

Date: Mon, 25 Aug 2003 16:41:55 +0100
From: Apache <psipred@cs.ucl.ac.uk>
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Subject: sod1 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: 9851599985898579999999973899848999998438986068998568997665022
Pred: CCHHH
AA: MATKAVCVLKGDPVQGIINFQKESNGPVKVWGSIKGLTEGLHGFHVHEFGDNTAGCTS
10 20 30 40 50 60

Conf: 045447667877878788877654565255685259999985870247877666517999
Pred: CCEEEEE
AA: AGPHFNPLSRKHGGPKDEERHVGDLGNVTADKDGADVSIEDSVISLSGDHCCIIGRTLTVV
70 80 90 100 110 120

Conf: 6388756778873201368888603788854219
Pred: ECCCECC
AA: HEKADDLGKGGNEESTKTGNAGSRLACGVIGIAQ
130 140 150

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=106182581927342>