EBI > Databases > Structure Databases > ProFunc



Nest analysis results for ao15

Nests are structural motifs that are often found in functionally important regions of protein structures.

6 nests were located in this chain, as shown below.

N4	C	Danidora unuma	View in		Ramachandran		01-44	Depth in	
Nest	Score	Residue range	Kaswol	Residue	region	accessibility	Cleft	CIETT	conservation
1.	3.97	Cys552-Leu554	&	Cys552	RIGHT	0.00%	•	25.57	1.00
				Gly553	LEFT	0.00%	-	-	1.00
				Leu554	-	0.00%	-	-	0.91
2.	3.97	Lys533-Ala536	æ	Lys533	RIGHT	0.00%	-	-	1.00
				Gly534	LEFT	0.00%	-	-	1.00
				Glu535	RIGHT	0.00%	•	26.19	0.87
				Ala536	-	0.00%	-	-	1.00
3.	1.00	Gly71-Ala73	æ	Gly71	LEFT	0.00%	-	-	1.00
				Gln72	RIGHT	0.00%	-	-	1.00
				Ala73	-	0.00%	-	-	1.00
4.	1.00	Asp131-Lys133	æ	Asp131	LEFT	0.00%	-	-	1.00
				Glu132	RIGHT	0.00%	-	-	1.00
				Lys133	-	0.00%	-	-	1.00
5.	1.00	Arg174-Glu176	&	Arg174	RIGHT	0.00%	-	-	1.00
				His175	LEFT	0.00%	-	-	1.00
				Glu176	-	0.00%	-	-	1.00
6.	0.49	Lys246-Asp248	æ	Lys246	RIGHT	0.00%	-	-	0.41
				His247	LEFT	0.00%	-	-	0.90
				Asp248	-	0.00%	-		0.17



References

- 1. Watson JD and Milner-White EJ (2002). A novel main-chain anion-binding site in proteins: the nest. A particular combination of phi,psi values in successive residues gives rise to anion-binding sites that occur commonly and are found often at functionally important regions. *J. Mol. Biol.*, **315**, 171-82.
- 2. Watson JD and Milner-White EJ (2002). The conformations of polypeptide chains where the main-chain parts of successive residues are enantiomeric. Their occurrence in cation and anion-binding regions of proteins. *J. Mol. Biol.*, **315**, 183-191.
- 3. Pal D, Suhnel J and Weiss MS (2002). New principles of protein structure: nests, eggs and what next? *Angew. Chem. Int. Ed.*, **41**, 4663-4665.

Terms of Use | EBI Funding | Contact EBI | © European Bioinformatics Institute 2010. EBI is an Outstation of the European Molecular Biology Laboratory