

Date: Wed, 17 Sep 2003 16:45:12 +0100
From: Apache <psipred@cs.ucl.ac.uk>
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Subject: kras2 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: 925899998179888889999974202433321211025778998636645899984355
Pred: CEEEEEEEECCCCCHHHHHHHHHCCCCCCCCCCCCEEEEEEEECCCCEEEEEEEECCC
AA: MTEYKLVVVGAGGVGKSALTIQLIQNHFVDEYDPTIEDSYRKQVVIDGETCLLDILDITAG
10 20 30 40 50 60

Conf: 433210124441378659998469864532568888888876235663047763221556
Pred: CCCCCHHHHHHHCCCCEEEEEEEECCCCCHHHHHHHHHHHHHCCCCCCCCCCCCCCCC
AA: QEEYSAMRDQYMRGTGEGFLCVFAINNTKSFEDIHHYREQIKRVKDSVDPMVLVGNKCDL
70 80 90 100 110 120

Conf: 52220389999999871984899207888577999999999972011257754345666
Pred: CCCCCHHHHHHHHHHHCCCCEEEEEEEECCCCCHHHHHHHHHHHHHHHCCCCCCCCCCCC
AA: PSRTVDTKQAQDLARSYGIPFIETSAKTRQGVDDAFYTLVREIRKHKEKMSKDGKKKKKK
130 140 150 160 170 180

Conf: 78866219
Pred: CCCCCCEC
AA: SKTKCVIM

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=106381301717938>