

Date: Tue, 19 Aug 2003 13:55:11 +0100
From: Apache <psipred@cs.ucl.ac.uk>
To: aprilb@biology2.wustl.edu
Subject: atp 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: 97640111067788889999999999999999999987015770011589999999999999999862
Pred: CCCC~~H~~HHHCCCHHHHHHHHHHHHHHHHHHHHHHHCCCCCCHHHHHHHHHHHHHHHHHHHHHHHHHHH
AA: MNENLFASFIAPTILGLPAAVLIILFPPLIPTSKYLINNRLITQOWLIKLTSMQMNTM
 10 20 30 40 50 60

[illegible][illegible][illegible]

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=106129730815339>