Phyre2

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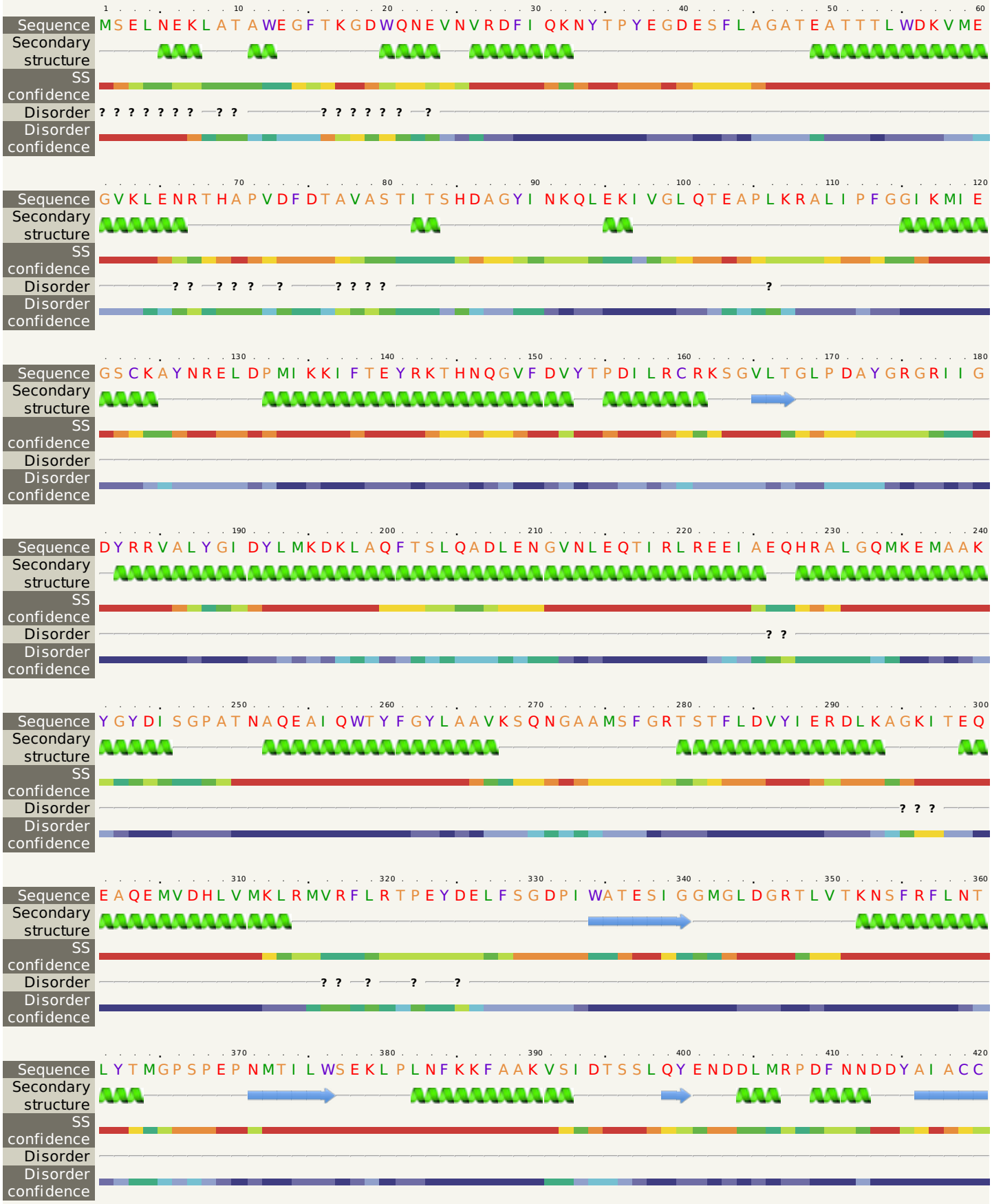
Description	P09373
Date	Thu Jan 5 11:02:15 GMT 2012
Unique Job ID	973cbb7f09cb6b84

Secondary structure and disorder prediction



Sequence
Secondary structure
SS confidence
Disorder
Disorder confidence
430 440 450 460 470 480
VSPMI VGKQMQFF GARANLAKTMLYAI NGGVDEKLKMQVGPKSEPI KGDVLNYDEVMERM
490 500 510 520 530 540
DHFMDWL AKQYI TALNI I HY MHDKYSYEASLMALHDRDVI RTMACGI AGL SVAADSLSAI
550 560 570 580 590 600
KYAKVKPI RDEDGLAI DFEI EGEYPQFGNNDPRVDDLAVDLVERFMKKI QKLHTYRDAI P
610 620 630 640 650 660
TQSVLTI TSNVVYGKKTGNTPDGRRAGAPF GPGANPMHGRDQKGAVASLTSVAKLPFAYA
670 680 690 700 710 720
KDGI SYTFSI VPNALGKDDEVRKTNLAGLMDGYFHHEASI EGGQHLNVNVMNREMLLDAM
730 740 750 760
ENPEKYPQLTI RVSGYAVRF NSLTKEQQQDVI TRTFTQSM
Confidence Key
High(9)
Low (0)
? Disordered
Alpha helix
Beta strand