

Table 1. Possible values for IATYPE

	C	CH	CH2	CH3	N	NH	NH2	NH3	O	OH	S	SH	P	FE	CU	CA	MG	MN	ZN
IATYPE	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19

Table 2. Protein atoms

0																					
	N	CA	C	CB	CG	CD	CE	CZ	CH	N	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
ALA	6	2	1	9	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	9
ARG	6	2	1	9	3	3	3	0	1	0	0	6	0	7	0	0	0	0	0	0	9
ASN	6	2	1	9	3	1	0	0	0	0	7	0	0	0	0	9	0	0	0	0	9
ASP	6	2	1	9	3	1	0	0	0	0	0	0	0	0	0	9	0	0	0	0	9
CYS	6	2	1	9	3	0	0	0	0	0	0	0	0	0	0	0	0	0	11	0	9
GLN	6	2	1	9	3	3	1	0	0	0	7	0	0	0	0	9	0	0	0	0	9
GLU	6	2	1	9	3	3	1	0	0	0	0	0	0	0	0	9	0	0	0	0	9
GLY	6	3	1	9	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	9
HIS	6	2	1	9	3	1	2	2	0	0	5	6	0	0	0	0	0	0	0	0	9
ILE	6	2	1	9	2	-1	4	0	0	0	0	0	0	0	0	0	0	0	0	0	9
LEU	6	2	1	9	3	2	4	0	0	0	0	0	0	0	0	0	0	0	0	0	9
LYS	6	2	1	9	3	3	3	3	0	0	0	8	0	0	0	0	0	0	0	0	9
MET	6	2	1	9	3	3	0	4	0	0	0	0	0	0	0	0	0	0	0	12	9
PHE	6	2	1	9	3	1	2	2	2	0	0	0	0	0	0	0	0	0	0	0	9
PRO	5	2	1	9	3	3	3	0	0	0	0	0	0	0	0	0	0	0	0	0	9
SER	6	2	1	9	3	0	0	0	0	0	0	0	0	0	10	0	0	0	0	0	9
THR	6	2	1	9	2	4	0	0	0	0	0	0	0	0	10	0	0	0	0	0	9
TRP	6	2	1	9	3	1	1	2	2	2	0	6	0	0	0	0	0	0	0	0	9
TYR	6	2	1	9	3	1	2	2	1	0	0	0	0	0	0	0	0	10	0	0	9
VAL	6	2	1	9	2	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	9
	N	CA	C	CB	CG	CD	CE	CZ	CH	N	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
0																					

f\_invacuo(q=0) = sqrt(I\_invacuo(q=0)/4)  
I\_invacuo: 3rd column in the crysol \*.int file  
pdb file: contains two atoms (having same ATYPE/RTYPE, but different coordinates: 0,0,0 and 50,0,0)  
Values below: f\_invacuo(q=0) ~ total valence electrons within an atomic group  
(actual Atomic Group, actual IATYPE)

ALA	N	CA*	C	CB*	O	CG*	CD	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	8.9991	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH3,4)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)
ARG	N	CA*	C	CB*	O	CG	CD	CE*	CZ	CH	ND	NE	NZ	NH*	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	7.9991	7.9994	7.9991	7.9991	5.9992	5.9992	5.9992	6.9946	7.9945	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH2,3)	(O,9)	(CH2,3)	(CH2,3)	(C,1)	(C,1)	(C,1)	(N,5)	(NH,6)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)
ASN	N	CA*	C	CB*	O	CG*	CD	CE	CZ	CH	ND*	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	7.9991	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH2,3)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)

ASP	N	CA*	C	CB*	0	CG*	CD	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
CYS	N	CA*	C	CB*	0	CG*	CD	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG*	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	16.9998 (SH,12)	15.9998 (S,11)	7.9994 (0,9)
GLN	N	CA*	C	CB*	0	CG	CD*	CE	CZ	CH	ND	NE*	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	7.9991 (CH2,3)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
GLU	N	CA*	C	CB*	0	CG	CD*	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	7.9991 (CH2,3)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
GLY	N	CA*	C	CB*	0	CG	CD	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	5.9992 (C,1)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
HIS	N	CA*	C	CB*	0	CG*	CD	CE*	CZ*	CH	ND	NE*	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
ILE	N	CA*	C	CB*	0	CG*	CD	CE*	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	6.9992 (CH,2)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
LEU	N	CA*	C	CB*	0	CG*	CD*	CE*	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	6.9992 (CH,2)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
LYS	N	CA*	C	CB*	0	CG	CD	CE	CZ*	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	7.9991 (CH2,3)	7.9991 (CH2,3)	7.9991 (CH2,3)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	9.9945 (NH3,8)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
MET	N	CA*	C	CB*	0	CG	CD*	CE*	CZ*	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD*	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	7.9991 (CH2,3)	5.9992 (C,1)	8.9991 (CH3,4)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
PHE	N	CA*	C	CB*	0	CG*	CD	CE*	CZ	CH*	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9992 (CH,2)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
PRO	N	CA*	C	CB*	0	CG	CD	CE*	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	6.9946 (N,5)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	7.9991 (CH2,3)	7.9991 (CH2,3)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)
SER	N	CA*	C	CB*	0	CG*	CD	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945 (NH,6)	12.7783 (?,?)	5.9992 (C,1)	7.9991 (CH2,3)	7.9994 (0,9)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	5.9992 (C,1)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	6.9946 (N,5)	8.9993 (OH,10)	7.9994 (0,9)	7.9994 (0,9)	7.9994 (0,9)	15.9998 (S,11)	15.9998 (S,11)	7.9994 (0,9)

THR	N	CA*	C	CB*	O	CG*	CD*	CE	CZ	CH	ND	NE	NZ	NH	OG*	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	6.9992	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH,2)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)
TRP	N	CA*	C	CB*	O	CG*	CD	CE	CZ*	CH*	ND	NE*	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	7.9991	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH2,3)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)
TYR	N	CA*	C	CB*	O	CG*	CD	CE*	CZ*	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	7.9991	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	8.9993	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH2,3)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)
VAL	N	CA*	C	CB*	O	CG*	CD*	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	6.9992	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH,2)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)

\*IATYPE different from Table 2  
 According to crysol.txt, if IATYPE = 0, actual IATYPE is assigned from the columns 13-14.

===== myTest with two-atom models: RTYPE – DAM (10Dec2013)

DAM	N	CA	C	CB	O	CG	CD	CE	CZ	CH	ND	NE	NZ	NH	OG	OD	OE	OH	SG	SD	OX
	7.9945	12.7783	5.9992	7.9991	7.9994	5.9992	5.9992	5.9992	5.9992	5.9992	6.9946	6.9946	6.9946	6.9946	7.9994	7.9994	7.9994	7.9994	15.9998	15.9998	7.9994
	(NH,6)	(?,?)	(C,1)	(CH2,3)	(O,9)	(C,1)	(C,1)	(C,1)	(C,1)	(C,1)	(N,5)	(N,5)	(N,5)	(N,5)	(O,9)	(O,9)	(O,9)	(O,9)	(S,11)	(S,11)	(O,9)