

Phyre2

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Description	RVBD2490c_(PE_PGRS43)_2801264_2806246
Date	Wed Aug 7 12:50:12 BST 2019
Unique Job ID	d674b45713b82c63

Secondary structure and disorder prediction



430 440 450 460 470 480
Sequence PGLGGAGGAG GASGGAGARA GANGLAAGND GPVSGGNGGK GGNGAHAPVA GGHGGNGGAG
Secondary structure
SS
confidence
Disorder
Disorder confidence

490 500 510 520 530 540
Sequence GNGGLVGDGG AGGHGGDGAA GAGYADMTAI FLGSSGTPGE DGGNGGAGGA GGAGGAHAGD
Secondary structure
SS
confidence
Disorder
Disorder confidence

550 560 570 580 590 600
Sequence GGAGGGAGGNG GAGGAGGNGA HGFNAVLVSD GGNGGDGGAG GRGGDGGAGG AGGDAPAGRA
Secondary structure
SS
confidence
Disorder
Disorder confidence

610 620 630 640 650 660
Sequence GSQGVGGDGG AGGAGGAPGN GGSGRGDMA FKDDGGAGG DGGDPGAGGK GGAGGAGATE
Secondary structure
SS
confidence
Disorder
Disorder confidence

670 680 690 700 710 720
Sequence GVTGATGATV HSGGNGGKGG NGADATVAGA NNGKGGAGGN GGLVGDGGAG GDGGSGAAGA
Secondary structure
SS
confidence
Disorder
Disorder confidence

730 740 750 760 770 780
Sequence NGANVGEDGA DGTLSGQPGE GSEANGGQGG VGGGGAGGAG GDGGAGSSAL GSGGNGGRGD
Secondary structure
SS
confidence
Disorder
Disorder confidence

790 800 810 820 830 840
Sequence AGQAGGAGGA GGAGGAGGSV SGGGGP GKG GAGGAGGAGA SGGGGK GAS GADSAEAVGG
Secondary structure
SS
confidence
Disorder
Disorder confidence

850 860 870 880 890 900
Sequence AGGKGGDGGV GGVGDDGGPG GDGGAGGAAP AGQVGS HGVG GVGDDGGLGG AGGNGGDGGH
Secondary structure
SS
confidence
Disorder
Disorder confidence

910 920 930 940 950 960
Sequence GSDGGDGGDG GDPGAGGLGG LGGDSGNGTR AASGV DASH GPSSGNGGN GGNGAQASVA
Secondary structure

Secondary structure
SS
confidence
Disorder
Disorder confidence

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Confidence Key
 High(9) [Color scale from red to blue] Low (0)
 ? Disordered (95%)
 [Alpha helix icon] Alpha helix (5%)
 [Beta strand icon] Beta strand (0%)