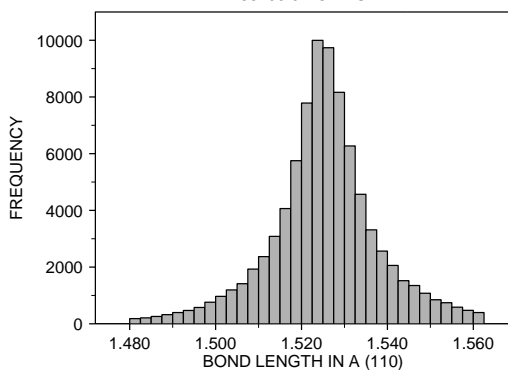
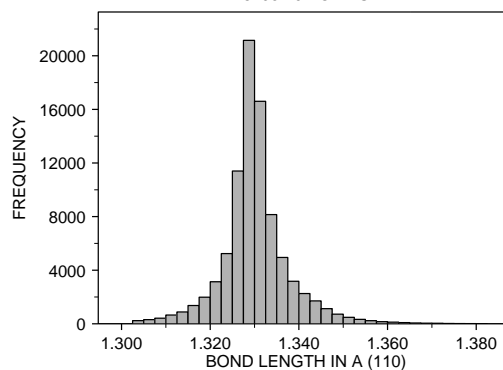


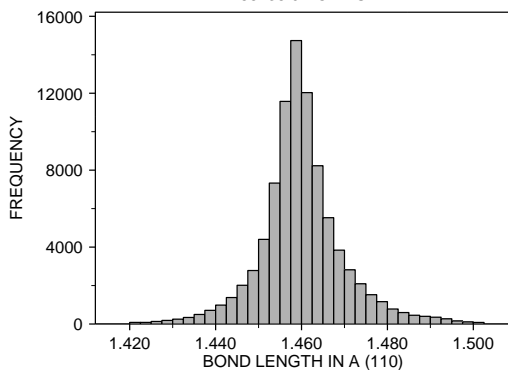
<2 : X-RAY RESOLUTION OF A (35)
ALA:CB:CA : BOND TYPE IN A (109)
88409.0 POINTS



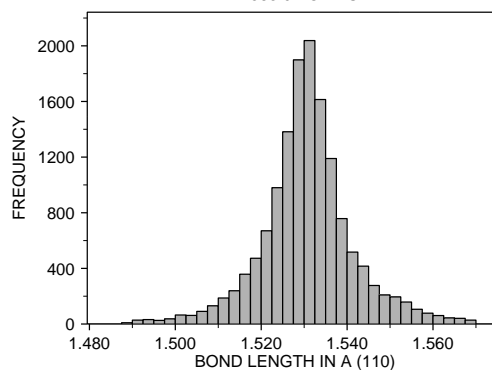
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ALA:C:N : BOND TYPE IN A (109)
87952.0 POINTS



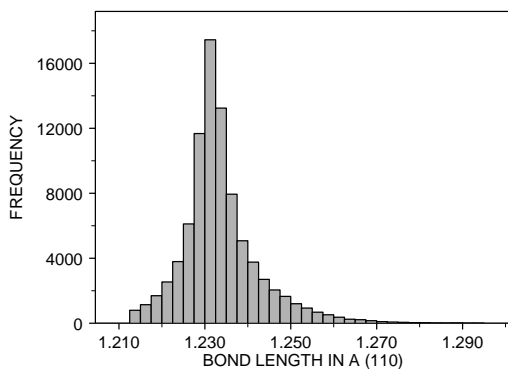
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ALA:N:CA : BOND TYPE IN A (109)
88438.0 POINTS



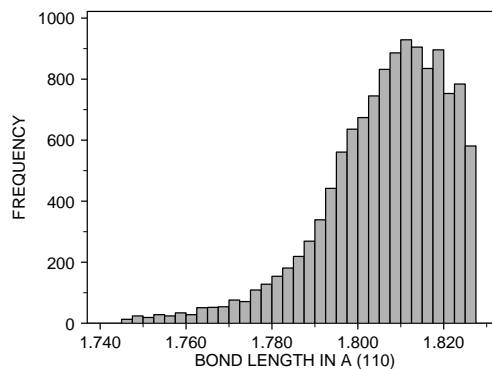
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CYS:CB:CA : BOND TYPE IN A (109)
14583.0 POINTS



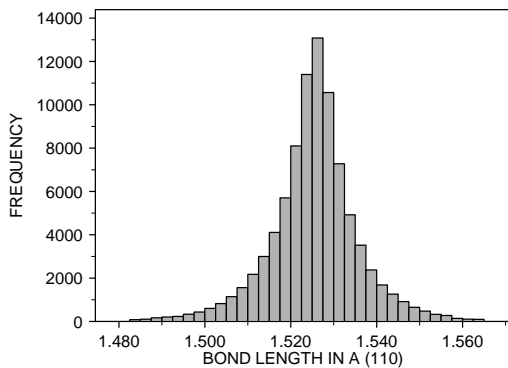
<2 : X-RAY RESOLUTION OF A (35)
ALA:O:C : BOND TYPE IN A (109)
88429.0 POINTS



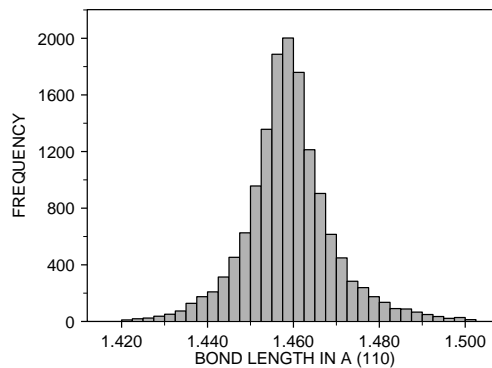
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CYS:SG:CB : BOND TYPE IN A (109)
14568.0 POINTS



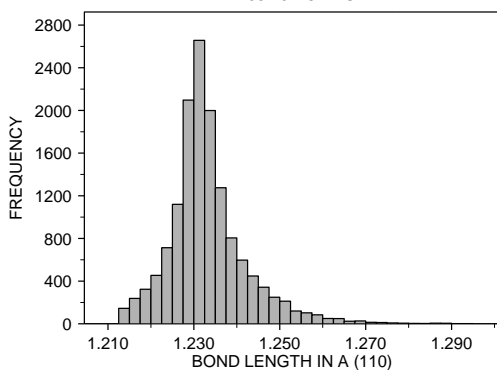
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ALA:C:CA : BOND TYPE IN A (109)
88439.0 POINTS



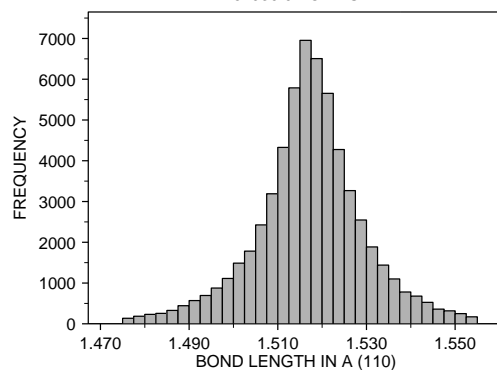
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CYS:N:CA : BOND TYPE IN A (109)
14585.0 POINTS



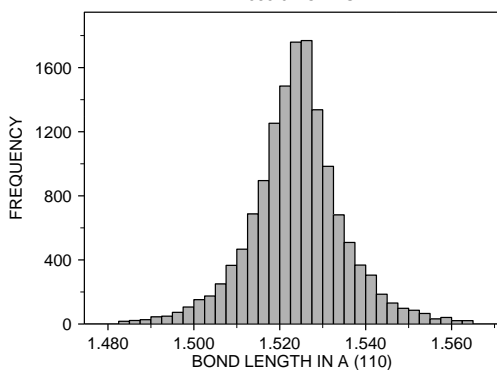
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CYS:O:C : BOND TYPE IN A (109)
14584.0 POINTS



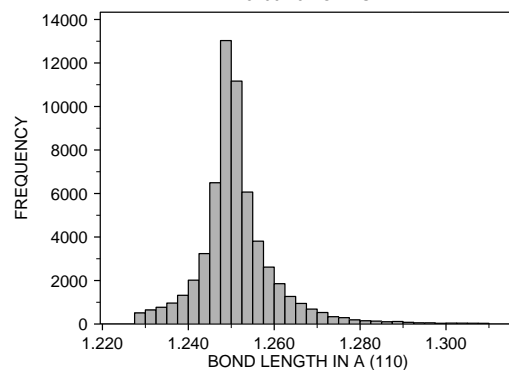
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ASP:CG:CB : BOND TYPE IN A (109)
61930.0 POINTS



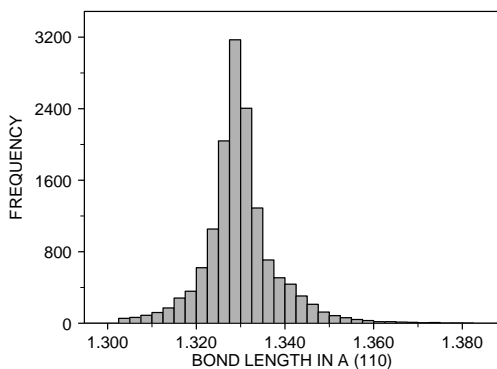
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CYS:C:CA : BOND TYPE IN A (109)
14585.0 POINTS



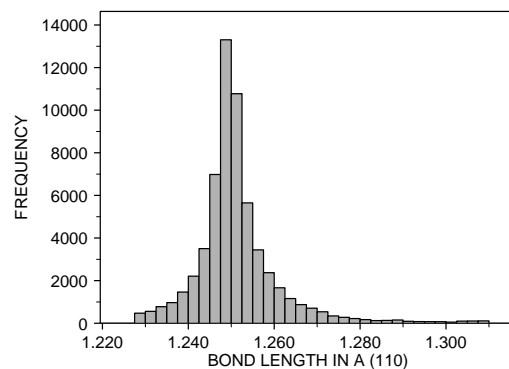
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ASP:OD1:CG : BOND TYPE IN A (109)
61902.0 POINTS



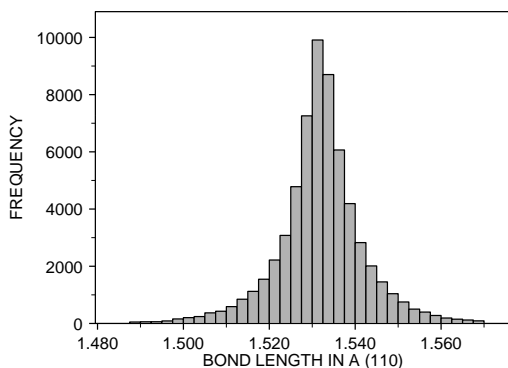
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CYS:C:+N : BOND TYPE IN A (109)
14481.0 POINTS



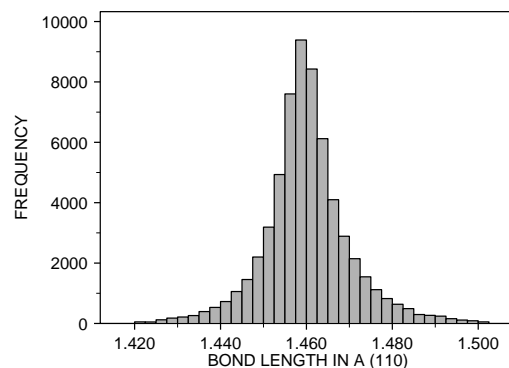
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ASP:OD2:CG : BOND TYPE IN A (109)
61899.0 POINTS



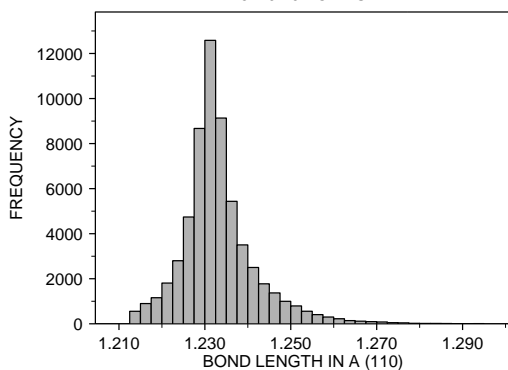
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ASP:CB:CA : BOND TYPE IN A (109)
62241.0 POINTS



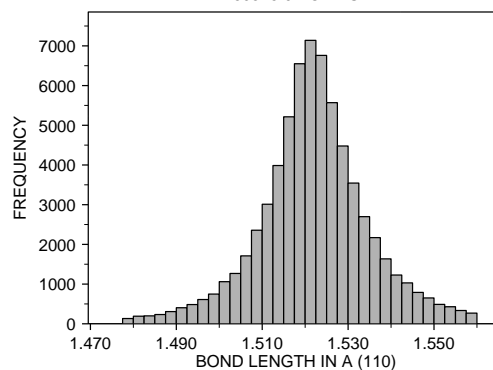
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ASP:N:CA : BOND TYPE IN A (109)
62290.0 POINTS



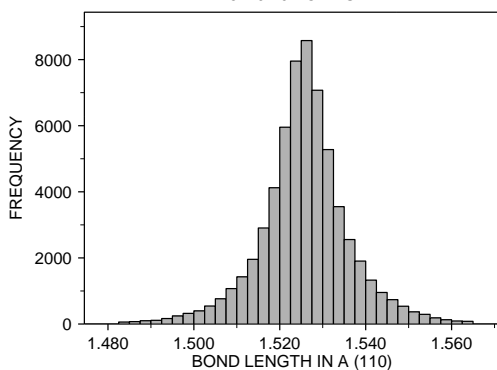
<2 : X-RAY RESOLUTION OF A (35)
ASP:O:C : BOND TYPE IN A (109)
62291.0 POINTS



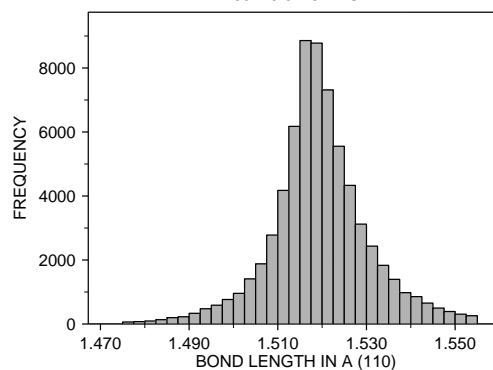
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GLU:CG:CB : BOND TYPE IN A (109)
69545.0 POINTS



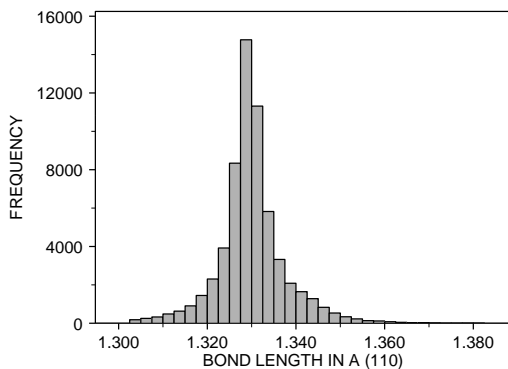
<2 : X-RAY RESOLUTION OF A (35)
ASP:C:CA : BOND TYPE IN A (109)
62291.0 POINTS



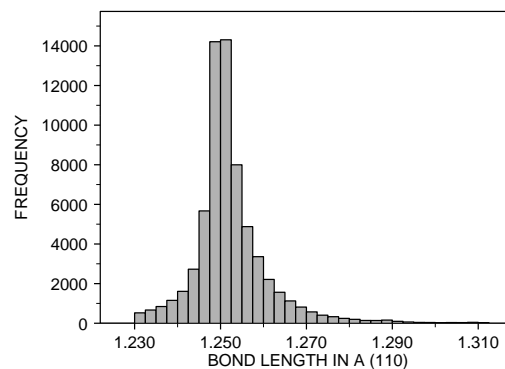
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GLU:CD:CG : BOND TYPE IN A (109)
69249.0 POINTS



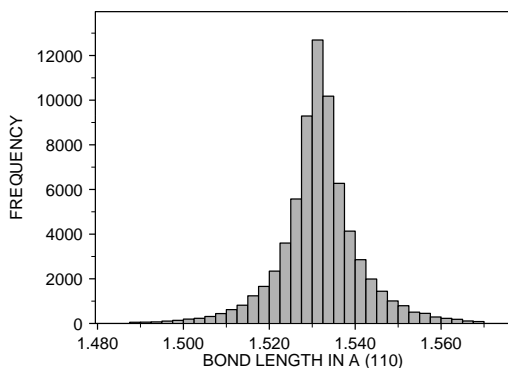
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ASP:C:+N : BOND TYPE IN A (109)
62029.0 POINTS



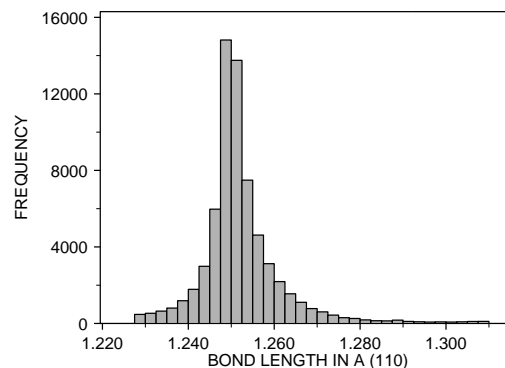
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GLU:OE1:CD : BOND TYPE IN A (109)
69221.0 POINTS

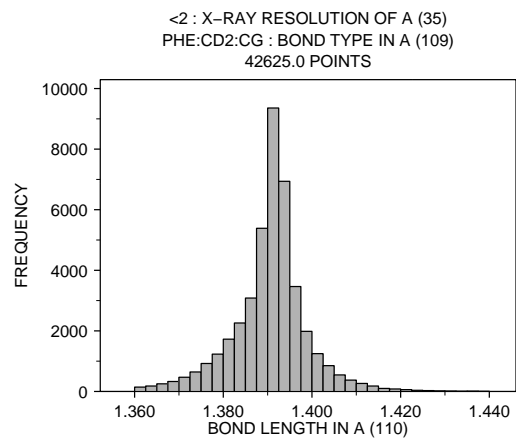
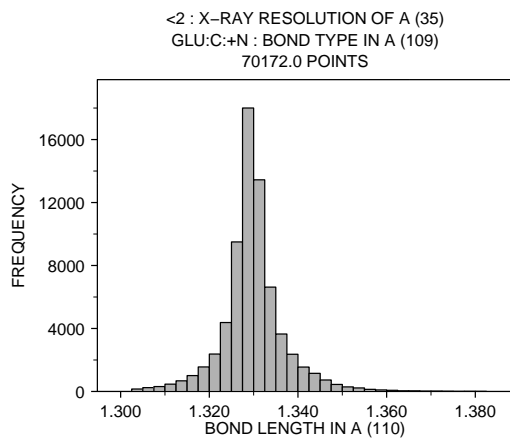
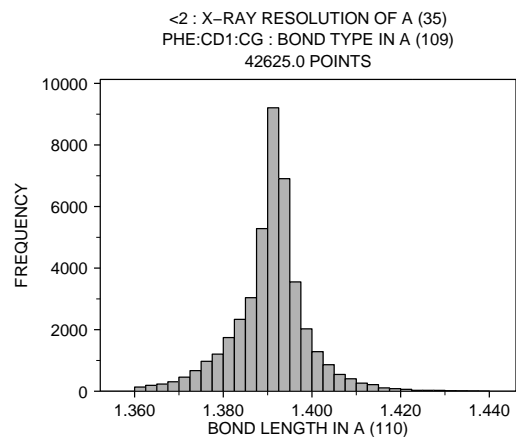
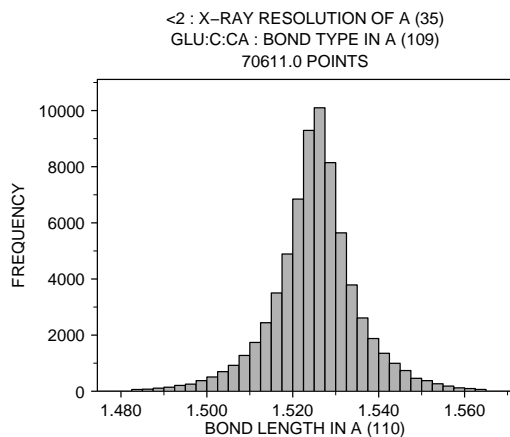
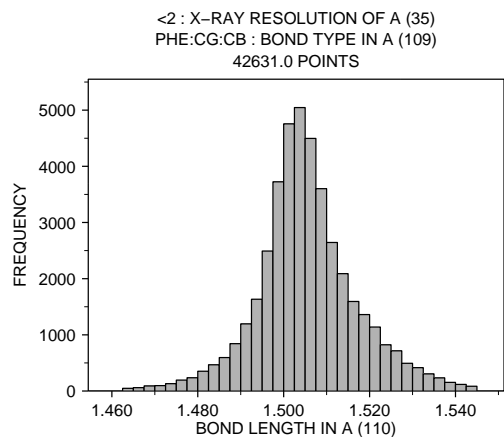
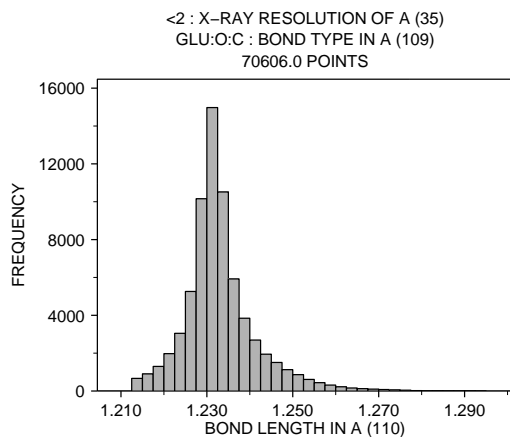
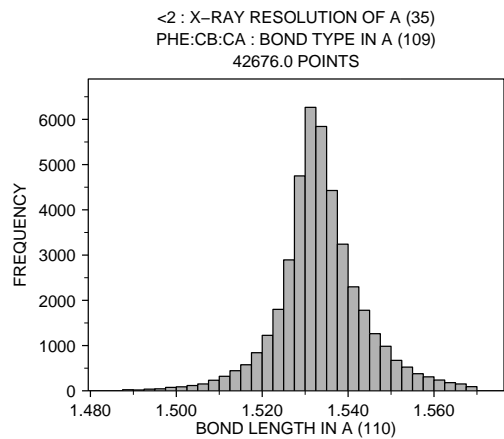
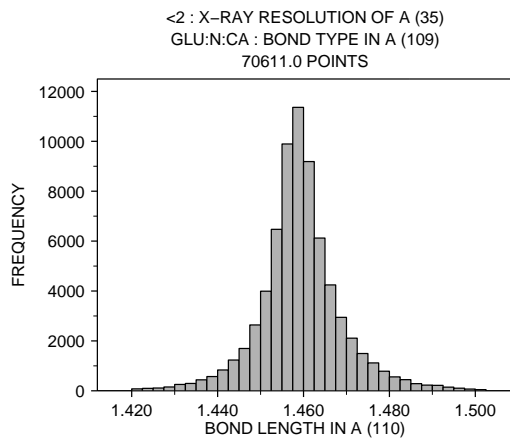


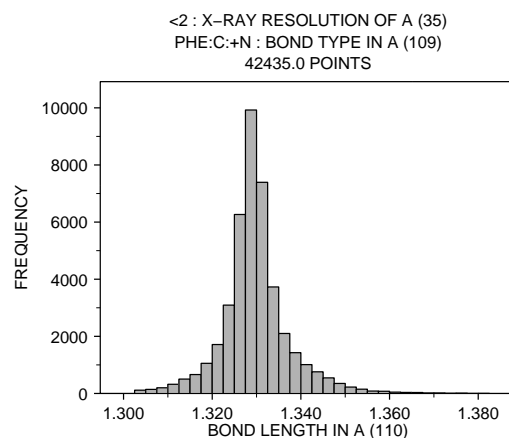
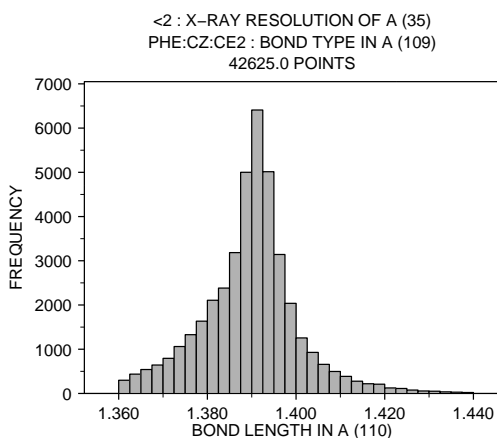
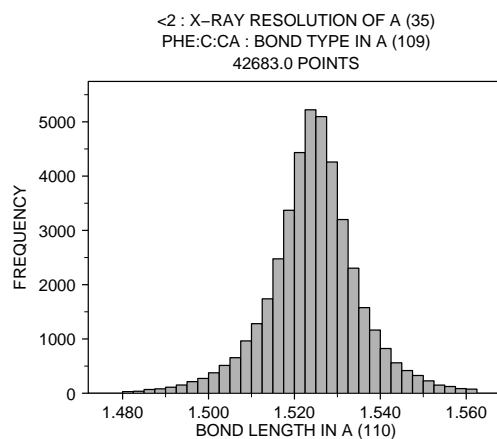
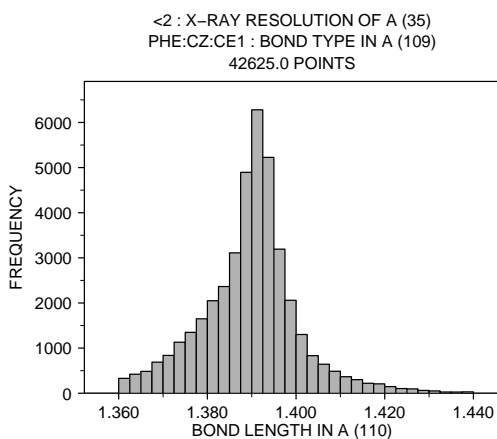
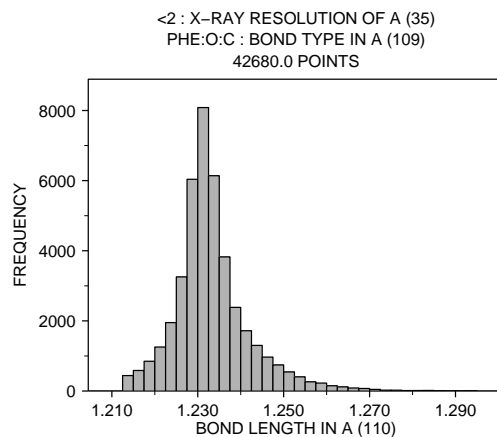
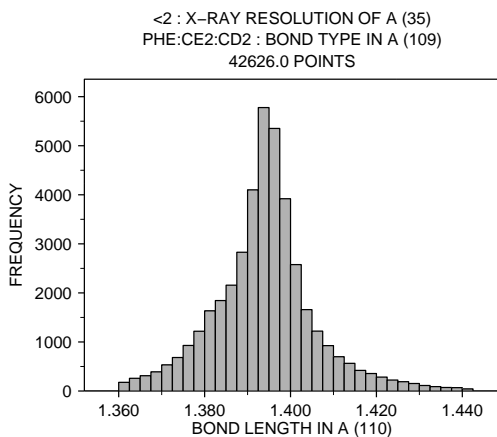
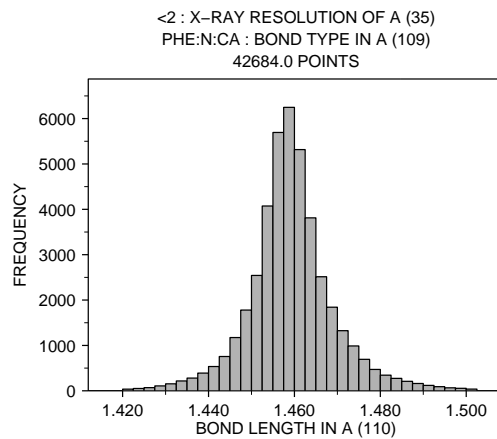
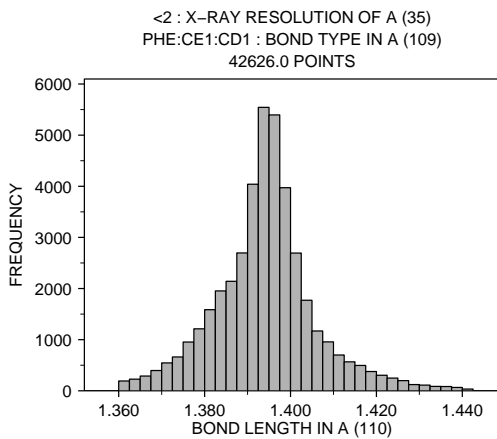
<2 : X-RAY RESOLUTION OF A (35)
GLU:CB:CA : BOND TYPE IN A (109)
70536.0 POINTS



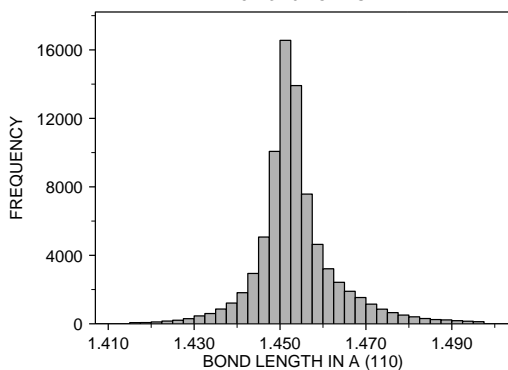
<2 : X-RAY RESOLUTION OF A (35)
GLU:OE2:CD : BOND TYPE IN A (109)
69212.0 POINTS



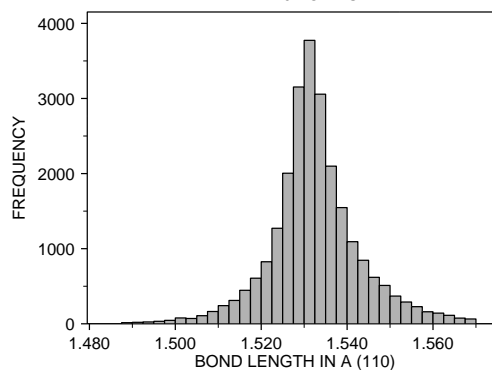




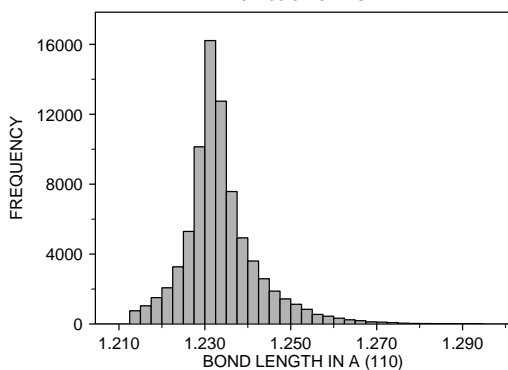
<2 : X-RAY RESOLUTION OF A (35)
GLY:N:CA : BOND TYPE IN A (109)
81137.0 POINTS



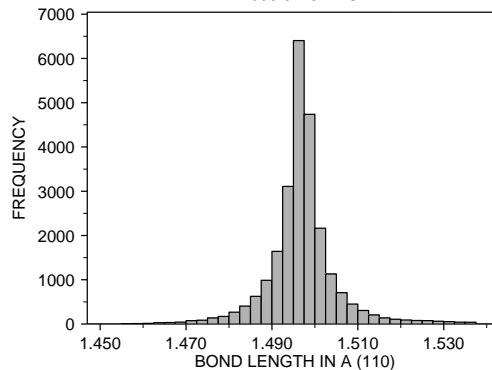
<2 : X-RAY RESOLUTION OF A (35)
HIS:CB:CA : BOND TYPE IN A (109)
24744.0 POINTS



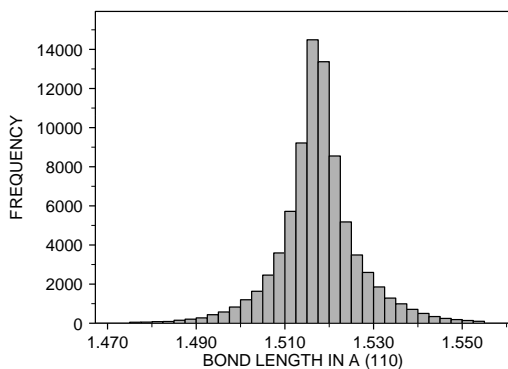
<2 : X-RAY RESOLUTION OF A (35)
GLY:O:C : BOND TYPE IN A (109)
81135.0 POINTS



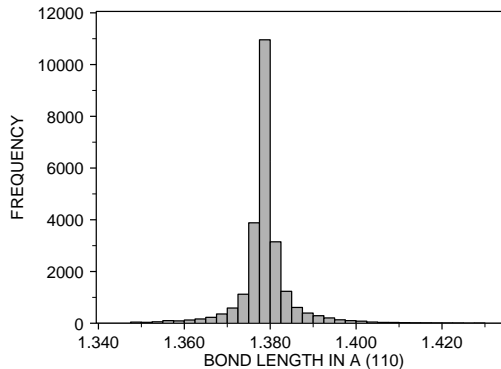
<2 : X-RAY RESOLUTION OF A (35)
HIS:CG:CB : BOND TYPE IN A (109)
24638.0 POINTS



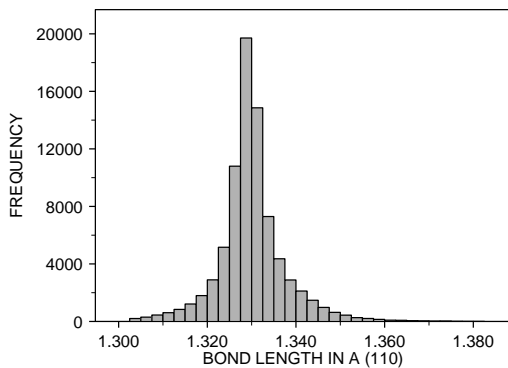
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GLY:C:CA : BOND TYPE IN A (109)
81139.0 POINTS



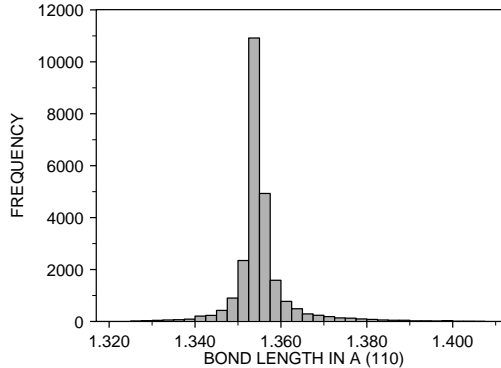
<2 : X-RAY RESOLUTION OF A (35)
HIS:ND1:CG : BOND TYPE IN A (109)
24630.0 POINTS



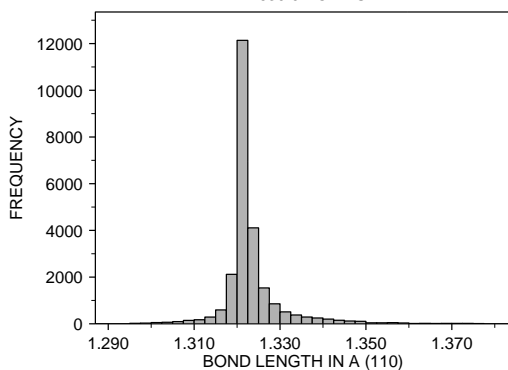
<2 : X-RAY RESOLUTION OF A (35)
GLY:C:+N : BOND TYPE IN A (109)
80708.0 POINTS



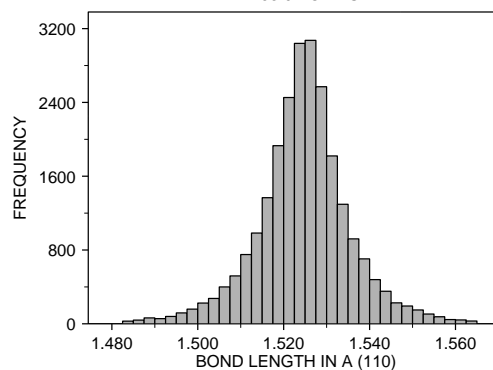
<2 : X-RAY RESOLUTION OF A (35)
HIS:CD2:CG : BOND TYPE IN A (109)
24630.0 POINTS



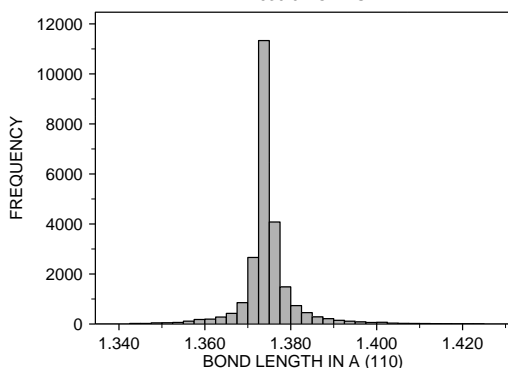
<2 : X-RAY RESOLUTION OF A (35)
HIS:CE1:ND1 : BOND TYPE IN A (109)
24630.0 POINTS



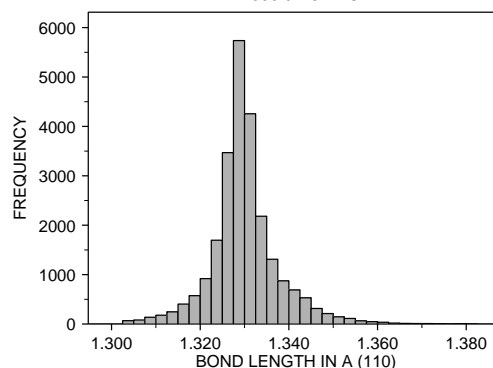
<2 : X-RAY RESOLUTION OF A (35)
HIS:C:CA : BOND TYPE IN A (109)
24753.0 POINTS



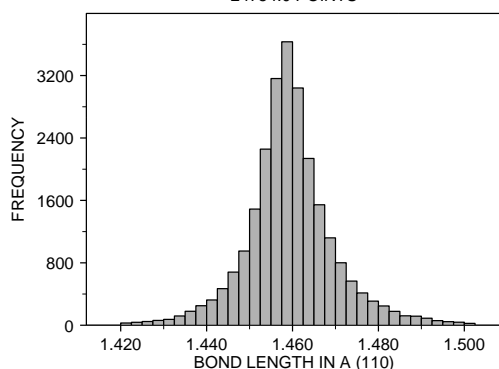
<2 : X-RAY RESOLUTION OF A (35)
HIS:NE2:CD2 : BOND TYPE IN A (109)
24630.0 POINTS



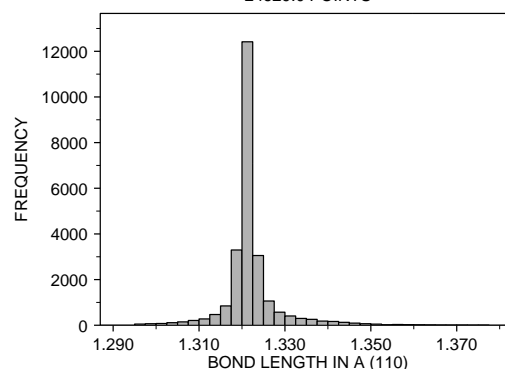
<2 : X-RAY RESOLUTION OF A (35)
HIS:C:+N : BOND TYPE IN A (109)
24586.0 POINTS



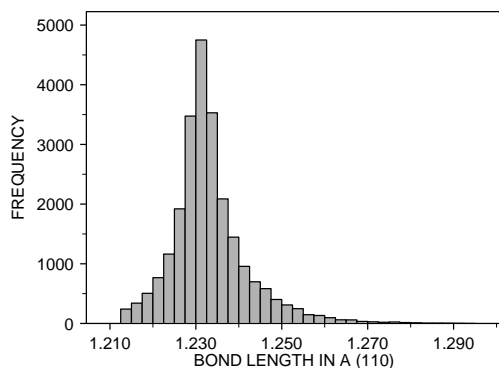
<2 : X-RAY RESOLUTION OF A (35)
HIS:N:CA : BOND TYPE IN A (109)
24754.0 POINTS



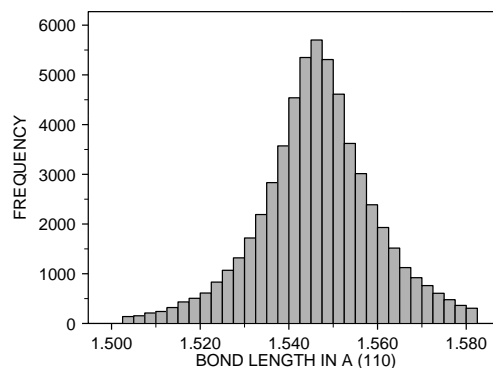
<2 : X-RAY RESOLUTION OF A (35)
HIS:NE2:CE1 : BOND TYPE IN A (109)
24629.0 POINTS

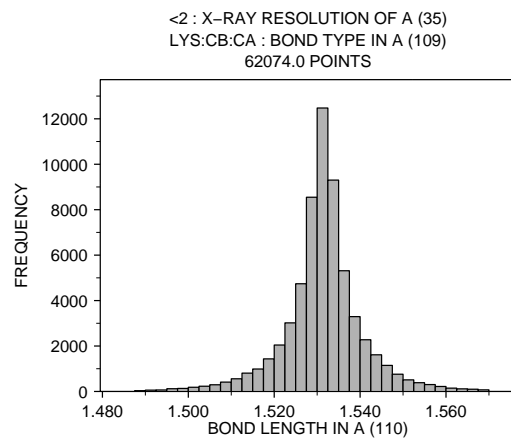
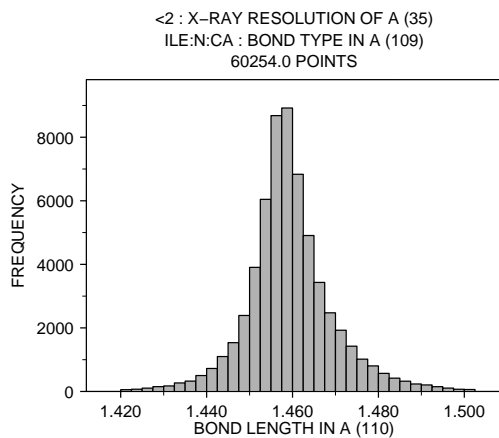
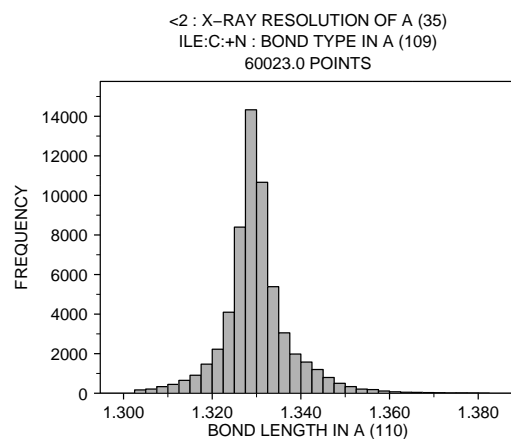
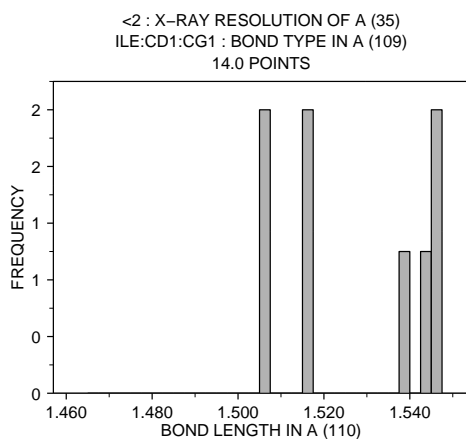
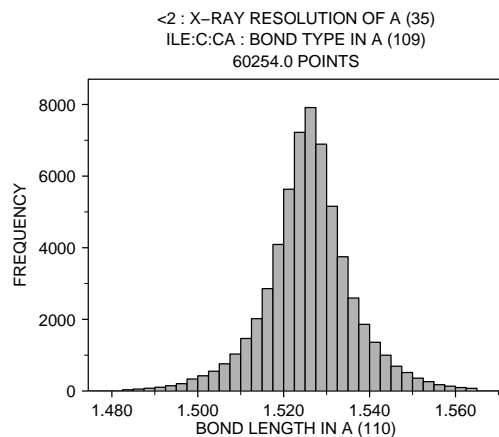
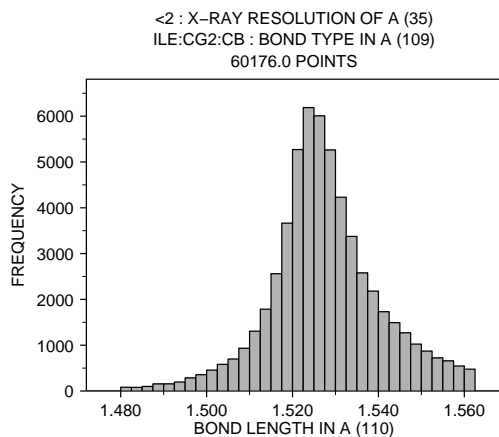
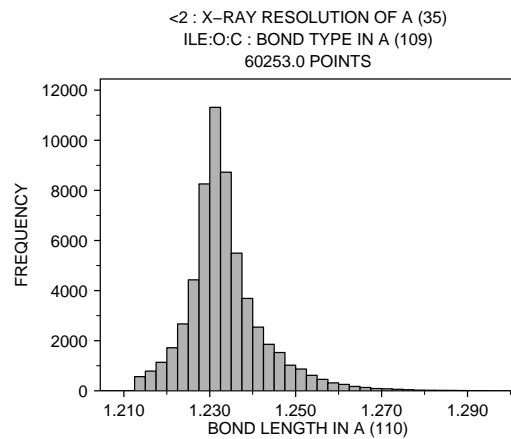
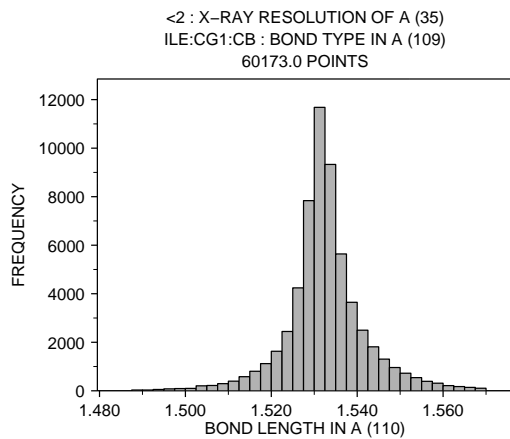


<2 : X-RAY RESOLUTION OF A (35)
HIS:O:C : BOND TYPE IN A (109)
24752.0 POINTS

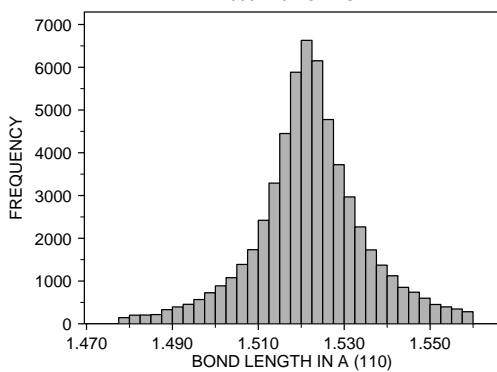


<2 : X-RAY RESOLUTION OF A (35)
ILE:CB:CA : BOND TYPE IN A (109)
60248.0 POINTS

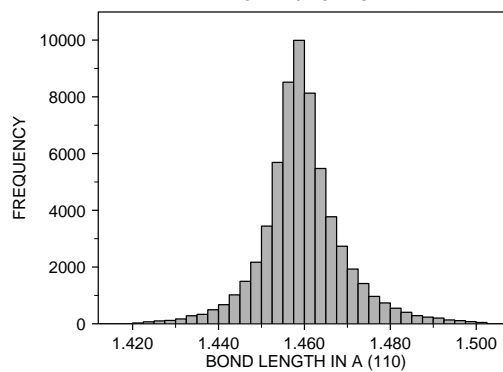




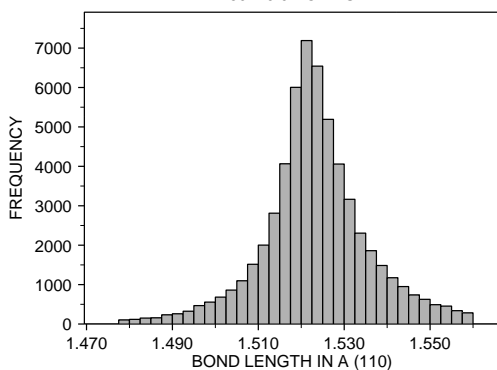
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LYS:CG:CB : BOND TYPE IN A (109)
60911.0 POINTS



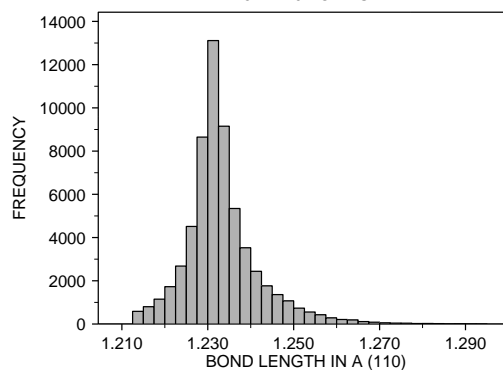
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LYS:N:CA : BOND TYPE IN A (109)
62121.0 POINTS



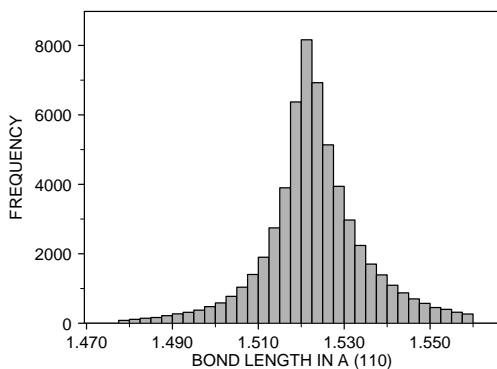
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LYS:CD:CG : BOND TYPE IN A (109)
60470.0 POINTS



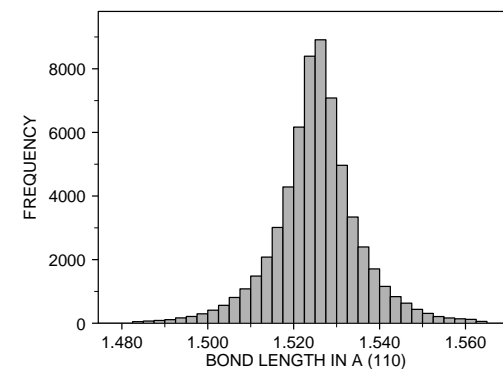
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LYS:O:C : BOND TYPE IN A (109)
62121.0 POINTS



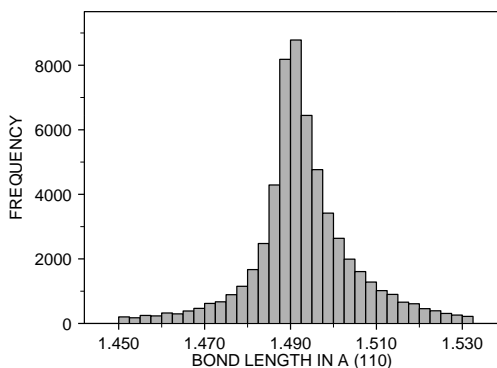
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LYS:CE:CD : BOND TYPE IN A (109)
60083.0 POINTS



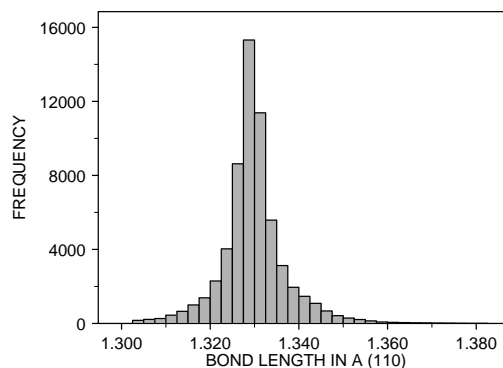
<2 : X-RAY RESOLUTION OF A (35)
LYS:C:CA : BOND TYPE IN A (109)
62121.0 POINTS



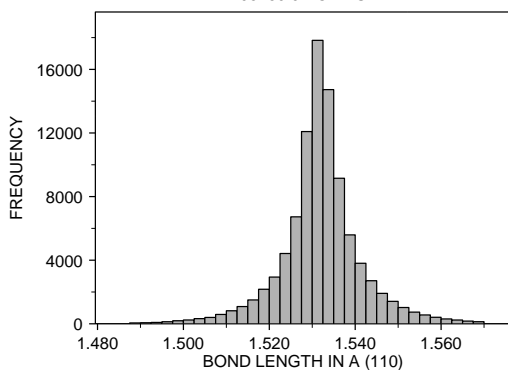
<2 : X-RAY RESOLUTION OF A (35)
LYS:NZ:CE : BOND TYPE IN A (109)
59969.0 POINTS



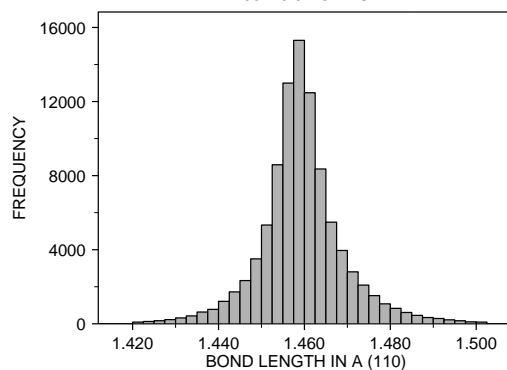
<2 : X-RAY RESOLUTION OF A (35)
LYS:C:+N : BOND TYPE IN A (109)
61631.0 POINTS



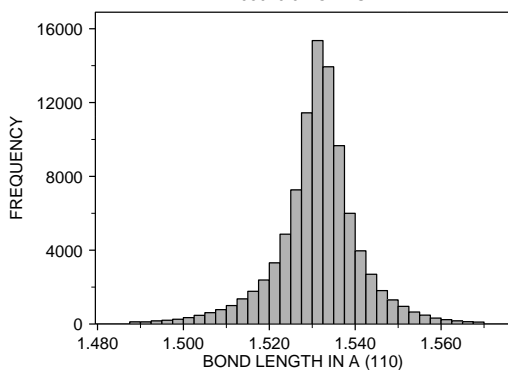
<2 : X-RAY RESOLUTION OF A (35)
LEU:CB:CA : BOND TYPE IN A (109)
95160.0 POINTS



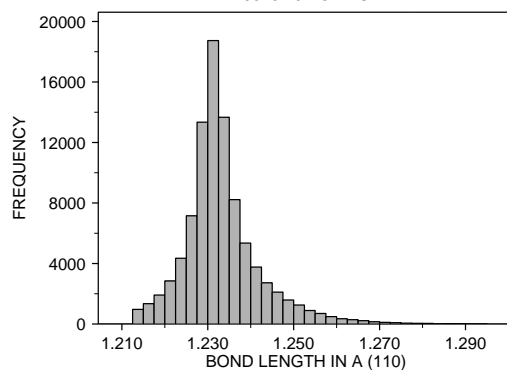
<2 : X-RAY RESOLUTION OF A (35)
LEU:N:CA : BOND TYPE IN A (109)
95170.0 POINTS



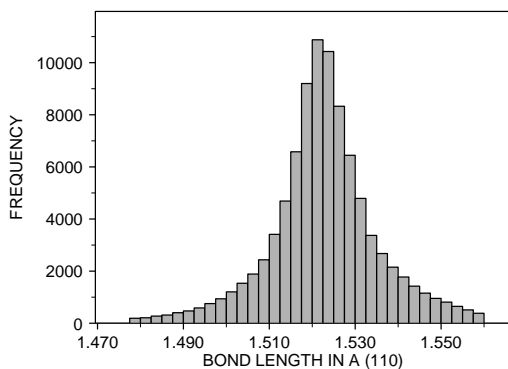
<2 : X-RAY RESOLUTION OF A (35)
LEU:CG:CB : BOND TYPE IN A (109)
95020.0 POINTS



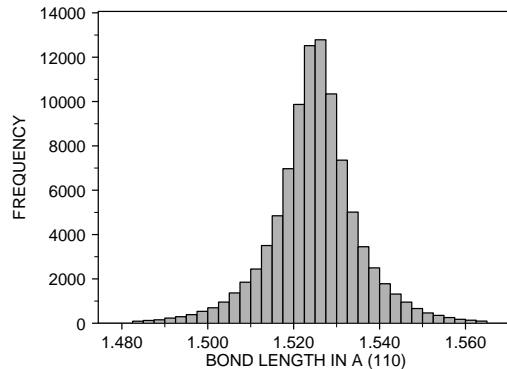
<2 : X-RAY RESOLUTION OF A (35)
LEU:O:C : BOND TYPE IN A (109)
95161.0 POINTS



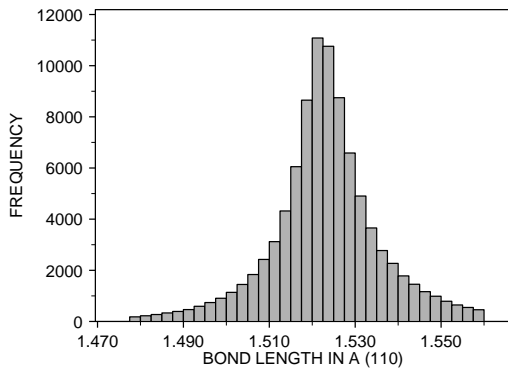
<2 : X-RAY RESOLUTION OF A (35)
LEU:CD1:CG : BOND TYPE IN A (109)
95011.0 POINTS



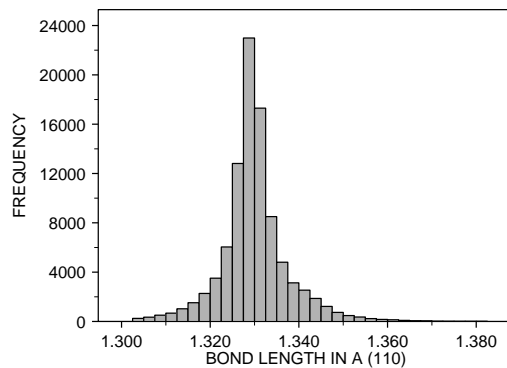
<2 : X-RAY RESOLUTION OF A (35)
LEU:C:CA : BOND TYPE IN A (109)
95169.0 POINTS

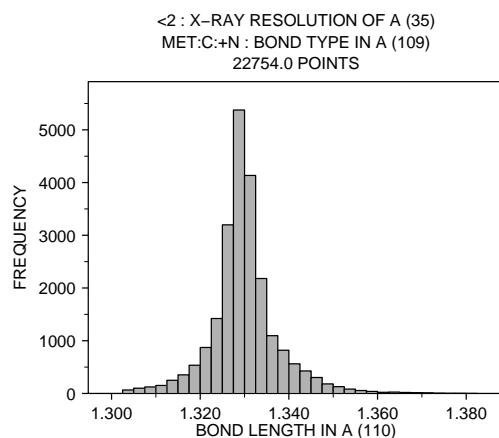
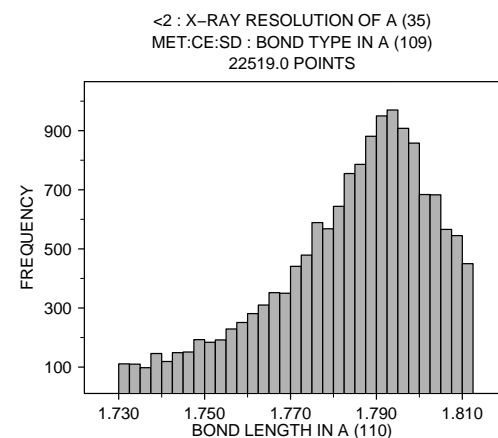
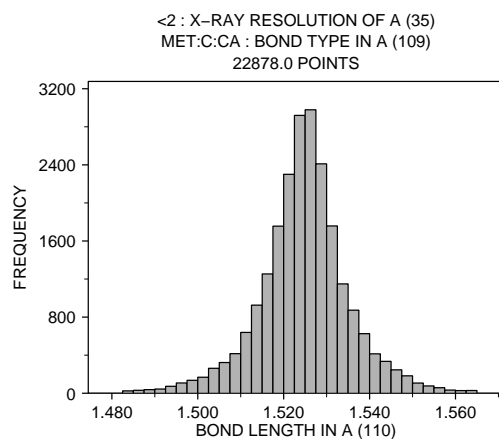
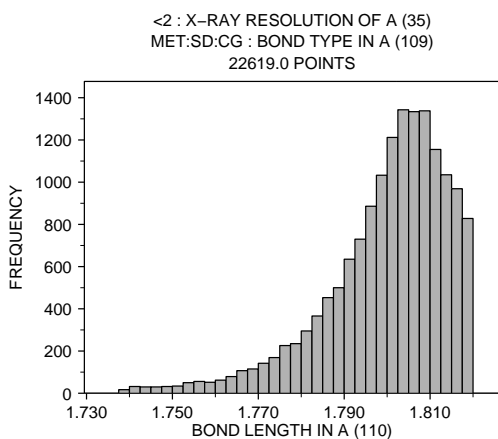
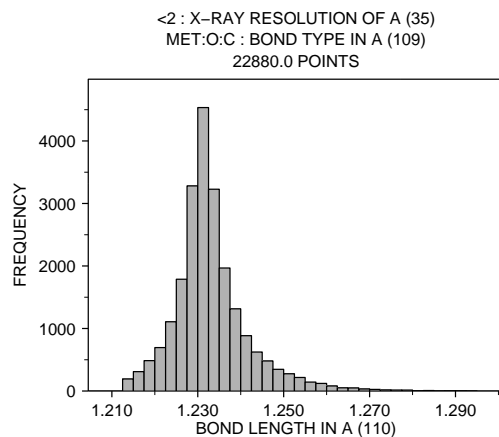
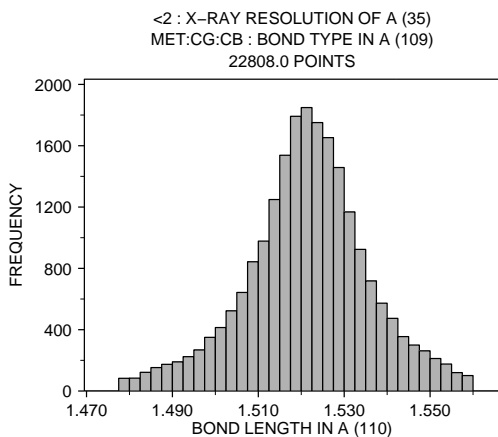
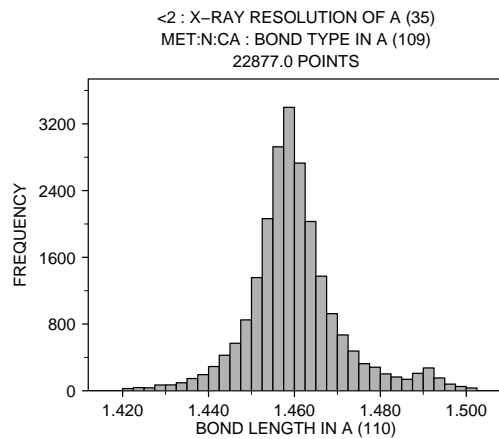
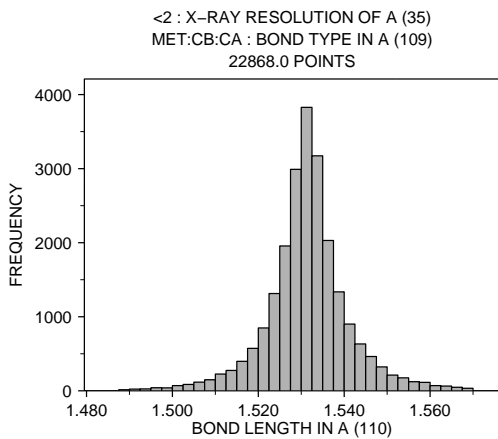


<2 : X-RAY RESOLUTION OF A (35)
LEU:CD2:CG : BOND TYPE IN A (109)
95014.0 POINTS

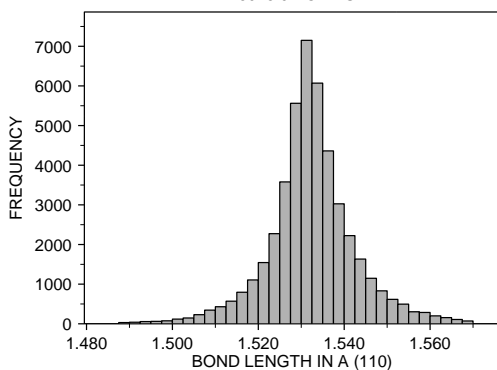


<2 : X-RAY RESOLUTION OF A (35)
LEU:C:+N : BOND TYPE IN A (109)
94618.0 POINTS

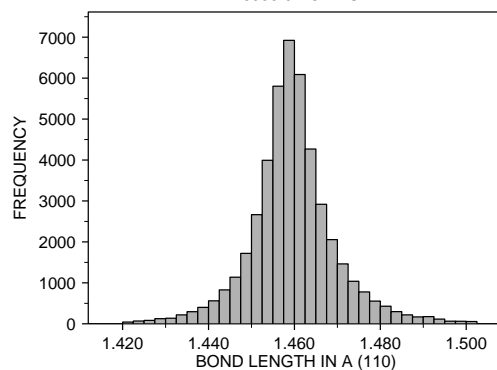




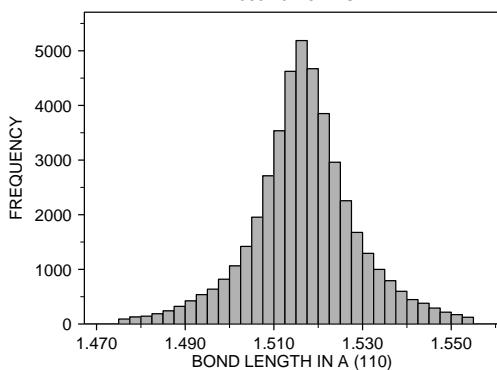
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ASN:CB:CA : BOND TYPE IN A (109)
46016.0 POINTS



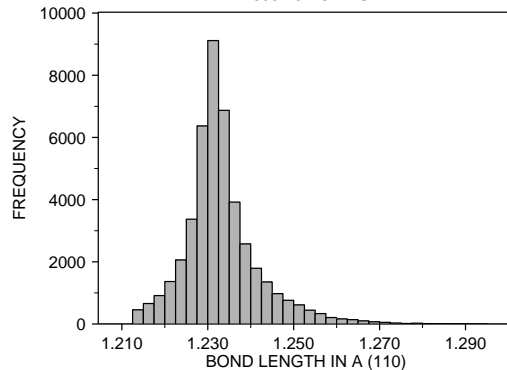
<2 : X-RAY RESOLUTION OF A (35)
ASN:N:CA : BOND TYPE IN A (109)
46033.0 POINTS



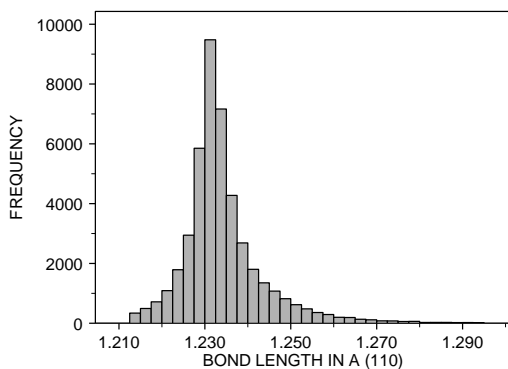
<2 : X-RAY RESOLUTION OF A (35)
ASN:CG:CB : BOND TYPE IN A (109)
45804.0 POINTS



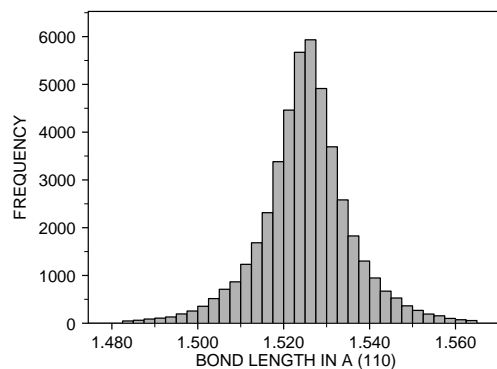
<2 : X-RAY RESOLUTION OF A (35)
ASN:O:C : BOND TYPE IN A (109)
46032.0 POINTS



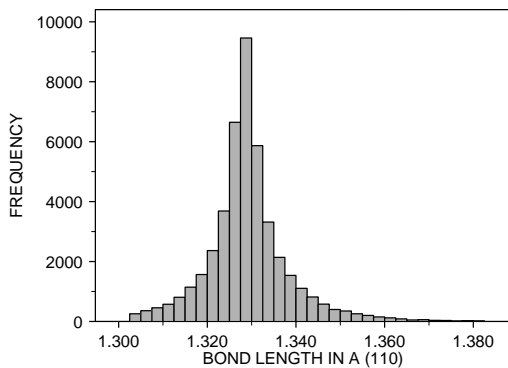
<2 : X-RAY RESOLUTION OF A (35)
ASN:OD1:CG : BOND TYPE IN A (109)
45788.0 POINTS



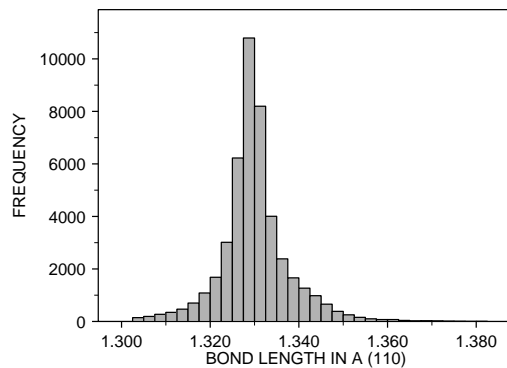
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ASN:C:CA : BOND TYPE IN A (109)
46033.0 POINTS



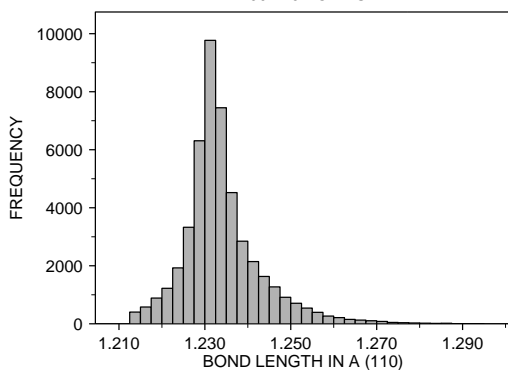
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ASN:ND2:CG : BOND TYPE IN A (109)
45786.0 POINTS



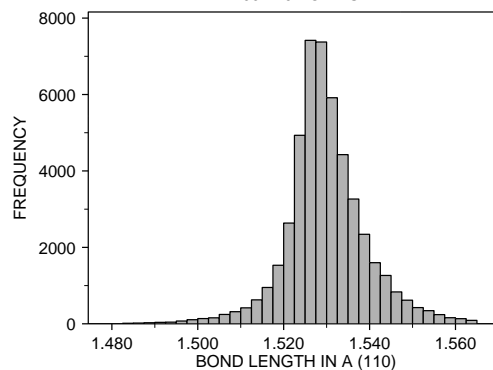
<2 : X-RAY RESOLUTION OF A (35)
ASN:C:+N : BOND TYPE IN A (109)
45754.0 POINTS



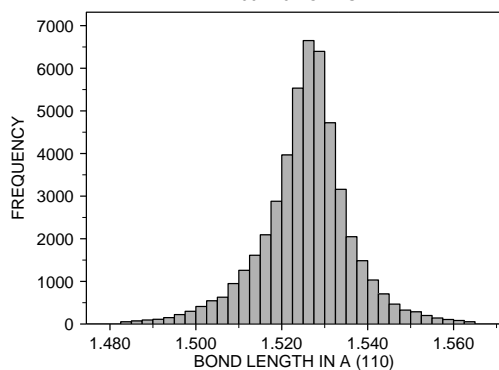
<2 : X-RAY RESOLUTION OF A (35)
PRO:O:C : BOND TYPE IN A (109)
49041.0 POINTS



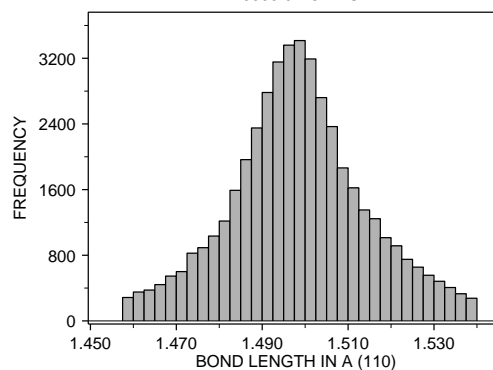
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PRO:CA:CB : BOND TYPE IN A (109)
49041.0 POINTS



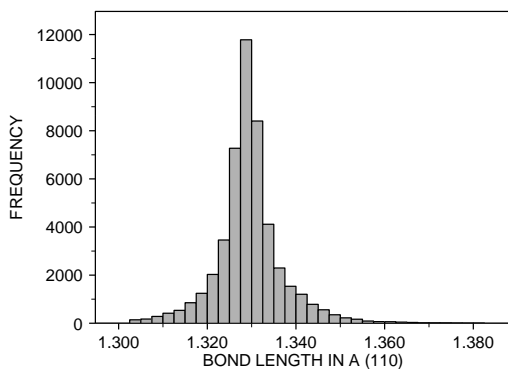
<2 : X-RAY RESOLUTION OF A (35)
PRO:C:CA : BOND TYPE IN A (109)
49047.0 POINTS



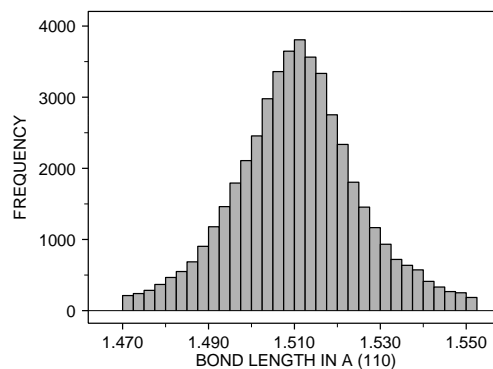
<2 : X-RAY RESOLUTION OF A (35)
PRO:CB:CG : BOND TYPE IN A (109)
49003.0 POINTS



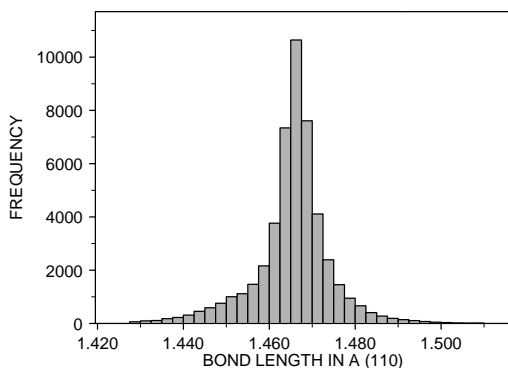
<2 : X-RAY RESOLUTION OF A (35)
PRO:C:+N : BOND TYPE IN A (109)
48677.0 POINTS



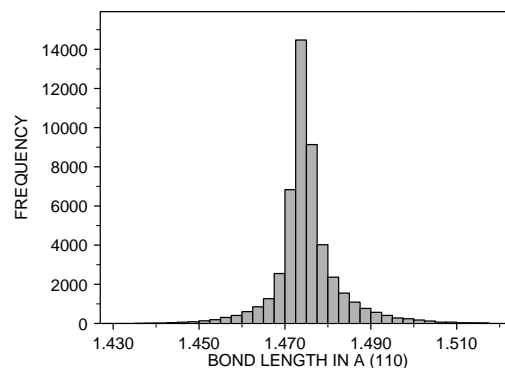
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PRO:CG:CD : BOND TYPE IN A (109)
49003.0 POINTS



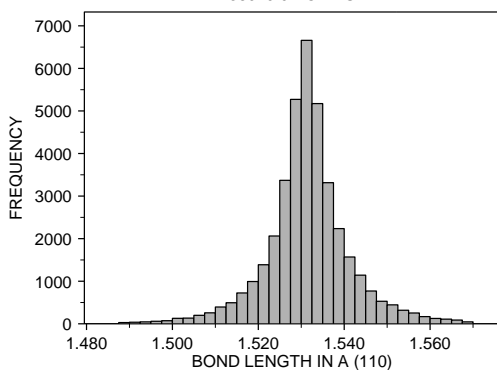
<2 : X-RAY RESOLUTION OF A (35)
PRO:N:CA : BOND TYPE IN A (109)
49047.0 POINTS



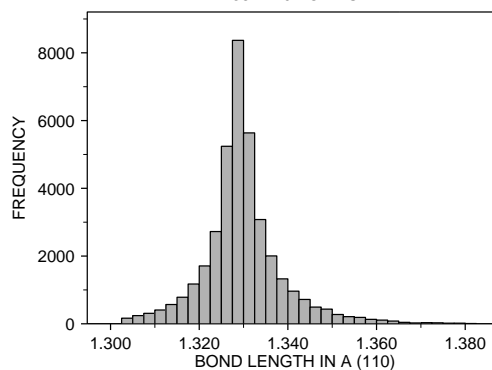
<2 : X-RAY RESOLUTION OF A (35)
PRO:N:CD : BOND TYPE IN A (109)
49004.0 POINTS



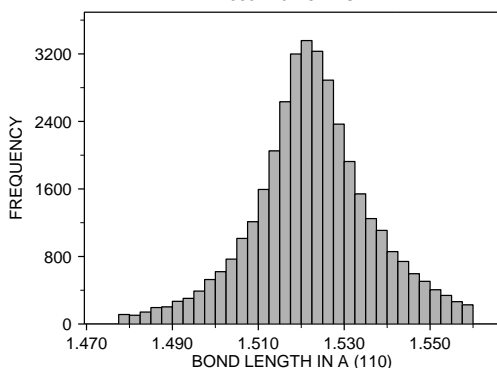
<2 : X-RAY RESOLUTION OF A (35)
GLN:CB:CA : BOND TYPE IN A (109)
38919.0 POINTS



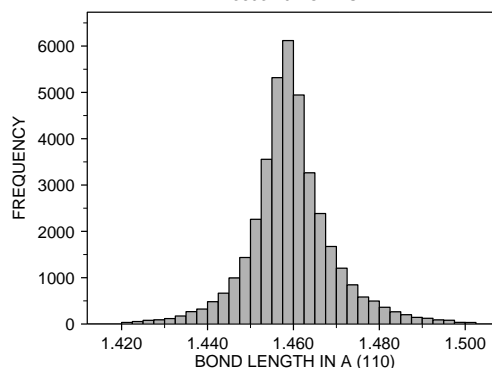
<2 : X-RAY RESOLUTION OF A (35)
GLN:NE2:CD : BOND TYPE IN A (109)
38477.0 POINTS



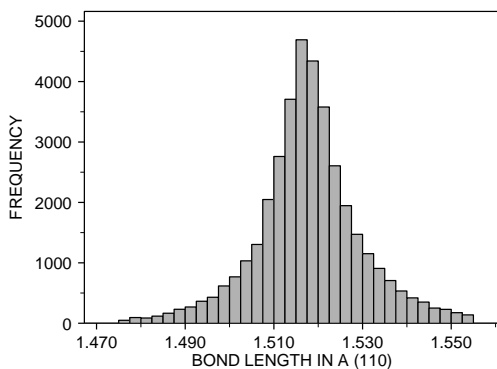
<2 : X-RAY RESOLUTION OF A (35)
GLN:CG:CB : BOND TYPE IN A (109)
38577.0 POINTS



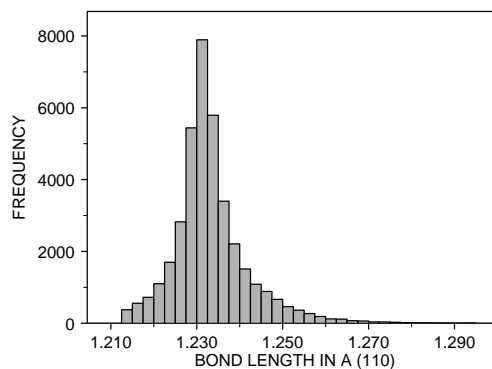
<2 : X-RAY RESOLUTION OF A (35)
GLN:N:CA : BOND TYPE IN A (109)
38937.0 POINTS



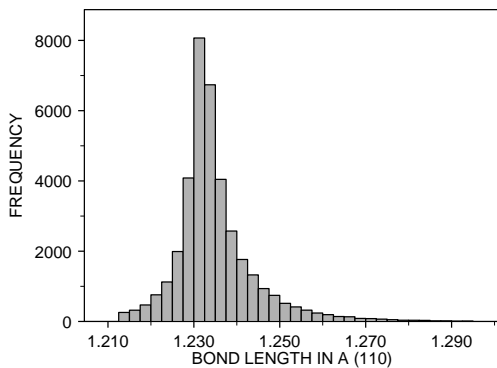
<2 : X-RAY RESOLUTION OF A (35)
GLN:CD:CG : BOND TYPE IN A (109)
38501.0 POINTS



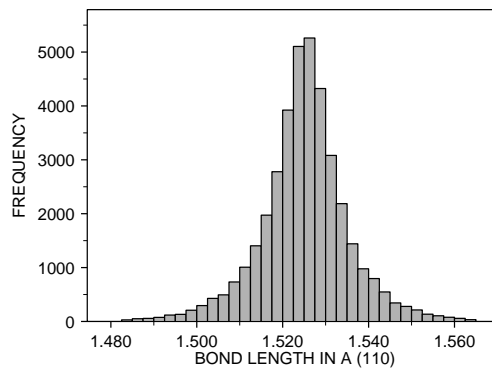
<2 : X-RAY RESOLUTION OF A (35)
GLN:O:C : BOND TYPE IN A (109)
38938.0 POINTS

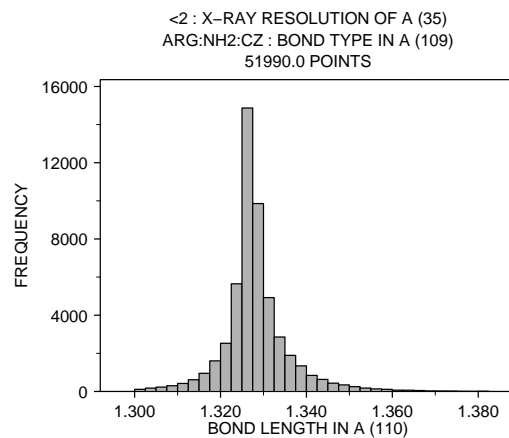
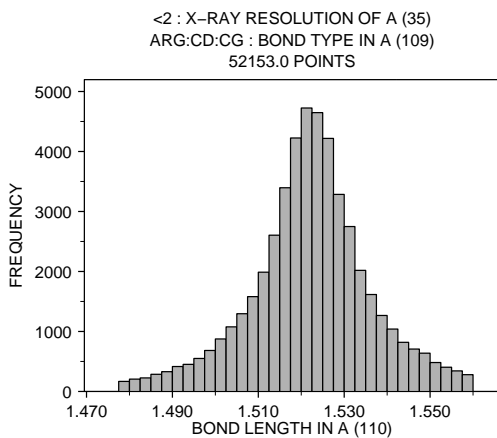
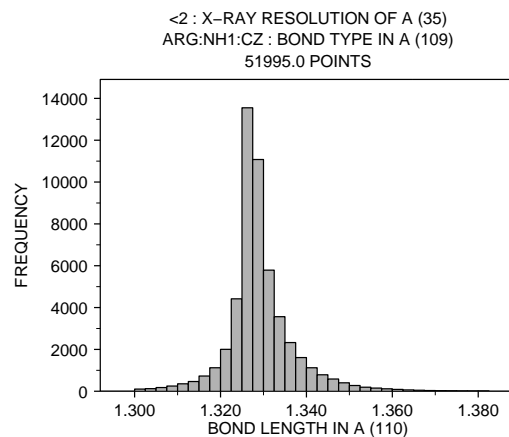
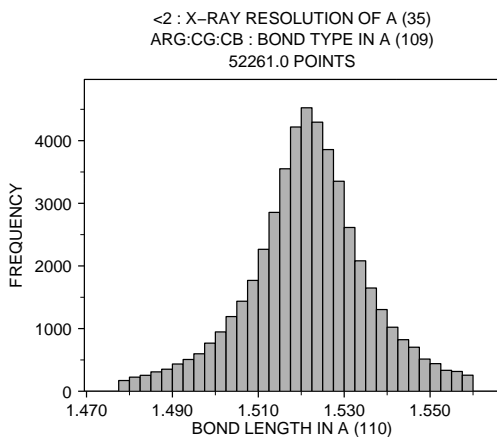
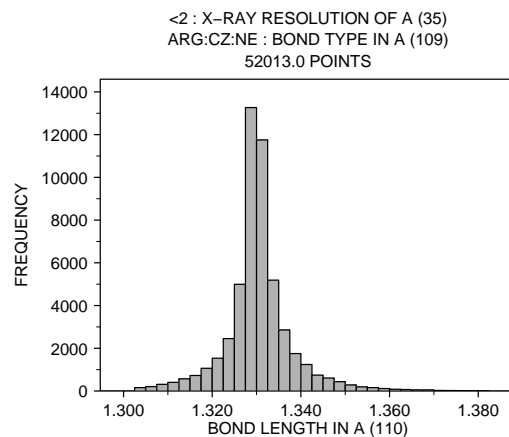
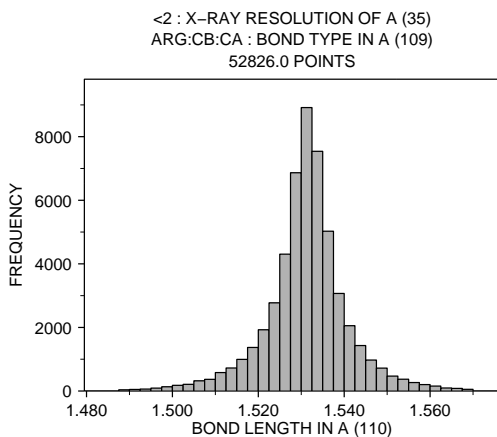
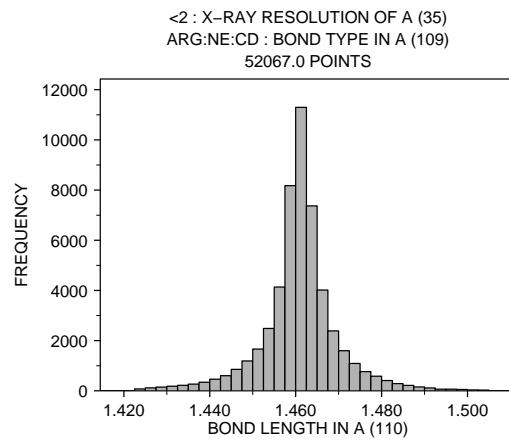
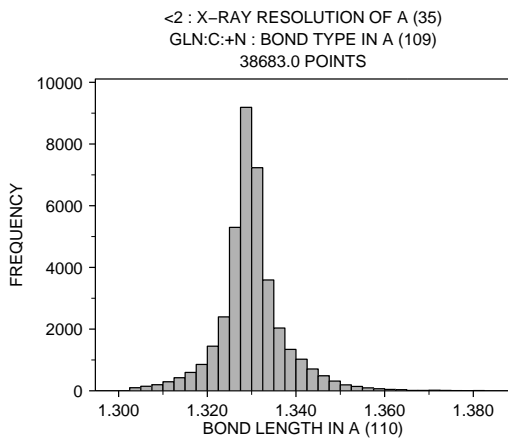


<2 : X-RAY RESOLUTION OF A (35)
GLN:OE1:CD : BOND TYPE IN A (109)
38478.0 POINTS

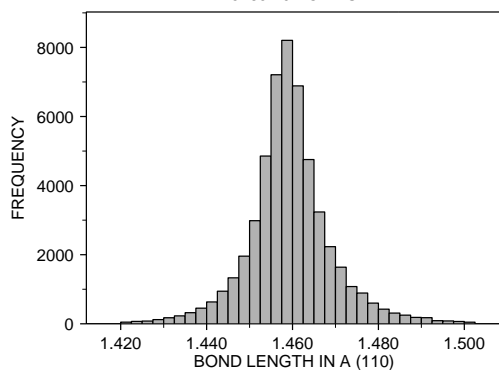


<2 : X-RAY RESOLUTION OF A (35)
GLN:C:CA : BOND TYPE IN A (109)
38938.0 POINTS

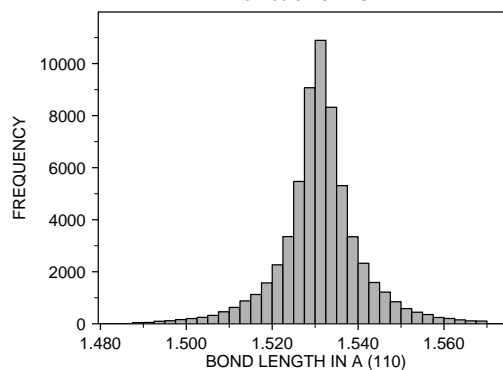




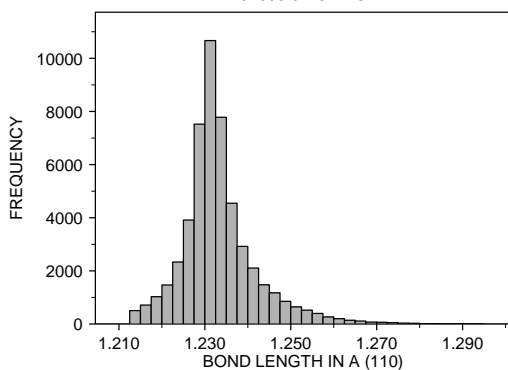
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ARG:N:CA : BOND TYPE IN A (109)
52857.0 POINTS



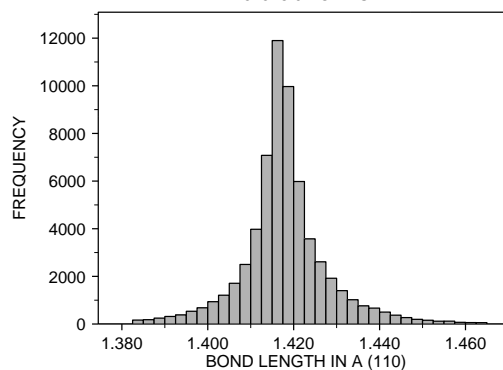
<2 : X-RAY RESOLUTION OF A (35)
SER:CB:CA : BOND TYPE IN A (109)
62709.0 POINTS



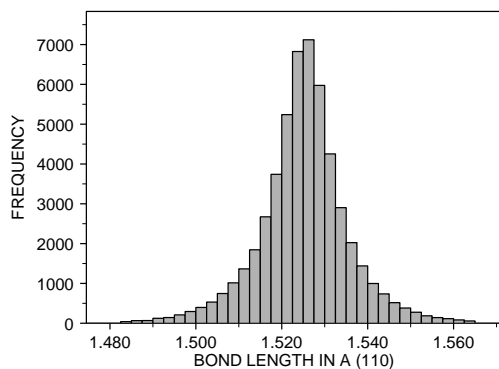
<2 : X-RAY RESOLUTION OF A (35)
ARG:O:C : BOND TYPE IN A (109)
52855.0 POINTS



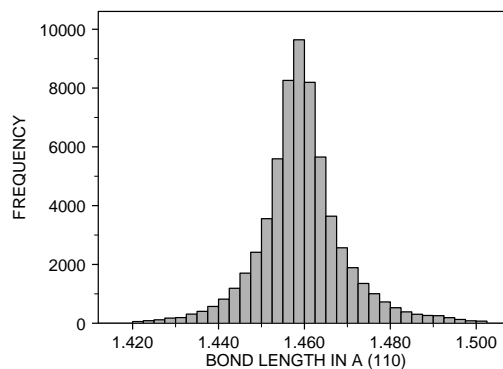
<2 : X-RAY RESOLUTION OF A (35)
SER:OG:CB : BOND TYPE IN A (109)
62528.0 POINTS



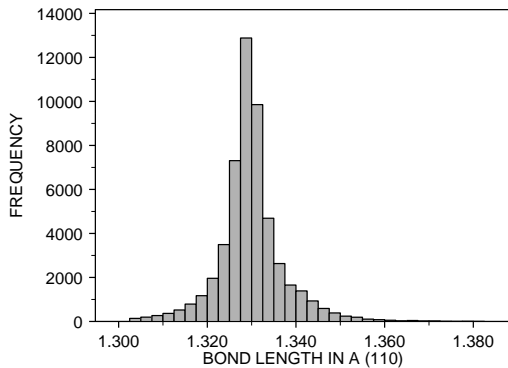
<2 : X-RAY RESOLUTION OF A (35)
ARG:C:CA : BOND TYPE IN A (109)
52857.0 POINTS



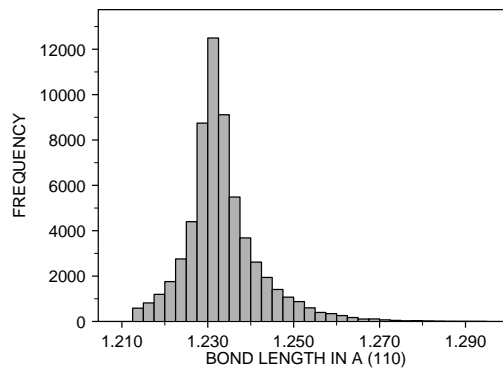
<2 : X-RAY RESOLUTION OF A (35)
SER:N:CA : BOND TYPE IN A (109)
62735.0 POINTS



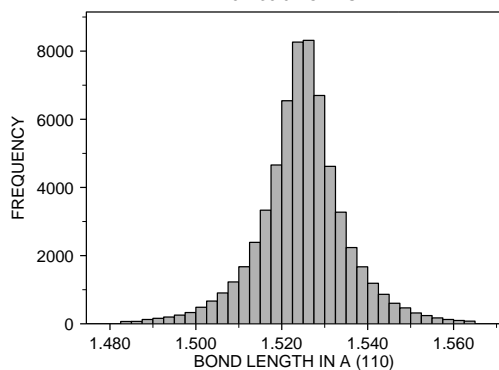
<2 : X-RAY RESOLUTION OF A (35)
ARG:C:+N : BOND TYPE IN A (109)
52511.0 POINTS



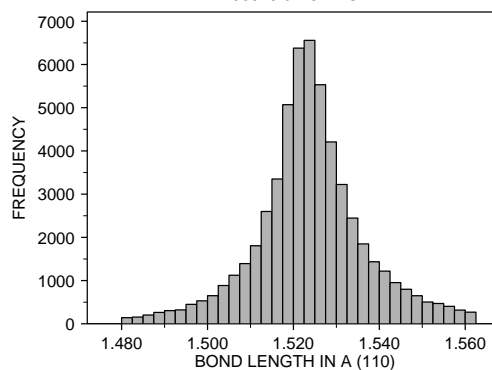
<2 : X-RAY RESOLUTION OF A (35)
SER:O:C : BOND TYPE IN A (109)
62729.0 POINTS



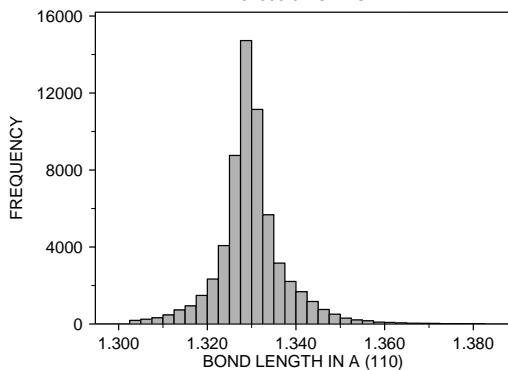
<2 : X-RAY RESOLUTION OF A (35)
SER:C:CA : BOND TYPE IN A (109)
62736.0 POINTS



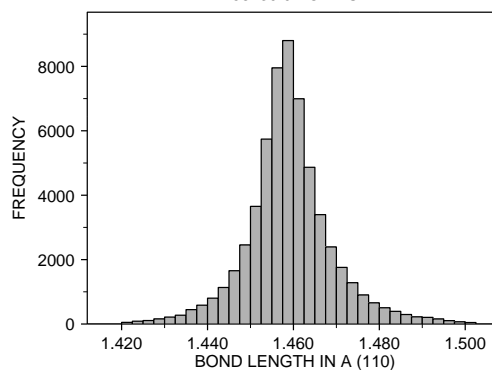
<2 : X-RAY RESOLUTION OF A (35)
THR:CG2:CB : BOND TYPE IN A (109)
58618.0 POINTS



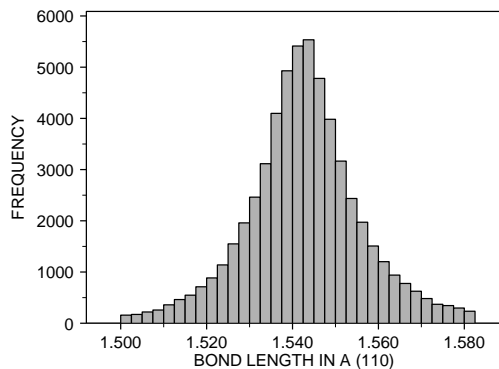
<2 : X-RAY RESOLUTION OF A (35)
SER:C:+N : BOND TYPE IN A (109)
62309.0 POINTS



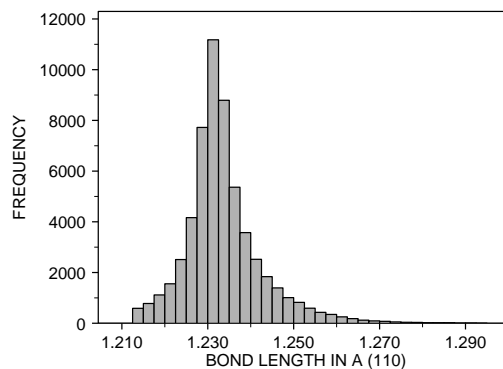
<2 : X-RAY RESOLUTION OF A (35)
THR:N:CA : BOND TYPE IN A (109)
58739.0 POINTS



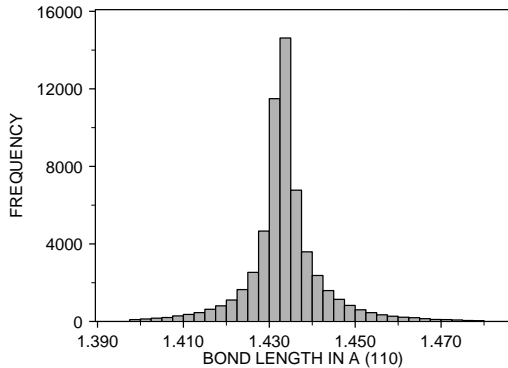
<2 : X-RAY RESOLUTION OF A (35)
THR:CB:CA : BOND TYPE IN A (109)
58725.0 POINTS



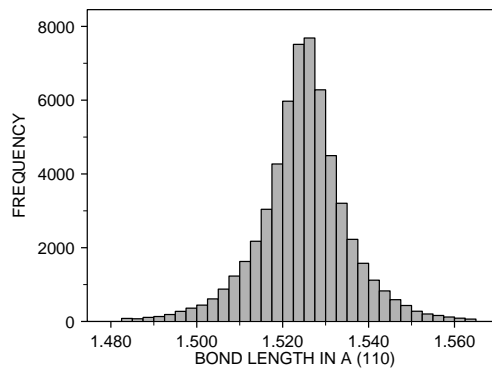
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THR:O:C : BOND TYPE IN A (109)
58740.0 POINTS

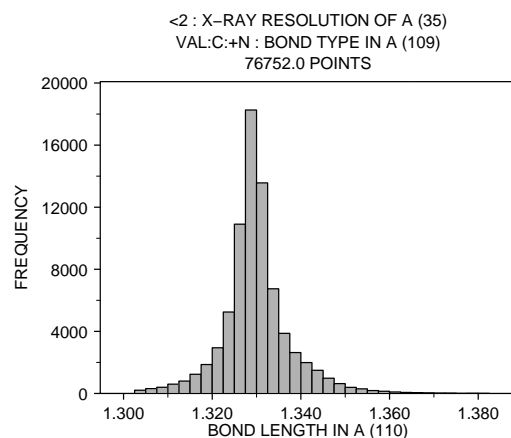
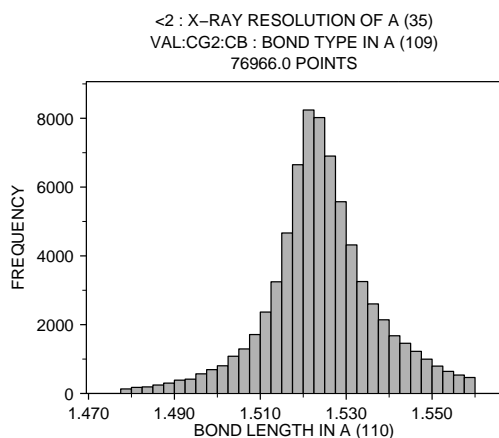
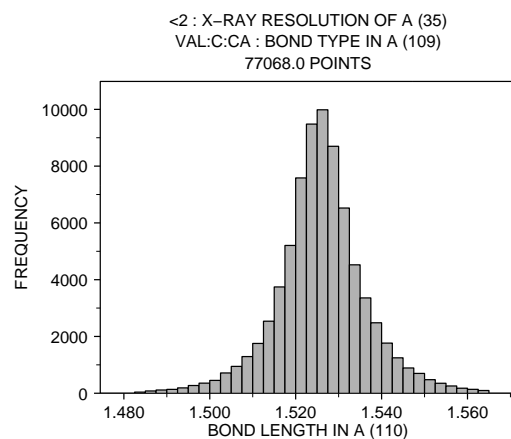
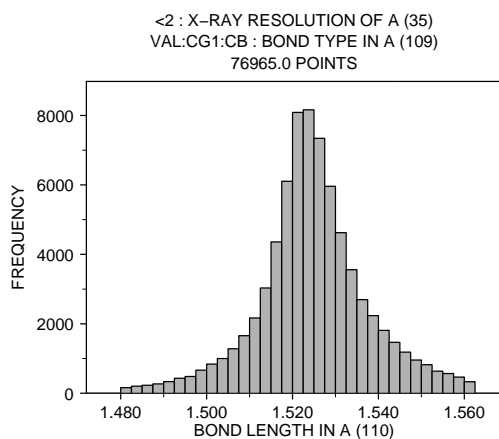
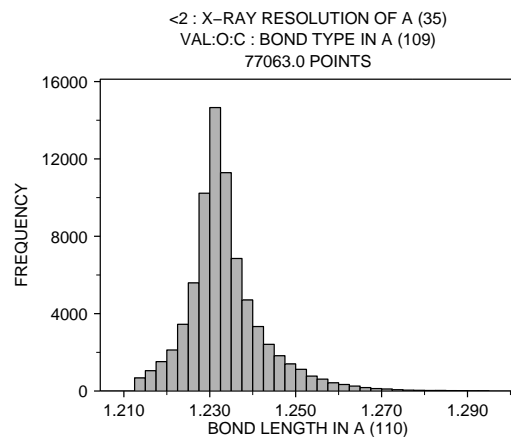
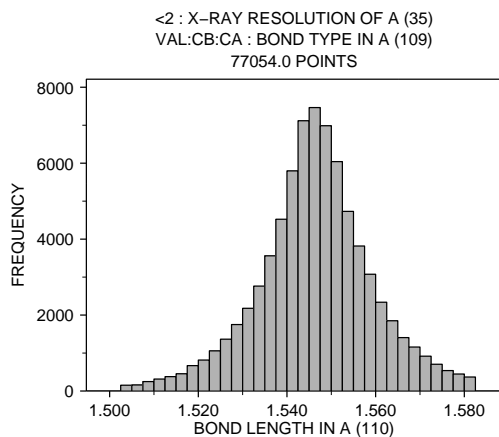
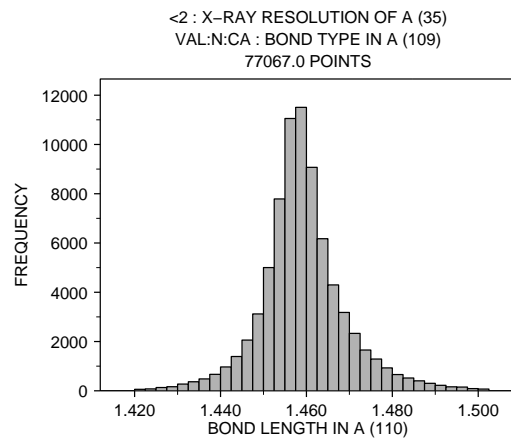
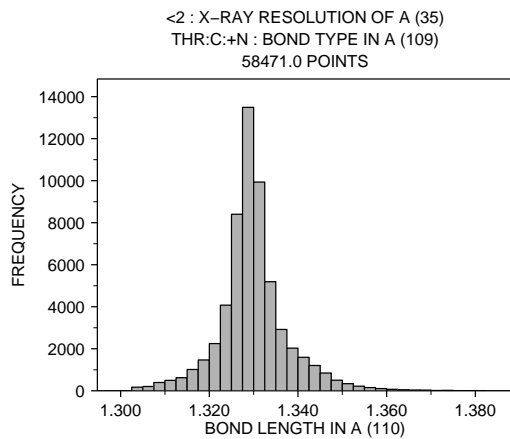


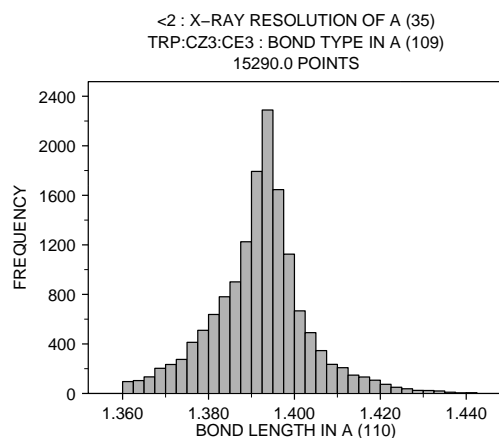
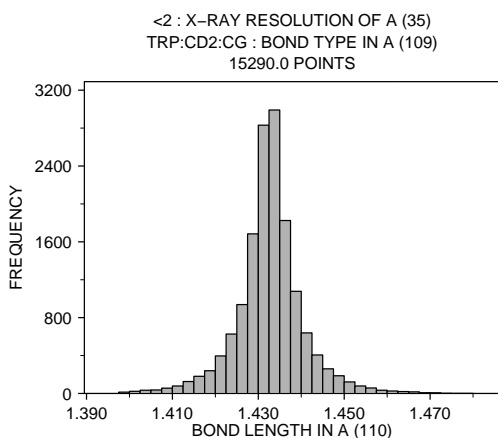
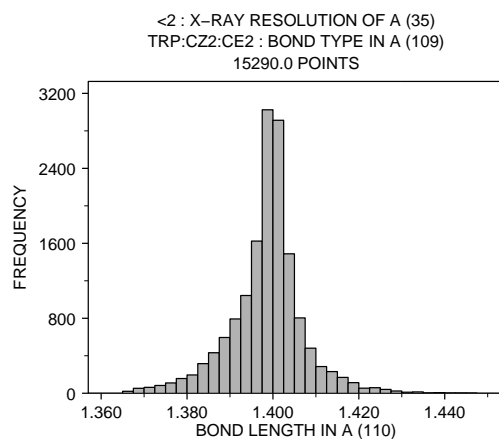
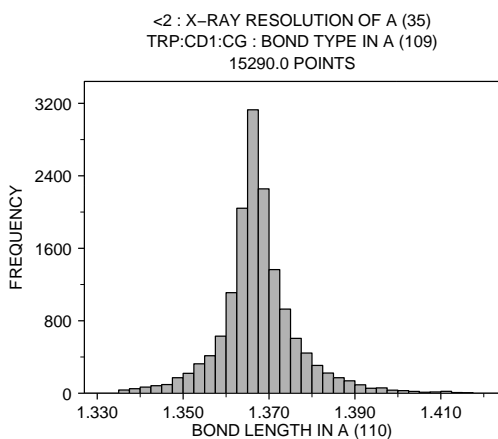
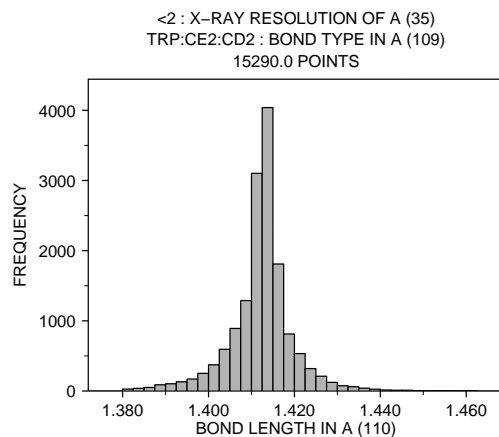
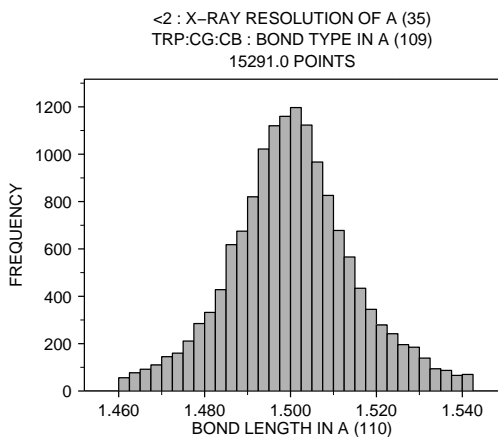
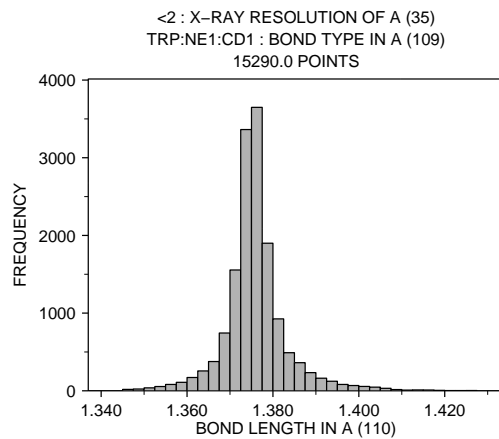
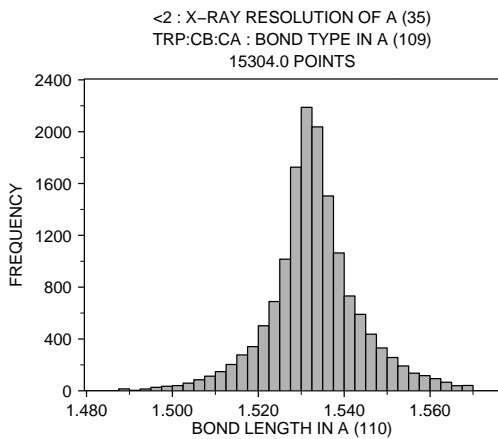
<2 : X-RAY RESOLUTION OF A (35)
THR:OG1:CB : BOND TYPE IN A (109)
58628.0 POINTS

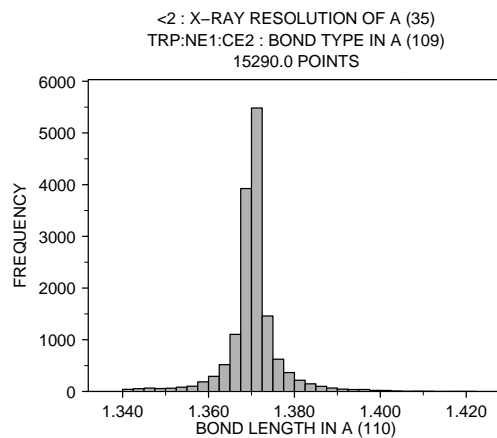
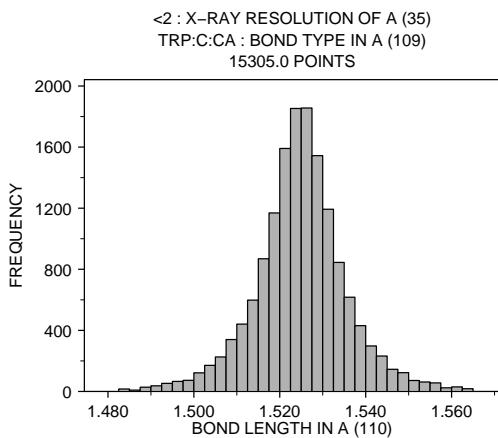
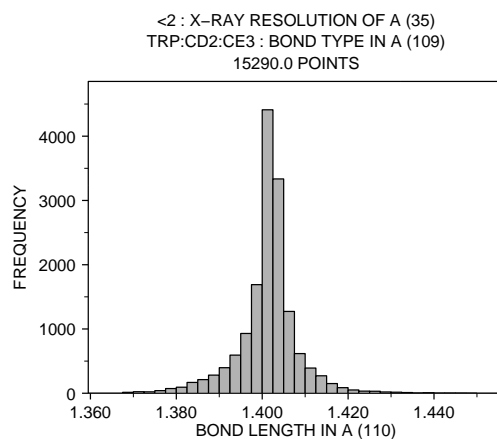
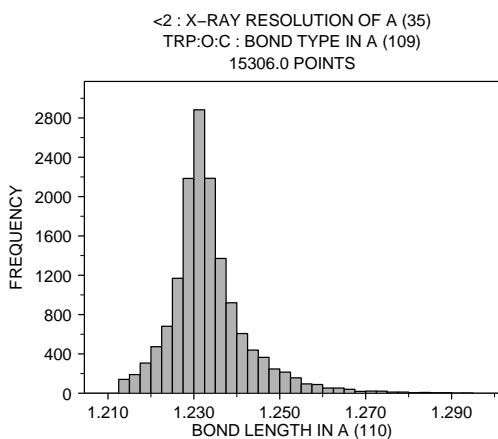
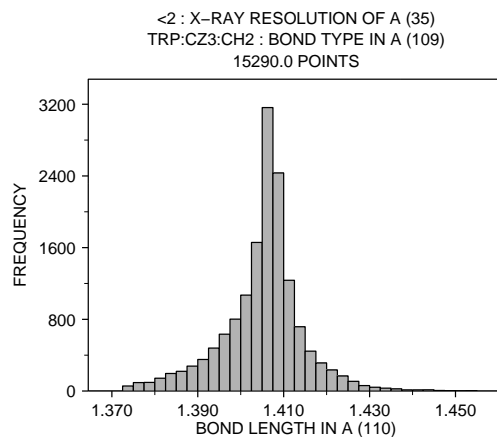
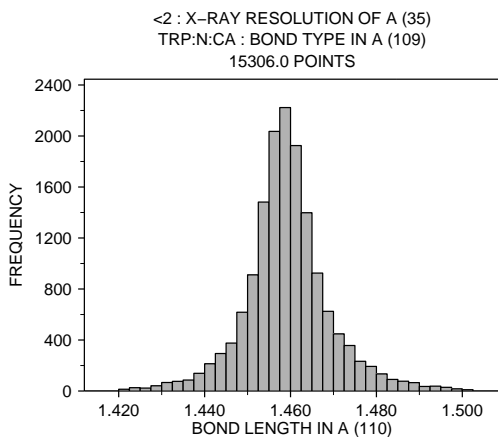
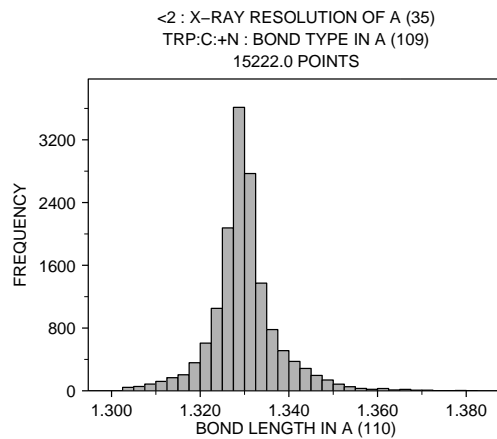
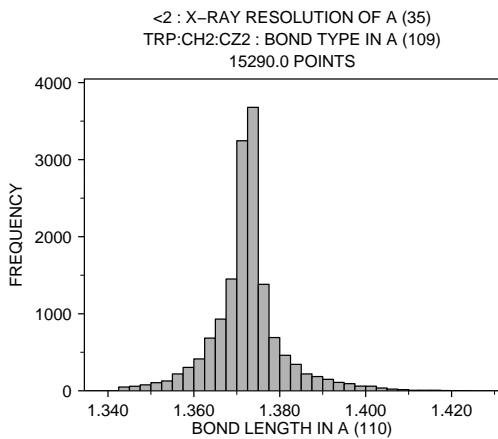


<2 : X-RAY RESOLUTION OF A (35)
THR:C:CA : BOND TYPE IN A (109)
58739.0 POINTS

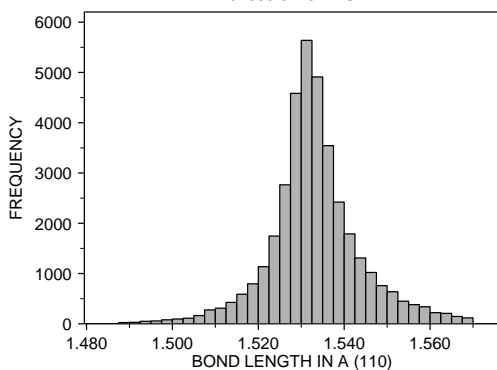




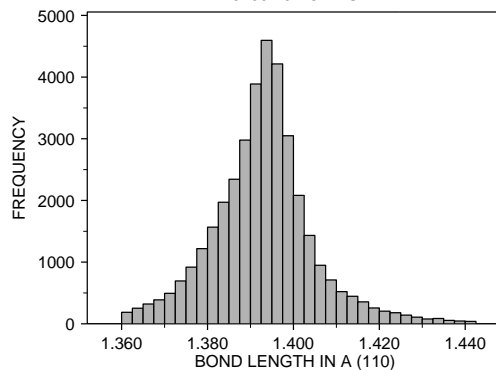




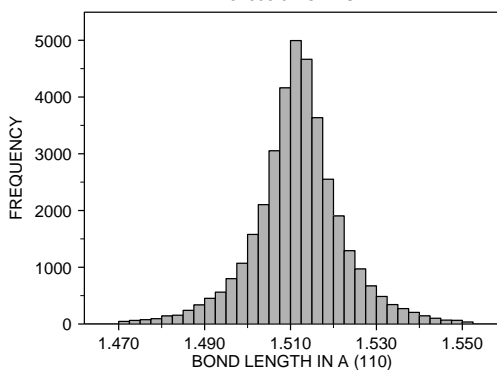
<2 : X-RAY RESOLUTION OF A (35)
TYR:CB:CA : BOND TYPE IN A (109)
37659.0 POINTS



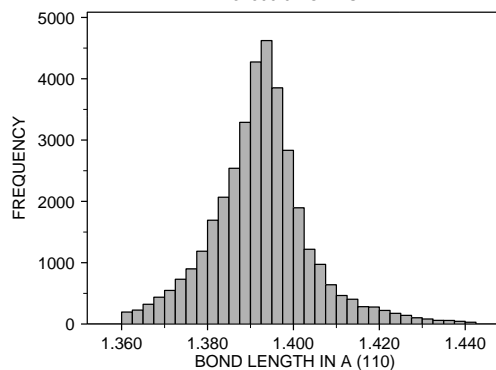
<2 : X-RAY RESOLUTION OF A (35)
TYR:CE1:CD1 : BOND TYPE IN A (109)
37601.0 POINTS



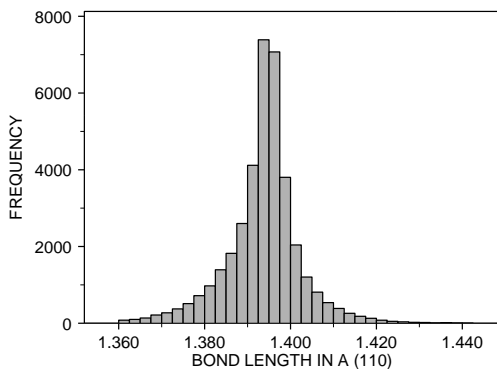
<2 : X-RAY RESOLUTION OF A (35)
TYR:CG:CB : BOND TYPE IN A (109)
37606.0 POINTS



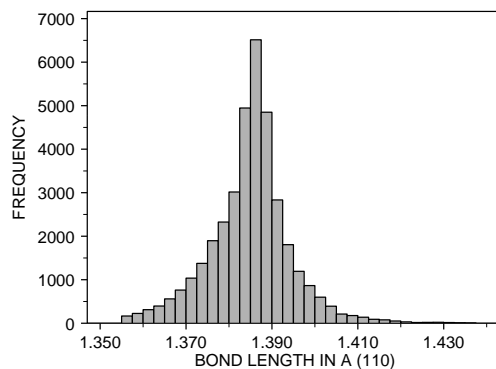
<2 : X-RAY RESOLUTION OF A (35)
TYR:CE2:CD2 : BOND TYPE IN A (109)
37600.0 POINTS



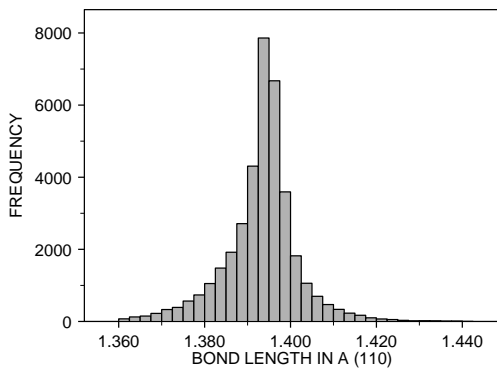
<2 : X-RAY RESOLUTION OF A (35)
TYR:CD1:CG : BOND TYPE IN A (109)
37601.0 POINTS



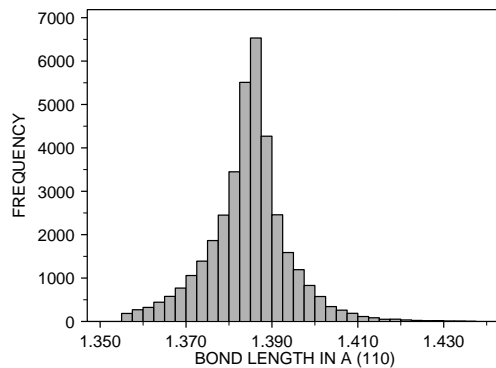
<2 : X-RAY RESOLUTION OF A (35)
TYR:CZ:CE1 : BOND TYPE IN A (109)
37601.0 POINTS



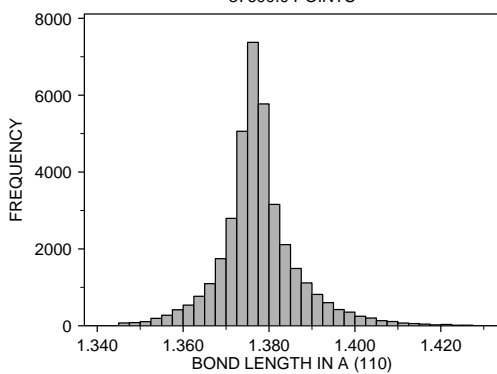
<2 : X-RAY RESOLUTION OF A (35)
TYR:CD2:CG : BOND TYPE IN A (109)
37602.0 POINTS



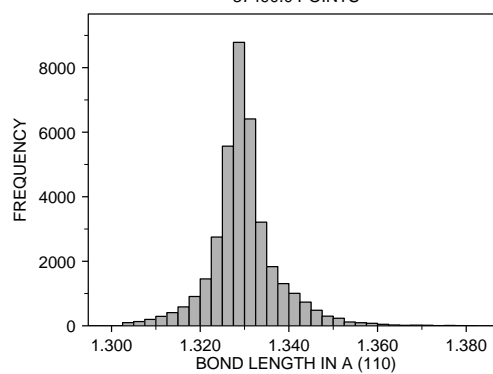
<2 : X-RAY RESOLUTION OF A (35)
TYR:CZ:CE2 : BOND TYPE IN A (109)
37600.0 POINTS



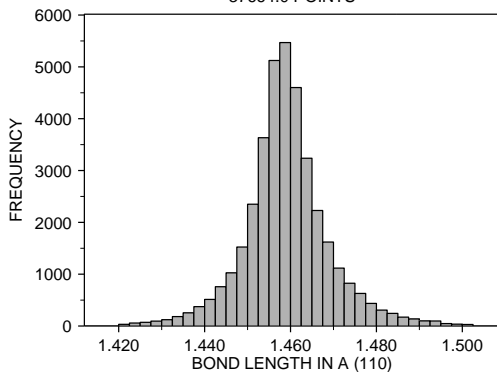
<2 : X-RAY RESOLUTION OF A (35)
TYR:OH:CZ : BOND TYPE IN A (109)
37600.0 POINTS



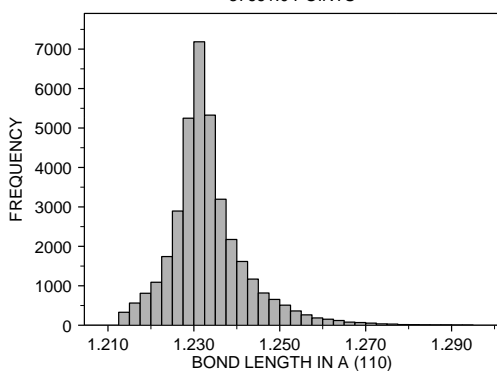
<2 : X-RAY RESOLUTION OF A (35)
TYR:C:+N : BOND TYPE IN A (109)
37490.0 POINTS



<2 : X-RAY RESOLUTION OF A (35)
TYR:N:CA : BOND TYPE IN A (109)
37664.0 POINTS



<2 : X-RAY RESOLUTION OF A (35)
TYR:O:C : BOND TYPE IN A (109)
37661.0 POINTS



<2 : X-RAY RESOLUTION OF A (35)
TYR:C:CA : BOND TYPE IN A (109)
37663.0 POINTS

