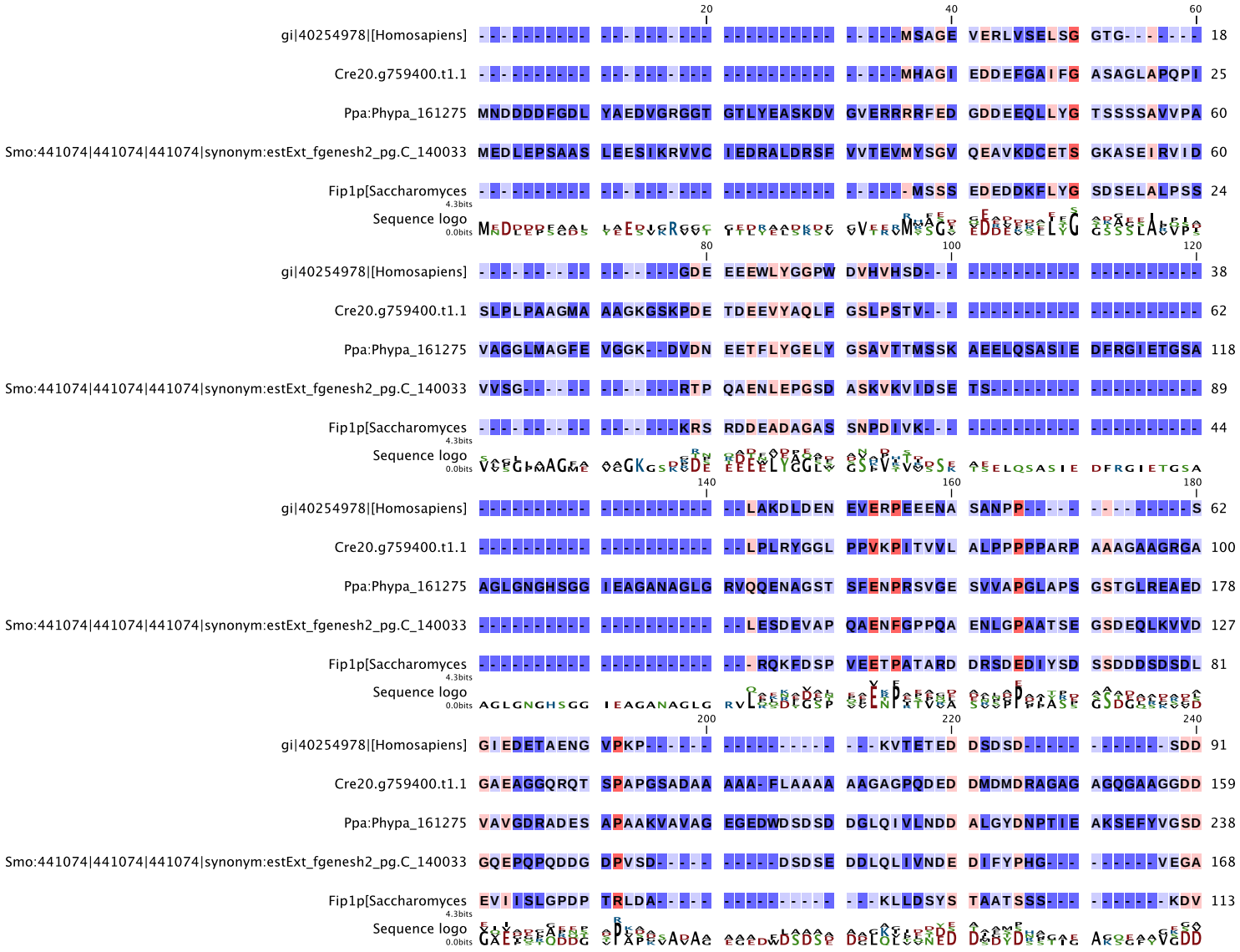
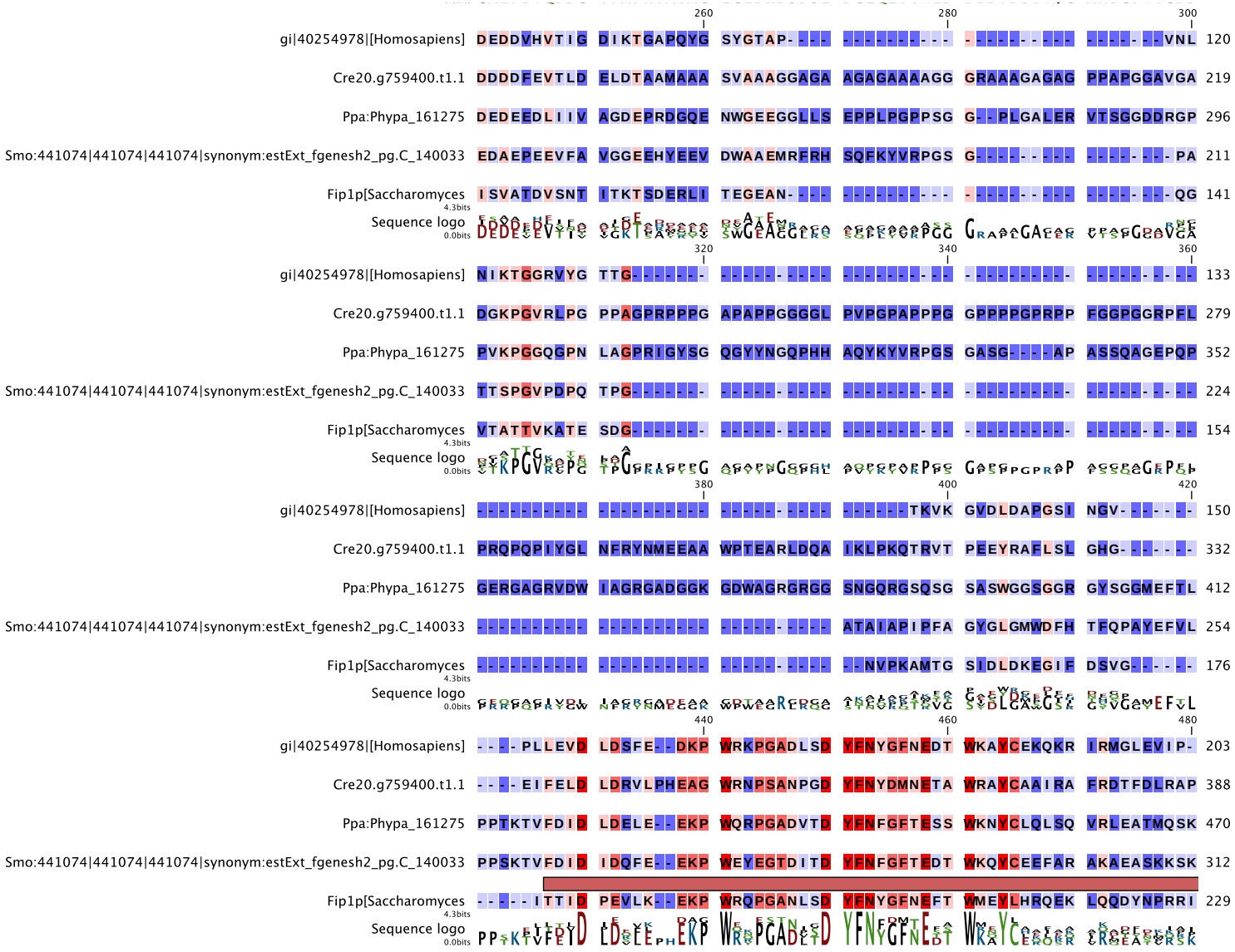
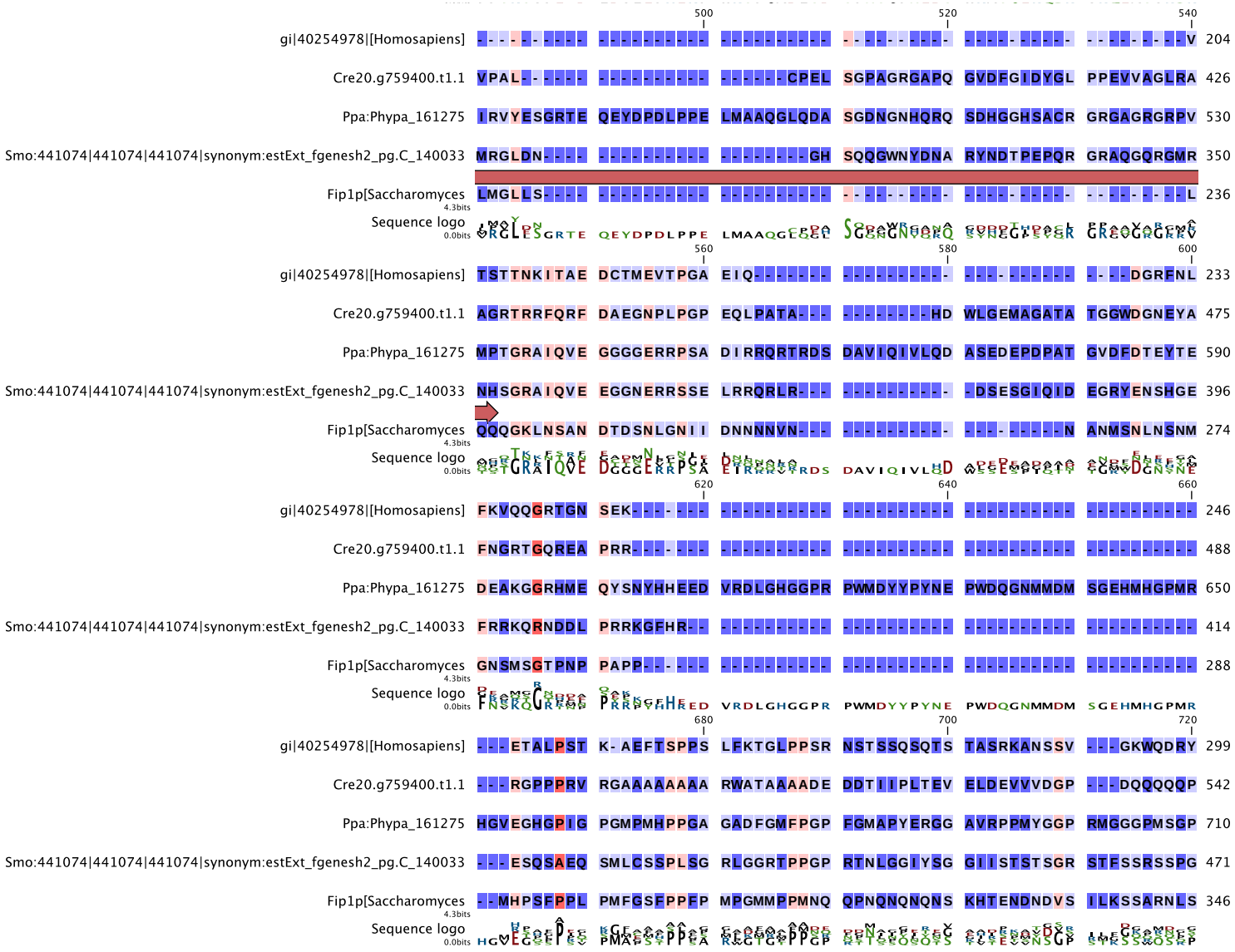
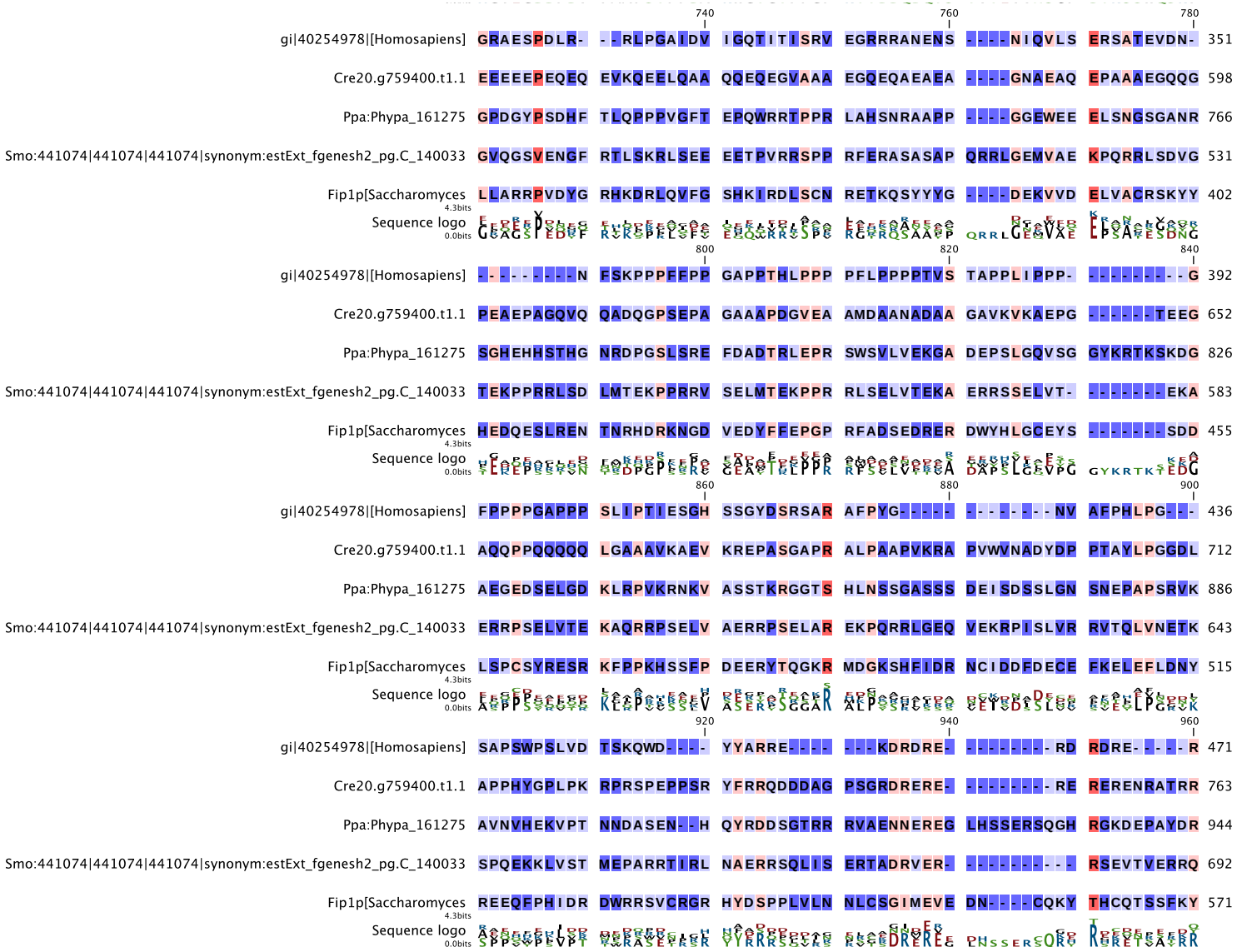
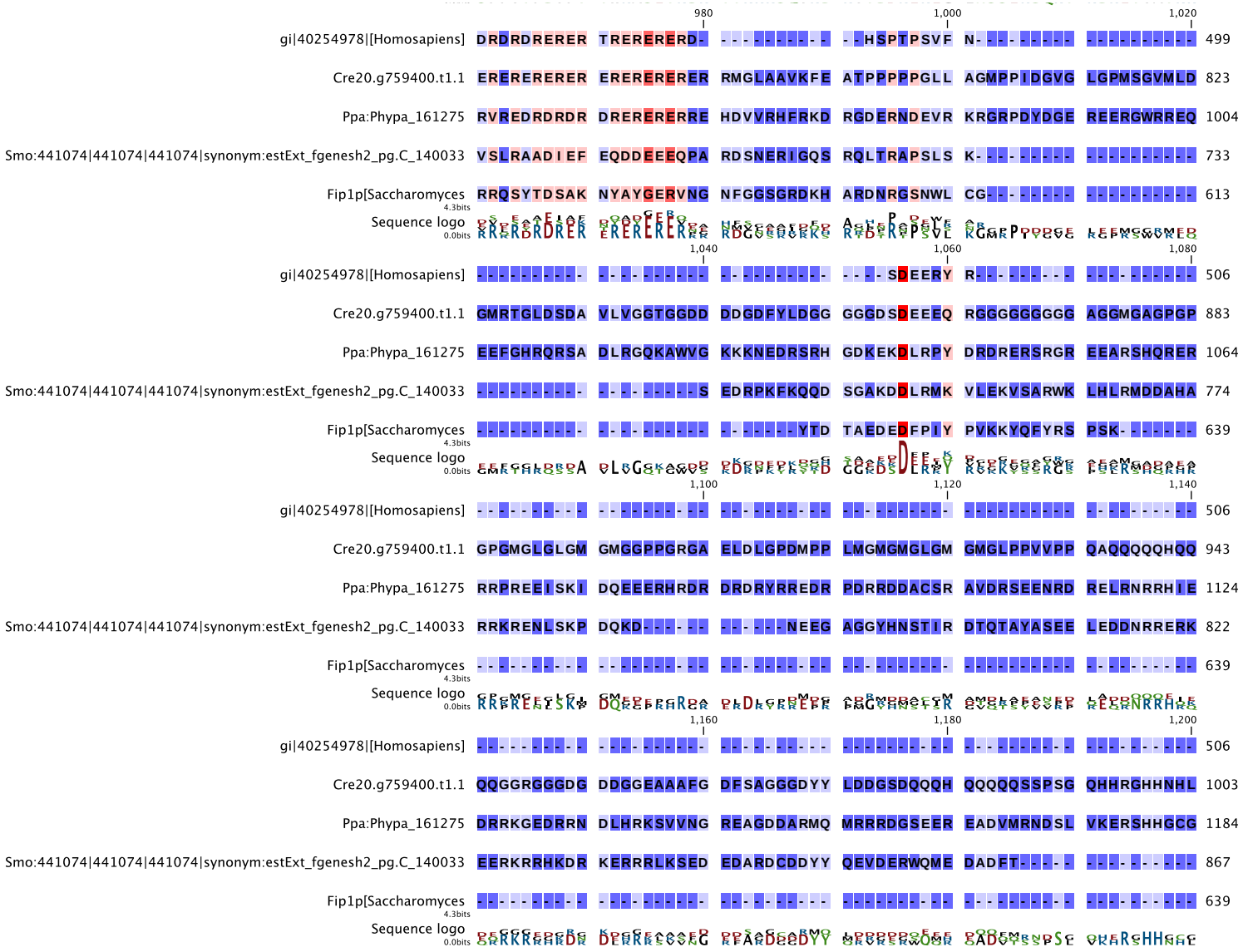
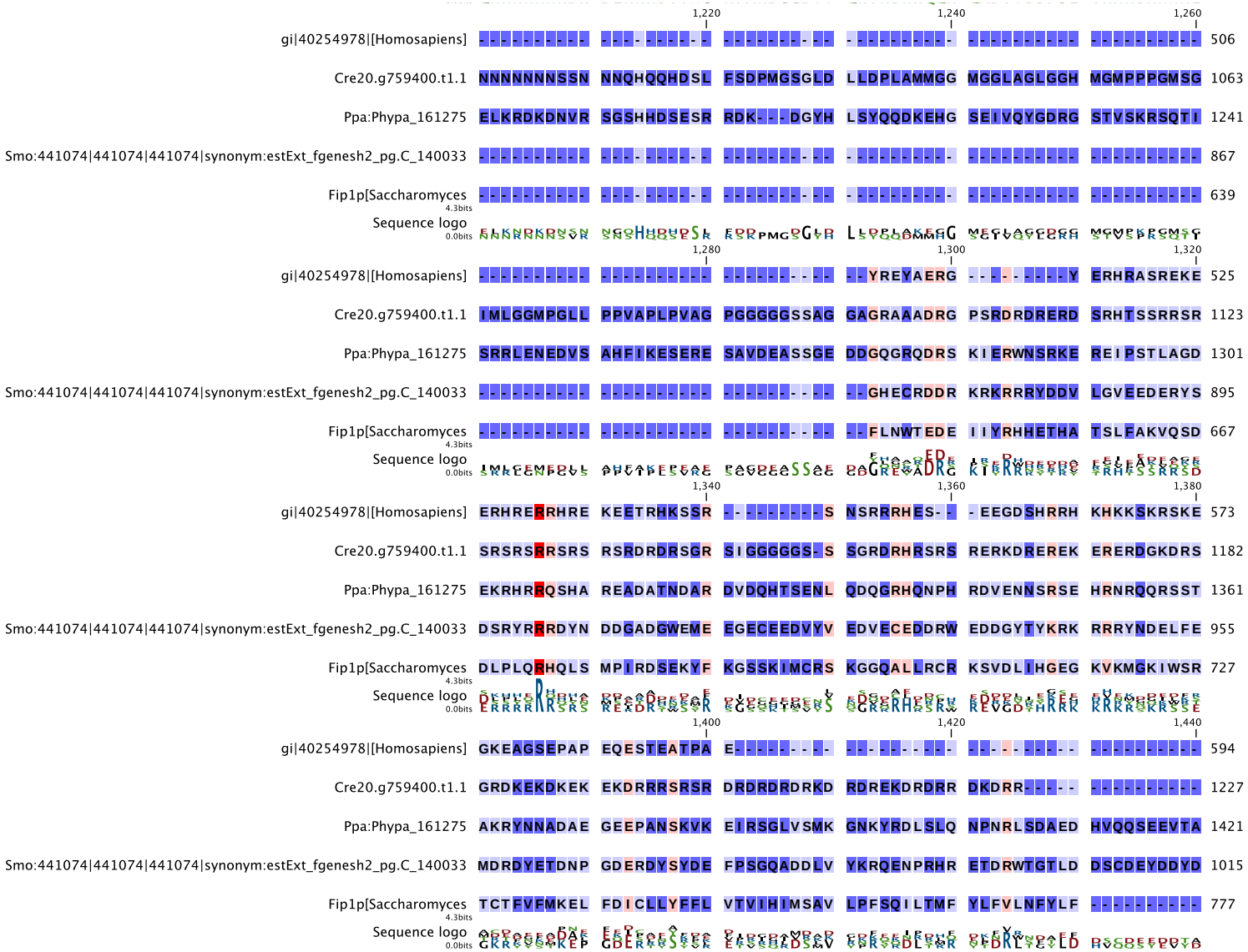

gi|40254978|[Homosapiens]
Cre20.g759400.t1.1
Ppa:Phypa_161275
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033
Fip1p[Saccharomyces
Sequence logo
4.3bits
0.0bits

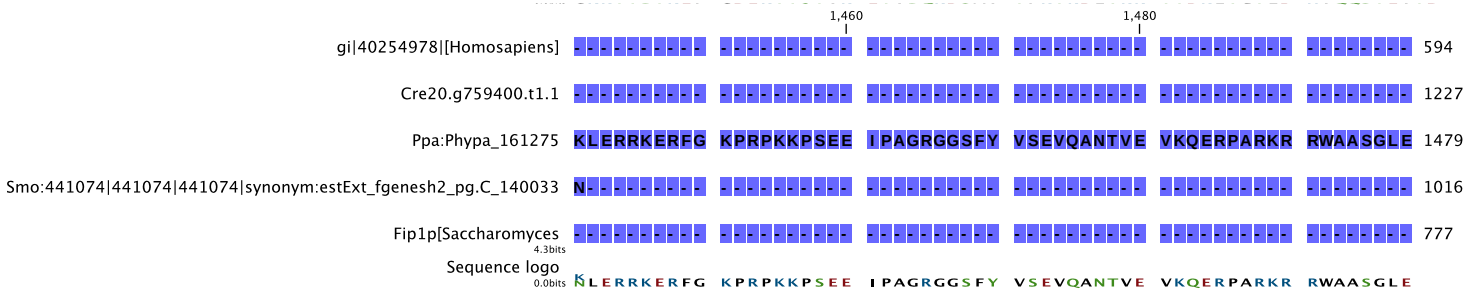

gi|40254978|[Homosapiens]
Cre20.g759400.t1.1
Ppa:Phypa_161275
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033
Fip1p[Saccharomyces
Sequence logo
4.3bits
0.0bits


500 520 540
gi|40254978|[Homosapiens] 204
Cre20.g759400.t1.1 426
Ppa:Phypa_161275 530
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 350
Fip1p[Saccharomyces 236
4.3bits
Sequence logo
0.0bits
560 580 600
gi|40254978|[Homosapiens] 233
Cre20.g759400.t1.1 475
Ppa:Phypa_161275 590
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 396
Fip1p[Saccharomyces 274
4.3bits
Sequence logo
0.0bits
620 640 660
gi|40254978|[Homosapiens] 246
Cre20.g759400.t1.1 488
Ppa:Phypa_161275 650
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 414
Fip1p[Saccharomyces 288
4.3bits
Sequence logo
0.0bits
680 700 720
gi|40254978|[Homosapiens] 299
Cre20.g759400.t1.1 542
Ppa:Phypa_161275 710
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 471
Fip1p[Saccharomyces 346
4.3bits
Sequence logo
0.0bits


740 760 780
gi|40254978|[Homosapiens] GRAESPDLR- --RLPGAIDV IGQTITISRV EGRRRANENS ----NIQVLS ERSATEVDN- 351
Cre20.g759400.t1.1 EEEEEPEQEQ EVKQEELQAA QQEQEGVAAA EGQEQAEAEA ----GNAEAQ EPAAAEGQQG 598
Ppa:Phypa_161275 GPDGYPSDHF TLQPPPVGFT EPQWRRTPPR LAHSNRAAPP ----GGEWEE ELSNGSGANR 766
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 GVQGSVENGF RTLSKRLSEE EETPVRRSPP RFERASASAP QRRLGEMVAE KPQRRLSDVG 531
Fip1p[Saccharomyces LLARRPVDYG RHKDRLQVFG SHKIRDLSCN RETKQSYYYG ----DEKVVD ELVACRSKYY 402
4.3bits
Sequence logo
0.0bits
800 820 840
gi|40254978|[Homosapiens] ---------N FSKPPPFFPP GAPPTHLPPP PFLPPPPTVS TAPPLIPPP- -------G 392
Cre20.g759400.t1.1 PEAEPAGQVQ QADQGPSEPA GAAAPDGVEA AMDAANADAA GAVKVKAEPG ------TEEG 652
Ppa:Phypa_161275 SGHEHHSTHG NRDPGSLSRE FDADTRLEPR SWSVLVEKGA DEPSLGQVSG GYKRTKSKDG 826
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 TEKPPRRLSD LMTEKPPRRV SELMTEKPPR RLSELVTEKA ERRSSELVT- -------EKA 583
Fip1p[Saccharomyces HEDQESLREN TNRHDRKNGD VEDYFFEPGP RFADSEDRER DWYHLGCEYS -------SDD 455
4.3bits
Sequence logo
0.0bits
860 880 900
gi|40254978|[Homosapiens] FPPPPGAPPP SLIPTIESGH SSGYDSRSAR AFPYG----- --------NV AFPHLPG--- 436
Cre20.g759400.t1.1 AQQPPQQQQQ LGAAAVKAEV KREPASGAPR ALPAAPVKRA PVWVNADYDP PTAYLPGGDL 712
Ppa:Phypa_161275 AEGEDSELGD KLRPVKRNKV ASSTKRGGTS HLNSSGASSS DEISDSSLGN SNEPAPSRVK 886
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 ERRPSELVTE KAQRRPSELV AERRPSELAR EKPQRRLGEQ VEKRPISLVR RVTQLVNETK 643
Fip1p[Saccharomyces LSPCSYRESR KFPPKHSSFP DEERYTQGKR MDGKSHFIDR NCIDDFDECE FKELEFLDNY 515
4.3bits
Sequence logo
0.0bits
920 940 960
gi|40254978|[Homosapiens] SAPSWPSLVD TSKQWD---- YYARRE---- ---KDRDRE- --------RD RDRE----R 471
Cre20.g759400.t1.1 APPHYGPLPK RPRSPEPPSR YFRRQDDDAG PSGRDRERE- --------RE RERENRATRR 763
Ppa:Phypa_161275 AVNVHEKVPT NNDASEN--H QYRDDSGTRR RVAENNEREG LHSSERSQGH RGKDEPAYDR 944
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 SPQEKKLVST MEPARRTIRL NAERRSQLIS ERTADRVER- --------- RSEVTVERRQ 692
Fip1p[Saccharomyces REEQFPHIDR DWRRSVCRGR HYDSPPLVLN NLCSGIMEVE DN----CQKY THCQTSSFKY 571
4.3bits
Sequence logo
0.0bits


980
1,000
1,020
gi|40254978|[Homosapiens] DRDRDRERER TRERERERD- ---------- -HSPTPSVF N-------- ---------- 499
Cre20.g759400.t1.1 ERERERERER ERERERERER RMGLAAVKFE ATPPPPPGLL AGMPPIDGVG LGPMSGVMLD 823
Ppa:Phypa_161275 RVREDRDRDR DRERERERRE HDVVRHFRKD RGDERNDEVR KRGRPDYDGE REERGWRREQ 1004
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 VSLRAADIEF EQDDEEEQPA RDSNERIGQS RQLTRAPSLS K-------- ---------- 733
Fip1p[Saccharomyces RRQSYTDSAK NYAYGERVNG NFGGSGRDKH ARDNRGSNWL CG-------- ---------- 613
4.3bits
Sequence logo
0.0bits
1,040
1,060
1,080
gi|40254978|[Homosapiens] ---------- ---------- ---------- ---SDEERY R------- ---------- 506
Cre20.g759400.t1.1 GMRTGLDSDA VLVGGTGGDD DDGDFYLDGG GGGDSDEEEQ RGGGGGGGGG AGGMGAGPGP 883
Ppa:Phypa_161275 EEFGHRQRSA DLRGQKAWVG KKKNEDRSRH GDKEKDLRFY DRDRERSRGR EEARSHQRER 1064
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 ---------- ---------- ---------- EDRPKFKQQD SGAKDDLRMK VLEKVSARWK LHLRMDDAHA 774
Fip1p[Saccharomyces ---------- ---------- ---------- ---------- TAEDEDFPIY PVKKYQFYRS PSK------- 639
4.3bits
Sequence logo
0.0bits
1,100
1,120
1,140
gi|40254978|[Homosapiens] ---------- ---------- ---------- ---------- ---------- ---------- 506
Cre20.g759400.t1.1 GPGMGLGLGM GMGGPPGRGA ELDLGPDMPP LMGMGMGLGM GMGLPPVVPP QAQQQQQHQQ 943
Ppa:Phypa_161275 RRPREEISKI DQEEERHRDR DRDRYRREDR PDRRDDACSR AVDRSEENRD RELRNRRHIE 1124
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 RRKRENLSKP DQKD------ ---------- NEEG AGGYHNSTIR DTQTAYASEE LEDDNRRERK 822
Fip1p[Saccharomyces ---------- ---------- ---------- ---------- ---------- ---------- 639
4.3bits
Sequence logo
0.0bits
1,160
1,180
1,200
gi|40254978|[Homosapiens] ---------- ---------- ---------- ---------- ---------- ---------- 506
Cre20.g759400.t1.1 QQGGRGGGDG DDGGEAAAFG DFSAGGGDYY LDDGSDQQQH QQQQQSSPSG QHHRGHHNHL 1003
Ppa:Phypa_161275 DRRKGEDRRN DLHRKSVVNG REAGDDARMQ MRRRDGSEER EADVMRNDSL VKERSHHGCG 1184
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 EERKRRHKDR KERRRLKSED EDARDCDDYY QEVDERWQME DADFT------ ---------- 867
Fip1p[Saccharomyces ---------- ---------- ---------- ---------- ---------- ---------- 639
4.3bits
Sequence logo
0.0bits


1,220
1,240
1,260
gi|40254978|[Homosapiens] 506
Cre20.g759400.t1.1 1063
Ppa:Phypa_161275 1241
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 867
Fip1p[Saccharomyces 639
4.3bits
Sequence logo
0.0bits
1,280
1,300
1,320
gi|40254978|[Homosapiens] 525
Cre20.g759400.t1.1 1123
Ppa:Phypa_161275 1301
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 895
Fip1p[Saccharomyces 667
4.3bits
Sequence logo
0.0bits
1,340
1,360
1,380
gi|40254978|[Homosapiens] 573
Cre20.g759400.t1.1 1182
Ppa:Phypa_161275 1361
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 955
Fip1p[Saccharomyces 727
4.3bits
Sequence logo
0.0bits
1,400
1,420
1,440
gi|40254978|[Homosapiens] 594
Cre20.g759400.t1.1 1227
Ppa:Phypa_161275 1421
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 1015
Fip1p[Saccharomyces 777
4.3bits
Sequence logo
0.0bits


1,460
1,480
gi|40254978|[Homosapiens] ---------- ---------- ---------- ---------- ---------- -------- 594
Cre20.g759400.t1.1 ---------- ---------- ---------- ---------- ---------- -------- 1227
Ppa:Phypa_161275 KLERRKERFG KPRPKKPSEE IPAGRGGSFY VSEVQANTVE VKQERPARKR RWAASGLE 1479
Smo:441074|441074|441074|synonym:estExt_fgenesh2_pg.C_140033 N--------- ---------- ---------- ---------- ---------- -------- 1016
Fip1p[Saccharomyces ---------- ---------- ---------- ---------- ---------- -------- 777
4.3bits
Sequence logo
0.0bits
KLERRKERFG KPRPKKPSEE IPAGRGGSFY VSEVQANTVE VKQERPARKR RWAASGLE