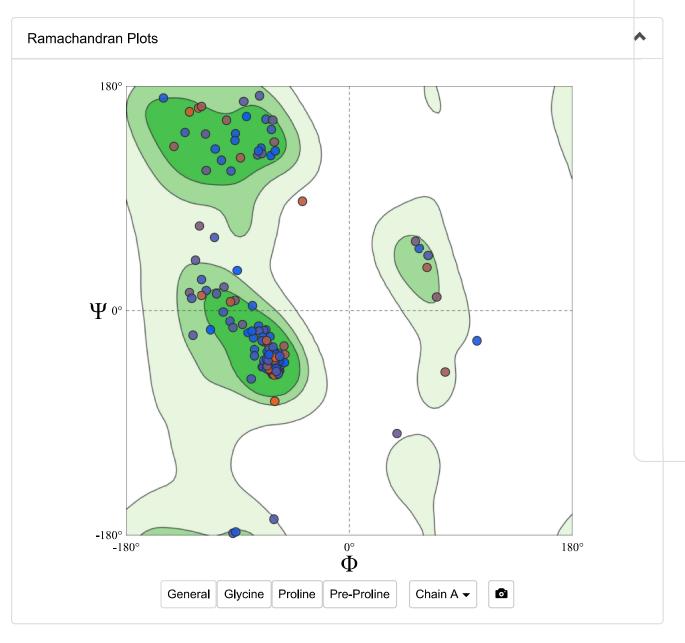
Structure Assessment

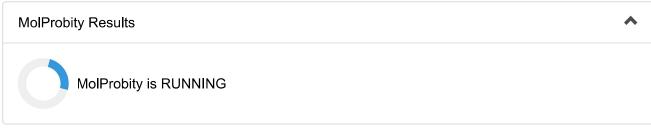
Help Examples ▼

Receptor de Ecdisona na pose Antagonista; Model 01; 🎍 🖺 ธ ե 🗆



Created: Fri 05 Mar 2021, 21:19;



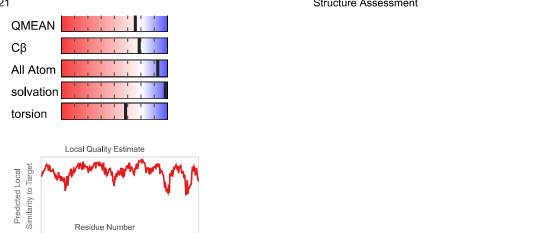


Quality Estimate

Residue Number

Protein Size (Residues)

Comparison with Non-redundant Set of PDB Structures





-0.43

-0.14 1.27

1.34

-1.16

QMEAN		40
Chain:A	PLSYNQLAVIYKLIWYQDGYEQPSEEDLKRIMSSPDENES	
<pre>template_upload.1.A</pre>	PISPEQEELIHRLVYFQNEYEQPSDEDLKRISNTPSEGED	42
QMEAN		
Chain:A	QHDVSFRHITEITILTVQLIVEFAKGLPAFTKIPQEDQIT	80
template_upload.1.A	QSDLNFRHITEITILTVQLIVEFAKRLPGFDKLLREDQIA	82
0.177.17		
QMEAN		
Chain:A	LLKACSSEVMMLRMARRYDHNSDSIFFANNRSYTRDSYKM	120
template_upload.1.A	LLKACSSEVMMLRMARRYDVGSDSILFANNQPYTRDSYSL	122
QMEAN		
Chain:A	AGMADNIEDLLHFCRQMYSMKVDNVEYALLTAIVIFSDRP	160
	AGMGETVDDLLRFCRQMYGMKVDNAEYALLTAIVIFSERP	
QMEAN		
Chain:A	GLEEAELVEAIQSYYIDTLRIYILNRHCGDPMSLVFFAKL	200
	SLIEGWKVEKIQEIYLEALKVYVDNRRKPASGTIFAKL	
QMEAN		
Chain:A	LSILTELRTLGNQNAEMCFSLKLKNRKLPKFLEEIWDV	238
template_upload.1.A	LSVLTELRTLGNLNSEMCFSLKLKNKKLPPFLAEIWDV	238