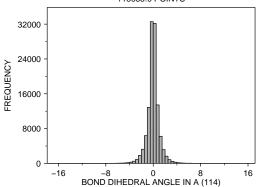
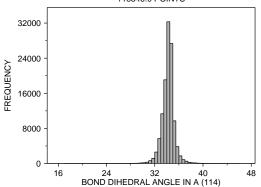
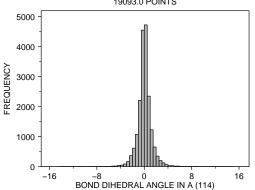
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ALA:C:CA:+N:O : BOND DIHEDRAL ANGLE TYPE IN A (113)
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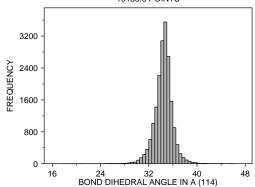
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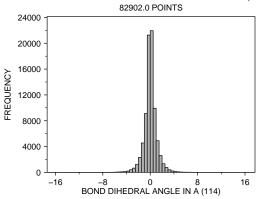
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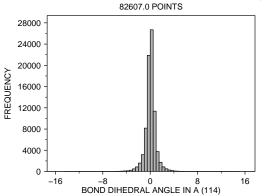
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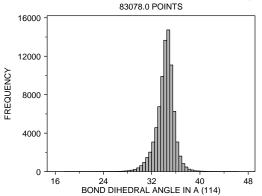
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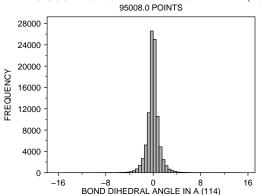
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<2 : X-RAY RESOLUTION OF A (35)
ASP:CA:N:C:CB : BOND DIHEDRAL ANGLE TYPE IN A (113)

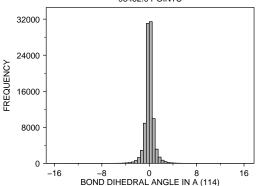


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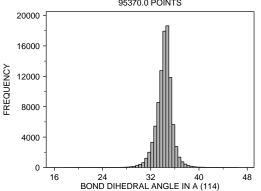


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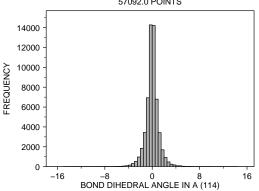
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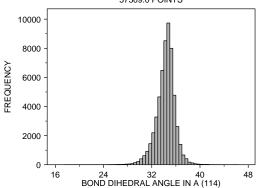
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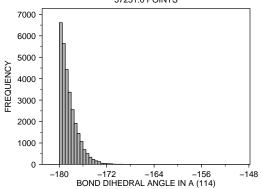
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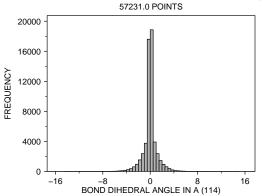
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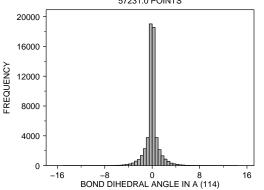
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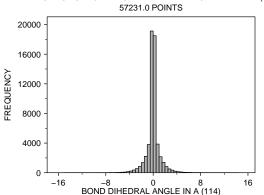
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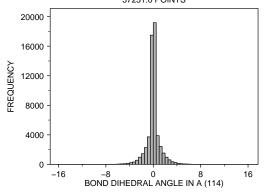


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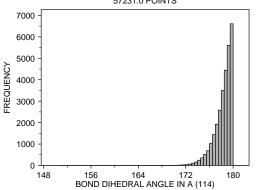


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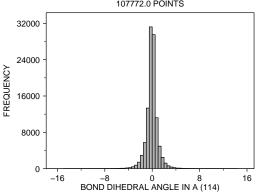
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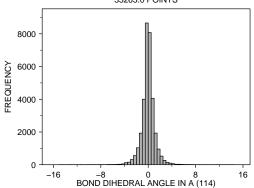
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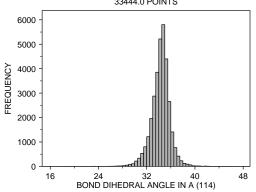
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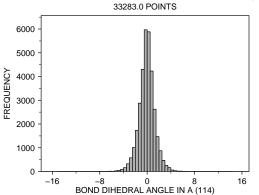
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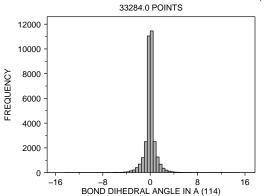
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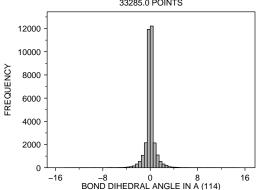
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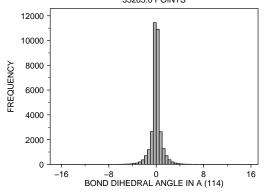


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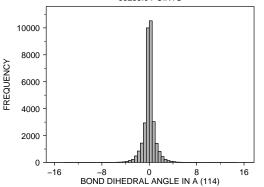


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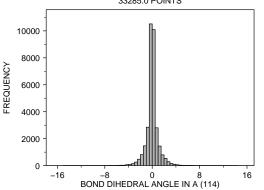
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HIS:CE1:NE2:CD2:CG : BOND DIHEDRAL ANGLE TYPE IN A (113)
33285.0 POINTS



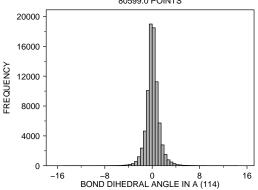
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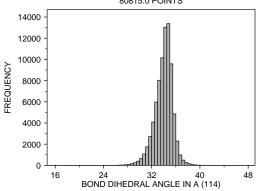
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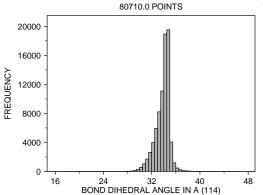
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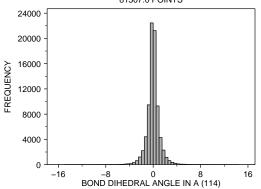
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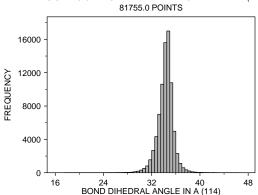
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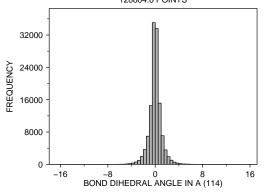


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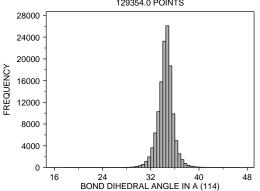


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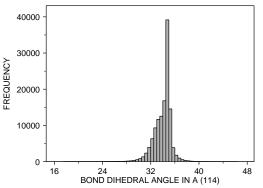
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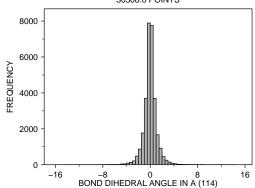
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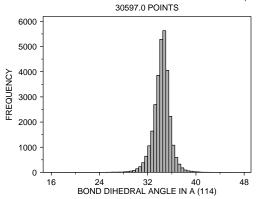
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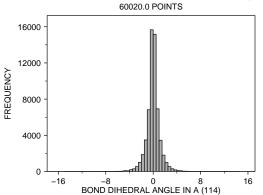
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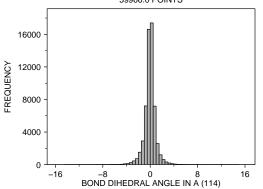
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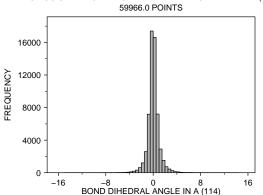
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ASN:C:CA:+N:O : BOND DIHEDRAL ANGLE TYPE IN A (113)



<2 : X-RAY RESOLUTION OF A (35)
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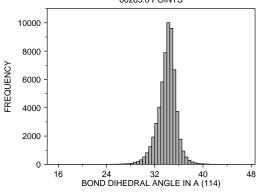


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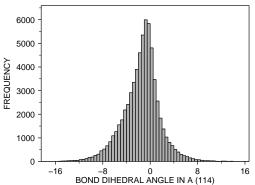


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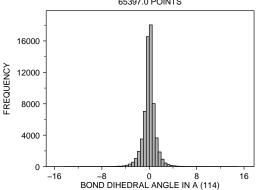
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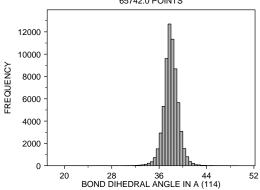
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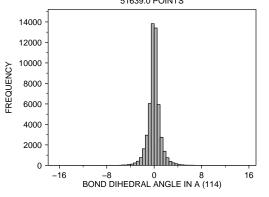
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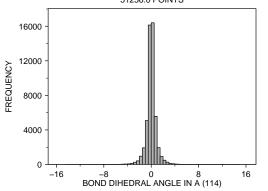
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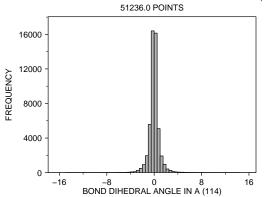
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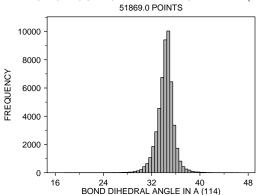
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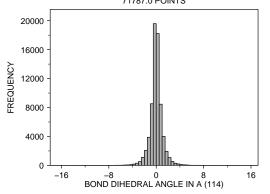


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GLN:CA:N:C:CB : BOND DIHEDRAL ANGLE TYPE IN A (113)



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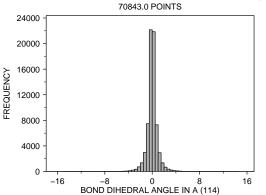
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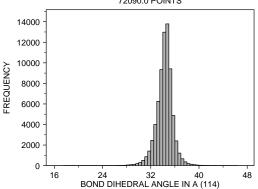
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ARG:CZ:NH1:NH2:NE : BOND DIHEDRAL ANGLE TYPE IN A (113)

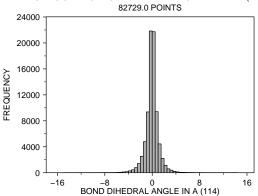
70843 0 POINTS



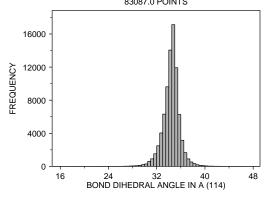
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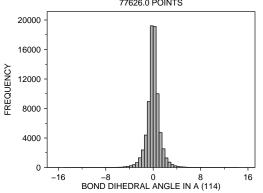
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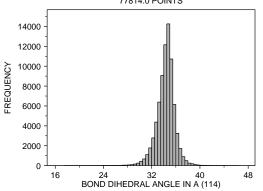
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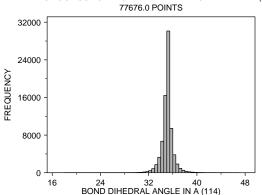
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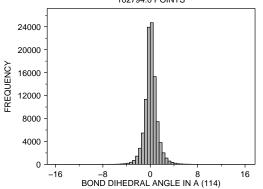


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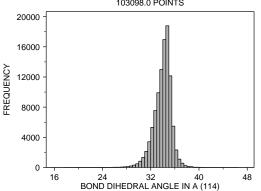


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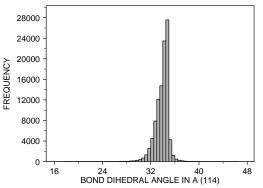
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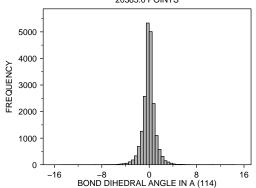
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103098.0 POINTS



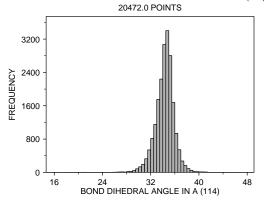
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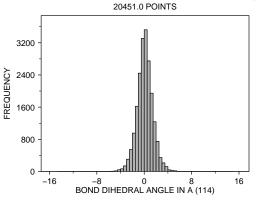
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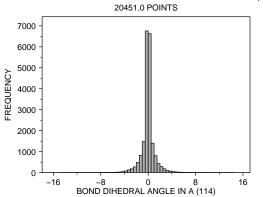
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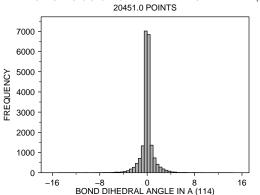
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<2 : X-RAY RESOLUTION OF A (35) TRP:CE3:CD2:CE2:CZ2 : BOND DIHEDRAL ANGLE TYPE IN A (113)

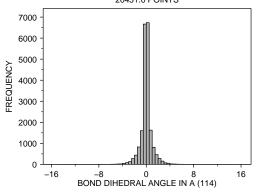


<2 : X-RAY RESOLUTION OF A (35)
TRP:CZ2:CH2:CZ3:CE3 : BOND DHEDRAL ANGLE TYPE IN A (113)

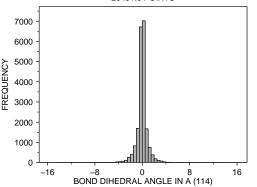


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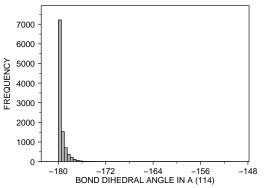
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TRP:CG:CD1:NE1:CE2 : BOND DIHEDRAL ANGLE TYPE IN A (113)
20451.0 POINTS



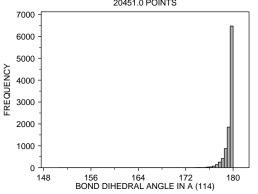
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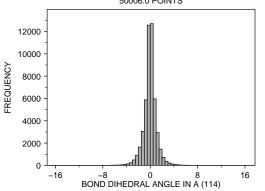
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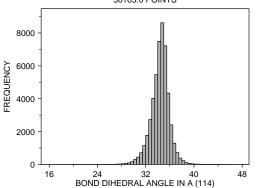
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TRP:CG:CD2:CE3:CZ3 : BOND DIHEDRAL ANGLE TYPE IN A (113)
20451.0 POINTS



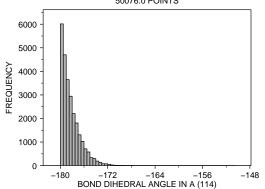
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TYR:C:CA:+N:O : BOND DIHEDRAL ANGLE TYPE IN A (113)
50006.0 POINTS



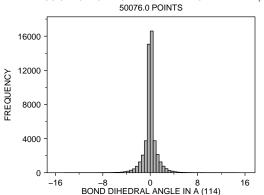
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TYR:CA:N:C:CB : BOND DIHEDRAL ANGLE TYPE IN A (113)
50163.0 POINTS



<2 : X-RAY RESOLUTION OF A (35)
TYR:CB:CG:CD1:CE1 : BOND DIHEDRAL ANGLE TYPE IN A (113)
50076.0 POINTS

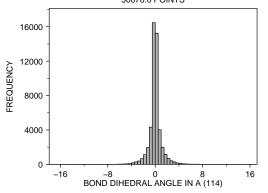


<2 : X-RAY RESOLUTION OF A (35)</p>
TYR:CG:CD1:CE1:CZ : BOND DIHEDRIAL ANGLE TYPE IN A (113)

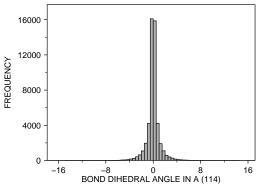


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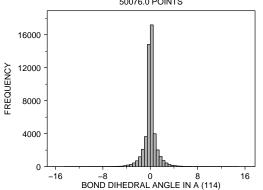
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TYR:CD1:CE1:CZ:CE2 : BOND DIHEDRAL ANGLE TYPE IN A (113)
50076.0 POINTS



<2 : X-RAY RESOLUTION OF A (35)
TYR:CE1:CZ:CE2:CD2 : BOND DIHEDRAL ANGLE TYPE IN A (113)
50076.0 POINTS

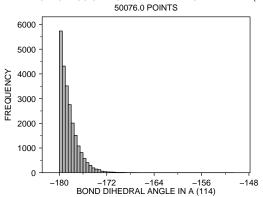


<2 : X-RAY RESOLUTION OF A (35)
TYR:CZ:CE2:CD2:CG : BOND DIHEDRAL ANGLE TYPE IN A (113)
50076.0 POINTS

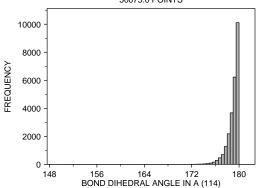


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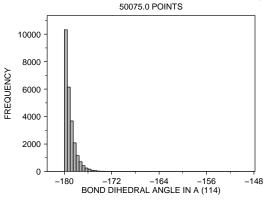
TYR:CE2:CD2:CG:CB : BOND DIHEDRAL ANGLE TYPE IN A (113)



<2 : X-RAY RESOLUTION OF A (35)
TYR:CD1:CE1:CZ:OH : BOND DIHEDRAL ANGLE TYPE IN A (113)
50075.0 POINTS



<2 : X-RAY RESOLUTION OF A (35)
TYR:CD2:CE2:CZ:OH : BOND DIHEDRIAL ANGLE TYPE IN A (113)



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