

Model details



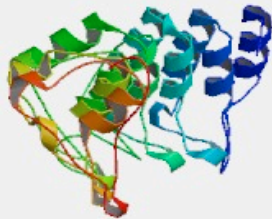
Sequence Information

Primary Database Link [Q8TDY4](#)
Original Database ID [EN ENSP00000338769](#)
Organism [Homo sapiens](#)
Annotation up-regulated in liver cancer 1 (development and differentiation enhancing factor-like 1)
Sequence Length 903

Model Information

Perform action on this model [?](#) :

Sequence Model Coverage [?](#)



[?](#) Sequence Identity 21.00%
[?](#) E-Value 1.4e-11
[?](#) Model Score 0.84
[?](#) Target Region 598-846
[?](#) Protein Length 903
[?](#) Template PDB Code [1n11A](#)
[?](#) Template Region 413-683
[?](#) MPQS 0.32
[?](#) z-Dope 0.6
[?](#) Dataset human_4-2007
[?](#) ModPipe Version ModPipe2.0

Filtered models for current sequence ([Show all models](#))



Cross-references

Template Structure

PDB	1n11	d34 region of human ankyrin-r and linker ; PFAM: PF00023 ; SCOP: 79778
DBALI	1n11A	
CATH	1.25.40.20	Domain: 1n11A0 Serine Threonine Protein Phosphatase 5, Tetratricopeptide repeat (66%)
Jena Image Library	1n11	

Target Sequence

TrEMBL	Q8TDY4	Up-regulated in liver cancer 1 (Development and differentiation enhancing factor-like 1)
UniProt	Q8TDY4	
InterPro	Q8TDY4	
PFAM	Q8TDY4	