Residue properties

23_128_72

a. Absolute deviation from mean Chi-1 value (excl. Pro)

b. Absolute deviation from mean of omega torsion

c. C-alpha chirality: abs. deviation of zeta torsion

d. Secondary structure & estimated accessibility

Highlighted residues are those that deviate by more than 2.0 st. devs. from ideal

e. Sequence & Ramachandran regions

f. Max. deviation (see listing)

g. G-factors
Residue properties
23_128_72

a. Absolute deviation from mean Chi-1 value (excl. Pro)

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Highlighted residues are those that deviate by more than 2.0 st. devs. from ideal

e. Sequence & Ramachandran regions

Most favoured ▲ Allowed ■ Generous □ Disallowed

f. Max. deviation (see listing)

g. G-factors
Residue properties
23_128_72

a. Absolute deviation from mean Chi-1 value (excl. Pro)

b. Absolute deviation from mean of omega torsion

c. C-alpha chirality: abs. deviation of zeta torsion

Highlighted residues are those that deviate by more than 2.0 st. devs. from ideal

d. Secondary structure & estimated accessibility

Key:
- Helix
- Beta strand
- Random coil

Accessibility shading:
- Buried
- Accessible

e. Sequence & Ramachandran regions

f. Max. deviation (see listing)

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<th>Phi-pi</th>
<th>Chi-chi1</th>
<th>Chi1 only</th>
<th>Chi3 &amp; chi4</th>
<th>Omega</th>
<th>Dihedrals</th>
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</table>
Residue properties

Model_12

a. Absolute deviation from mean Chi-1 value (excl. Pro)

b. Absolute deviation from mean of omega torsion

c. C-alpha chirality: abs. deviation of zeta torsion

d. Secondary structure & estimated accessibility

e. Sequence & Ramachandran regions

f. Max. deviation (see listing)

g. G-factors
Residue properties

Model_12

a. Absolute deviation from mean Chi-1 value (excl. Pro)

b. Absolute deviation from mean of omega torsion

c. C-alpha chirality: abs. deviation of zeta torsion

d. Secondary structure & estimated accessibility

Highlighted residues are those that deviate by more than 2.0 st. devs. from ideal

e. Sequence & Ramachandran regions

f. Max. deviation (see listing)

g. G-factors
Histograms showing RMS distances of planar atoms from best-fit plane. Black bars indicate large deviations from planarity: RMS dist > 0.03 for rings, and > 0.02 otherwise.
Histograms showing RMS distances of planar atoms from best-fit plane.
Black bars indicate large deviations from planarity: RMS dist > 0.03 for rings, and > 0.02 otherwise.

> signifies data points off the graph in the direction shown.