

Appendix

This appendix is essentially a visual representation of the information encoded in the coarse-grain .cgc files, showing how each of the amino acids as well as a POPC lipid is divided into coarse-grained beads. Excerpts from the standard CHARMM topology file are shown with the Martini bead assignments overlaid.

```

RESI ALA      0.00
GROUP
ATOM N      NH1 -0.47 ! |
ATOM HN     H   0.31 ! |
ATOM CA     CT1 0.07 ! |
ATOM HA     HB  0.09 ! |
GROUP
ATOM CB     CT3 -0.27 ! |
ATOM HB1    HA  0.09 ! |
ATOM HB2    HA  0.09 ! |
ATOM HB3    HA  0.09 ! |
GROUP
ATOM C       C   0.51 ! |
ATOM O       O  -0.51 ! |
BOND CB CA N HN N CA
BOND C CA C +N CA HA CB HB1 CB HB2 CB HB3
DOUBLE O C

```

BAS = P4

```

RESI ASP     -1.00
GROUP
ATOM N      NH1 -0.47 ! |
ATOM HN     H   0.31 ! |
ATOM CA     CT1 0.07 ! |
ATOM HA     HB  0.09 ! |
GROUP
ATOM CB     CT2 -0.28 ! |
ATOM HB1    HA  0.09 ! |
ATOM HB2    HA  0.09 ! |
ATOM CG     CC  0.62 ! |
ATOM OD1    OC -0.76 ! |
ATOM OD2    OC -0.76 ! |
GROUP
ATOM C       C   0.51 ! |
ATOM O       O  -0.51 ! |
BOND CB CA CG CB OD2 CG
BOND N HN N CA C CA C +N
BOND CA HA CB HB1 CB HB2
DOUBLE O C CG OD1

```

BAS = P5

```

RESI ARG      1.00
GROUP
ATOM N      NH1 -0.47 ! |
ATOM HN     H   0.31 ! |
ATOM CA     CT1 0.07 ! |
ATOM HA     HB  0.09 ! |
GROUP
ATOM CB     CT2 -0.18 ! |
ATOM HB1    HA  0.09 ! |
ATOM HB2    HA  0.09 ! |
GROUP
ATOM CG     CT2 -0.18 ! |
ATOM HG1    HA  0.09 ! |
ATOM HG2    HA  0.09 ! |
GROUP
ATOM CD     CT2  0.20 ! |
ATOM HD1    HA  0.09 ! |
ATOM HD2    HA  0.09 ! |
ATOM NE     NC2 -0.70 ! |
ATOM HE     C   0.44 ! |
ATOM CZ     C   0.64 ! |
ATOM NH1    NC2 -0.80 ! |
ATOM HH11   HC  0.46 ! |
ATOM HH12   HC  0.46 ! |
ATOM NH2    NC2 -0.80 ! |
ATOM HH21   HC  0.46 ! |
ATOM HH22   HC  0.46 ! |
GROUP
ATOM C       C   0.51 ! |
ATOM O       O  -0.51 ! |
BOND CB CA CG CB CD CG NE CD CZ NE
BOND NH2 CZ N HN N CA
BOND C CA C +N CA HA CB HB1
BOND CB HB2 CG HG1 CG HG2 CD HD1 CD HD2
BOND NE HE NH1 HH11 NH1 HH12 NH2 HH21 NH2 HH22
DOUBLE O C CZ NH1

```

BAS = P5

SI1 = N0r

SI2 = Qdr

```

RESI CYS      0.00
GROUP
ATOM N      NH1 -0.47 ! |
ATOM HN     H   0.31 ! |
ATOM CA     CT1 0.07 ! |
ATOM HA     HB  0.09 ! |
GROUP
ATOM CB     CT2 -0.11 ! |
ATOM HB1    HA  0.09 ! |
ATOM HB2    HA  0.09 ! |
ATOM SG     S  -0.23 ! |
ATOM HG1    HS  0.16 ! |
GROUP
ATOM C       C   0.51 ! |
ATOM O       O  -0.51 ! |
BOND CB CA SG CB N HN N CA
BOND C CA C +N CA HA CB HB1
BOND CB HB2 SG HG1
DOUBLE O C

```

BAS = P5

```

RESI ASN      0.00
GROUP
ATOM N      NH1 -0.47 ! |
ATOM HN     H   0.31 ! |
ATOM CA     CT1 0.07 ! |
ATOM HA     HB  0.09 ! |
GROUP
ATOM CB     CT2 -0.18 ! |
ATOM HB1    HA  0.09 ! |
ATOM HB2    HA  0.09 ! |
GROUP
ATOM CG     CC  0.55 ! |
ATOM OD1    O  -0.55 ! |
GROUP
ATOM ND2    NH2 -0.62 ! |
ATOM HD21   H   0.32 ! |
ATOM HD22   H   0.30 ! |
GROUP
ATOM C       C   0.51 ! |
ATOM O       O  -0.51 ! |
BOND CB CA CG CB ND2 CG
BOND N HN N CA C CA C +N
BOND CA HA CB HB1 CB HB2 ND2 HD21 ND2 HD22
DOUBLE O C CG OD1

```

BAS = P5

SID = P5n

```

RESI GLN      0.00
GROUP
ATOM N      NH1 -0.47 ! |
ATOM HN     H   0.31 ! |
ATOM CA     CT1 0.07 ! |
ATOM HA     HB  0.09 ! |
GROUP
ATOM CB     CT2 -0.18 ! |
ATOM HB1    HA  0.09 ! |
ATOM HB2    HA  0.09 ! |
GROUP
ATOM CG     CT2 -0.18 ! |
ATOM HG1    HA  0.09 ! |
ATOM HG2    HA  0.09 ! |
GROUP
ATOM CD     CC  0.55 ! |
ATOM OE1    O  -0.55 ! |
GROUP
ATOM NE2    NH2 -0.62 ! |
ATOM HE21   H   0.32 ! |
ATOM HE22   H   0.30 ! |
GROUP
ATOM C       C   0.51 ! |
ATOM O       O  -0.51 ! |
BOND CB CA CG CB CD CG NE2 CD
BOND N HN N CA C CA
BOND C +N CA HA CB HB1 CB HB2 CG HG1
BOND CG HG2 NE2 HE21 NE2 HE22
DOUBLE O C CD OE1

```

BAS = P5

SID = P4q

```

RESI GLU      -1.00
GROUP
ATOM N        NH1    -0.47
ATOM HN       H       0.31
ATOM CA       CT1     0.07
ATOM HA       HB       0.09
GROUP
ATOM CB       CT2    -0.18
ATOM HB1      HA       0.09
ATOM HB2      HA       0.09
GROUP
ATOM CG       CT2    -0.28
ATOM HG1      HA       0.09
ATOM HG2      HA       0.09
ATOM CD       CC       0.62
ATOM OE1      OC      -0.76
ATOM OE2      OC      -0.76
GROUP
ATOM C        C        0.51
ATOM O        O       -0.51
BOND CB CA CG CB CD CG OE2 CD
BOND N HN N CA C CA
BOND C +N CA HA CB HB1 CB HB2 CG HG1
BOND CG HG2
DOUBLE O C CD OE1

```

BAS = P5

SID = Qae

```

RESI GLY      0.00
GROUP
ATOM N        NH1    -0.47
ATOM HN       H       0.31
ATOM CA       CT2    -0.02
ATOM HA1      HB       0.09
ATOM HA2      HB       0.09
GROUP
ATOM C        C        0.51
ATOM O        O       -0.51
BOND N HN N CA C CA
BOND C +N CA HA1 CA HA2
DOUBLE O C

```

BAS = P5

```

RESI HSD      0.00 ! neutral HIS, proton on ND1
GROUP
ATOM N        NH1    -0.47
ATOM HN       H       0.31
ATOM CA       CT1     0.07
ATOM HA       HB       0.09
GROUP
ATOM CB       CT2    -0.09
ATOM HB1      HA       0.09
ATOM HB2      HA       0.09
ATOM ND1      NR1    -0.36
ATOM HD1      H       0.32
ATOM CG       CPH1   -0.05
GROUP
ATOM CE1      CPH2    0.25
ATOM HE1      HR1     0.13
ATOM NE2      NR2    -0.70
ATOM CD2      CPH1    0.22
ATOM HD2      HR3     0.10
GROUP
ATOM C        C        0.51
ATOM O        O       -0.51
BOND CB CA CG CB ND1 CG CE1 ND1
BOND NE2 CD2 N HN N CA
BOND C +N CA HA CB HB1
BOND CB HB2 ND1 HD1 CD2 HD2 CE1 HE1
DOUBLE O C CG CD2 CE1 NE2

```

SID = SP1h

SID = SC4h

BAS = P5

SID = SP1h

```

RESI HSE      0.00 ! neutral His, proton on NE2
GROUP
ATOM N        NH1    -0.47
ATOM HN       H       0.31
ATOM CA       CT1     0.07
ATOM HA       HB       0.09
GROUP
ATOM CB       CT2    -0.08
ATOM HB1      HA       0.09
ATOM HB2      HA       0.09
ATOM ND1      NR2    -0.70
ATOM CG       CPH1    0.22
ATOM CE1      CPH2    0.25
ATOM HE1      HR1     0.13
GROUP
ATOM NE2      NR1    -0.36
ATOM HE2      H       0.32
ATOM CD2      CPH1   -0.05
ATOM HD2      HR3     0.09
GROUP
ATOM C        C        0.51
ATOM O        O       -0.51
BOND CB CA CG CB ND1 CG
BOND NE2 CD2 N HN N CA
BOND C +N NE2 CE1 CA HA CB HB1
BOND CB HB2 NE2 HE2 CD2 HD2 CE1 HE1
DOUBLE O C CD2 CG CE1 ND1

```

SID = SP1h

SID = SC4h

BAS = P5

SID = SP1h

```

RESI HSP      1.00 ! Protonated His
GROUP
ATOM N        NH1    -0.47
ATOM HN       H       0.31
ATOM CA       CT1     0.07
ATOM HA       HB       0.09
GROUP
ATOM CB       CT2    -0.05
ATOM HB1      HA       0.09
ATOM HB2      HA       0.09
ATOM CD2      CPH1    0.19
ATOM HD2      HR1     0.13
ATOM CG       CPH1    0.19
GROUP
ATOM NE2      NR3    -0.51
ATOM HE2      H       0.44
ATOM ND1      NR3    -0.51
ATOM HD1      H       0.44
ATOM CE1      CPH2    0.32
ATOM HE1      HR2     0.18
GROUP
ATOM C        C        0.51
ATOM O        O       -0.51
BOND CB CA CG CB ND1 CG CE1 ND1
BOND NE2 CD2 N HN N CA
BOND C +N CA HA CB HB1
BOND CB HB2 ND1 HD1 NE2 HE2 CD2 HD2 CE1 HE1
DOUBLE O C CD2 CG NE2 CE1

```

SID = SP1h

SID = SC4h

BAS = P5

SID = SQdh

```

RESI ILE      0.00
GROUP
ATOM N        NH1    -0.47
ATOM HN       H       0.31
ATOM CA       CT1     0.07
ATOM HA       HB       0.09
GROUP
ATOM CB       CT1    -0.09
ATOM HB       HA       0.09
GROUP
ATOM CG2      CT3    -0.27
ATOM HG21     HA       0.09
ATOM HG22     HA       0.09
ATOM HG23     HA       0.09
GROUP
ATOM CG1      CT2    -0.18
ATOM HG11     HA       0.09
ATOM HG12     HA       0.09
GROUP
ATOM CD       CT3    -0.27
ATOM HD1      HA       0.09
ATOM HD2      HA       0.09
ATOM HD3      HA       0.09
GROUP
ATOM C        C        0.51
ATOM O        O       -0.51
BOND CB CA CG1 CB CG2 CB CD CG1
BOND N HN N CA C CA C +N
BOND CA HA CB HB CG1 HG11 CG1 HG12 CG2 HG21
BOND CG2 HG22 CG2 HG23 CD HD1 CD HD2 CD HD3
DOUBLE O C

```

BAS = P5

SID = AC1i

```

RESI LEU      0.00
GROUP
ATOM N      NH1 -0.47 !
ATOM HN     H    0.31 !
ATOM CA     CT1 0.07 !
ATOM HA     HB   0.09 !
GROUP
ATOM CB     CT2 -0.18 !
ATOM HB1    HA   0.09 !
ATOM HB2    HA   0.09 !
GROUP
ATOM CG     CT1 -0.09 !
ATOM HG     HA   0.09 !
GROUP
ATOM CD1    CT3 -0.27 !
ATOM HD11   HA   0.09 !
ATOM HD12   HA   0.09 !
ATOM HD13   HA   0.09 !
GROUP
ATOM CD2    CT3 -0.27 !
ATOM HD21   HA   0.09 !
ATOM HD22   HA   0.09 !
ATOM HD23   HA   0.09 !
GROUP
ATOM C      C    0.51 !
ATOM O      O    -0.51 !
BOND CB CA CG CB CD1 CG CD2 CG
BOND N HN N CA C CA C +N
BOND CA HA CB HB1 CB HB2 CG HG CD1 HD11
BOND CD1 HD12 CD1 HD13 CD2 HD21 CD2 HD22 CD2 HD23
DOUBLE O C

```

Diagram illustrating the structure of LEU (Leucine) with highlighted regions: BAS = P5 (blue dashed box) and SID = AC11 (red dashed box).

```

RESI LYS      1.00
GROUP
ATOM N      NH1 -0.47 !
ATOM HN     H    0.31 !
ATOM CA     CT1 0.07 !
ATOM HA     HB   0.09 !
GROUP
ATOM CB     CT2 -0.18 !
ATOM HB1    HA   0.09 !
ATOM HB2    HA   0.09 !
GROUP
ATOM CG     CT2 -0.18 !
ATOM HG1    HA   0.09 !
ATOM HG2    HA   0.09 !
GROUP
ATOM CD     CT2 -0.18 !
ATOM HD1    HA   0.09 !
ATOM HD2    HA   0.09 !
GROUP
ATOM CE     CT2 0.21 !
ATOM HE1    HA   0.05 !
ATOM HE2    HA   0.05 !
ATOM NZ     NH3 -0.30 !
ATOM HZ1    HC   0.33 !
ATOM HZ2    HC   0.33 !
ATOM HZ3    HC   0.33 !
GROUP
ATOM C      C    0.51 !
ATOM O      O    -0.51 !
BOND CB CA CG CB CD CG CE CD NZ CE
BOND N HN N CA C CA
BOND C +N CA HA CB HB1 CB HB2 CG HG1
BOND CG HG2 CD HD1 CD HD2 CE HE1 CE HE2
DOUBLE O C
BOND NZ HZ1 NZ HZ2 NZ HZ3

```

Diagram illustrating the structure of LYS (Lysine) with highlighted regions: BAS = P5 (blue dashed box), SI1 = C3k (red dashed box), and SI2 = Qdk (red dashed box).

```

RESI MET      0.00
GROUP
ATOM N      NH1 -0.47 !
ATOM HN     H    0.31 !
ATOM CA     CT1 0.07 !
ATOM HA     HB   0.09 !
GROUP
ATOM CB     CT2 -0.18 !
ATOM HB1    HA   0.09 !
ATOM HB2    HA   0.09 !
GROUP
ATOM CG     CT2 -0.14 !
ATOM HG1    HA   0.09 !
ATOM HG2    HA   0.09 !
ATOM SD     S    -0.09 !
ATOM CE     CT3 -0.22 !
ATOM HE1    HA   0.09 !
ATOM HE2    HA   0.09 !
ATOM HE3    HA   0.09 !
GROUP
ATOM C      C    0.51 !
ATOM O      O    -0.51 !
BOND CB CA CG CB SD CG CE SD
BOND N HN N CA C CA C +N
BOND CA HA CB HB1 CB HB2 CG HG1 CG HG2
BOND CE HE1 CE HE2 CE HE3
DOUBLE O C

```

Diagram illustrating the structure of MET (Methionine) with highlighted regions: BAS = P5 (blue dashed box) and SID = C5m (red dashed box).

```

RESI PRO      0.00
GROUP
ATOM N      N    -0.29 !
ATOM CD     CP3 0.00 !
ATOM HD1    HA   0.09 !
ATOM HD2    HA   0.09 !
ATOM CA     CP1 0.02 !
ATOM HA     HB   0.09 !
GROUP
ATOM CB     CP2 -0.18 !
ATOM HB1    HA   0.09 !
ATOM HB2    HA   0.09 !
GROUP
ATOM CG     CP2 -0.18 !
ATOM HG1    HA   0.09 !
ATOM HG2    HA   0.09 !
GROUP
ATOM C      C    0.51 !
ATOM O      O    -0.51 !
BOND C CA C +N
BOND N CA CA CB CB CG CG CD N CD
BOND HA CA HG1 CG HG2 CG HD1 CD HD2 CD HB1 CB HB2 CB
DOUBLE O C

```

Diagram illustrating the structure of PRO (Proline) with highlighted regions: SID = AC2p (red dashed box) and BAS = P5 (blue dashed box).

```

RESI PHE      0.00
GROUP
ATOM N      NH1 -0.47 !
ATOM HN     H    0.31 !
ATOM CA     CT1 0.07 !
ATOM HA     HB   0.09 !
GROUP
ATOM CB     CT2 -0.18 !
ATOM HB1    HA   0.09 !
ATOM HB2    HA   0.09 !
GROUP
ATOM CG     CA   0.00 !
GROUP
ATOM CD1    CA  -0.115 !
ATOM HD1    HP   0.115 !
GROUP
ATOM CE1    CA  -0.115 !
ATOM HE1    HP   0.115 !
GROUP
ATOM CZ     CA  -0.115 !
ATOM HZ     HP   0.115 !
GROUP
ATOM CD2    CA  -0.115 !
ATOM HD2    HP   0.115 !
GROUP
ATOM CE2    CA  -0.115 !
ATOM HE2    HP   0.115 !
GROUP
ATOM C      C    0.51 !
ATOM O      O    -0.51 !
BOND CB CA CG CB CD2 CG CE1 CD1
BOND CZ CE2 N HN
BOND N CA C CA C +N CA HA
BOND CB HB1 CB HB2 CD1 HD1 CD2 HD2 CE1 HE1
DOUBLE O C CD1 CG CZ CE1 CE2 CD2
BOND CE2 HE2 CZ HZ

```

Diagram illustrating the structure of PHE (Phenylalanine) with highlighted regions: SI1 = SC4f (red dashed box), SI2 = SC4f (red dashed box), SI3 = SC4f (red dashed box), and BAS = P5 (blue dashed box).

```

RESI SER          0.00
GROUP
ATOM N      NH1   -0.47
ATOM HN     H      0.31
ATOM CA     CT1   0.07
ATOM HA     HB      0.09
GROUP
ATOM CB     CT2   0.05
ATOM HB1    HA     0.09
ATOM HB2    HA     0.09
ATOM OG     OH1   -0.66
ATOM HG1    H      0.43
GROUP
ATOM C      C      0.51
ATOM O      O     -0.51
BOND CB CA  OG CB N HN N CA
BOND C CA  C +N CA HA CB HB1
BOND CB HB2 OG HG1
DOUBLE O C

```

BAS = P5

SID = P1s

```

RESI THR          0.00
GROUP
ATOM N      NH1   -0.47
ATOM HN     H      0.31
ATOM CA     CT1   0.07
ATOM HA     HB      0.09
GROUP
ATOM CB     CT1   0.14
ATOM HB     HA     0.09
ATOM OG1    OH1   -0.66
ATOM HG1    H      0.43
GROUP
ATOM CG2    CT3   -0.27
ATOM HG21   HA     0.09
ATOM HG22   HA     0.09
ATOM HG23   HA     0.09
GROUP
ATOM C      C      0.51
ATOM O      O     -0.51
BOND CB CA  OG1 CB CG2 CB N HN
BOND N CA  C CA C +N CA HA
BOND CB HB OG1 HG1 CG2 HG21 CG2 HG22 CG2 HG23
DOUBLE O C

```

BAS = P5

SID = P1t

```

RESI TRP          0.00
GROUP
ATOM N      NH1   -0.47
ATOM HN     H      0.31
ATOM CA     CT1   0.07
ATOM HA     HB      0.09
GROUP
ATOM CB     CT2   -0.18
ATOM HB1    HA     0.09
ATOM HB2    HA     0.09
GROUP
ATOM CG     CY     -0.03
ATOM CD1    CA     0.035
ATOM HD1    HP     0.115
ATOM NE1    NY     -0.61
ATOM HE1    H      0.38
ATOM CE2    CPT    0.13
ATOM CD2    CPT    -0.02
GROUP
ATOM CE3    CA     -0.115
ATOM HE3    HP     0.115
GROUP
ATOM CZ3    CA     -0.115
ATOM HZ3    HP     0.115
GROUP
ATOM CZ2    CA     -0.115
ATOM HZ2    HP     0.115
GROUP
ATOM CH2    CA     -0.115
ATOM HH2    HP     0.115
GROUP
ATOM C      C      0.51
ATOM O      O     -0.51
BOND CB CA  CG CB CD2 CG NE1 CD1
BOND CZ2 CE2
BOND N HN  N CA C CA C +N
BOND CZ3 CH2 CD2 CE3 NE1 CE2 CA HA CB HB1
BOND CB HB2 CD1 HD1 NE1 HE1 CE3 HE3 CZ2 HZ2
BOND CZ3 HZ3 CH2 HH2
DOUBLE O C CD1 CG CE2 CD2 CZ3 CE3 CH2 CZ2

```

SI1 = SC4w

BAS = P5

SI2 = SP1w

SI3 = SC4w

SI4 = SC4w

```

RESI TYR          0.00
GROUP
ATOM N      NH1   -0.47
ATOM HN     H      0.31
ATOM CA     CT1   0.07
ATOM HA     HB      0.09
GROUP
ATOM CB     CT2   -0.18
ATOM HB1    HA     0.09
ATOM HB2    HA     0.09
GROUP
ATOM CG     CA     0.00
GROUP
ATOM CD1    CA     -0.115
ATOM HD1    HP     0.115
GROUP
ATOM CE1    CA     -0.115
ATOM HE1    HP     0.115
GROUP
ATOM CZ     CA     0.11
ATOM OH     OH1   -0.54
ATOM HH     H      0.43
GROUP
ATOM CD2    CA     -0.115
ATOM HD2    HP     0.115
GROUP
ATOM CE2    CA     -0.115
ATOM HE2    HP     0.115
GROUP
ATOM C      C      0.51
ATOM O      O     -0.51
BOND CB CA  CG CB CD2 CG CE1 CD1
BOND CZ CE2 OH CZ
BOND N HN  N CA C CA C +N
BOND CA HA CB HB1 CB HB2 CD1 HD1 CD2 HD2
BOND CE1 HE1 CE2 HE2 OH HH
DOUBLE O C CD1 CG CE1 CZ CE2 CD2

```

SI1 = SC4y

BAS = P5

SI3 = SP1y

SI2 = SC4y

```

RESI VAL          0.00
GROUP
ATOM N      NH1   -0.47
ATOM HN     H      0.31
ATOM CA     CT1   0.07
ATOM HA     HB      0.09
GROUP
ATOM CB     CT1   -0.09
ATOM HB     HA     0.09
GROUP
ATOM CG1    CT3   -0.27
ATOM HG11   HA     0.09
ATOM HG12   HA     0.09
ATOM HG13   HA     0.09
GROUP
ATOM CG2    CT3   -0.27
ATOM HG21   HA     0.09
ATOM HG22   HA     0.09
ATOM HG23   HA     0.09
GROUP
ATOM C      C      0.51
ATOM O      O     -0.51
BOND CB CA  CG1 CB CG2 CB N HN
BOND N CA  C CA C +N CA HA
BOND CB HB CG1 HG11 CG1 HG13 CG2 HG21
BOND CG2 HG22 CG2 HG23
DOUBLE O C

```

BAS = P5

SID = AC3v

```

graph TD
    NH3 --> PO4
    PO4 --> GL1
    GL1 --> C1B
    GL1 --> GL2
    C1B --> C2B
    C2B --> D3B
    D3B --> C4B
    C4B --> C5B
    GL2 --> C1A
    C1A --> C2A
    C2A --> C3A
    C3A --> C4A

```

GROUP			
ATOM C35	CTL2	-0.18	!
ATOM H5X	HAL2	0.09	!
ATOM H5Y	HAL2	0.09	!
GROUP			
ATOM C36	CTL2	-0.18	!
ATOM H6X	HAL2	0.09	!
ATOM H6Y	HAL2	0.09	!
GROUP			
ATOM C37	CTL2	-0.18	!
ATOM H7X	HAL2	0.09	!
ATOM H7Y	HAL2	0.09	!
GROUP			
ATOM C38	CTL2	-0.18	!
ATOM H8X	HAL2	0.09	!
ATOM H8Y	HAL2	0.09	!
GROUP			
ATOM C39	CTL2	-0.18	!
ATOM H9X	HAL2	0.09	!
ATOM H9Y	HAL2	0.09	!
GROUP			
ATOM C310	CTL2	-0.18	!
ATOM H10X	HAL2	0.09	!
ATOM H10Y	HAL2	0.09	!
GROUP			
ATOM C311	CTL2	-0.18	!
ATOM H11X	HAL2	0.09	!
ATOM H11Y	HAL2	0.09	!
GROUP			
ATOM C312	CTL2	-0.18	!
ATOM H12X	HAL2	0.09	!
ATOM H12Y	HAL2	0.09	!
GROUP			
ATOM C313	CTL2	-0.18	!
ATOM H13X	HAL2	0.09	!
ATOM H13Y	HAL2	0.09	!
GROUP			
ATOM C314	CTL2	-0.18	!
ATOM H14X	HAL2	0.09	!
ATOM H14Y	HAL2	0.09	!
GROUP			
ATOM C315	CTL2	-0.18	!
ATOM H15X	HAL2	0.09	!
ATOM H15Y	HAL2	0.09	!
GROUP			
ATOM C316	CTL3	-0.27	!
ATOM H16X	HAL3	0.09	!
ATOM H16Y	HAL3	0.09	!
ATOM H16Z	HAL3	0.09	!