

Date: Wed, 6 Aug 2003 22:49:09 +0100
From: Apache <psipred@cs.ucl.ac.uk>
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Subject: PSIPRED Results

PSIPRED PREDICTION RESULTS

Key

Conf: Confidence (0=low, 9=high)
Pred: Predicted secondary structure (H=helix, E=strand, C=coil)
AA: Target sequence

Conf: 988887245325056436887768857886752888705458731488875888731456
Pred: CCCCCCEEEECCEEECC
AA: MGPWGKLRWTVALLLAAAGTAVGDR CERNEFQCQDGK CISYKWVCDGSAECQDGSDSEQ
10 20 30 40 50 60

Conf: 52135655665554788787655787778888677875466754654667641247871
Pred: CCECCCCC
AA: ETCLSVTCKSGDFSCGGRVNR CIPQFWRC DGQVDCDNGSDEQGCPPKTC SQDEF RCHDGK
70 80 90 100 110 120

Conf: 238870514557655776665540147875322058874003664557754244425750
Pred: CCCCCCEEECCC
AA: CISRQFVCDSDRDCLDGSDEASCPVLTCGPASFQCNSSTCIPQLWACDNDPDCEDGSDEW
130 140 150 160 170 180

Conf: 135567643267775156764447886020685025787403677765565566554666
Pred: CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCEEECCCCCCCCCCCCCCCCCCCCCCCCCCCCC
AA: PQRCRGLYVFQGDSSPCSAFEFHCLSGECIHSSWRCDGGPDCKDKSDEENCAVATCRPDE
190 200 210 220 230 240

Conf: 634672002785202554212243554101123210257777523302440576314422
Pred: CCCCCCCCCCEEECCCCCEEECCCCCEEECCCCCCCCCCCCCCCCCEEECCCCCCCCCCC
AA: FQCSDGNCIHGSRQC DREYDCKDMSDEVGCVNVTLC EGP NKFCHSGECITL DKVCNMAR
250 260 270 280 290 300

Conf: 456652255221133114677520387333424320452688505158870344521468
Pred: CCCCCCCCCCEEEEEECCCCCCCCCCCCCCCCCCCCCEEECCCCCEEECCCCCEEEEEECCCC
AA: DCRDWSDEPIKECGTNECLDNNGGCSHVCNDLKIGYECLCPDGFQLVAQRRCEDIDECQD
310 320 330 340 350 360

Conf: 877887166269966785512320688888765345565101010002222215786224
Pred: CCCCCCEEECCCCCEEEEEECCCCCCCCCCCCCCCCCCCCCHHCEEEEEEEEEEECCCCCEE
AA: PDTCSQLCVNLEGGYKCQCEEGFQLDPHTKACKAVGSIAYLFFTNRHEVRKMTLDRSEYT
370 380 390 400 410 420

Conf: 422688707786777518759998537883899862788677516898824655443158

Pred: EECCCCCEEEEEEECCCEEEEEEECCCEEEEEEECCCCCCCCCEEEEEEECCCCCCEEE
AA: SLIPNLRNVVALDTEVASNRIFYWSDLSQRMICSTQLDRAHGVSSYDTVISRDIQAPDGLA
430 440 450 460 470 480

Conf: 860589799981789726777258977899982788888507981579889997179986
Pred: EECCCCCEEEEEEECCCEEEEEEECCCEEEEEEECCCCCCCCCEEEEEEECCCCCCEEE
AA: VDWIHSNIYWTDSVLGTVSVADTKGVKRKTLFRENGSKPRAIVVDPVHGFMYWTDWGTPA
490 500 510 520 530 540

Conf: 168862388840899942788883479823898799981787789998538997289972
Pred: EEEEEEECCCCCEEEEEEECCCCCCCCCEEEEEEECCCCCEEEEEEECCCCCEEEEEEE
AA: KIKKGGLNGVDIYSLVTENIQWPNGITLDLLSGRLYWVDSKLHSISSIDVNGGNRKILE
550 560 570 580 590 600

Conf: 688688851799718889998379983899965789643798326888744899962016
Pred: CCCCCCCCCCEEEEEEECCCEEEEEEECCCEEEEEEECCCCCEEEEEEECCCCCEEEEEEE
AA: DEKRLAHPFSLAVFEDKVFWDIINEAIFSANRLTGSDVNLLAENLLSPEDMVLFFHNLTO
610 620 630 640 650 660

Conf: 644543223678888864354106888888886268768986844568635666210101
Pred: CCCCCCCCCCCCCCCCCCEEEEECCCCCCCCCCCCCEEECCCCCECCCCCCCCCCCCCEE
AA: PRGVNWCERTTSLNGGCQYLCPLPAPQINPHSPKFTCACPDGMLLARDMRSCLTEAEAAVA
670 680 690 700 710 720

Conf: 24566553131578768777664236768704225666530146753640125512566
Pred: CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCEEECCCCCCCCCEEEEECCCCCEEEEEEECCC
AA: TQETSTVRLKVSSTAVRTQHTTTRPVPDTSRLPGATPGLTTVEIVTMSHQALGDVAGRGN
730 740 750 760 770 780

Conf: 778776401220276158998403035666540157885013788854345601024305
Pred: CCCCCCCCCCEEECCCEEEEEEEECCHHHHHHHCCCCCCCCCCCCCCCCCCCCCEEEEC
AA: EKKPSSVRALSIVLPIVLLVFLCLGVFLWKNWRLKNINSINFDPVYQKTTEDEVHICH
790 800 810 820 830 840

Conf: 88872421021004433469
Pred: CCCCCECHHHEECCCCCCC
AA: NQDGYSYPSRQMVSLDDVA
850 860

Calculate PostScript, PDF and JPEG graphical output for this result
using:
<http://bioinf.cs.ucl.ac.uk/cgi-bin/psipred/gra/nph-view2.cgi?id=106020452318838>