Phyre^2

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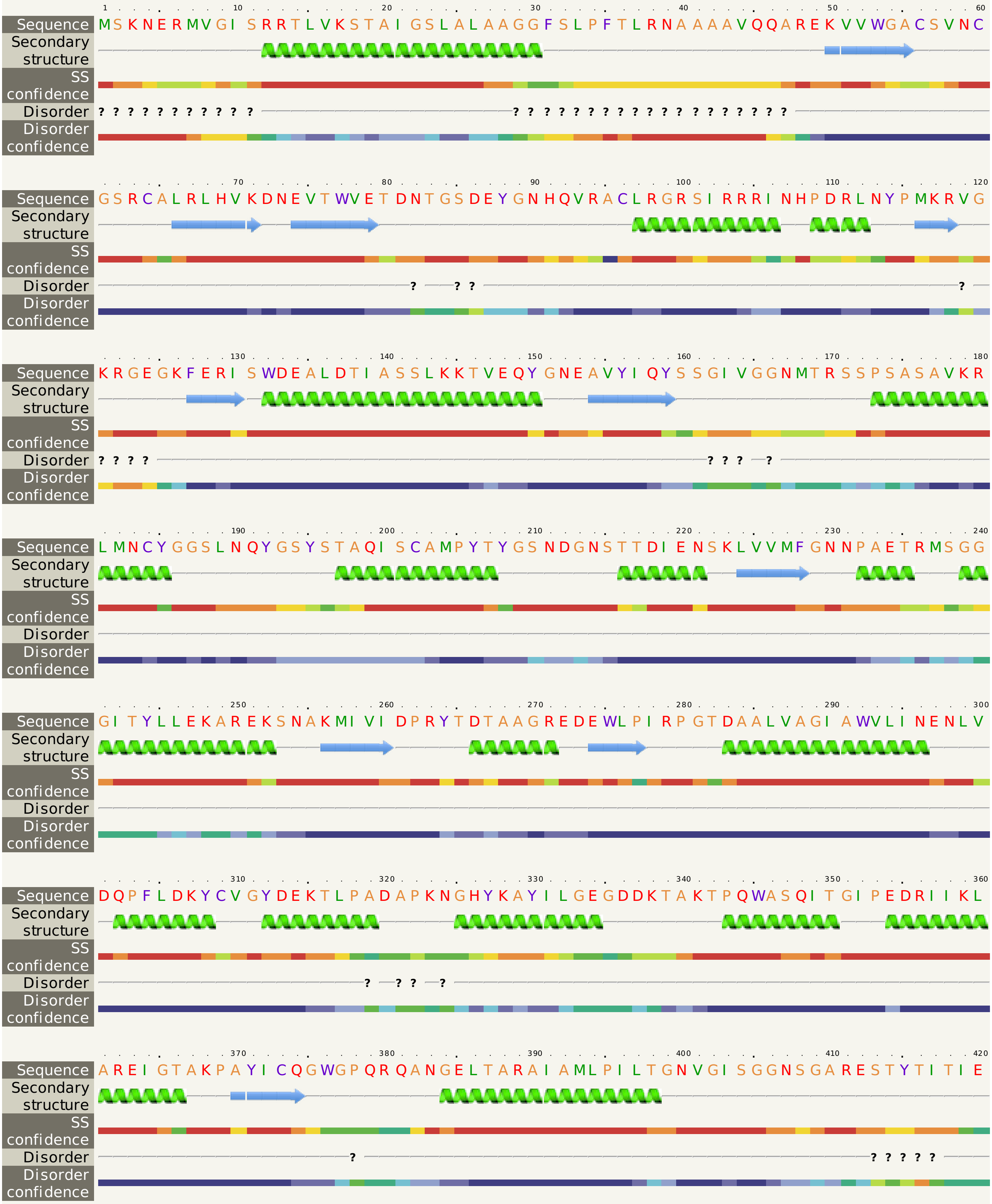
Description	P77374
Date	Thu Jan 5 12:28:16 GMT 2012
Unique Job ID	f80e8aa8888f794b

Secondary structure and disorder prediction




430 440 450 460 470 480
Sequence RLPVLDNPVKTSISCFSWTDAIDHGPQMTAIRDGVRGKDKLDVPIKFIWNYAGNTLVNQH
Secondary structure
SS confidence
Disorder ? ? ? ? ?
Disorder confidence
490 500 510 520 530 540
Sequence SDINKTHEILQDESKCEMIVVIENFMTSSAKYADILLPDLMTVEQEDIIPNDYAGNMGYL
Secondary structure
SS confidence
Disorder
Disorder confidence
550 560 570 580 590 600
Sequence IFLQPVTSEKFERKPIYWILSEVAKRLGPDVYQKFTEGRTQEQWLQHLYAKMLAKDPALP
Secondary structure
SS confidence
Disorder ? ? ?
Disorder confidence
610 620 630 640 650 660
Sequence SYDELKKMGIYKRKDPNGHFVAYKAFRDDPEANPLKTPSGKIEIYSSRLAEIARTWELEK
Secondary structure
SS confidence
Disorder
Disorder confidence
670 680 690 700 710 720
Sequence DEVISPLPVYASTFEGWNSPERRTFPLQLFGFHYKSRTHSTYGNIDLLKAACRQEVWINP
Secondary structure
SS confidence
Disorder ? ? ? ? ?
Disorder confidence
730 740 750 760 770 780
Sequence IDAQKRGIANGDMVRVFNHRGEVRLPAKVTPRILPGVSAMGQGAWHEANMSGDKIDHGGC
Secondary structure
SS confidence
Disorder ? ? ? ? ? ? ?
Disorder confidence
790 800
Sequence VNTLTTLRPSPLAKGNPQHTNLVEIEKI
Secondary structure
SS confidence
Disorder ? ?
Disorder confidence
Confidence Key
High(9)
Low (0)
? Disordered
Alpha helix
Beta strand