

Date: Thu, 25 Sep 2003 22:24:23 +0100  
From: Apache <psipred@cs.ucl.ac.uk>  
To: aprilb@biology2.wustl.edu  
Subject: cypla1 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

[illegible]

Pred: CCCCCHHHHHHHHHHHHHHHHHHHHHCCCCCCCCCCCCCCCCCCCHHHHHHHHCCCHN

AA: MLFPISMSATEFLLASVIFCLVFWVIRASRPQVPKGLKNPPGPWGWPLIGHMLTLGKNPH  
10 20 30 40 50 60

Conf: 999999984488568997893479841767889999626742246764432010367762

Pred: HHHHHHHHCCCCEEEECCCCEEEECCHHHHHHHHHCCCCCCCCCCCCCHCCCCCCC

AA: LALSRSQQYGDVLQIRIGSTPVVVL~~S~~GLDTIRQALVRQGD~~D~~FKGRPDLYTFTLISNGQS

Conf: 0214775279999987666523302443334565433679999999999999985058

Pred: EEECCCCHHHHHHHHHHHHHCCCHHHCCC

AA: MSFSPDSGPVWAARRRLAQNLKSFSIASDPASSTSCYLEEHVSKEAEVLISLTQLQELMAG  
130 140 150 160 170 180

Conf: 97204689998878998889874266553025789999999999985124534322100

Pred: CCCCCHHHHHHHHHHHHHHHHHHHHCCCCCCCCHHHHHHHHHHHHHHHHHHHHCCCCCCCCCCCC

AA: PGHFNPYRYVVSVTNVICAICFGRRYDHNHQELLSLVNLNNNFGEVVGSGNPADFIPIIL  
190 200 210 220 230 240

[illegible]

Pred: CCCCCHHHHHHHHHHHHHHHHHHHHHHHHHHHHHCCCCCHHHHHHHHHHHHHHHHHHHHHCCC

AA: RYLPNPSLNAFKDLNEKFYSFMQKMKVEHYKTFEKGHIRDITDSLIEHCQEKQLDENANV  
250 260 270 280 290 300

Conf: 68889999999999731135789999999998308743689999999871577876445

Pred: C C C H H H H H H H H H H H H C C H H H H H H H H H H H H H H C C C C H H H H H H H H H H H H C C C C C C C H H

AA: QLSDEKIINIVLDFGAGFDTVTTAISWSLMYLMVNPVRQRKIQEELDTVIGRSRRPRLS

Conf: 65136026899988887406876542115541203885847887266213542165202

Pred: HHHCCCHHHHHHHHHHHHHCCCCCCCCCCCCCCCCCECCCCCECCCCCEEECHHHHCCCCC

AA: DRSHLPYMEAFILETFRHSSFVPFTIPHSTTRDTSLKGFIYPKGRCVFVNQWQINHDQKL

Conf: 58865355655676532010158864244787343023205899999999997501267

Pred: CCCCCCECCCCCCCCCHHCCCCCCCCCCCCCCCCCHHHHHHHHHHHHHHHHCCEE  
AA: WVNPFSEFLPERFLTPDGAIDKVLSEKVIIFGMGKRKCIGETVARWEVFLFLAILLQRFVEF  
430 440 450 460 470 480

Conf: 60778878750110057468996058998569  
Pred: ECCCCCCCCCEEEEEEECCCCCEEEEEEECC  
AA: SVPLGVKVDMTPIYGLTMKHACCEHFQMLRS  
490 500 510

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=106452363620584>