## AcriiA4 Anti-CRISPR Protein – 5vw1.pdb

ACHIA4 model to be judged has an orange/shver/white backbone.	AcrIIA4 model to be judged has	an orange/silver/white backbone.
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What is Displayed	How it is Displayed	Why it is Important
N Terminus	Blue Endcap	Amino terminus – beginning of the protein
		chain
C Terminus	Red Endcap	Carboxyl terminus – end of the protein chain
Primary Structure	Green strings	Primary structure determines the sequence
		of amino acids in a protein
Beta Pleated Sheets	Silver painted	Secondary structures with hydrogen bonding
	backbone	between parallel peptides
Alpha Helices	White painted	Secondary structures with hydrogen bonding
	backbone	between amine groups and carboxyl groups,
		with 3.6 residues per turn and a separation of
		1.5 Å
3 <sub>10</sub> Helix	Dark blue colored	Secondary structures like alpha helices, with
	backbone	3 residues per turn and a separation of 2.0 Å
Lys18	Amino acid sidechain	Prevents entrance of non-complimentary
		DNA in RuvC active site
Asn25, Ser26	Blue LEGO bricks	Prevents Cutting of non-target DNA by

blocking RuvC active site

Asp14, Asn36	Paper Clips	Prevents PAM nucleotide stabilization & PAM
		recognition
Asn39, Asp69	3D Printed Sidechain	Prevents verification of non-target DNA
Glu70	3D Printed Sidechain	Prevents PAM recognition
Asp37	Red Push pin	Prevents PAM recognition; interacts with
		both Topo and CTD
Glu40	White Push pin	Prevents PAM recognition; interacts with CTD
RuvC Domain of	Yellow alpha-carbon	AcrIIA4 binds to the RuvC domain of the
CRISPR Cas9	backbone	CRISPR Cas9 Protein to prevent cleavage of
(residues 960-1021)		DNA.
Topo Domain of	Red alpha-carbon	AcrIIA4 binds to the Topo domain to prevent
CRISPR Cas9	backbone	PAM recognition by the Cas9 protein.
(residues 1102-1149)		
CTD Domain of	Light blue alpha-	AcrIIA4 binds to the CTD domain to prevent
CRISPR Cas9	carbon backbone	PAM recognition by the Cas9 protein.
(residues 1200-1247)		