Chi1-Chi2 plots

Arg (60)  Asn (16)  Asp (48)

Cys (-)  Gln (26)  Glu (34)

His (20)  Ile (30)  Leu (80)

Numbers of residues are shown in brackets. Those in unfavourable conformations (score < -3.00) are labelled. Shading shows favourable conformations as obtained from an analysis of 163 structures at resolution 2.0A or better.
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