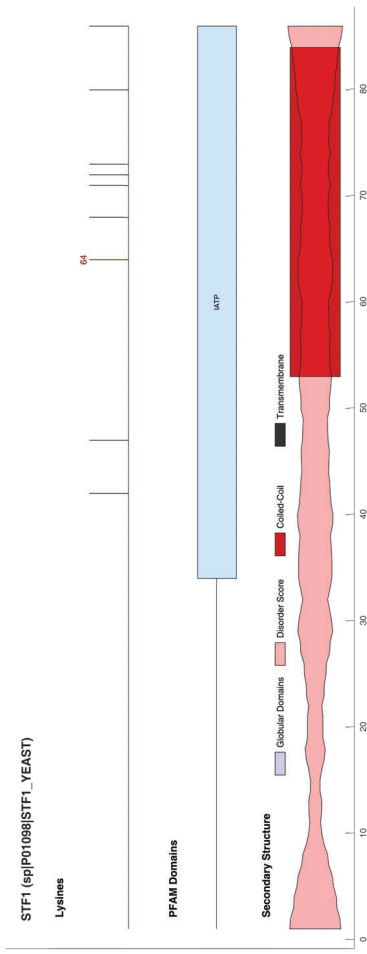
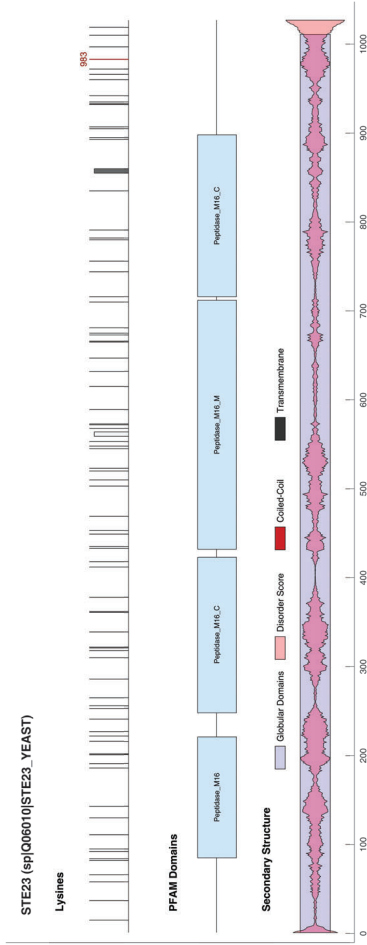
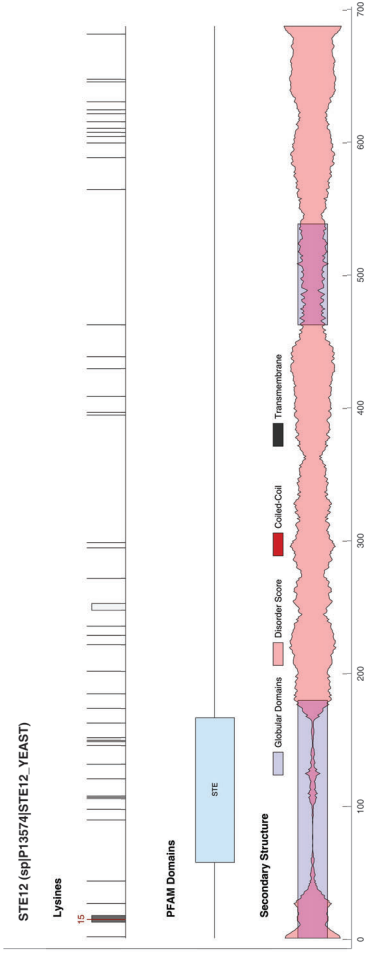
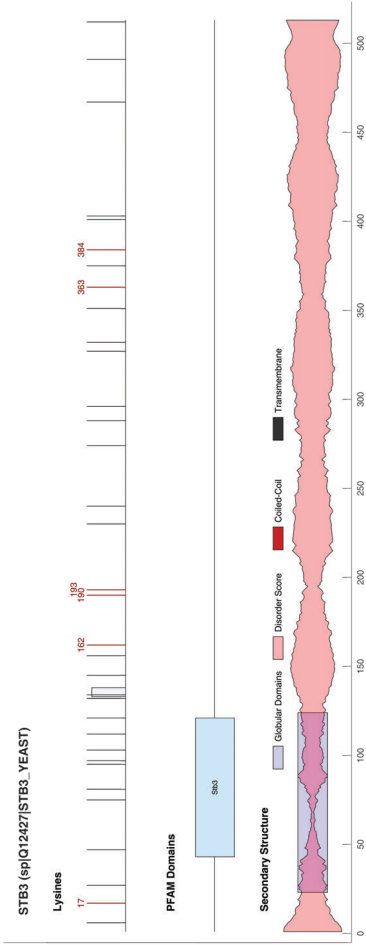


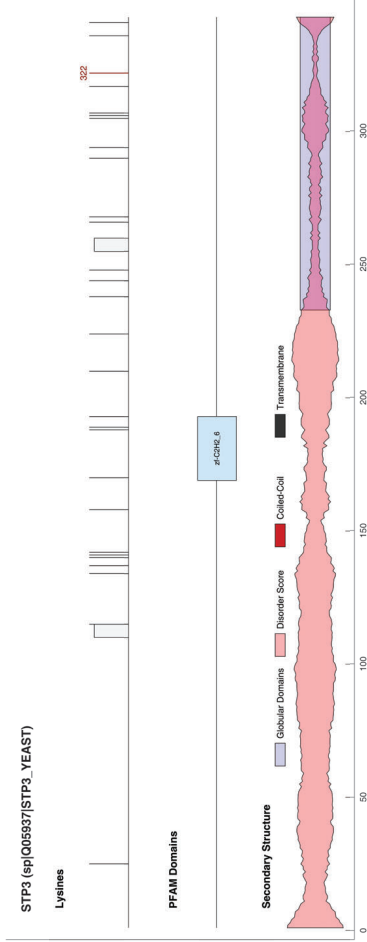
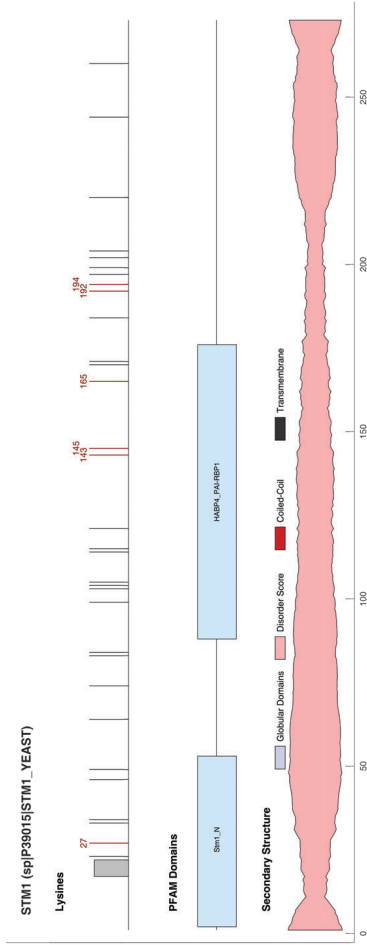
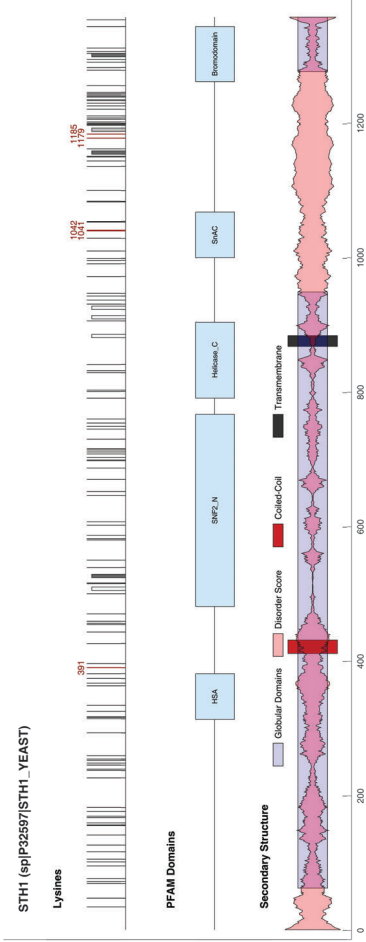
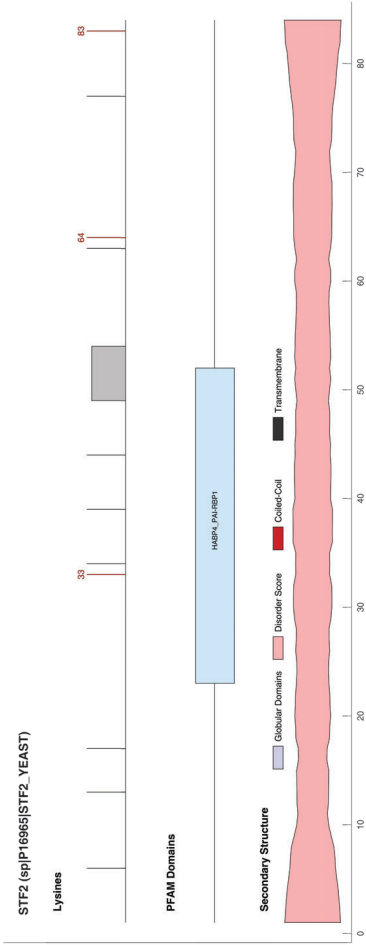




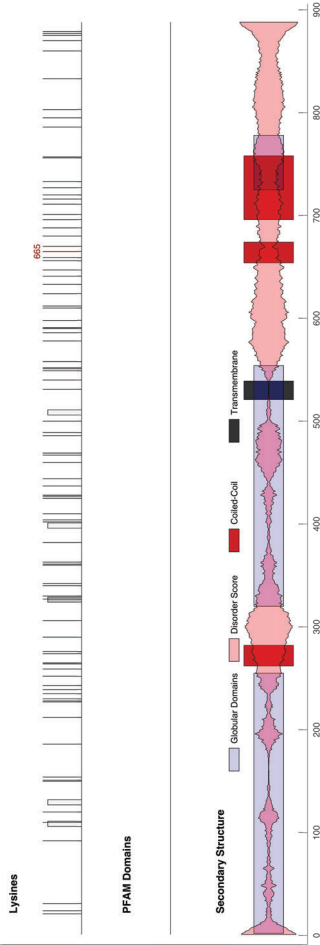




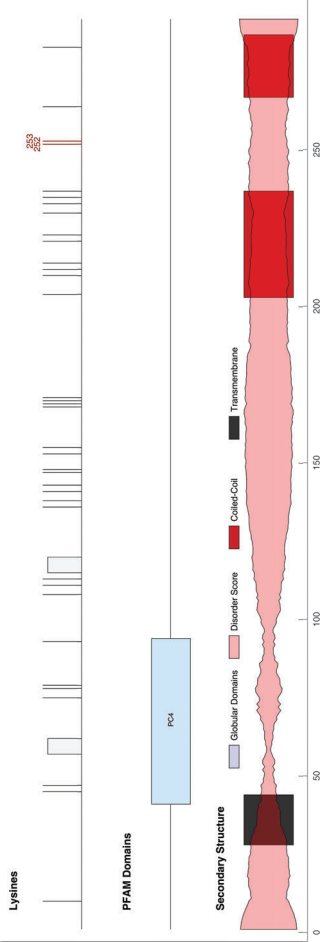




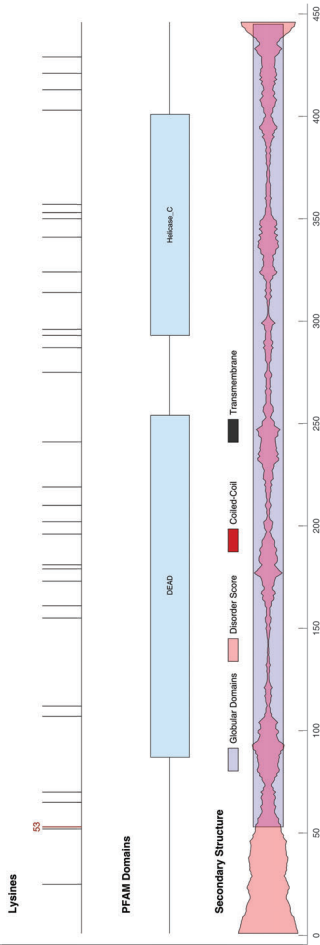
STU2 (sp|P46675|STU2_YEAST)



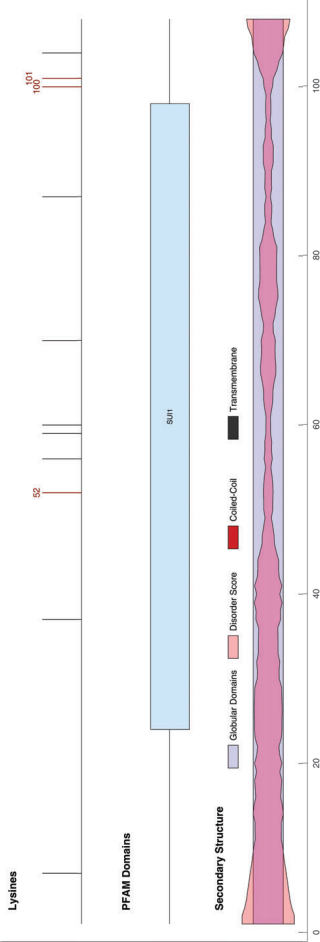
SUB1 (sp|P54000|SUB1_YEAST)



SUB2 (sp|Q07478|SUB2_YEAST)



SU1 (sp|P32911|SU1_YEAST)



SUM1 (sp|P46676|SUM1_YEAST)

Lysines



PFAM Domains



SWC3 (sp|P31376|SWC3_YEAST)

Lysines



PFAM Domains



SWC4 (sp|P53201|SWC4_YEAST)

Lysines

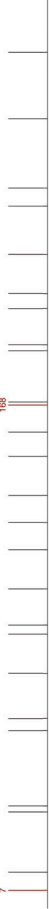


PFAM Domains



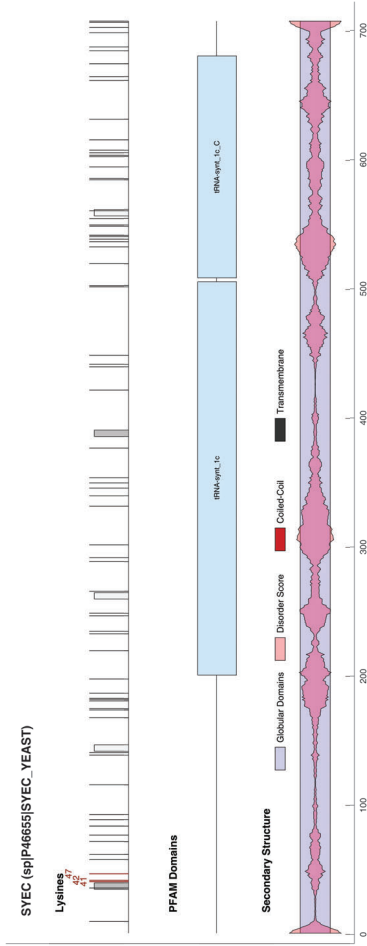
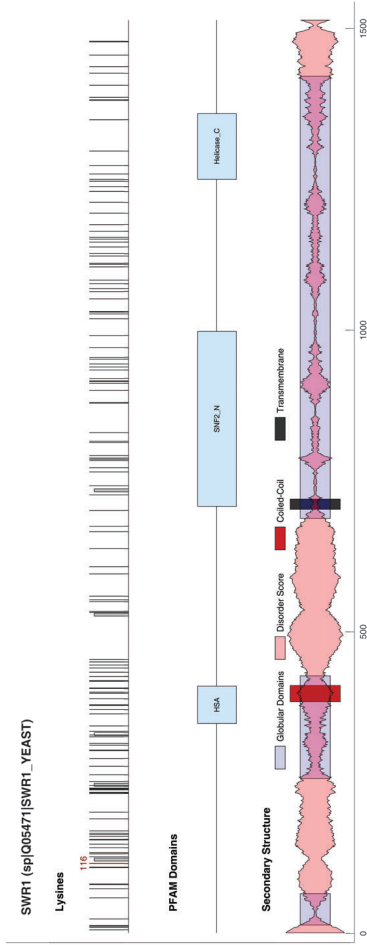
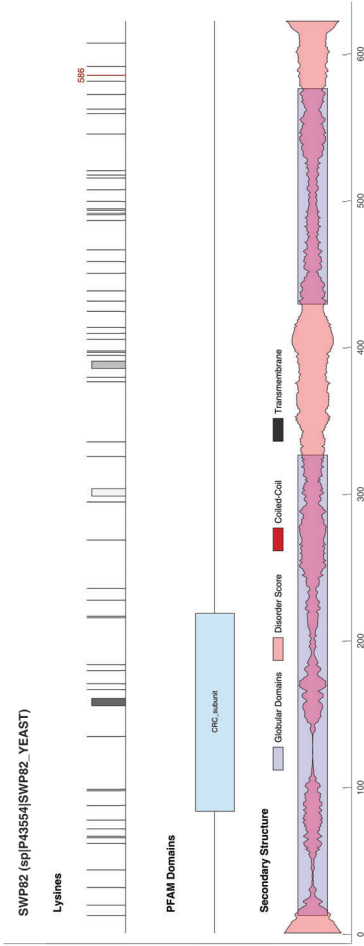
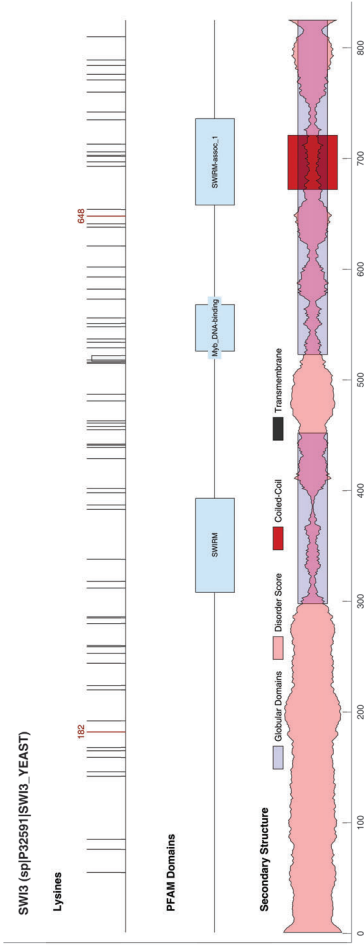
SWC5 (sp|P38326|SWC5_YEAST)

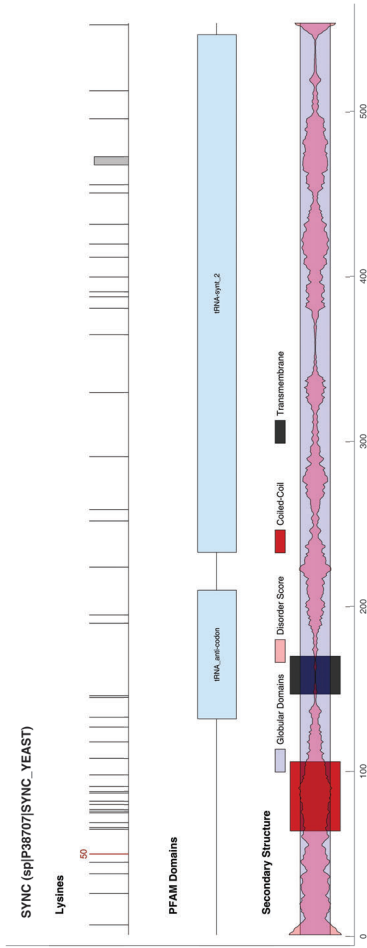
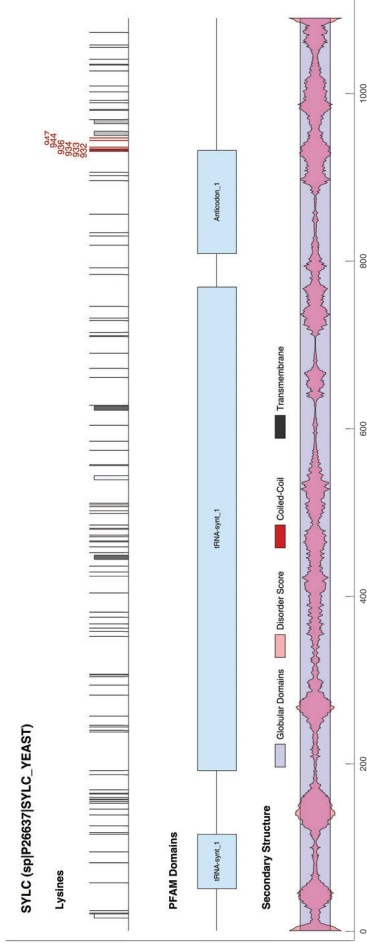
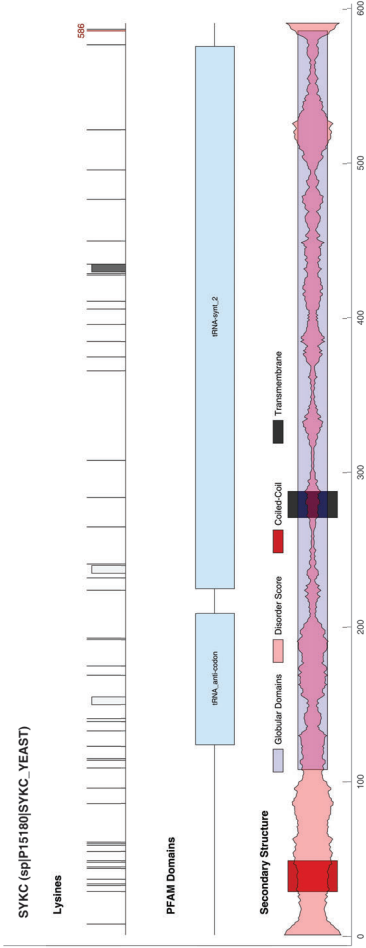
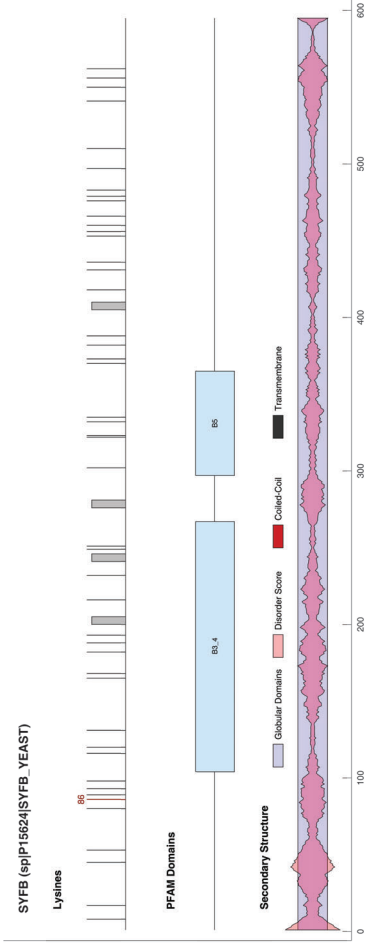
Lysines



PFAM Domains







SYYC (sp|P36421|SYYC_YEAST)

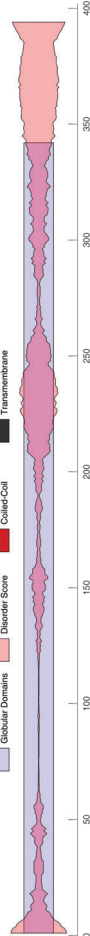
Lysines



PFAM Domains



Secondary Structure



T2EA (sp|P36100|T2EA_YEAST)

Lysines



PFAM Domains



Secondary Structure



T2EB (sp|P36145|T2EB_YEAST)

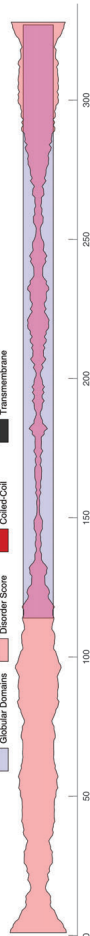
Lysines



PFAM Domains



Secondary Structure



T2FA (sp|P41895|T2FA_YEAST)

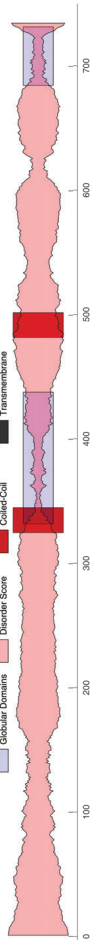
Lysines



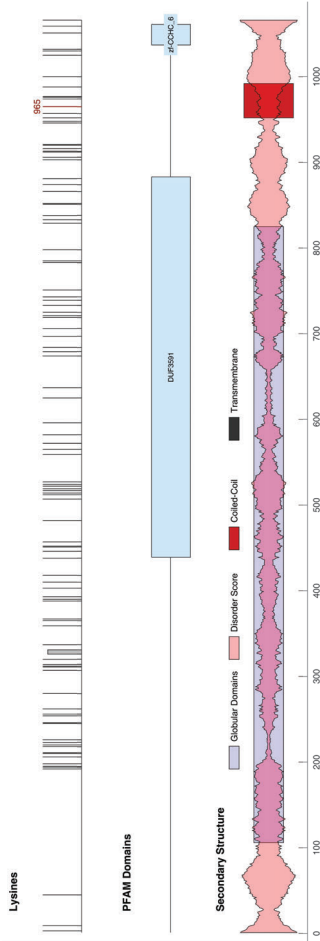
PFAM Domains



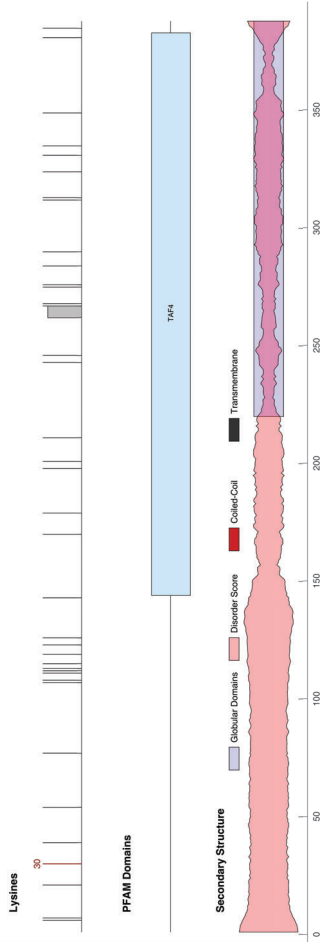
Secondary Structure



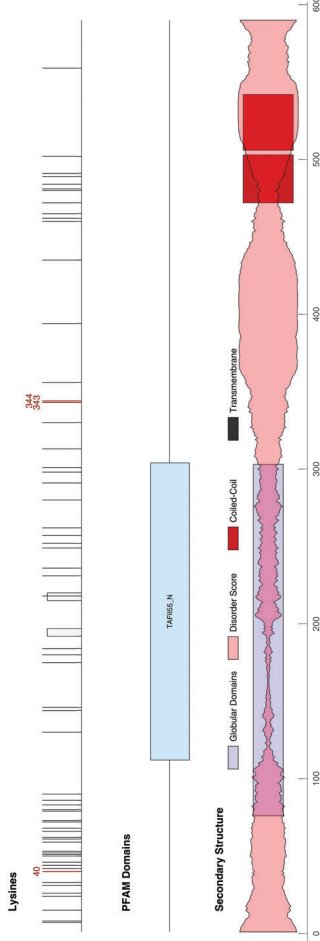
TAF1 (sp|P46677|TAF1_YEAST)



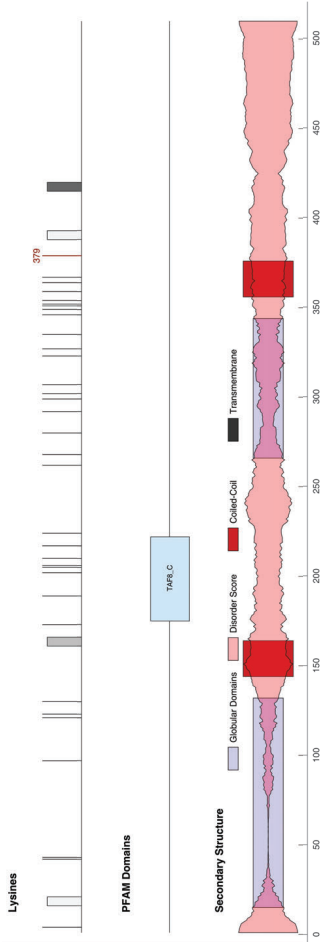
TAF4 (sp|P50105|TAF4_YEAST)

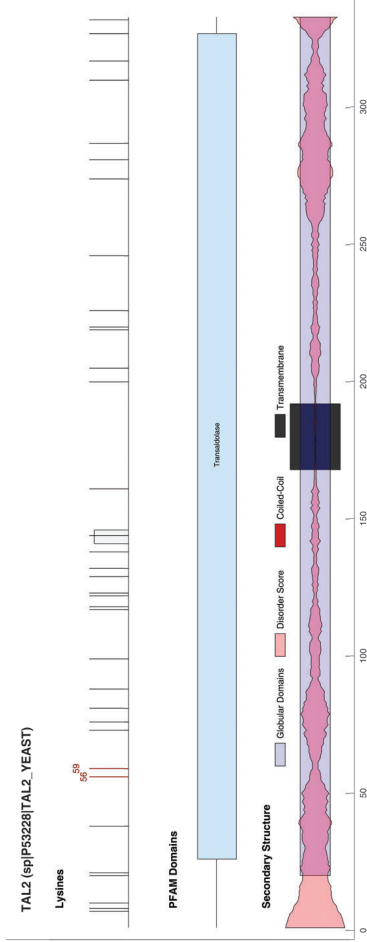
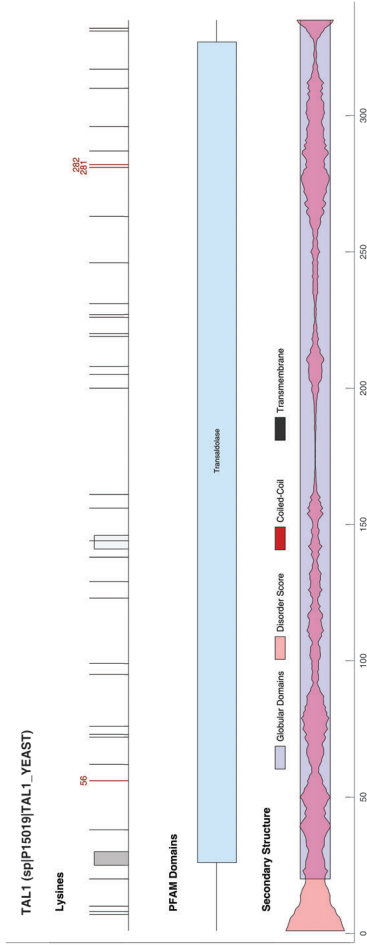
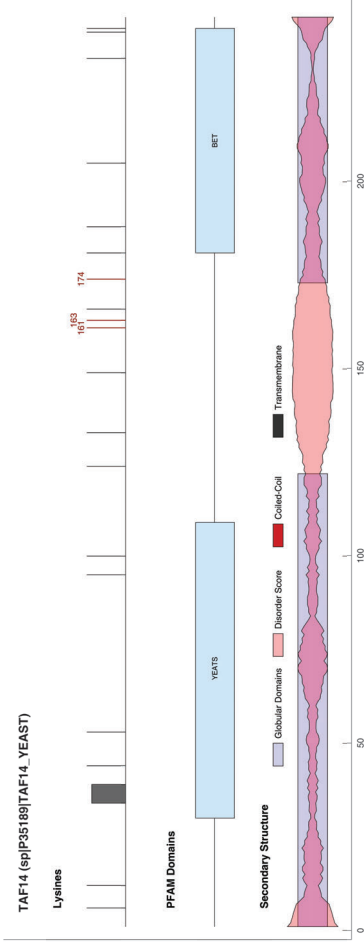
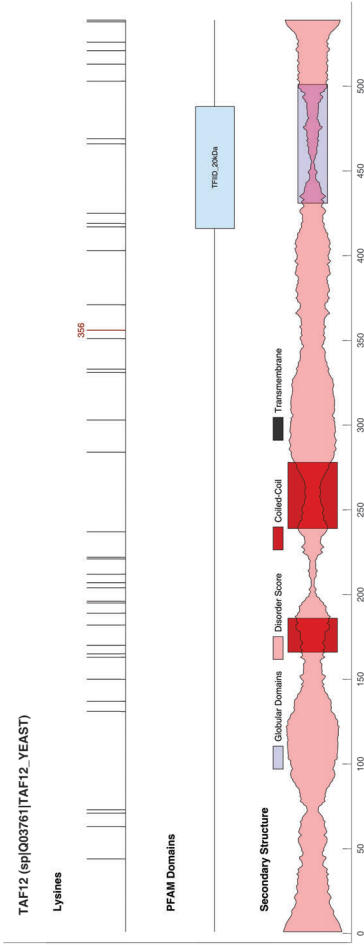


TAF7 (sp|Q05021|TAF7_YEAST)

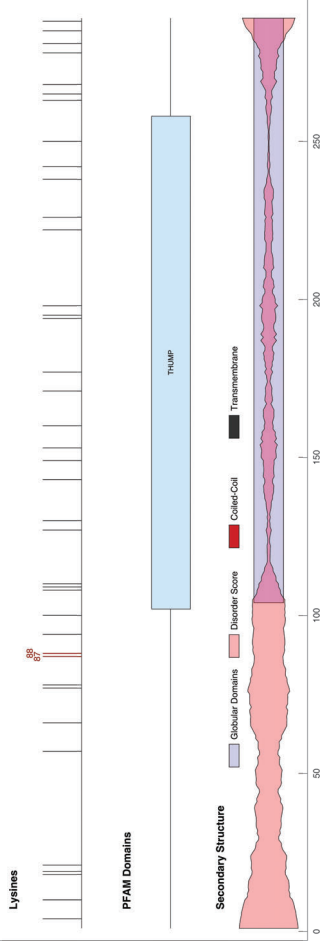


TAF8 (sp|Q03750|TAF8_YEAST)

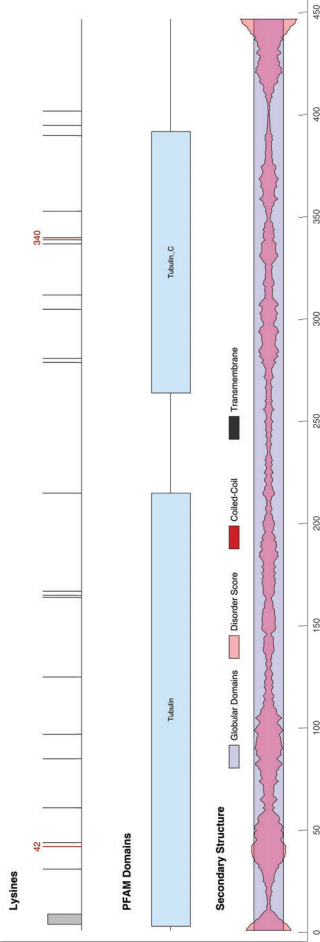




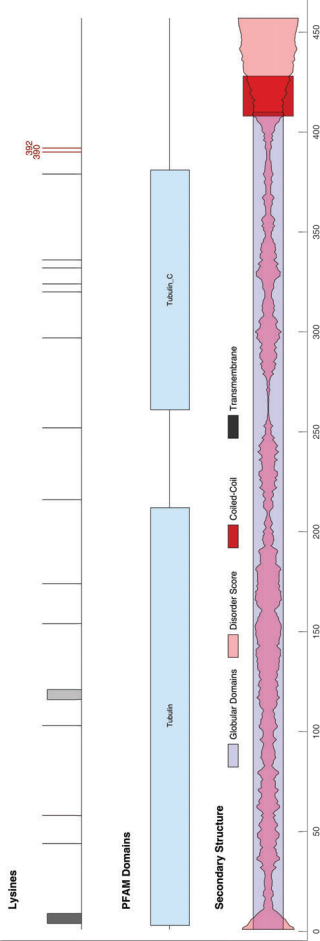
TAN1 (sp|P53072|TAN1_YEAST)



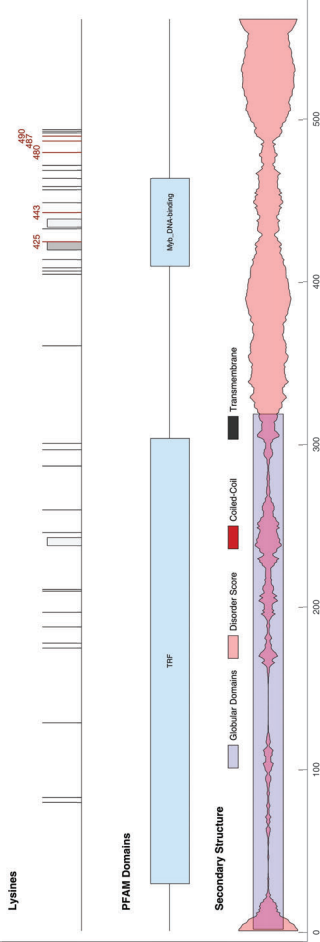
TBA1 (sp|P09733|TBA1_YEAST)

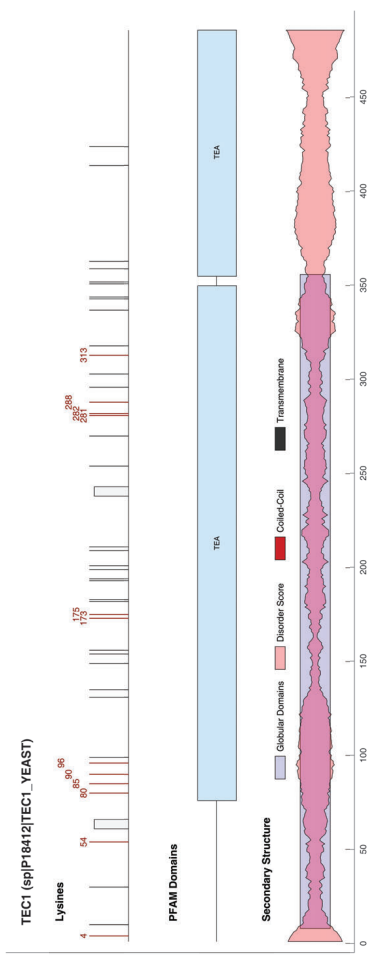
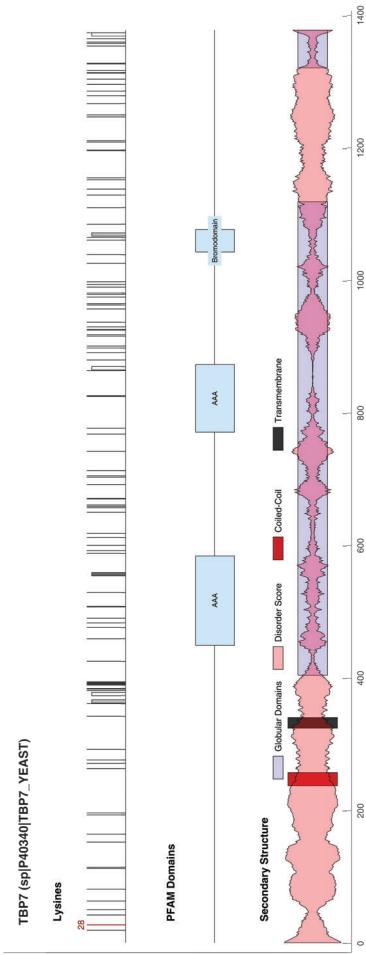
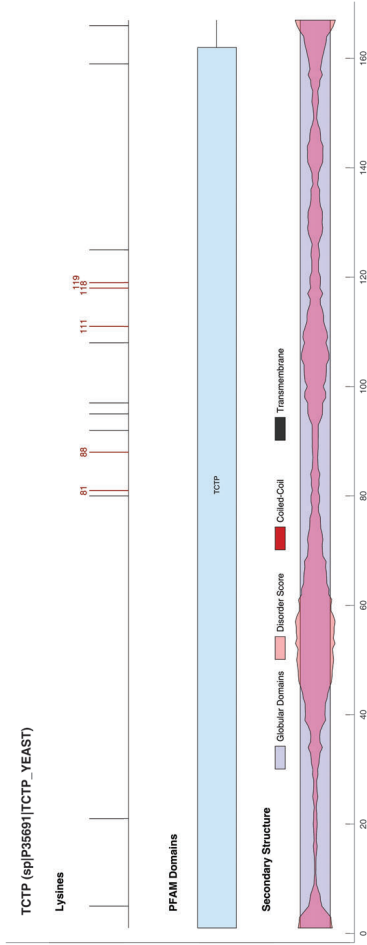
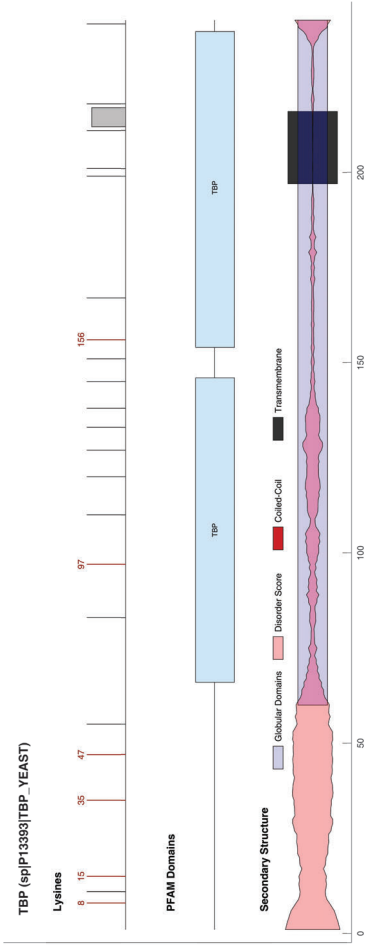


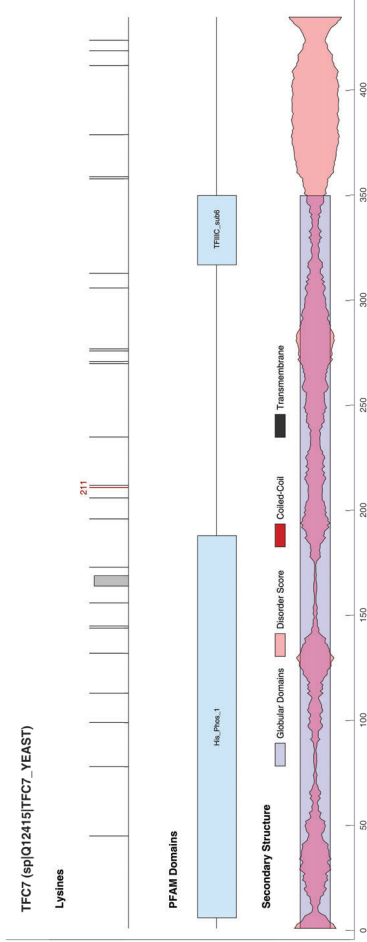
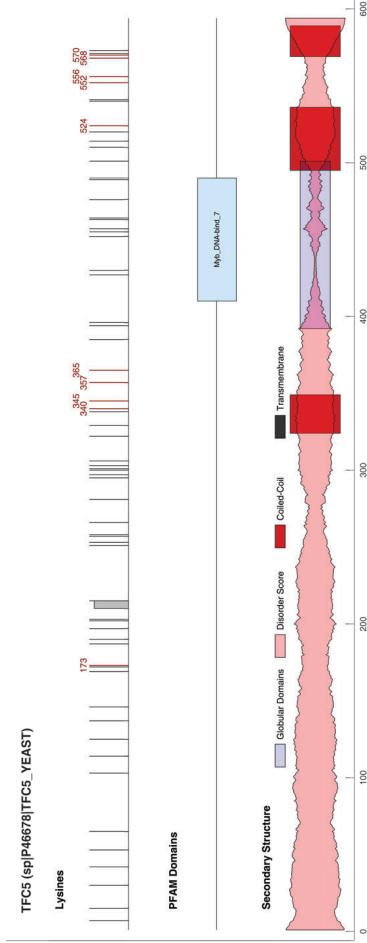
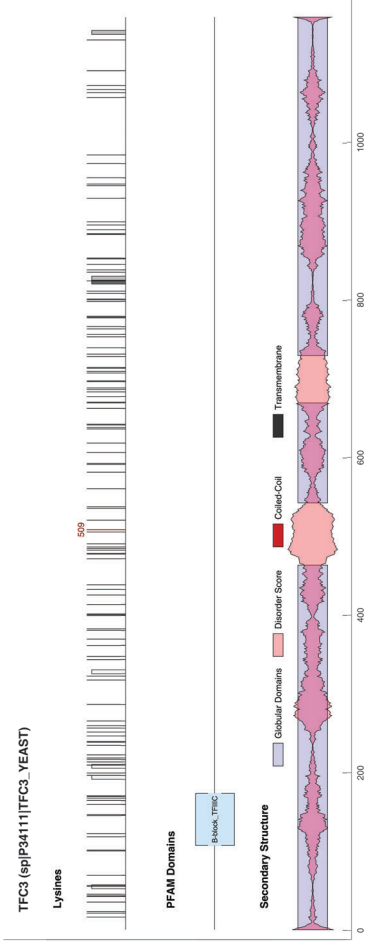
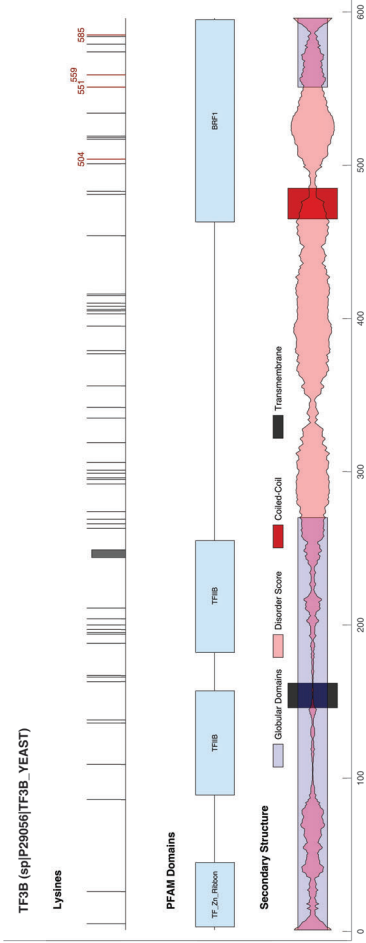
TBB (sp|P02557|TBB_YEAST)



TBF1 (sp|Q02457|TBF1_YEAST)

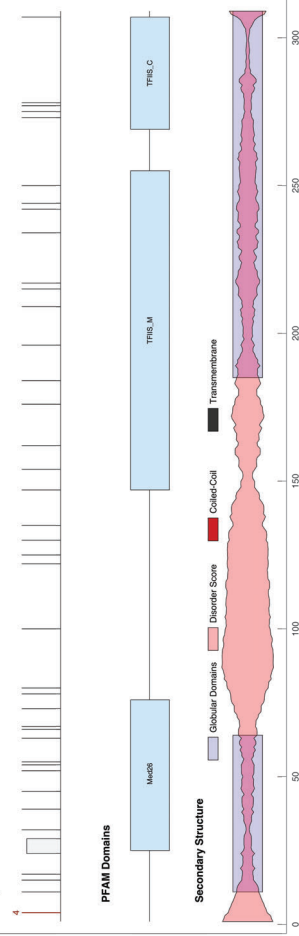






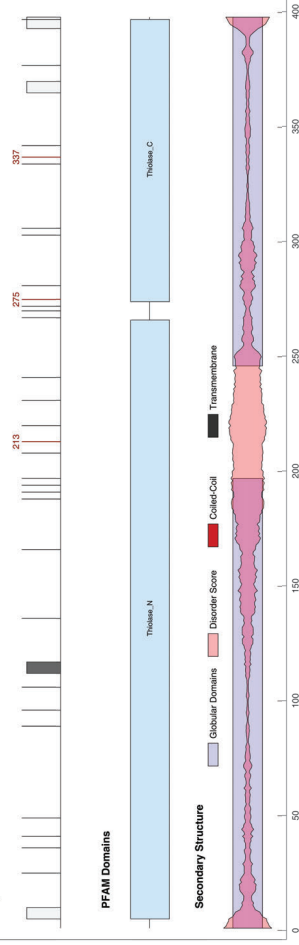
TFS2 (sp|P07273|TFS2_YEAST)

Lysines



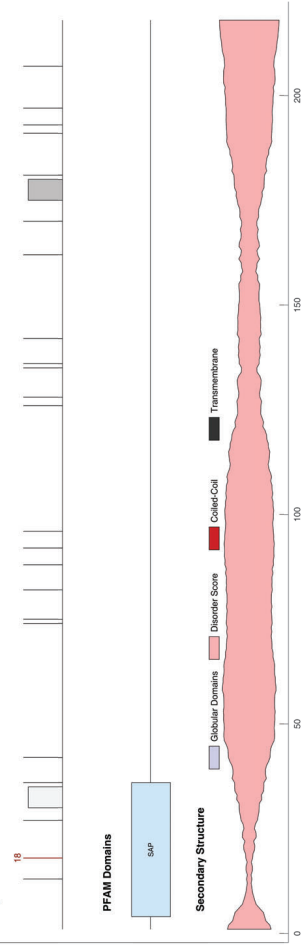
THIL (sp|P41338|THIL_YEAST)

Lysines



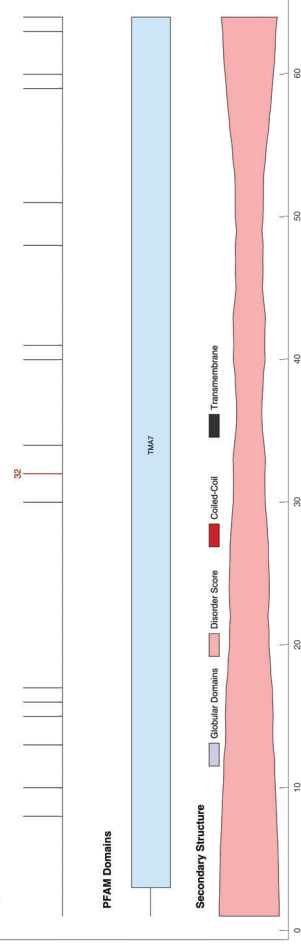
THO1 (sp|P40040|THO1_YEAST)

Lysines

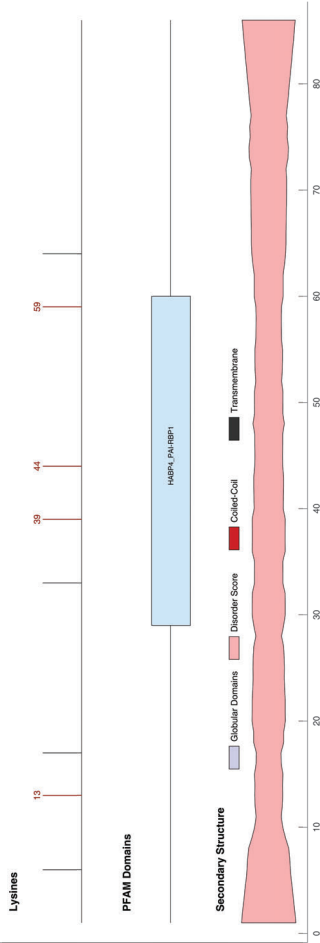


TMA7 (sp|Q3E764|TMA7_YEAST)

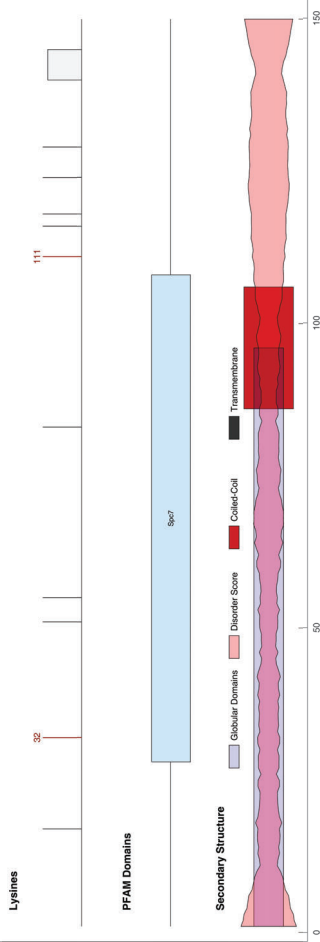
Lysines



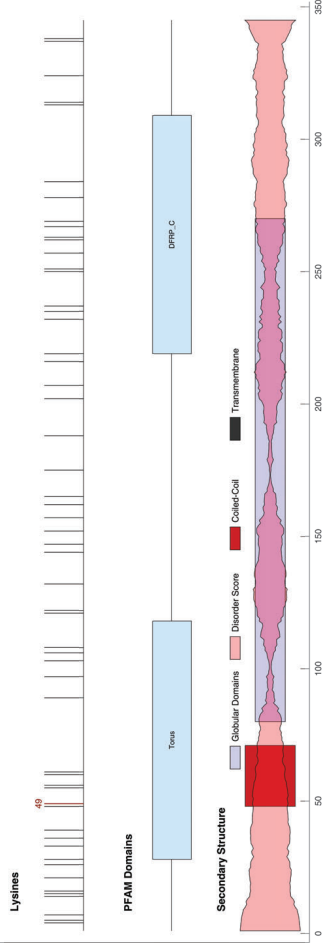
TMA10 (ep|Q06177|TMA10_YEAST)



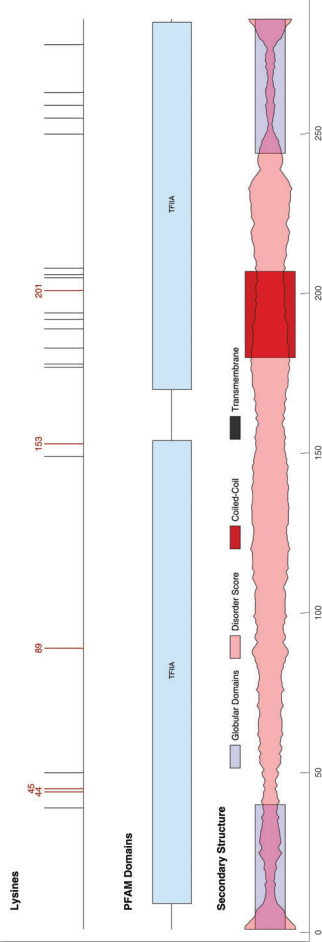
TMA17 (ep|Q12513|TMA17_YEAST)

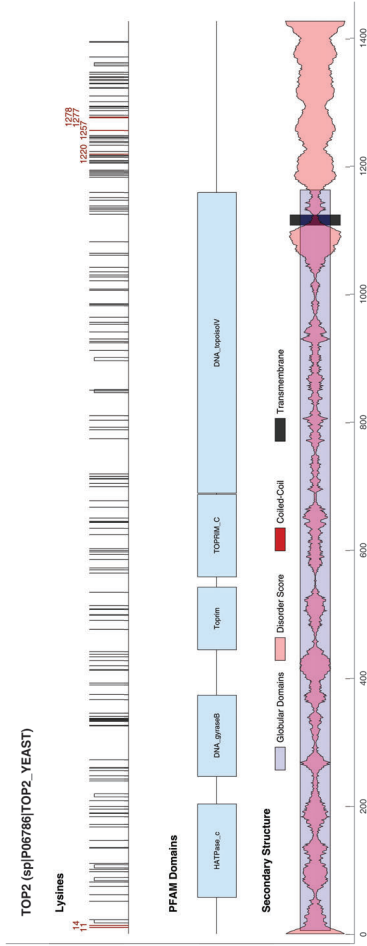
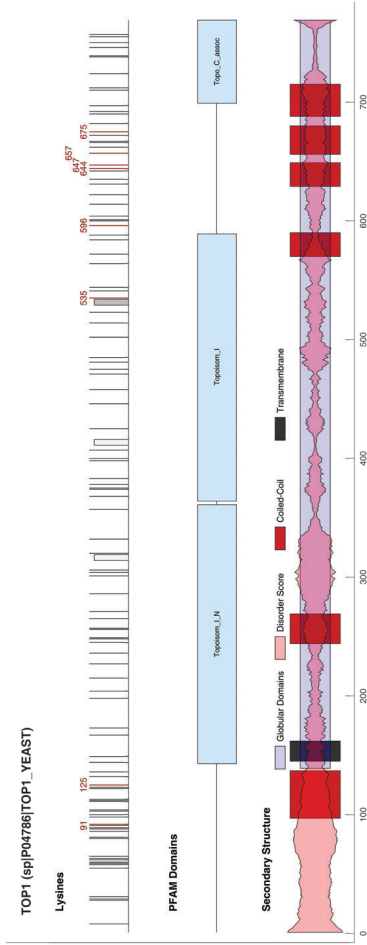
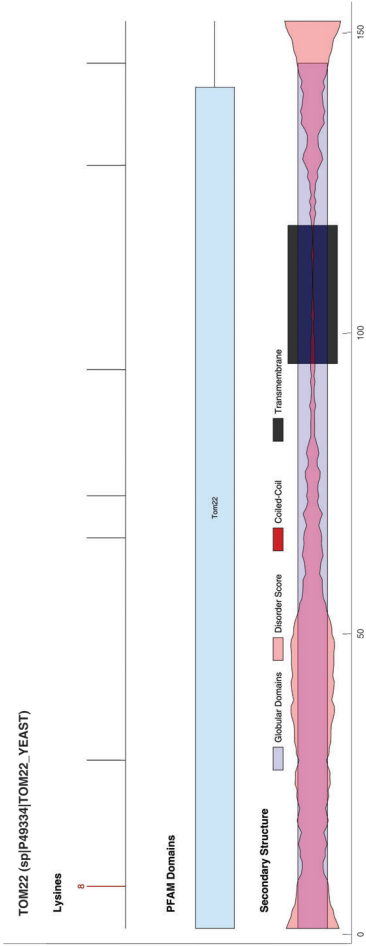
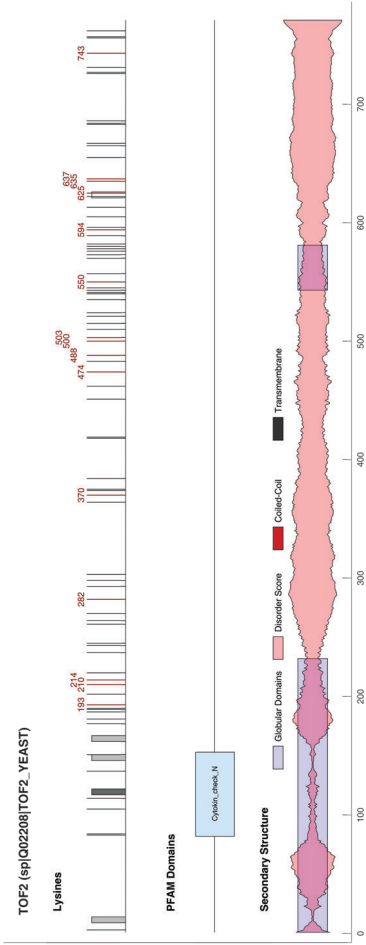


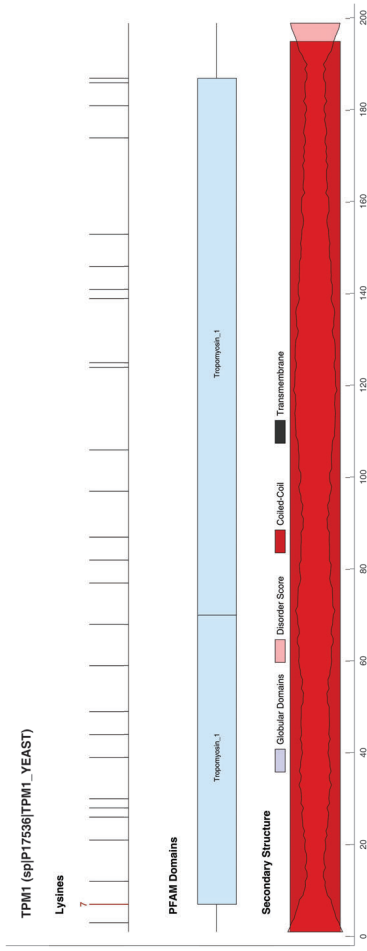
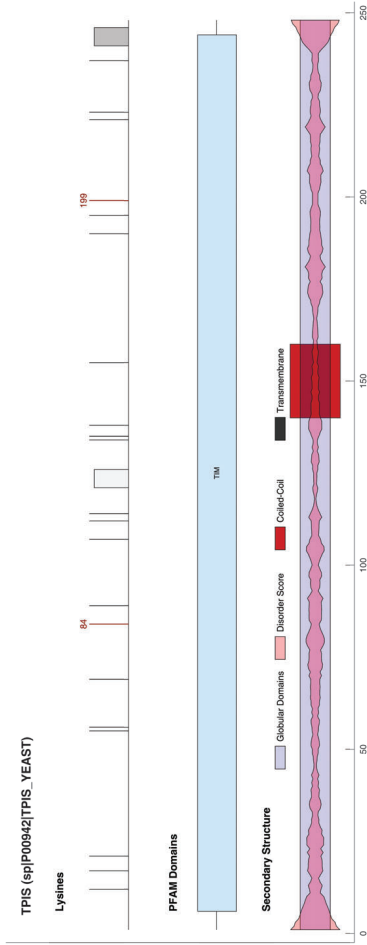
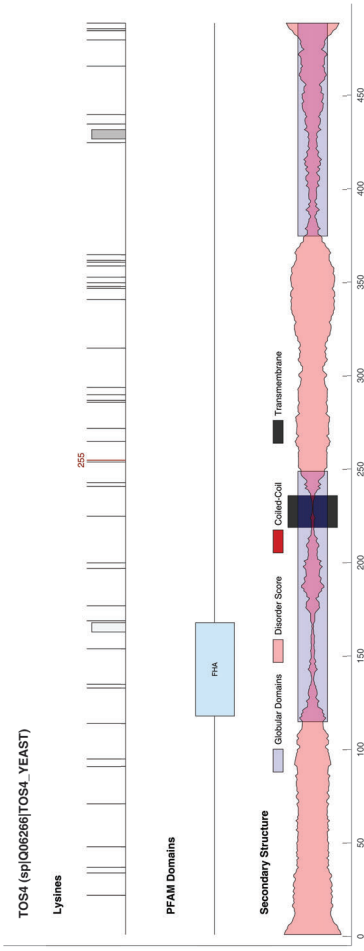
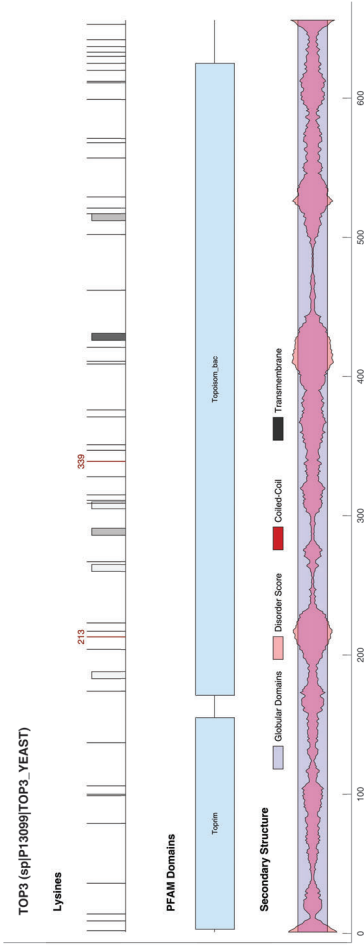
TMA46 (ep|Q12000|TMA46_YEAST)



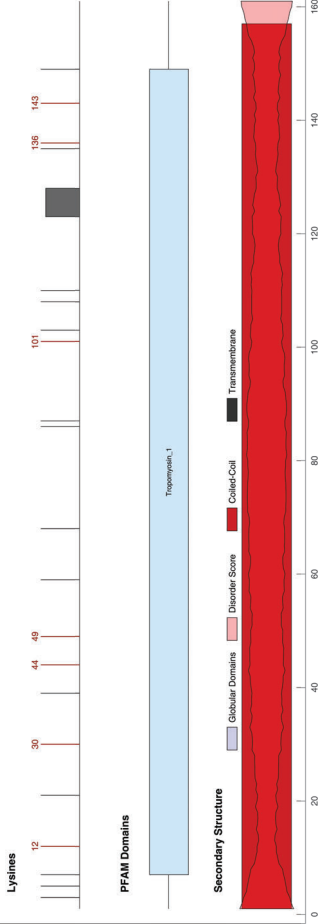
TOA1 (sp|P32773|TOA1_YEAST)



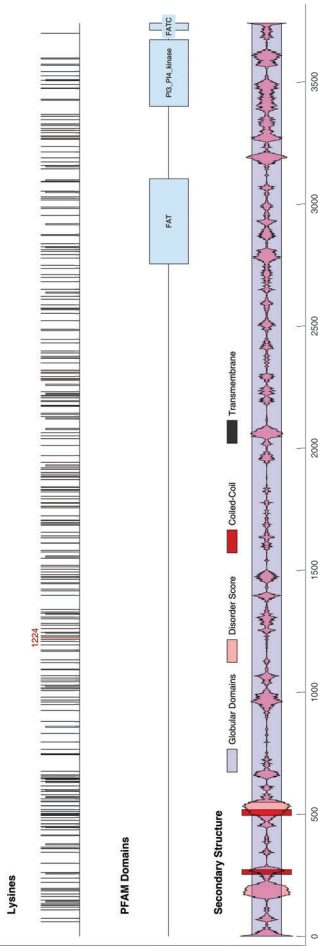




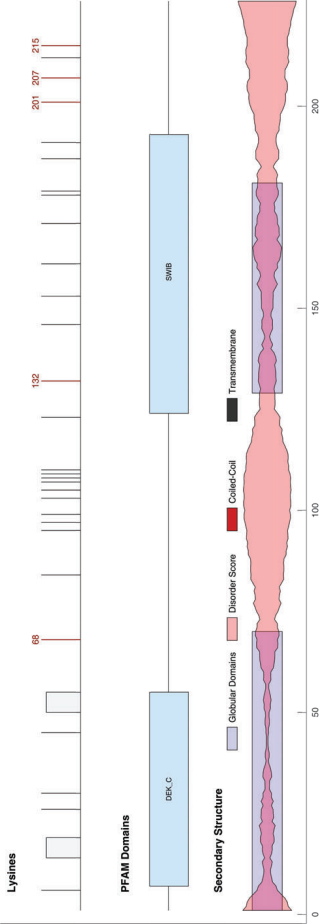
TPM2 (sp|P40414|TPM2_YEAST)



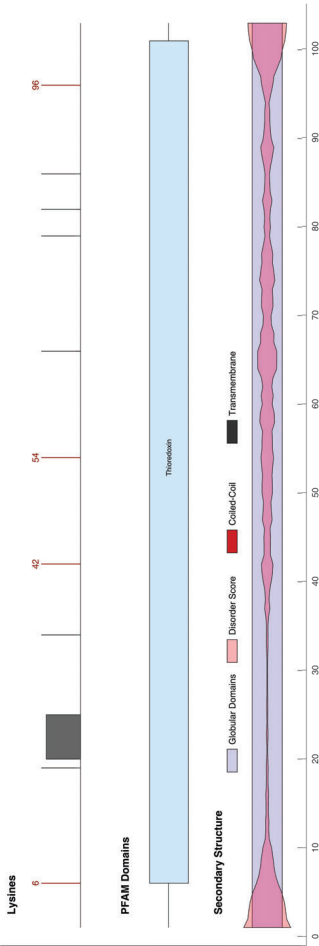
TRA1 (sp|P38811|TRA1_YEAST)



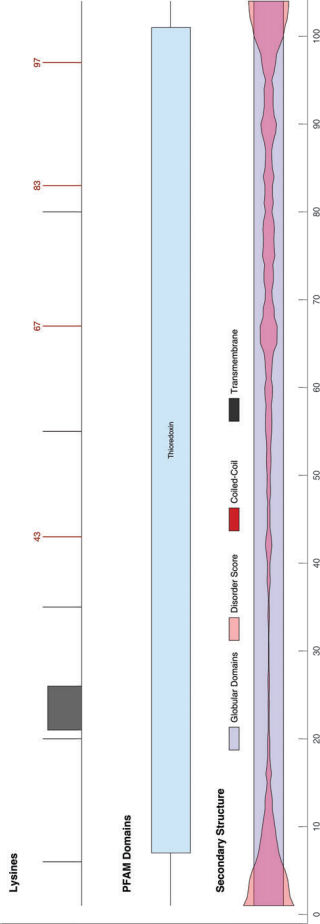
TRI1 (sp|Q05024|TRI1_YEAST)



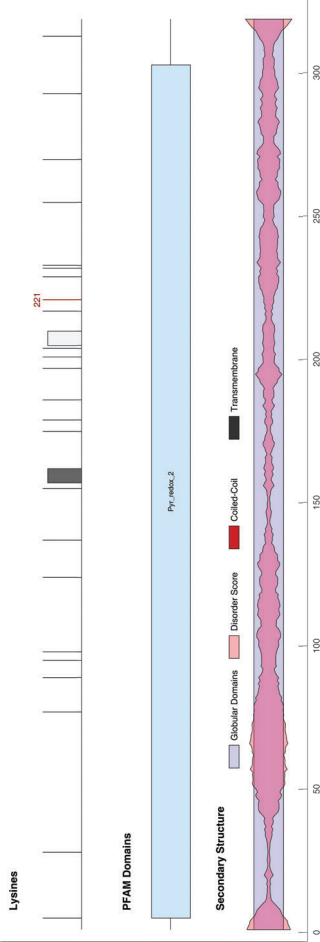
TRX1 (sp|P22217|TRX1_YEAST)



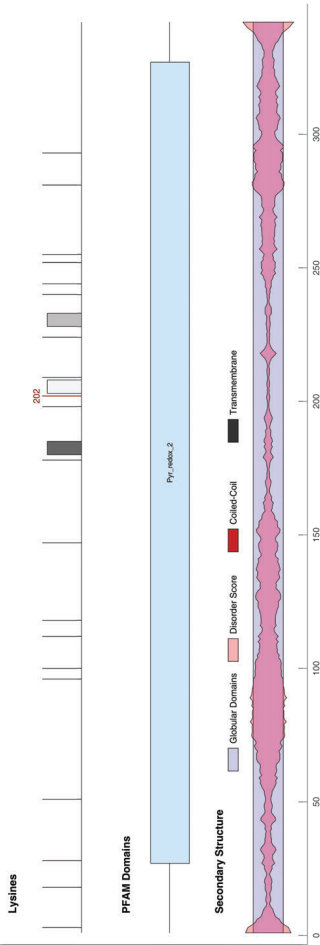
TRX2 (sp|P22803|TRX2_YEAST)



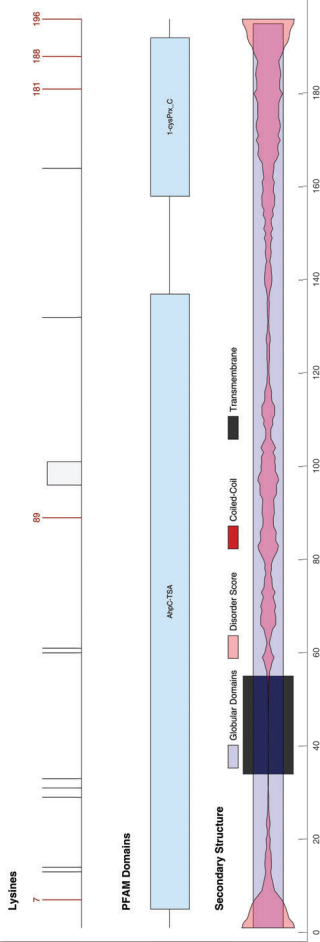
TRXB1 (sp|P2509|TRXB1_YEAST)

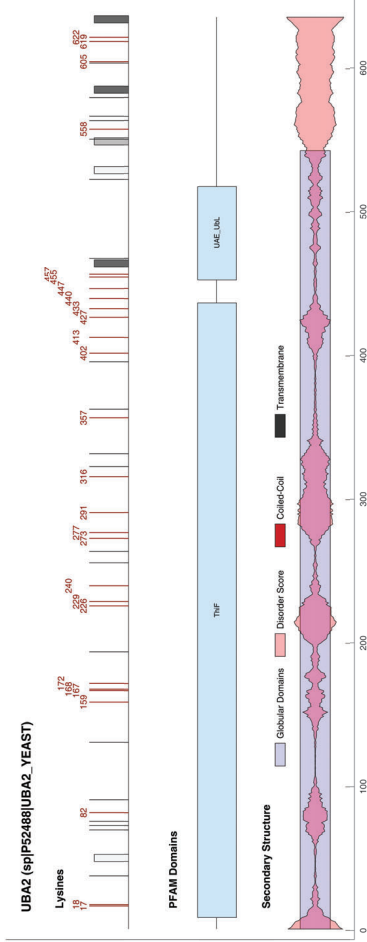
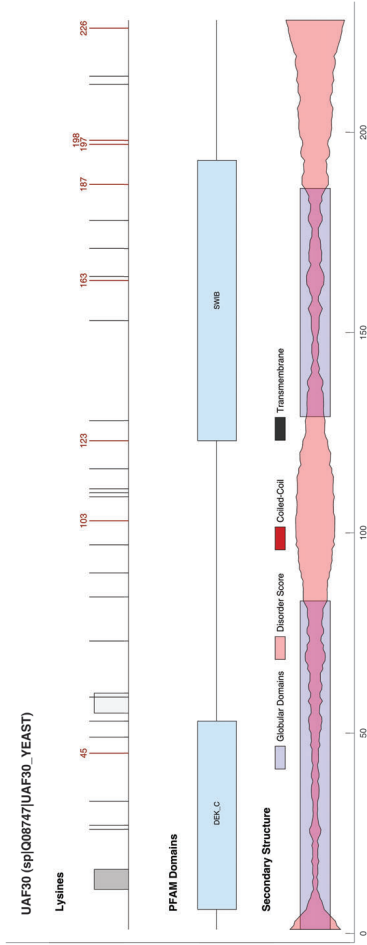
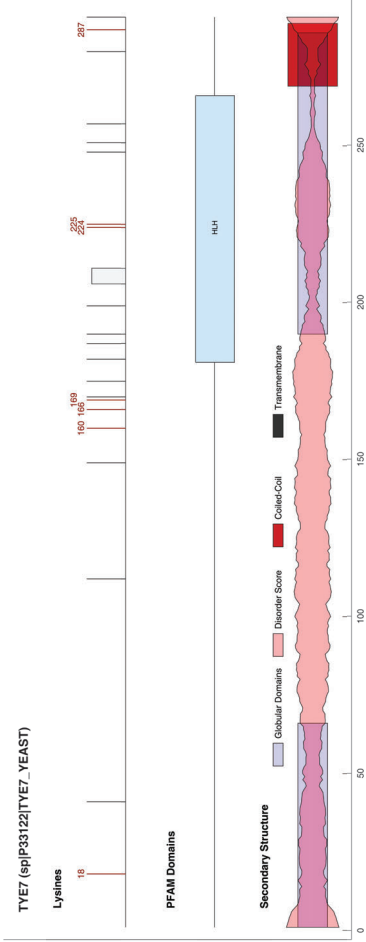
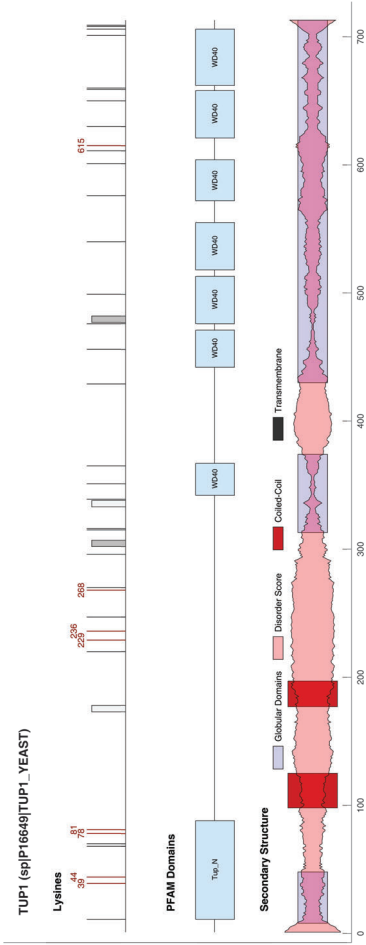


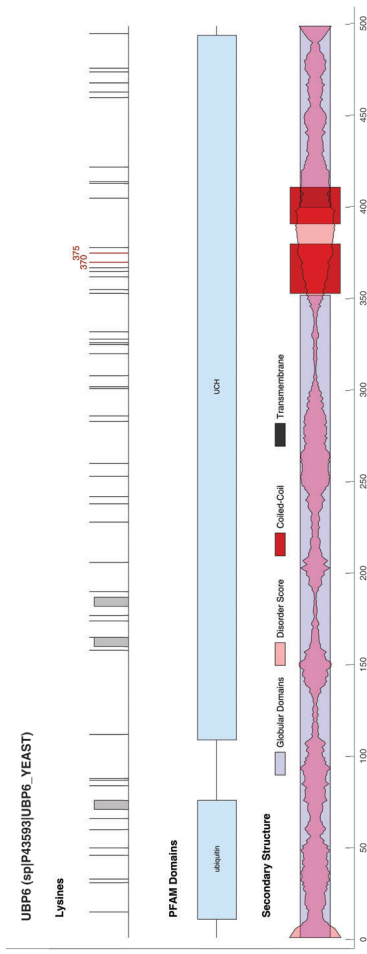
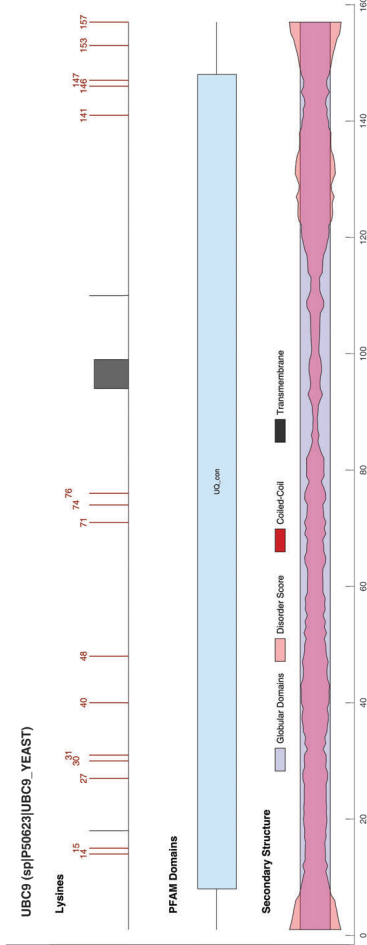
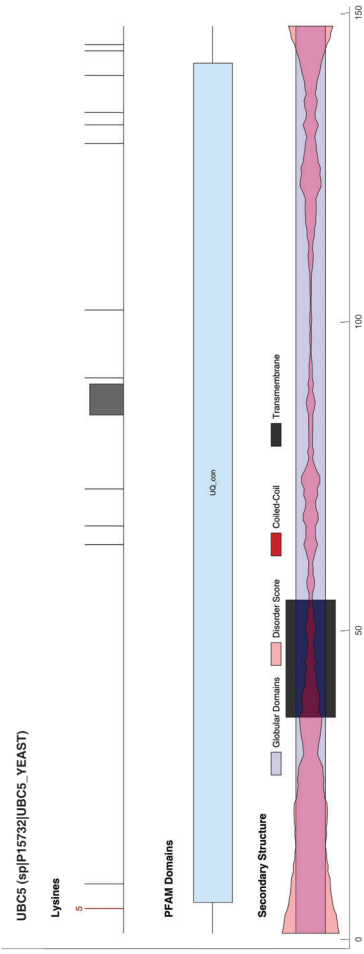
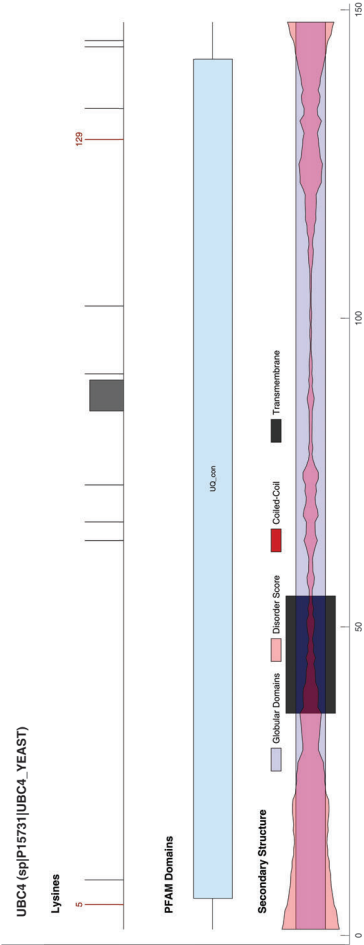
TRXB2 (sp|P38816|TRXB2_YEAST)

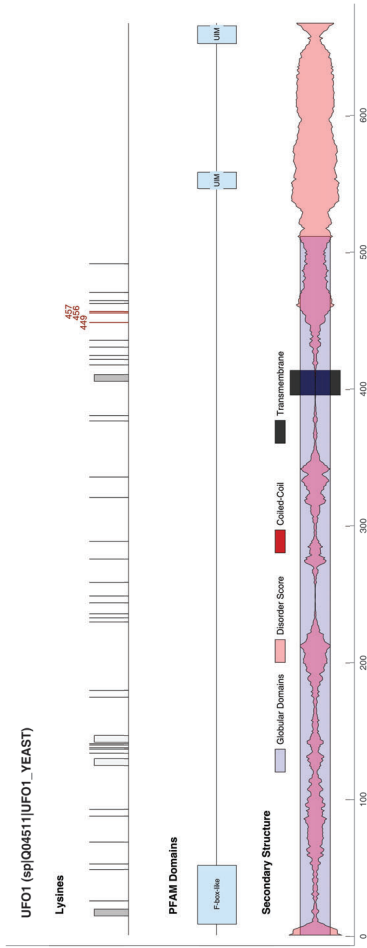
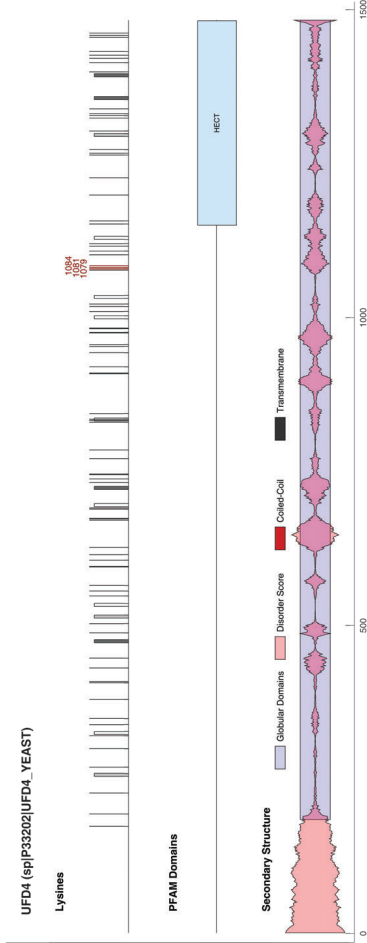
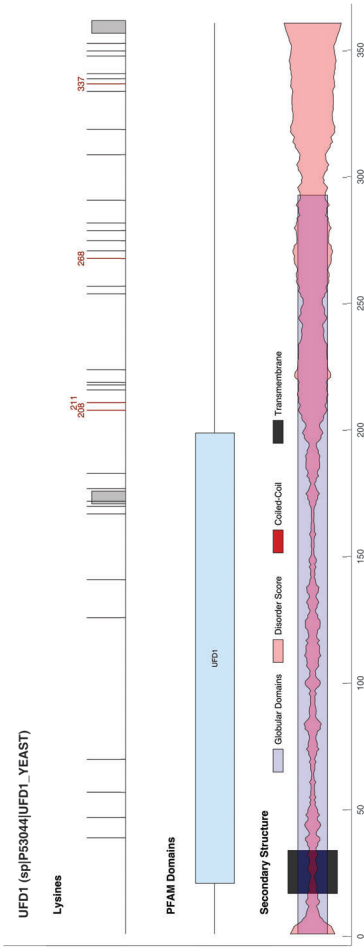
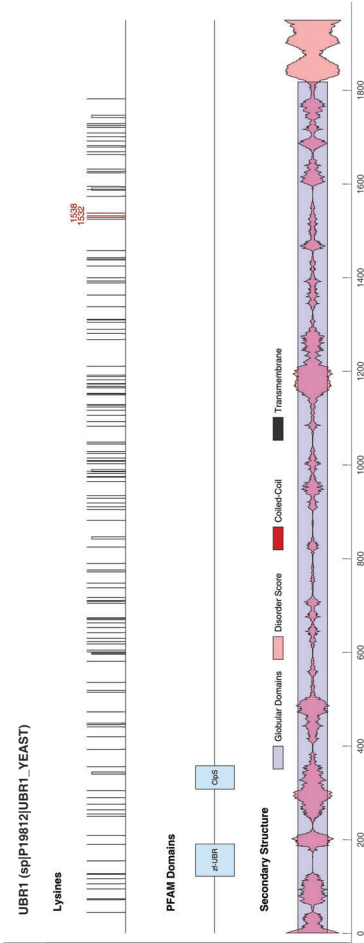


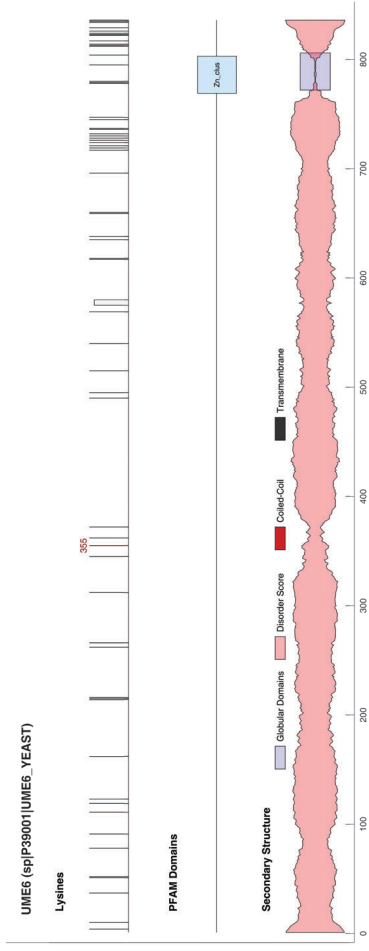
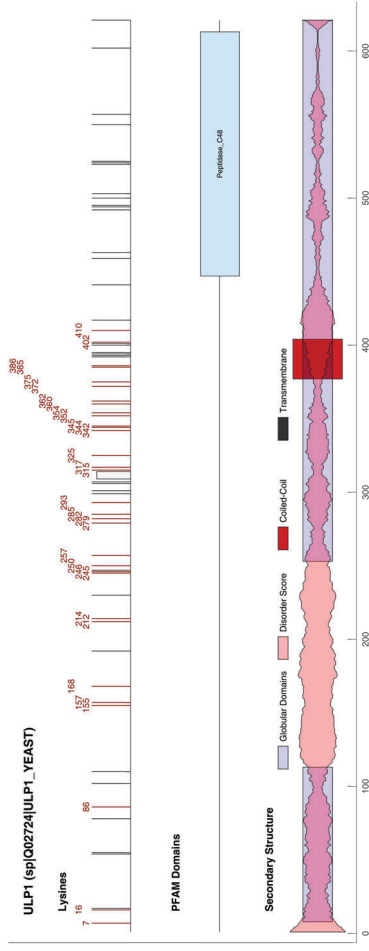
TSA1 (sp|P34760|TSA1_YEAST)

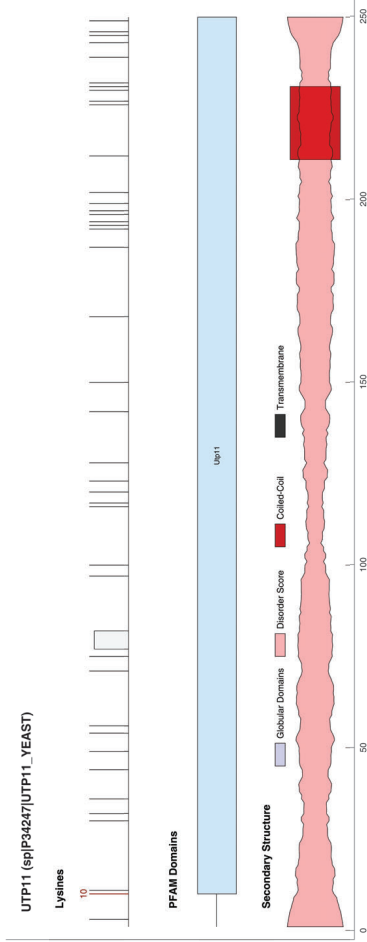
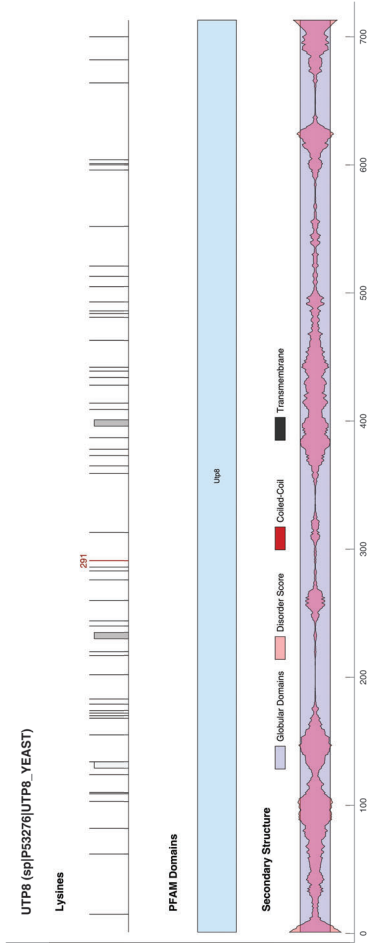
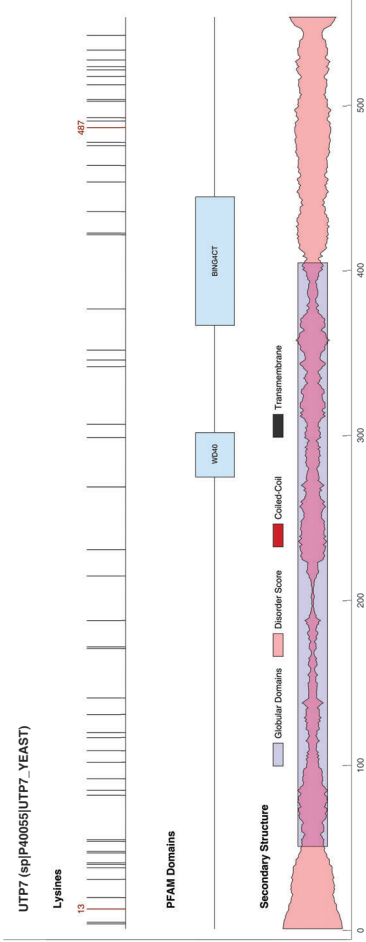
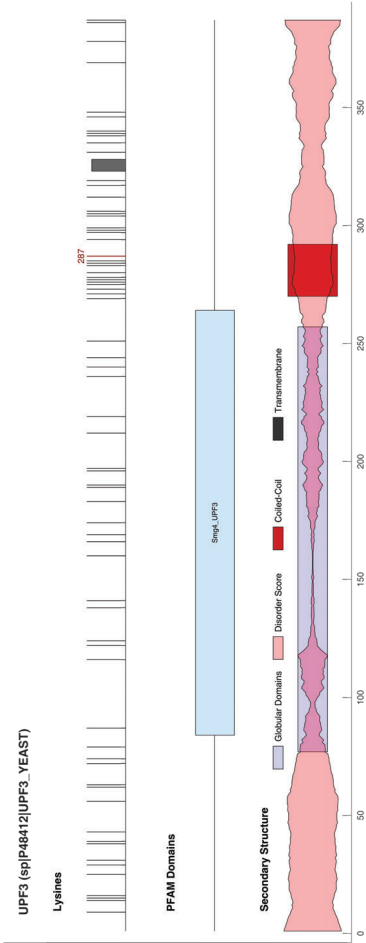


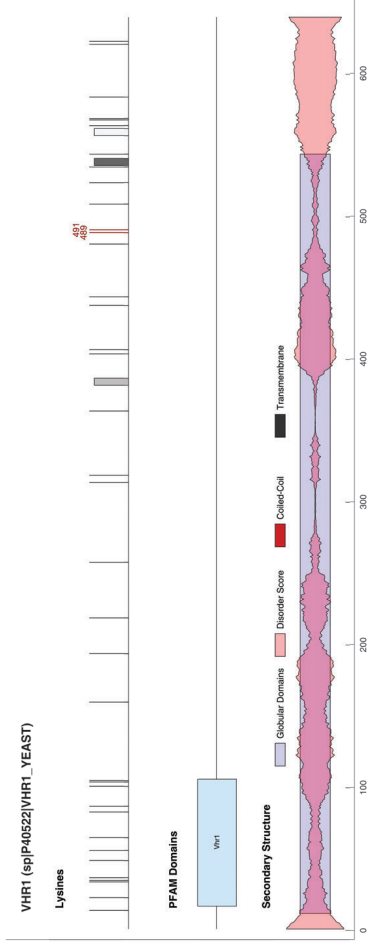
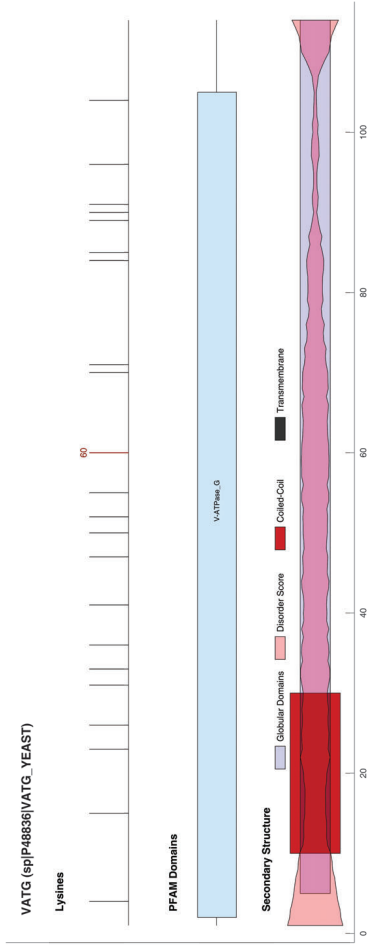
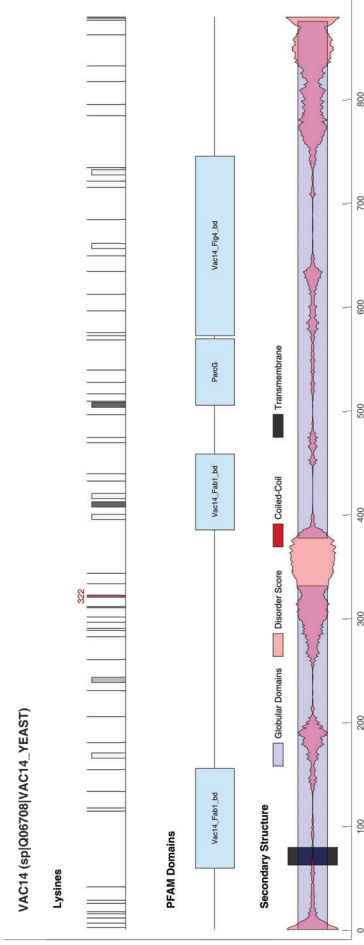
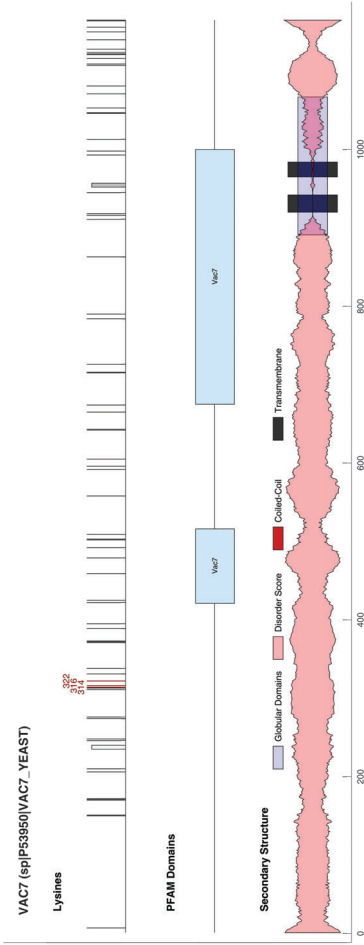


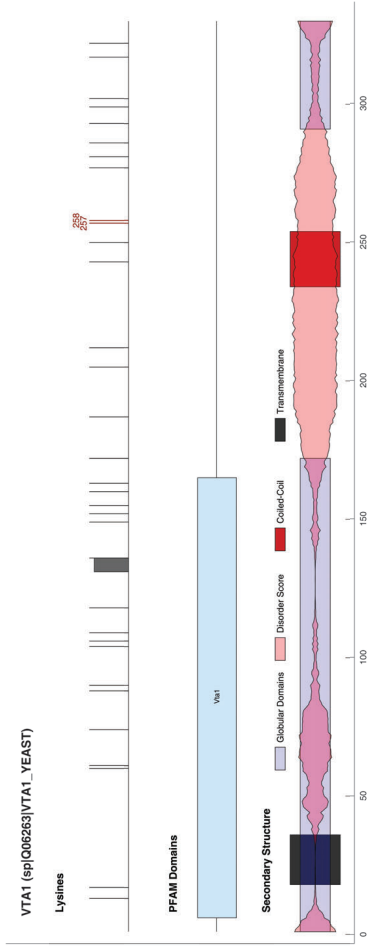
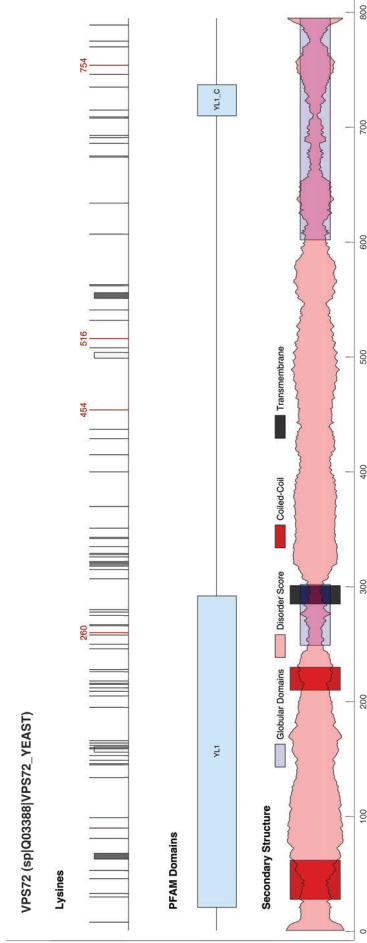
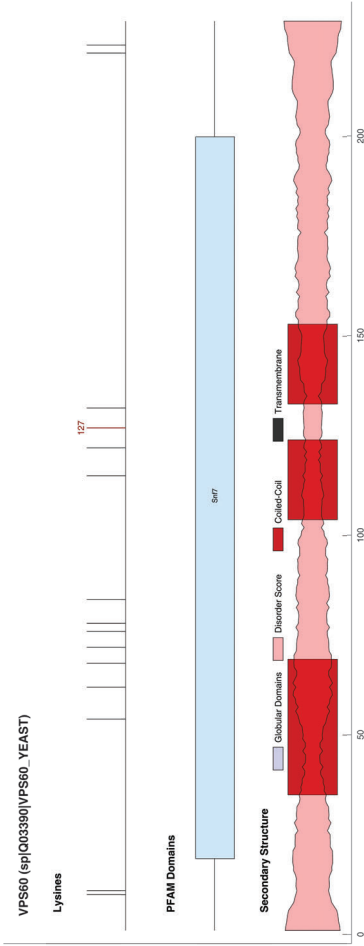
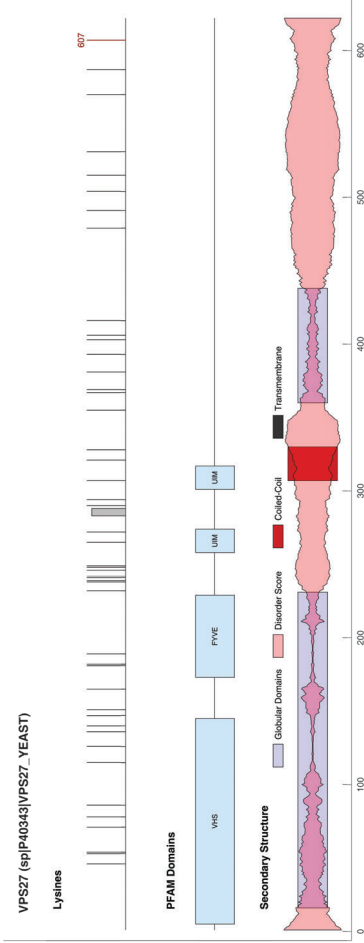


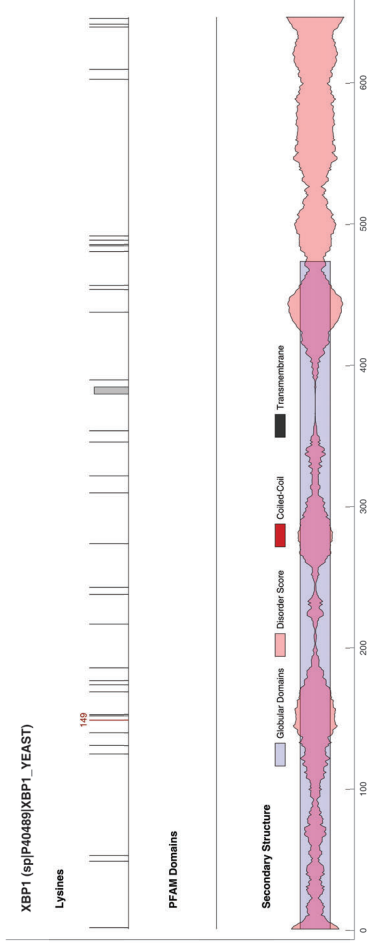
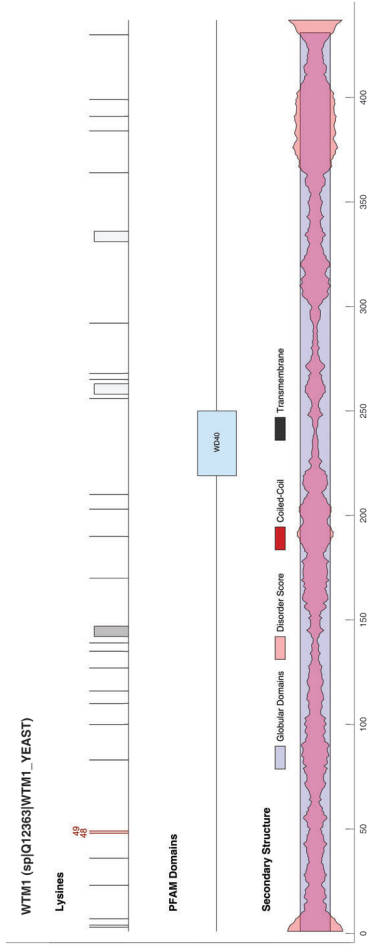
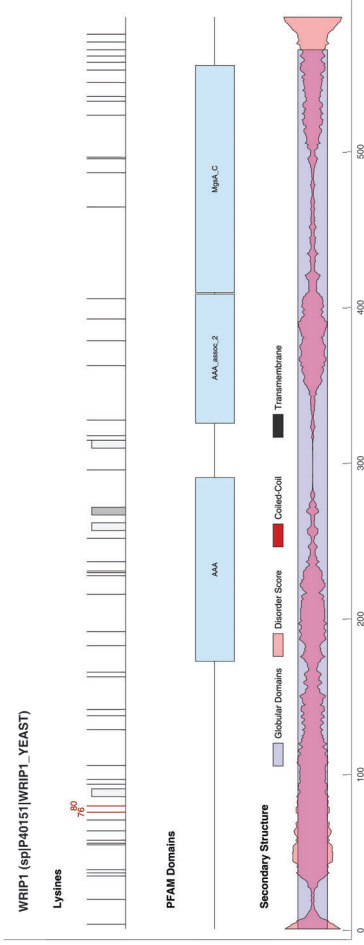
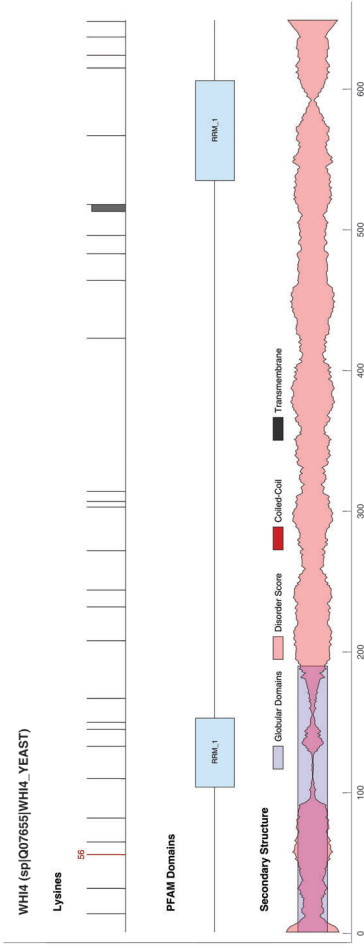




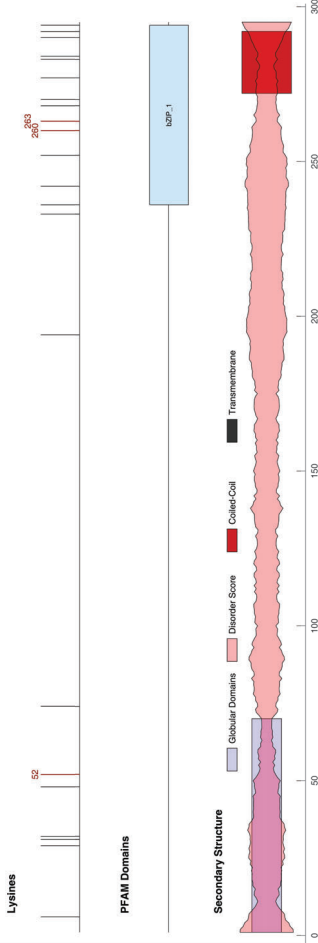




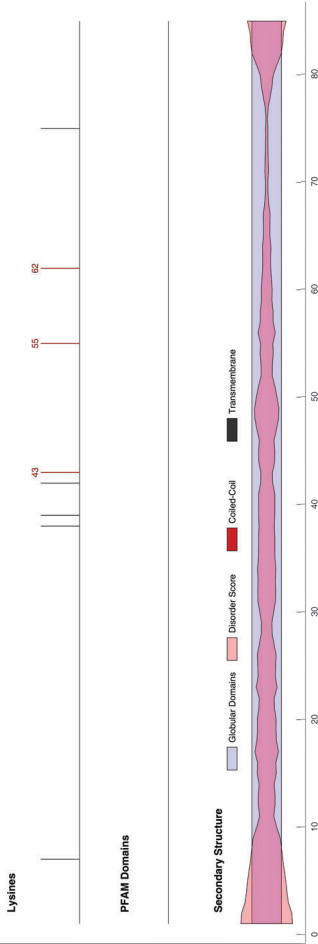




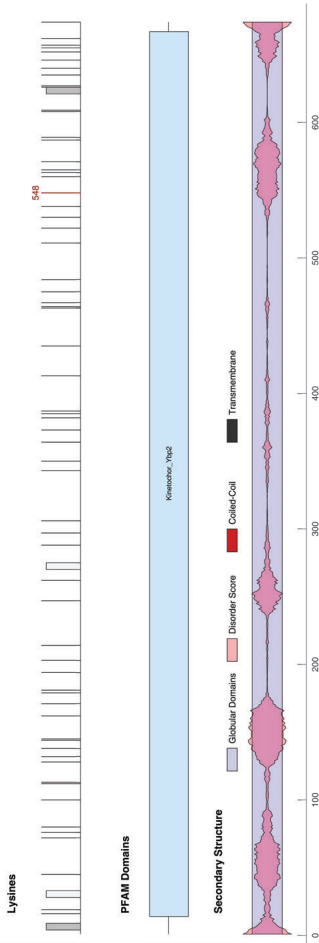
YAP4 (sp|P40917|YAP4_YEAST)



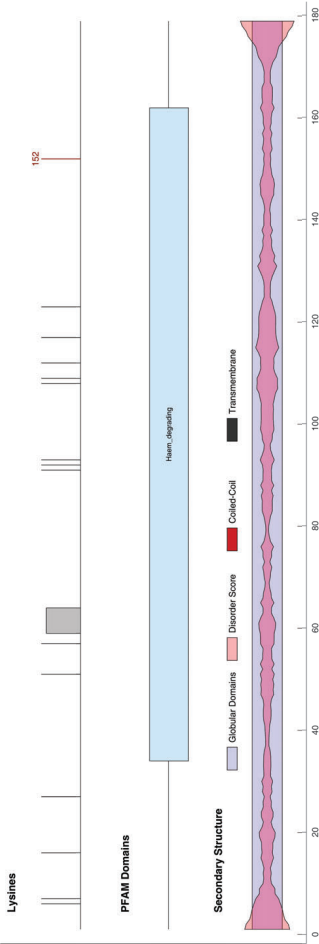
YB085 (sp|C43137|YB085_YEAST)



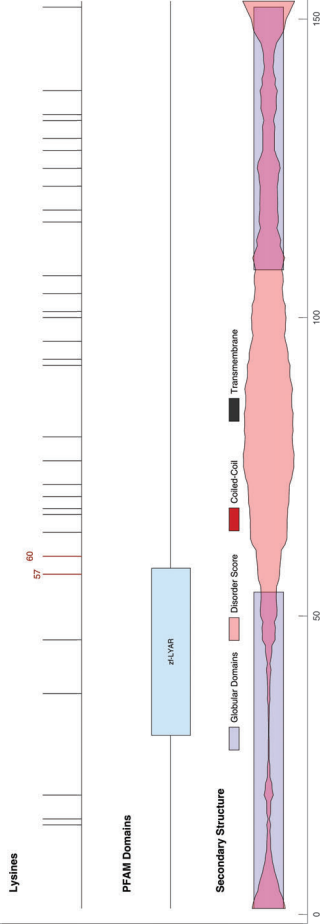
YBP1 (sp|P38315|YBP1_YEAST)



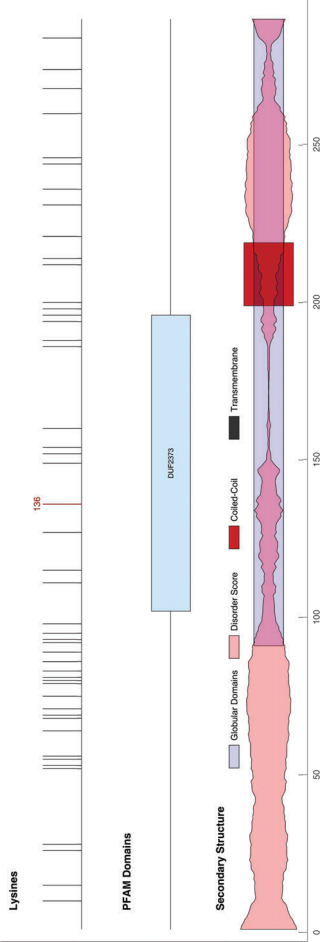
YBY7 (sp|P38276|YBY7_YEAST)



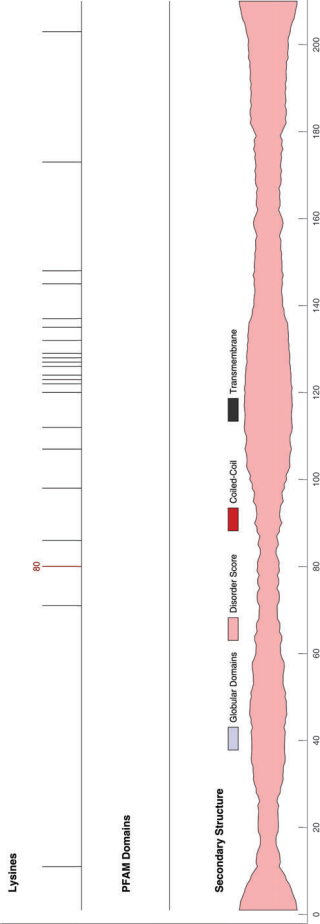
YC16 (sp|P37263|YC16_YEAST)



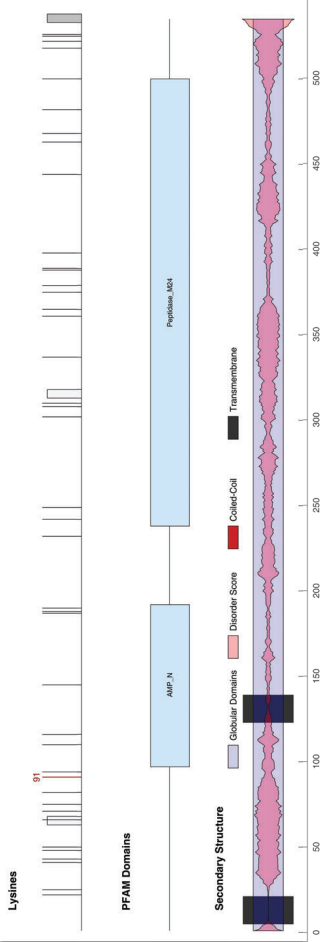
YC06 (sp|P25617|YC06_YEAST)

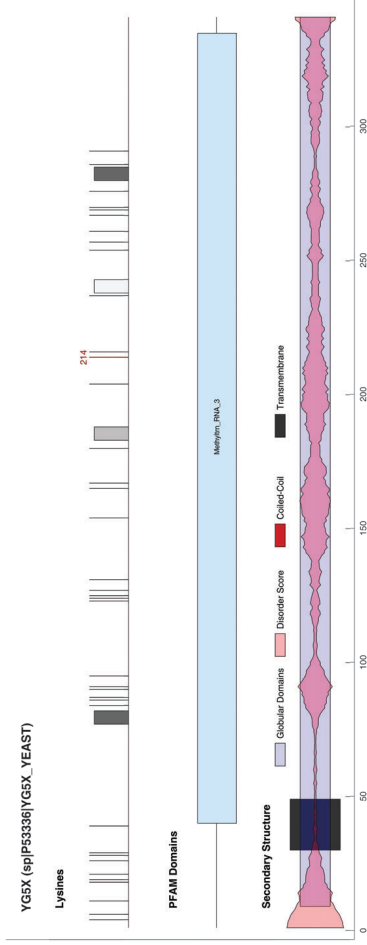
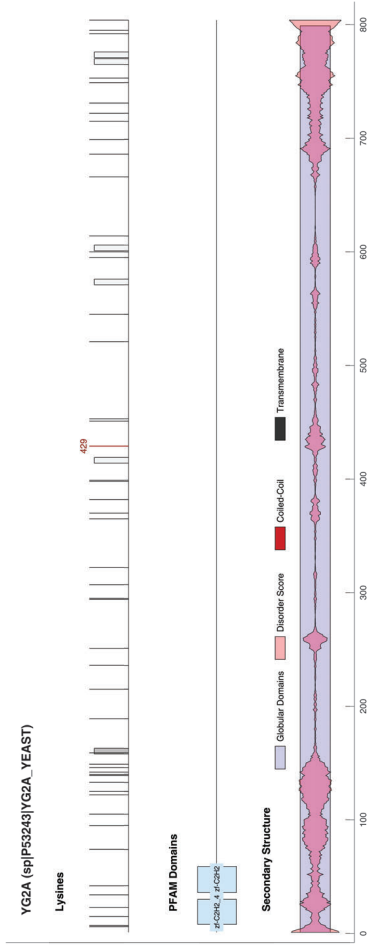
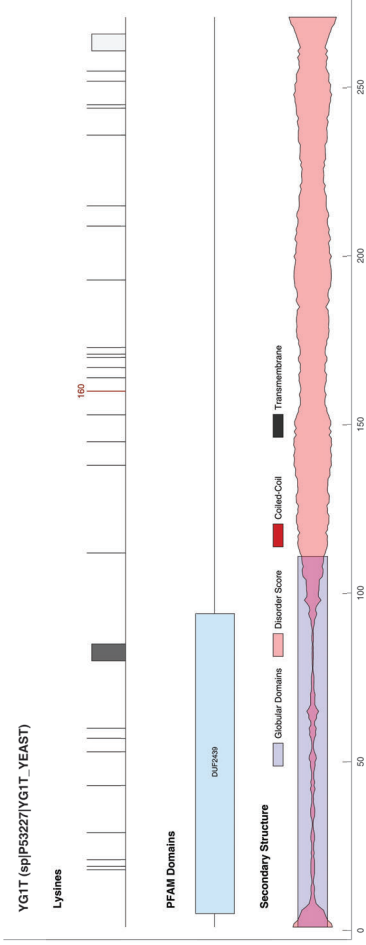
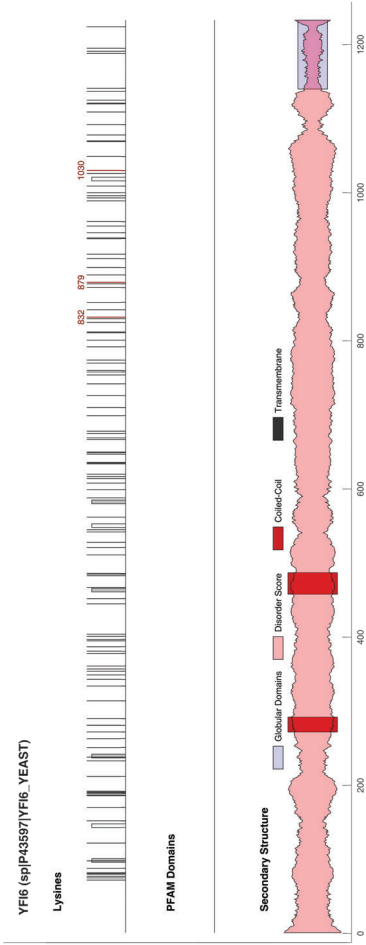


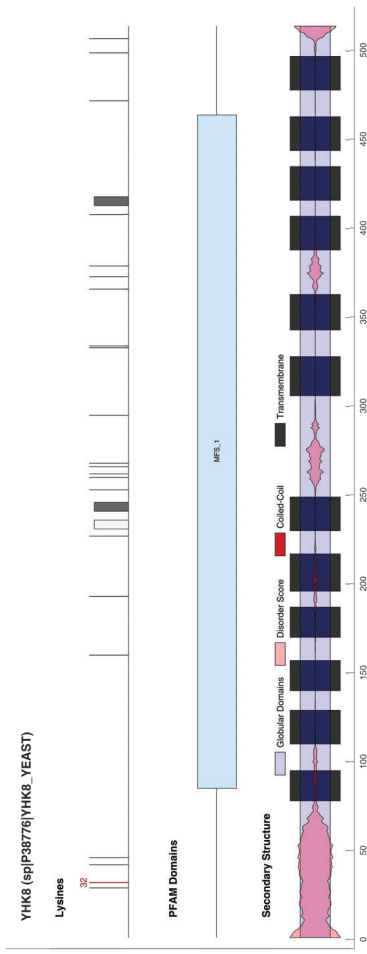
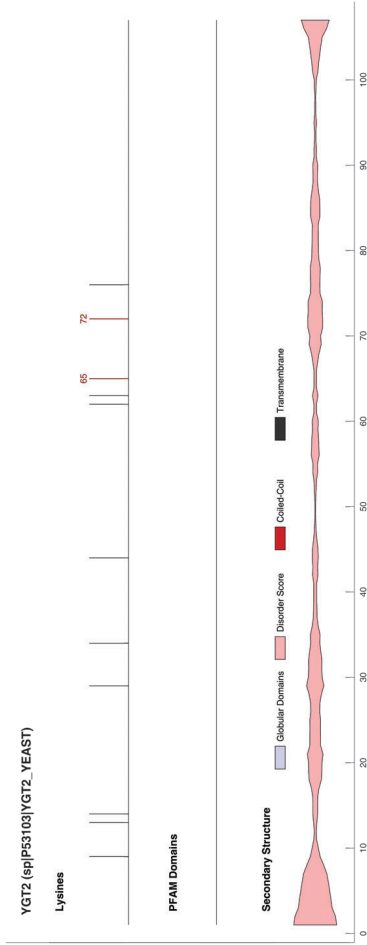
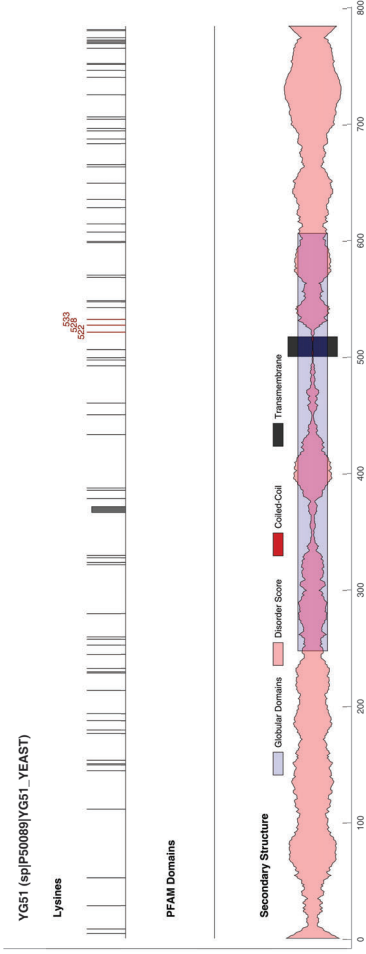
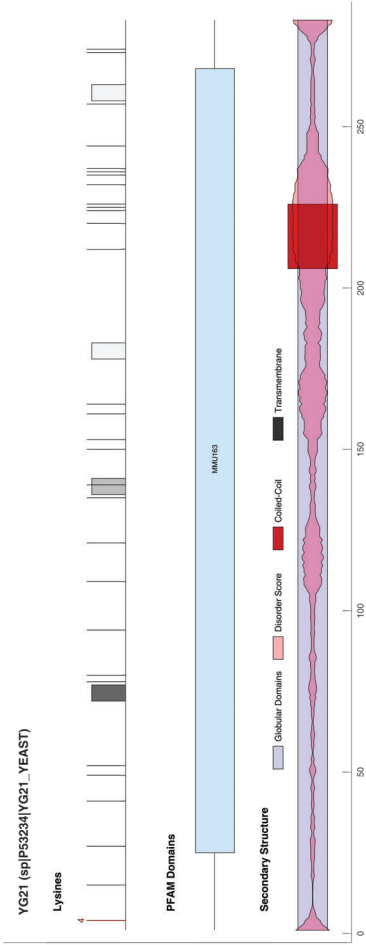
YEQ9 (sp|P40052|YEQ9_YEAST)

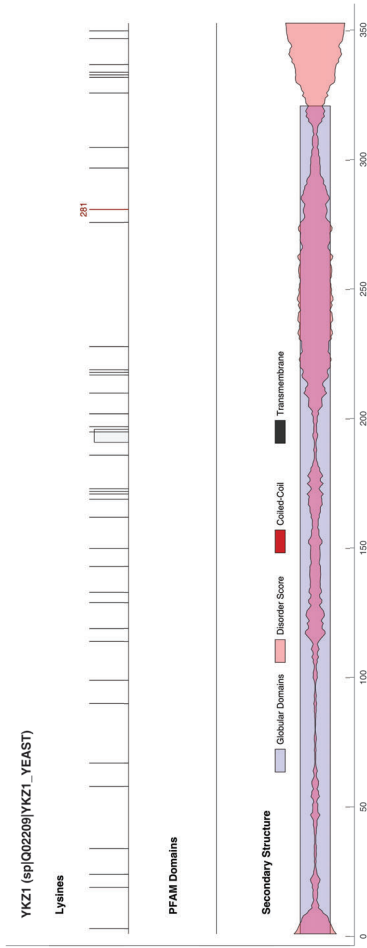
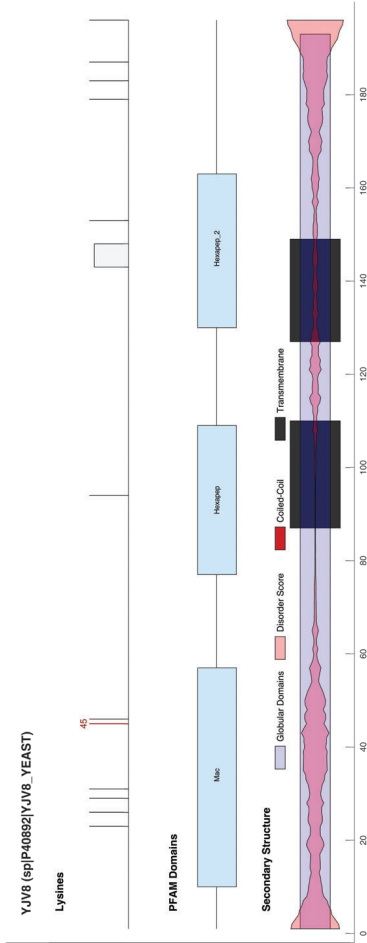
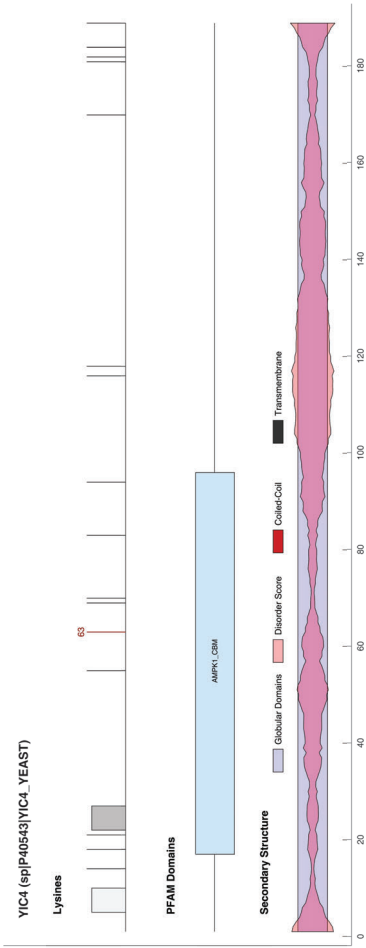
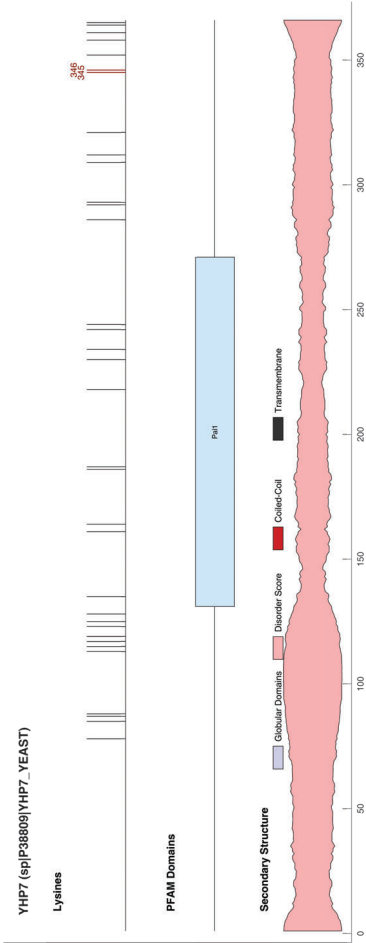


YFH6 (sp|P43590|YFH6_YEAST)

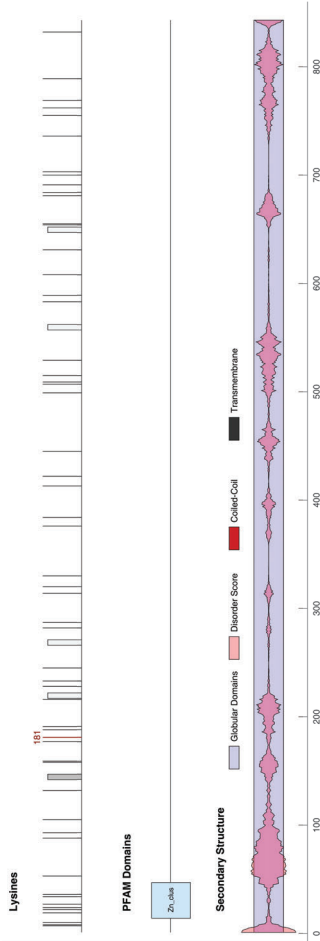




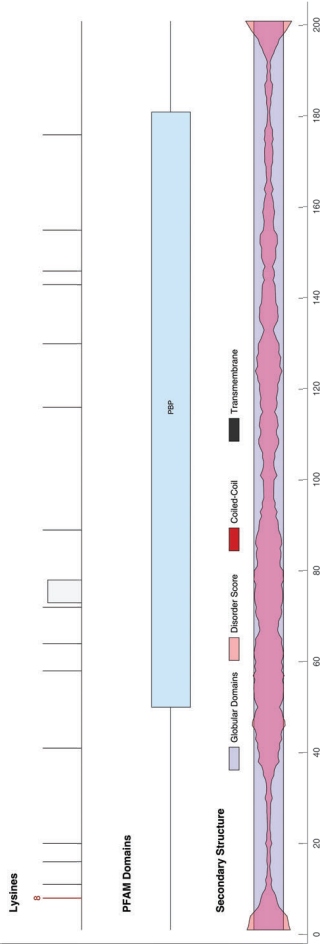




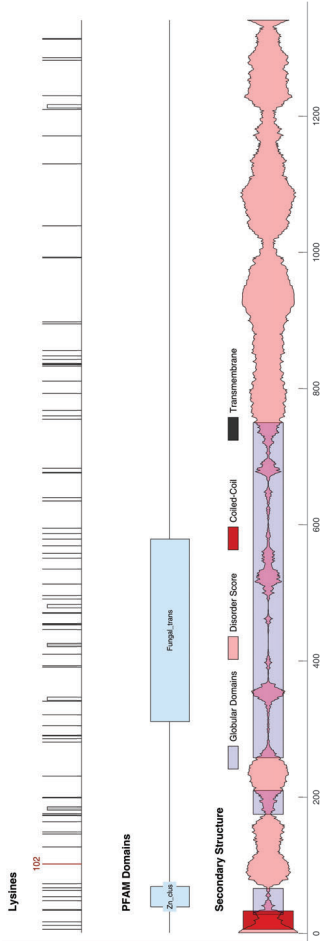
YL054 (sp|Q12244|YL054_YEAST)



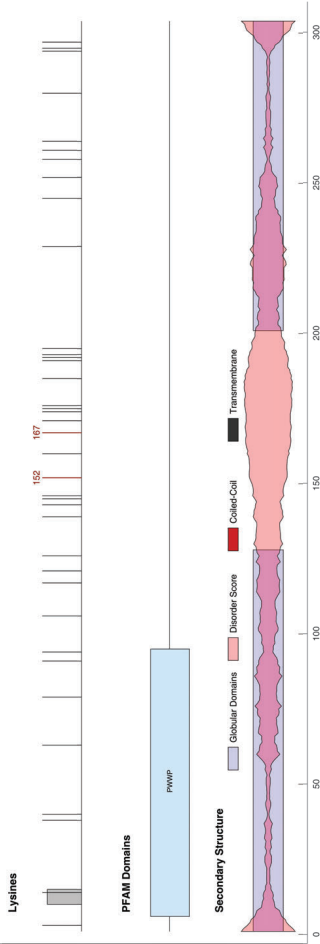
YL179 (sp|Q06252|YL179_YEAST)



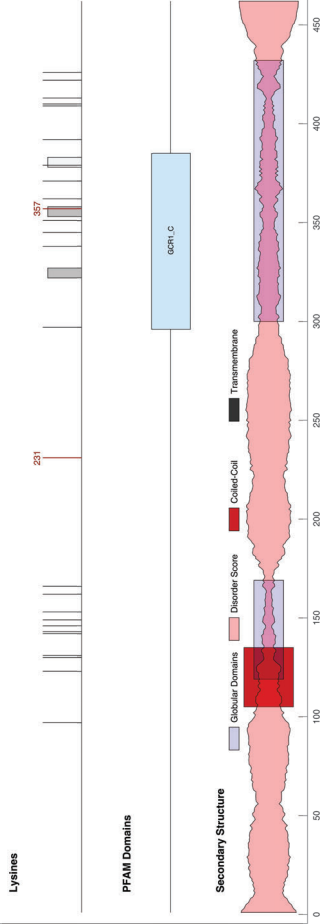
YL278 (sp|Q05854|YL278_YEAST)



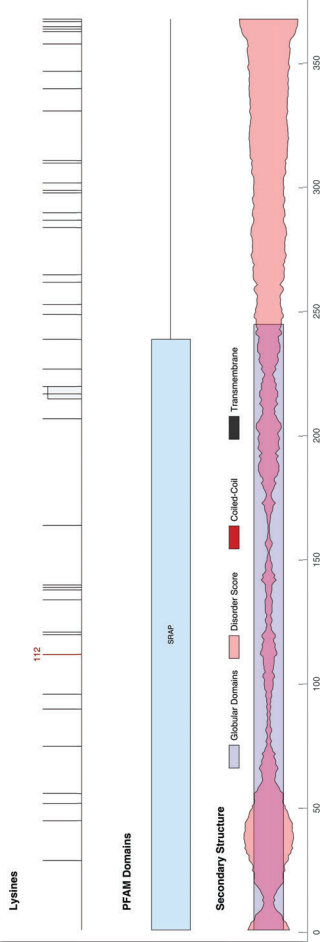
YL455 (sp|Q06188|YL455_YEAST)



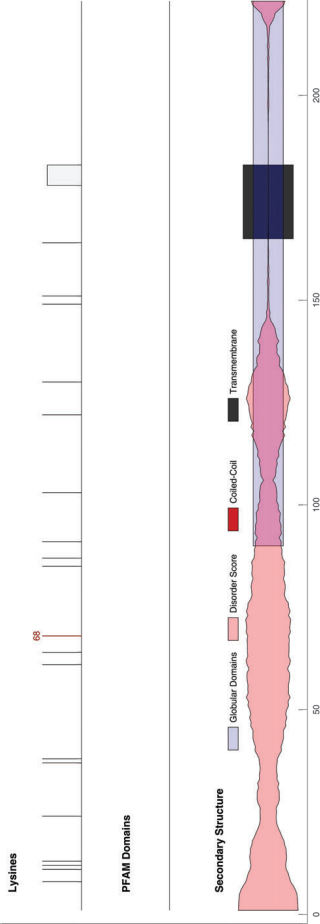
YM01 (sp|Q04461|YM01_YEAST)



YM04 (sp|Q04471|YM04_YEAST)



YM24 (sp|P40218|YM24_YEAST)



YN034 (sp|Q3E841|YN034_YEAST)

