BASA	**All**	**BASAT**	Total Interface Accessible Surface Area
	by Chain	**BASA1**	Interface Accessible Surface Area for Chain A
		%BASA1	Interface Accessible Surface Area for Chain A (%)
		BASA2	Interface Accessible Surface Area for Chain B
		%BASA2	Interface Accessible Surface Area for Chain B (%)
	%	**BASAN**	% Neutral contribution
		BASAP	% Polar contribution
		BASANP	% Non polar contribution
		BASAC	% Charged contribution
GAP Volume	**All**	**GAPV**	Gap Volume
SEGMENTS	**All**	**SEGT**	Nb Segments Total
	by Chain	**SEG1**	Nb Segments for Chain A
		SEG2	Nb Segments for Chain B
SSBONDS	**All**	**DisuInt**	Number of Disulfide bonds
HBONDS		**Hb30**	Number of hydrogen bonds
SALT BRIDGES		**SBAB**	Number of Salt Bridges
CONTACT	**All**	**NbCont**	Number of non-bonded contacts
	Atom (by Chain)	**ContCh**	Number of all non-bonded atoms
		ContN	Number of neutral non-bonded atoms
		ContP	Number of polar non-bonded atoms
		ContHyd	Number of hydrophobic non-bonded atoms
		ContC	Number of charged non-bonded atoms
	Resid (by Chain)	**ContRes**	Number of all non-bonded residues
		ContRN	Number of neutral non-bonded residues
		ContRP	Number of polar non-bonded residues
		ContRHyd	Number of hydrophobic non-bonded residues
		ContRC	Number of charged non-bonded residues

GENERAL PROPERTIES	**Atom (by Chain)**	**AIntaro**	% Aromatic
		AIntacid	% Acidic
		AIntbas	% Basic
		AIntacy	% Acyclic
		AIntcyc	% Cyclic
		AIntali	% Aliphatic
		AIntS	% Small
		AIntM	% Medium
		AIntL	% Large
		AIntH	% Helix
		AIntSh	% Sheet
		AIntCo	% Coil
	Resid (by Chain)	**RIntaro**	% Aromatic
		RIntacid	% Acidic
		RIntbas	% Basic
		RIntacy	% Acyclic
		RIntcyc	% Cyclic
		RIntali	% Aliphatic
		RIntS	% Small
		RIntM	% Medium
		RIntL	% Large
		RIntH	% Helix
		RintSh	% Sheet
		RIntCo	% Coil
ATOMIC PROPERTIES	**by Chain**	**AIntCar**	% Carbon
		AIntOx	% Oxygen
		AIntNit	% Nitrogen
		AIntSul	% Sulfur
		AIntH	% Hydrogen
		AIntPho	% Phosphorus
		AIntCF	% Fluor
		AIntNOPS	% (N+O+P+S)