

Boron-boron, carbon-carbon and nitrogen-nitrogen bonding in N-heterocyclic carbenes and their diazaboryl and triazole analogues; Wanzlick equilibria revisited

Katarzyna Młodzikowska,^{a†} Adam A. Rajkiewicz,^{a†} Karol Grela,^b and Bartosz Trzaskowski^{a*}

^a Centre of New Technologies, University of Warsaw, Banacha 2C, 02-097 Warszawa

^b Faculty of Chemistry, Biological and Chemical Research Centre, University of Warsaw, Żwirki i Wigury 101, 02-089 Warszawa

[†] These authors contributed equally to this work

Contents

Molecule partitioning scheme in functional SAPT0 calculations	2
Calculated structural and energetic parameters and partial charges	2
Table S1	2
Table S2	3
Table S3	4
Table S4	4
Table S5	5
Table S6	6
Table S7	6
Table S8	7
Geometries in xyz format	8

Molecule partitioning scheme in functional SAPT0 calculations

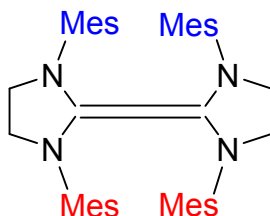


Figure S1. Functional SAPT0 partitioning scheme used throughout the study, taking dimer of **2** as an example. For each dimer the intramolecular interaction energy estimation consisted of two independent calculations. In the first one the interaction energy was calculated between the two side groups in blue, with the remainder of the molecule (in red and black) acting as the linker. Second calculations included interaction energy between the two side groups in red, with the remainder of the molecule (in blue and black) acting as the linker.

Calculated structural and energetic parameters and partial charges

Table S1. The comparison of calculated structural and energetic parameters of **1-6** and their dimers at the M06-2X level of theory.

parameter	1	2	2 dim	3	3 dim	4	4 dim	5	5 dim	6	6 dim
X1-X1' bond length [Å]	-	-	1.36	-	1.34	-	1.35	-	1.35	-	1.35
X1-X1' bond order (NBO)	2	-	2	-	2	-	2	-	2	-	2
more stable form ^a	dim	-	mon	-	dim	-	dim	-	mon	-	mon
ΔG DFT (kcal/mol) ^b	-	-	6.6	-	-1.3	-	-3.8	-	22.2	-	22.8
ΔG CCSD(T)(kcal/mol) ^b	-	-	8.0	-	-3.5	-	-4.2	-	21.9	-	24.2
HOMO-LUMO gap (eV)	2.91	7.96	5.93	9.53	7.83	9.31	7.79	9.31	6.65	9.35	6.62
Natural charges											
X1	-0.14	0.24	0.28	0.22	0.27	0.22	0.26	0.13	0.25	0.13	0.25
Imidazole	-0.34	-0.41	-0.35	-0.45	-0.40	-0.46	-0.42	-0.50	-0.43	-0.51	-0.44
R2	0.24	0.21	0.20	0.22	0.20	0.23	0.21	0.25	0.21	0.25	0.22
R3	0.24	0.21	0.16	0.22	0.20	0.23	0.21	0.25	0.21	0.25	0.22
X1'	-0.14	-	0.28	-	0.27	-	0.26	-	0.25	-	0.25
Imidazole'	-0.34	-	-0.36	-	-0.40	-	-0.42	-	-0.43	-	-0.44
R2'	0.24	-	0.20	-	0.20	-	0.21	-	0.21	-	0.22
R3'	0.24	-	0.16	-	0.20	-	0.21	-	0.21	-	0.22
Mulliken partial charges											
X1	-0.25	0.19	0.41	0.18	0.31	0.17	0.30	0.12	0.33	0.11	0.32

Imidazole	-0.15	-0.44	-0.35	-0.39	-0.32	-0.39	-0.33	-0.45	-0.35	-0.44	-0.35
R2	0.20	0.22	0.19	0.20	0.16	0.19	0.16	0.22	0.18	0.22	0.17
R3	0.20	0.22	0.16	0.20	0.16	0.19	0.16	0.22	0.18	0.22	0.18
X1'	-0.25	-	0.40	-	0.31	-	0.31	-	0.33	-	0.31
Imidazole'	-0.15	-	-0.35	-	-0.32	-	-0.32	-	-0.35	-	-0.35
R2'	0.20	-	0.19	-	0.16	-	0.16	-	0.18	-	0.18
R3'	0.20	-	0.16	-	0.16	-	0.16	-	0.18	-	0.17

^a dim - dimer, mon – monomer

Table S2. The comparison of calculated structural and energetic parameters of **1-6** and their dimers at the M06 level of theory.

parameter	1	2	2 dim	3	3 dim	4	4 dim	5	5 dim	6	6 dim
X1-X1' bond length [Å]	-	-	1.37	-	1.3	-	1.35	-	1.35	-	1.35
X1-X1' bond order (NBO)	2	-	2	-	2	-	2	-	2	-	2
HOMO-LUMO gap (eV)	2.91	5.92	4.11	7.18	5.86	7.00	5.85	6.93	4.74	6.98	4.72
Natural charges											
X1	-0.14	0.24	0.29	0.23	0.27	0.22	0.26	0.14	0.25	0.14	0.25
Imidazole	-0.34	-0.42	-0.36	-0.44	-0.40	-0.46	-0.42	-0.50	-0.42	-0.50	-0.44
R2	0.24	0.21	0.20	0.22	0.20	0.23	0.15	0.25	0.21	0.25	0.22
R3	0.24	0.21	0.16	0.22	0.20	0.23	0.15	0.25	0.21	0.25	0.22
X1'	-0.14	-	0.28	-	0.27	-	0.26	-	0.25	-	0.25
Imidazole'	-0.34	-	-0.36	-	-0.40	-	-0.42	-	-0.42	-	-0.44
R2'	0.24	-	0.20	-	0.20	-	0.15	-	0.21	-	0.22
R3'	0.24	-	0.16	-	0.20	-	0.15	-	0.21	-	0.22
Mulliken partial charges											
X1	-0.25	0.17	0.44	0.16	0.33	0.15	0.33	0.11	0.35	0.10	0.34
Imidazole	-0.15	-0.37	-0.37	-0.34	-0.30	-0.36	-0.30	-0.40	-0.32	-0.40	-0.33
R2	0.20	0.21	0.20	0.17	0.15	0.18	0.15	0.20	0.16	0.20	0.17
R3	0.20	0.21	0.17	0.17	0.15	0.18	0.15	0.20	0.16	0.20	0.16
X1'	-0.25	-	0.43	-	0.33	-	0.33	-	0.35	-	0.34
Imidazole'	-0.15	-	-0.37	-	-0.30	-	-0.30	-	-0.32	-	-0.33
R2'	0.20	-	0.20	-	0.15	-	0.15	-	0.16	-	0.17
R3'	0.20	-	0.17	-	0.15	-	0.15	-	0.16	-	0.16

Table S3. The comparison of calculated structural and energetic parameters of **7-9**, **16** and **17** and their dimers at the M06-2X level of theory.

parameter	7	7 dim	8	8 dim	9	9 dim	16	16 dim	17	17 dim
X1-X1' bond length [Å]	-	1.35	-	1.35	-	1.35	-	1.38	-	1.37
X1-X1' bond order (NBO)	-	2	-	2	-	2	-	2	-	2
more stable form ^a	-	dim	-	dim	-	dim	-	mon	-	mon
ΔG DFT (kcal/mol)	-	-5.4	-	-5.9	-	-0.6	-	54.8	-	33.6
ΔG	-	-4.4	-	-2.9	-	0.7	-	55.9	-	37.7
CCSD(T)(kcal/mol)										
HOMO-LUMO gap	7.89	6.28	7.27	5.33	7.71	5.80	5.01	3.81	6.20	4.76
				Natural charges						
X1	0.22	0.30	0.22	0.27	0.22	0.27	0.24	0.29	0.24	0.29
imidazole	-0.42	-0.34	-0.41	-0.36	-0.42	-0.37	-0.42	-0.36	-0.42	-0.36
R2	0.21	0.19	0.20	0.18	0.20	0.17	0.21	0.20	0.21	0.20
R3	0.20	0.18	0.21	0.18	0.22	0.19	0.21	0.16	0.21	0.16
X1'	-	0.26	-	0.27	-	0.27	-	0.28	-	0.28
imidazole'	-	-0.38	-	-0.36	-	-0.37	-	-0.36	-	-0.36
R2'	-	0.19	-	0.18	-	0.17	-	0.20	-	0.20
R3'	-	0.16	-	0.18	-	0.19	-	0.16	-	0.16
				Mulliken partial charges						
X1	0.19	0.35	0.19	0.37	0.18	0.37	0.17	0.44	0.17	0.44
imidazole	-0.39	-0.32	-0.38	-0.32	-0.40	-0.33	-0.42	-0.37	-0.42	-0.37
R2	0.22	0.16	0.22	0.16	0.21	0.16	0.21	0.20	0.21	0.20
R3	0.17	0.14	0.16	0.16	0.19	0.17	0.21	0.17	0.21	0.17
X1'	-	0.34	-	0.37	-	0.37	-	0.43	-	0.43
imidazole'	-	-0.32	-	-0.32	-	-0.33	-	-0.37	-	-0.37
R2'	-	0.16	-	0.18	-	0.16	-	0.20	-	0.20
R3'	-	0.14	-	0.16	-	0.17	-	0.17	-	0.17

^a dim - dimer, mon – monomer

Table S4. The comparison of calculated structural and energetic parameters of **7-9**, **16** and **17** and their dimers at the M06 level of theory.

parameter	7	7 dim	8	8 dim	9	9 dim	16	16 dim	17	17 dim
X1-X1' bond length [Å]	-	1.35	-	1.35	-	1.36	-	1.40	-	1.38
X1-X1' bond order (NBO)	-	2	-	2	-	2	-	2	-	2
HOMO-LUMO gap	5.81	4.42	5.15	3.43	5.66	4.03	3.33	2.37	4.54	3.16
				Natural charges						
X1	0.22	0.30	0.22	0.27	0.22	0.27	0.24	0.29	0.24	0.29
imidazole	-0.42	-0.37	-0.42	-0.36	-0.43	-0.36	-0.42	-0.36	-0.42	-0.36

R2	0.21	0.19	0.21	0.18	0.21	0.17	0.21	0.20	0.21	0.20
R3	0.21	0.18	0.21	0.18	0.22	0.19	0.21	0.16	0.21	0.16
X1'	-	0.26	-	0.27	-	0.27	-	0.28	-	0.28
imidazole'	-	-0.37	-	-0.36	-	-0.36	-	-0.36	-	-0.36
R2'	-	0.16	-	0.18	-	0.17	-	0.20	-	0.20
R3'	-	0.18	-	0.18	-	0.19	-	0.16	-	0.16
Mulliken partial charges										
X1	0.17	0.39	0.17	0.40	0.16	0.41	0.17	0.44	0.17	0.44
imidazole	-0.37	-0.30	-0.37	-0.33	-0.38	-0.34	-0.42	-0.37	-0.42	-0.37
R2	0.21	0.17	0.22	0.18	0.21	0.17	0.21	0.20	0.21	0.20
R3	0.16	0.13	0.15	0.15	0.17	0.17	0.21	0.17	0.21	0.17
X1'	-	0.36	-	0.40	-	0.40	-	0.43	-	0.43
imidazole'	-	-0.30	-	-0.34	-	-0.34	-	-0.37	-	-0.37
R2'	-	0.18	-	0.19	-	0.17	-	0.20	-	0.20
R3'	-	0.15	-	0.15	-	0.17	-	0.17	-	0.17

Table S5. The comparison of calculated structural and energetic parameters of systems **10** and **11** and their dimers at the M06-2X level of theory.

parameter	10	10A	10B	10C	11	11A	11B	11C
X1-X1' bond length [Å]	-	1.61	1.72	1.67	-	1.67	1.670	1.68
X1-X1' bond order (NBO)	-	1	1	1	-	1	1	1
more stable form ^a	-	mon	-	-	-	mon	-	-
ΔG DFT (kcal/mol) ^b	-	11.7	-	-	-	24.6	-	-
ΔG CCSD(T)(kcal/mol)	-	29.0	-	-	-	40.5	-	-
HOMO-LUMO gap	4.21	2.80	7.45	2.40	4.24	1.60	7.08	2.86
Natural charges								
X1	0.21	0.32	0.69	0.67	0.13	0.47	0.59	0.57
imidazole	-1.10	-0.77	-0.37	-0.50	-1.15	-0.62	-0.41	-0.52
R2	0.05	-0.11	0.19	-0.75	0.07	-0.21	0.20	-0.74
R3	0.05	-0.11	0.18	-0.81	0.07	-0.17	0.21	-0.74
X1'	-	0.32	0.69	0.67	-	0.47	0.59	0.57
imidazole'	-	-0.77	-0.37	-0.49	-	-0.62	-0.41	-0.52
R2'	-	-0.12	0.19	-0.74	-	-0.22	0.20	-0.75
R3'	-	-0.12	0.18	-0.70	-	-0.17	0.21	-0.74
Mulliken partial charges								
X1	-0.07	0.30	0.41	0.41	-0.11	0.32	0.36	0.34
imidazole	-0.92	-0.60	-0.32	-0.46	-0.96	-0.54	-0.36	-0.53
R2	-0.04	-0.20	0.16	-0.75	-0.02	-0.24	0.18	-0.73
R3	-0.05	-0.20	0.16	-0.81	-0.02	-0.21	0.18	-0.73
X1'	-	0.29	0.40	0.39	-	0.31	0.35	0.34
imidazole'	-	-0.60	-0.32	-0.50	-	-0.55	-0.37	-0.53

R2'	-	-0.20	0.16	-0.76	-	-0.25	0.19	-0.75
R3'	-	-0.20	0.16	-0.72	-	-0.21	0.18	-0.72

^a dim - dimer, mon – monomer ^b – in the 6-31G** basis set

Table S6. The comparison of calculated structural and energetic parameters of systems **10** and **11** and their dimers at the M06 level of theory.

parameter	10	10A	10B	10C	11	11A	11B	11C
X1-X1' bond length [Å]	-	1.63	1.71	1.670	-	1.67	1.69	1.68
X1-X1' bond order (NBO)	-	1	1	1	-	1	1	1
HOMO-LUMO gap	2.57	1.67	5.68	0.63	2.62	0.90	5.31	1.70
Natural charges								
X1	0.23	0.39	0.68	0.59	0.17	0.51	0.61	0.57
imidazole	-1.08	-0.72	-0.37	-0.60	-1.12	-0.61	-0.42	-0.55
R2	0.05	-0.13	0.19	-0.63	0.06	-0.23	0.20	-0.73
R3	0.04	-0.15	0.18	-0.77	0.06	-0.16	0.21	-0.71
X1'	-	0.40	0.68	0.58	-	0.51	0.61	0.57
imidazole'	-	-0.73	-0.37	-0.61	-	-0.61	-0.42	-0.57
R2'	-	-0.13	0.19	-0.64	-	-0.23	0.20	-0.73
R3'	-	-0.14	0.18	-0.75	-	-0.16	0.21	-0.70
Mulliken partial charges								
X1	-0.08	0.35	0.41	0.37	-0.11	0.33	0.36	0.34
imidazole	-0.88	-0.56	-0.32	-0.55	-0.77	-0.52	-0.35	-0.53
R2	-0.05	-0.21	0.16	-0.67	-0.18	-0.26	0.17	-0.74
R3	-0.07	-0.23	0.16	-0.78	-0.04	-0.21	0.17	-0.73
X1'	-	0.35	0.41	0.39	-	0.33	0.36	0.34
imidazole'	-	-0.57	-0.31	-0.56	-	-0.52	-0.35	-0.53
R2'	-	-0.21	0.15	-0.68	-	-0.27	0.18	-0.74
R3'	-	-0.22	0.16	-0.76	-	-0.21	0.17	-0.73

Table S7. The comparison of calculated structural and energetic parameters of systems **12** and **13** and their dimers at the M06-2X level of theory.

parameter	12	12A	12B	12C	13	13A	13B	13C
X1-X1' bond length [Å]	-	1.589	1.689	1.674	-	1.593	1.677	1.763
X1-X1' bond order (NBO)	-	2	1	1	-	2	1	1
more stable form ^a	-	mon	-	-	-	mon	-	-

ΔG DFT (kcal/mol) ^b	-	21.6	-	-	-	37.6	-	-
ΔG	-	36.0	-	-	-	55.6	-	-
CCSD(T)(kcal/mol)								
HOMO-LUMO gap	6.46	4.67	8.38	2.54	6.09	4.18	8.01	4.69
			Natural charges					
X1	0.14	0.12	0.67	0.07	0.06	0.15	0.57	0.60
imidazole	-1.23	-1.16	-0.40	-1.75	-1.27	-1.15	-0.45	-2.03
R2	0.11	0.08	0.20	-0.08	0.14	0.09	0.22	0.01
R3	0.11	0.08	0.20	-0.18	0.14	0.07	0.22	0.02
X1'	-	0.12	0.67	0.07	-	0.12	0.57	0.59
imidazole'	-	-1.16	-0.40	-1.75	-	-1.18	-0.45	-2.03
R2'	-	0.08	0.20	-0.08	-	0.08	0.22	0.01
R3'	-	0.08	0.20	-0.18	-	0.06	0.22	0.02
			Mulliken partial charges					
X1	-0.20	0.11	0.37	0.04	-0.25	0.09	0.32	0.23
imidazole	-1.02	-0.92	-0.29	-1.45	-1.07	-0.92	-0.35	-1.67
R2	0.01	-0.04	0.15	-0.22	0.04	-0.03	0.17	-0.16
R3	0.01	-0.04	0.15	-0.33	0.04	-0.04	0.17	-0.16
X1'	-	0.11	0.37	0.04	-	0.09	0.32	0.22
imidazole'	-	-0.92	-0.29	-1.45	-	-0.92	-0.35	-1.68
R2'	-	-0.04	0.15	-0.22	-	-0.03	0.17	-0.17
R3'	-	-0.04	0.15	-0.33	-	-0.06	0.17	-0.16

^a dim - dimer, mon – monomer ^b – in the 6-31G** basis set

Table S8. The comparison of calculated structural and energetic parameters of systems **12** and **13** and their dimers at the M06 level of theory.

parameter	12	12A	12B	12C	13	13A	13B	13C
X1-X1' bond length [Å]	-	1.59	1.69	1.67	-	1.59	1.68	1.76
X1-X1' bond order (NBO)	-	2	1	1	-	2	1	1
HOMO-LUMO gap	4.61	3.00	6.35	1.35	4.18	2.51	6.08	2.88
			Natural charges					
X1	0.15	0.16	0.66	0.18	0.09	0.18	0.59	0.60
imidazole	-1.22	-1.13	-0.39	-1.66	-1.26	-1.14	-0.44	-1.98
R2	0.11	0.06	0.20	-0.13	0.13	0.08	0.22	-0.02
R3	0.11	0.06	0.20	-0.22	0.13	0.05	0.22	0.00
X1'	-	0.16	0.66	0.18	-	0.16	0.59	0.59
imidazole'	-	-1.13	-0.39	-1.66	-	-1.12	-0.44	-1.99
R2'	-	0.06	0.20	-0.13	-	0.07	0.22	-0.01
R3'	-	0.06	0.20	-0.22	-	0.05	0.22	0.00

Mulliken partial charges

X1	-0.21	0.14	0.36	0.09	-0.24	0.13	0.32	0.23
imidazole	-0.98	-0.87	-0.25	-1.37	-1.02	-0.88	-0.30	-1.60
R2	-0.01	-0.06	0.13	-0.27	0.01	-0.04	0.15	-0.21
R3	-0.01	-0.06	0.13	-0.36	0.01	-0.07	0.15	0.19
X1'	-	0.14	0.36	0.09	-	0.10	0.32	0.21
imidazole'	-	-0.87	-0.25	-1.37	-	-0.86	-0.30	-1.60
R2'	-	-0.06	0.13	-0.27	-	-0.06	0.15	-0.20
R3'	-	-0.06	0.13	-0.36	-	-0.08	0.15	-0.20

Geometries in xyz format

```

130
D1
B1 5.6073 11.9795 3.8713
B2 4.1595 12.1489 3.9926
C1_1 7.0767 11.8340 3.7572
N2_1 8.0532 12.7190 4.1784
C3_1 9.3198 12.2280 3.8915
H3_1 10.2067 12.7945 4.1380
C4_1 9.1619 11.0283 3.2996
H4_1 9.8812 10.3077 2.9360
N5_1 7.7970 14.0576 3.2069
C1_2 2.6895 12.2861 4.1056
N2_2 1.8937 13.3505 3.7194
C3_2 0.5574 13.0920 3.9945
H3_2 -0.2088 13.8216 3.7728
C4_2 0.4901 11.8634 4.5428
H4_2 -0.3499 11.2762 4.8858
N5_2 1.7867 11.3721 4.6192
C1_3 7.8119 14.0576 4.6006
C2_3 7.9698 14.3694 5.9601
C3_3 7.8008 15.6994 6.3387
H3_3 7.9147 15.9808 7.3836
C4_3 7.4770 16.6705 5.3987
H4_3 7.3445 17.7037 5.7151
C5_3 7.3184 16.3326 4.0627
H5_3 7.0621 17.1026 3.3357
C6_3 7.4953 15.0179 3.6307
C1_4 8.2647 13.2816 6.9736
H1_4 8.8691 12.5119 6.4727
C2_4 6.9633 12.6137 7.4248
H2A_4 6.3270 13.3391 7.9519
H2B_4 7.1731 11.7833 8.1122
H2C_4 6.3863 12.2234 6.5747
C3_4 9.0578 13.7744 8.1776
H3A_4 9.9643 14.3148 7.8797
H3B_4 9.3572 12.9251 8.8029
H3C_4 8.4592 14.4414 8.8119
C1_5 7.3721 14.6762 2.1606
H1_5 7.4856 13.5895 2.0464
C2_5 8.4839 15.3395 1.3499
H2A_5 8.4175 16.4346 1.3984
H2B_5 8.4135 15.0483 0.2942
H2C_5 9.4760 15.0480 1.7175
C3_5 5.9907 15.0399 1.6273
H3A_5 5.2127 14.5405 2.2215
H3B_5 5.8887 14.7188 0.5815
H3C_5 5.8115 16.1231 1.6595
C1_6 7.2503 9.5332 2.8092
C2_6 7.1568 9.2466 1.4372
C3_6 6.6830 7.9879 1.0737
H3_6 6.5970 7.7281 0.0206
C4_6 6.3094 7.0615 2.0399
H4_6 5.9364 6.0855 1.7347
C5_6 6.4113 7.3708 3.3879
H5_6 6.1211 6.6340 4.1365
C6_6 6.9000 8.6102 3.8038
C1_7 7.5236 10.2902 0.4007
H1_7 8.3266 10.9123 0.8214
C2_7 6.3328 11.2127 0.1299
H2A_7 5.5052 10.6445 -0.3181
H2B_7 6.6153 12.0102 -0.5707
H2C_7 5.9561 11.6776 1.0517
C3_7 8.0444 9.6930 -0.9009
H3A_7 8.8586 8.9790 -0.7283
H3B_7 8.4215 10.4883 -1.5543
H3C_7 7.2516 9.1742 -1.4557
C1_8 7.0784 8.9012 5.2784
H1_8 7.4262 9.9377 5.3870
C2_8 8.1481 7.9920 5.8805
H2A_8 7.8518 6.9357 5.8300
H2B_8 8.3128 8.2407 6.9363
H2C_8 9.1048 8.0965 5.3526
C3_8 5.7552 8.7932 6.0265
H3A_8 5.0148 9.4711 5.5811
H3B_8 5.8887 9.0693 7.0814
H3C_8 5.3518 7.7719 5.9998
C1_9 2.3725 14.6226 3.2908
C2_9 2.2720 14.9493 1.9282
C3_9 2.6927 16.2193 1.5402
H3_9 2.6313 16.5111 0.4938
C4_9 3.2014 17.1154 2.4726
H4_9 3.5307 18.1009 2.1480
C5_9 3.2941 16.7642 3.8112
H5_9 3.6943 17.4765 4.5321
C6_9 2.8674 15.5115 4.2534
C1_10 1.7665 13.9338 0.9220
H1_10 1.0367 13.2886 1.4332
C2_10 2.9144 13.0373 0.4507
H2A_10 3.6611 13.6312 -0.0958
H2B_10 2.5405 12.2567 -0.2259
H2C_10 3.4289 12.5514 1.2915
C3_10 1.0599 14.5651 -0.2714

```


H3A_10	0.2721	15.2619	0.0385	H28	14.4043	8.8577	5.7468
H3B_10	0.6015	13.7856	-0.8911	H29	14.1922	7.1183	5.8714
H3C_10	1.7613	15.1118	-0.9153	C30	11.5086	7.1727	2.3975
C1_11	2.9145	15.1645	5.7259	C31	10.4566	6.2551	2.2650
H1_11	2.5763	14.1263	5.8469	C32	10.3690	5.5143	1.0903
C2_11	1.9608	16.0534	6.5218	H33	9.5575	4.7932	0.9803
H2A_11	2.2541	17.1100	6.4637	C34	11.2873	5.6693	0.0524
H2B_11	1.9616	15.7660	7.5805	C35	12.3263	6.5804	0.2192
H2C_11	0.9324	15.9709	6.1472	H36	13.0670	6.6976	-0.5733
C3_11	4.3400	15.2349	6.2608	C37	12.4548	7.3407	1.3810
H3A_11	4.9929	14.5720	5.6771	C38	9.4654	6.0620	3.3723
H3B_11	4.3704	14.9147	7.3112	H39	8.6816	5.3559	3.0798
H3C_11	4.7462	16.2543	6.2118	H40	8.9831	7.0080	3.6526
C1_12	2.0897	10.0188	4.9435	H41	9.9614	5.6889	4.2762
C2_12	2.1207	9.6453	6.2961	C42	11.1458	4.8799	-1.2149
C3_12	2.3427	8.3010	6.5873	H43	10.7662	3.8707	-1.0181
H3_12	2.3739	7.9718	7.6240	H44	12.1027	4.7858	-1.7394
C4_12	2.5364	7.3757	5.5685	H45	10.4402	5.3584	-1.9067
H4_12	2.7135	6.3304	5.8163	C46	13.5975	8.2978	1.5435
C5_12	2.5093	7.7747	4.2402	H47	14.4248	8.0358	0.8760
H5_12	2.6645	7.0401	3.4505	H48	13.9689	8.2989	2.5754
C6_12	2.2719	9.1058	3.8970	H49	13.3052	9.3298	1.3043
C1_13	1.9668	10.6853	7.3882	98			
H1_13	1.2977	11.4727	7.0114	D2 dim			
C2_13	3.3186	11.3401	7.6833	N1	13.4768	6.4266	5.8460
H2A_13	4.0215	10.5952	8.0831	N2	13.1455	5.5993	3.7330
H2B_13	3.2075	12.1374	8.4303	C3	13.5043	5.2339	5.0601
H2C_13	3.7714	11.7709	6.7793	C4	12.9725	7.5325	5.0359
C3_13	1.3501	10.1329	8.6673	H5	13.7803	8.2178	4.7228
H3A_13	0.4119	9.5992	8.4741	H6	12.2407	8.1216	5.6055
H3B_13	1.1395	10.9507	9.3664	C7	12.3547	6.8246	3.8379
H3C_13	2.0323	9.4428	9.1809	H8	11.2942	6.5876	4.0225
C1_14	2.1838	9.5135	2.4415	H9	12.4215	7.4014	2.9111
H1_14	2.0617	10.6040	2.3937	C10	13.3757	6.5979	7.2588
C2_14	0.9624	8.8849	1.7737	C11	12.1840	6.3118	7.9454
H2A_14	1.0317	7.7890	1.7613	C12	12.0740	6.6539	9.2924
H2B_14	0.8778	9.2247	0.7335	H13	11.1453	6.4280	9.8192
H2C_14	0.0361	9.1555	2.2964	C14	13.1156	7.2709	9.9795
C3_14	3.4700	9.1757	1.6973	C15	14.2822	7.5586	9.2778
H3A_14	4.3280	9.6505	2.1936	H16	15.1101	8.0463	9.7961
H3B_14	3.4206	9.5426	0.6628	C17	14.4297	7.2427	7.9289
H3C_14	3.6475	8.0926	1.6553	C18	11.0438	5.6327	7.2558
49				H19	10.1473	5.6261	7.8847
M2				H20	11.3032	4.5928	7.0186
N1	11.6403	8.4770	5.6571	H21	10.7891	6.1173	6.3041
N2	11.6076	7.9608	3.5776	C22	13.0074	7.5828	11.4417
C3	11.8183	7.4406	4.8075	H23	11.9634	7.6714	11.7595
C4	11.4464	9.7866	4.9979	H24	13.5213	8.5177	11.6934
H5	12.3725	10.3796	5.0662	H25	13.4660	6.7906	12.0493
H6	10.6428	10.3618	5.4725	C26	15.7006	7.6053	7.2233
C7	11.1271	9.3596	3.5758	H27	16.3979	8.0976	7.9109
H8	10.0473	9.3834	3.3574	H28	15.5212	8.2852	6.3813
H9	11.6375	9.9603	2.8139	H29	16.2003	6.7168	6.8189
C10	11.9391	8.4155	7.0467	C30	14.0308	5.5070	2.6112
C11	10.9023	8.5694	7.9751	C31	13.4506	5.1760	1.3705
C12	11.2133	8.5317	9.3327	C32	14.2670	5.0104	0.2554
H13	10.4089	8.6383	10.0624	H33	13.8056	4.7276	-0.6928
C14	12.5189	8.3447	9.7797	C34	15.6473	5.1728	0.3221
C15	13.5265	8.1817	8.8309	C35	16.1904	5.5841	1.5353
H16	14.5526	8.0209	9.1652	H36	17.2622	5.7826	1.5985
C17	13.2620	8.2176	7.4640	C37	15.4095	5.7906	2.6743
C18	9.4835	8.7442	7.5228	C38	11.9669	5.0393	1.2179
H19	8.7830	8.4607	8.3150	H39	11.7187	4.4337	0.3389
H20	9.2729	8.1335	6.6364	H40	11.4886	6.0197	1.0827
H21	9.2628	9.7873	7.2561	H41	11.5145	4.5751	2.0992
C22	12.8324	8.3348	11.2465	C42	16.5144	4.9035	-0.8704
H23	13.7542	7.7818	11.4567	H43	17.4706	5.4330	-0.8014
H24	12.0231	7.8774	11.8268	H44	16.0245	5.2064	-1.8031
H25	12.9692	9.3535	11.6322	H45	16.7422	3.8330	-0.9599
C26	14.3606	8.0261	6.4629	C46	16.0583	6.4145	3.8738
H27	15.3346	7.9529	6.9571	H47	17.1503	6.3955	3.7795

H48 15.7875 5.9330 4.8146
H49 15.7591 7.4685 3.9622
N50 14.2679 3.6098 6.7441
N51 13.6594 2.7571 4.7002
C52 13.7893 3.9671 5.4525
C53 13.8687 2.2189 6.9600
H54 12.8361 2.1683 7.3431
H55 14.5308 1.7391 7.6871
C56 13.9646 1.6187 5.5645
H57 14.9750 1.2128 5.3771
H58 13.2422 0.8100 5.3947
C59 15.5593 3.9748 7.2478
C60 16.7333 4.0134 6.4692
C61 17.9159 4.4632 7.0613
H62 18.8161 4.5141 6.4452
C63 17.9938 4.8121 8.4053
C64 16.8472 4.6597 9.1794
H65 16.8855 4.8894 10.2459
C66 15.6385 4.2434 8.6298
C67 16.8133 3.4862 5.0658
H68 17.7043 3.8708 4.5553
H69 15.9410 3.7244 4.4562
H70 16.8994 2.3905 5.0793
C71 19.2617 5.3450 9.0012
H72 19.2449 6.4418 9.0502
H73 20.1364 5.0607 8.4067
H74 19.4122 4.9804 10.0240
C75 14.4526 4.0487 9.5238
H76 14.5665 4.6182 10.4532
H77 14.3335 2.9919 9.8018
H78 13.5254 4.3612 9.0334
C79 12.8568 2.4711 3.5555
C80 13.5040 2.0190 2.3926
C81 12.7365 1.5838 1.3148
H82 13.2452 1.2454 0.4103
C83 11.3454 1.5708 1.3579
C84 10.7230 2.0085 2.5233
H85 9.6332 2.0003 2.5834
C86 11.4538 2.4573 3.6229
C87 14.9986 1.9936 2.2930
H88 15.3146 1.6450 1.3031
H89 15.4500 1.3309 3.0429
H90 15.4232 2.9930 2.4478
C91 10.5476 1.1353 0.1659
H92 11.0203 0.2915 -0.3498
H93 10.4562 1.9489 -0.5661
H94 9.5324 0.8373 0.4478
C95 10.7361 2.9261 4.8474
H96 9.6640 2.7109 4.7862
H97 10.8642 4.0080 4.9766
H98 11.1285 2.4559 5.7581
17

M3

N1 11.9140 8.7456 5.5251
N2 11.2262 8.1070 3.6057
C3 11.7655 7.6479 4.7543
C4 11.4148 9.9982 4.9444
H5 12.1861 10.7806 4.9704
H6 10.5506 10.3698 5.5170
C7 11.0451 9.5618 3.5293
H8 10.0130 9.8173 3.2514
H9 11.7077 9.9920 2.7620
C10 10.9732 7.2960 2.4489
H11 11.1914 6.2577 2.7085
H12 9.9243 7.3726 2.1268
C13 12.3749 8.7244 6.8842
H14 12.6750 7.7021 7.1244
H15 13.2359 9.3940 7.0262
H16 11.5859 9.0385 7.5849
H17 11.6080 7.5922 1.5994
34

D3 dim

N1 12.2823 5.5531 6.5113
N2 12.0888 4.9319 4.2518
C3 12.0279 6.0050 5.1857
C4 12.2514 4.0918 6.4368
H5 11.2318 3.7275 6.6352
H6 12.9206 3.6512 7.1840
C7 12.6526 3.8029 4.9933
H8 13.7485 3.7962 4.8901
H9 12.2693 2.8533 4.6041
C10 10.8438 4.6566 3.5504
H11 10.5301 5.5630 3.0220
H12 11.0078 3.8655 2.8100
C13 13.4663 6.1209 7.1384
H14 13.3675 7.2116 7.1544
H15 13.5380 5.7665 8.1730
N24 11.5674 8.3292 5.8044
N25 11.6463 7.7634 3.5233
C26 11.7637 7.2833 4.8585
C27 11.5777 9.5668 5.0233
H28 12.6012 9.9686 4.9707
H29 10.9390 10.3249 5.4897
C30 11.1058 9.1183 3.6434
H31 10.0062 9.0886 3.6013
H32 11.4600 9.7511 2.8222
C33 12.8531 7.6424 2.7190
H34 13.1511 6.5889 2.6878
H35 12.6450 7.9722 1.6948
C36 10.4207 8.1556 6.6835
H37 10.5326 7.2104 7.2254
H38 10.3938 8.9710 7.4153
H33 9.4568 8.1352 6.1420
H36 13.7002 8.2327 3.1149
H39 14.4036 5.8634 6.6111
H40 10.0245 4.3450 4.2246
23

M4

N1 11.9767 8.7690 5.4954
N2 11.1552 8.1095 3.6294
C3 11.7620 7.6634 4.7502
C4 11.3784 9.9974 4.9518
H5 12.0718 10.8454 5.0212
H6 10.4637 10.2654 5.5043
C7 11.0750 9.5739 3.5202
H8 10.0867 9.8957 3.1677
H9 11.8256 9.9556 2.8099
C10 10.9478 7.2698 2.4701
H11 10.8366 6.2474 2.8466
H12 9.9973 7.5513 1.9912
C13 12.4162 8.7078 6.8721
H14 13.0018 7.7871 6.9664
H15 13.0939 9.5529 7.0687
C16 12.0910 7.3407 1.4715
H17 11.9184 6.6677 0.6245
H18 12.2187 8.3526 1.0689
H19 13.0267 7.0484 1.9616
C20 11.2664 8.7038 7.8656
H21 11.6340 8.6193 8.8942
H22 10.6694 9.6214 7.8026
H23 10.6060 7.8530 7.6626
45

D4 dim

N1 12.3259 5.5545 6.5036
N2 12.0480 4.9215 4.2652
C3 12.0289 6.0033 5.1874
C4 12.3235 4.0926 6.4567
H5 11.3456 3.7017 6.7762
H6 13.0819 3.6826 7.1345
C7 12.5850 3.7670 4.9848
H8 13.6636 3.6633 4.7918
H9 12.0982 2.8427 4.6510

C10	10.8144	4.7025	3.5094	N16	11.5461	8.3949	5.7447
H11	10.5946	5.6377	2.9772	N17	11.6810	7.8783	3.4851
H12	11.0252	3.9446	2.7421	C18	11.8723	7.3779	4.8038
C13	13.4797	6.1832	7.1481	C19	11.1410	9.4991	4.9611
H14	13.2878	7.2645	7.1626	C20	11.2505	9.2133	3.6615
H15	13.4962	5.8516	8.1956	C21	10.6631	8.0091	6.8342
C16	9.6178	4.2846	4.3519	H22	10.4594	8.8846	7.4598
H17	8.7069	4.2305	3.7447	H23	9.7084	7.6020	6.4618
H18	9.7655	3.2981	4.8109	H24	11.1397	7.2396	7.4509
H19	9.4457	5.0083	5.1593	H25	11.0774	9.8433	2.7987
C20	14.8171	5.8960	6.4808	H26	10.7897	10.4028	5.4422
H21	15.6239	6.4702	6.9505	C27	12.7741	7.6650	2.5491
H22	15.0879	4.8340	6.5481	H28	13.7184	8.0989	2.9179
H23	14.7817	6.1669	5.4175	H29	12.9329	6.5924	2.3942
N24	11.5232	8.3247	5.7987	H30	12.5201	8.1248	1.5881
N25	11.6879	7.7775	3.5269	21			
C26	11.7634	7.2854	4.8587	M6			
C27	11.5037	9.5765	5.0425	N1	11.9745	8.8052	5.4684
H28	12.4895	10.0640	5.0842	N2	11.1380	8.1485	3.6410
H29	10.7711	10.2733	5.4674	C3	11.7622	7.6580	4.7547
C30	11.1700	9.1426	3.6138	C4	11.5041	9.9466	4.8326
H31	10.0823	9.1506	3.4462	C7	10.9784	9.5278	3.6594
H32	11.6283	9.7795	2.8476	C6	12.5475	8.8029	6.8027
C33	12.8828	7.5851	2.7043	H7	13.0811	7.8525	6.9020
H34	13.0902	6.5067	2.6880	H8	13.2874	9.6121	6.8715
H35	12.6274	7.8731	1.6751	H9	10.5166	10.0782	2.8499
C36	10.4113	8.1051	6.7244	H10	11.5796	10.9357	5.2657
H37	10.6193	7.1677	7.2574	C11	10.7774	7.3095	2.5120
H38	10.4400	8.9078	7.4743	H12	10.7367	6.2854	2.8958
C39	14.1109	8.3483	3.1788	H13	9.7676	7.5826	2.1762
H40	14.9935	8.0872	2.5837	C14	11.7785	7.4267	1.3789
H41	13.9725	9.4348	3.1017	H15	11.5033	6.7781	0.5406
H42	14.3263	8.1108	4.2286	H16	11.8364	8.4567	1.0064
C43	9.0398	8.0425	6.0675	H17	12.7750	7.1387	1.7308
H44	8.2643	7.7808	6.7964	C18	11.4838	8.9448	7.8744
H45	8.7573	9.0040	5.6186	H19	11.9273	8.9267	8.8754
H46	9.0292	7.2850	5.2729	H20	10.9337	9.8872	7.7639
15				H21	10.7640	8.1229	7.7962
M5				42			
N1	11.7583	8.5252	5.6212	D6 dim			
N2	11.4723	7.9609	3.6028	N1	11.4204	5.5916	6.5395
C3	11.8187	7.4225	4.8126	N2	11.9035	4.7692	4.4213
C4	11.3947	9.6841	4.9486	C3	11.8533	5.9446	5.2259
C7	11.2118	9.3234	3.6588	C4	11.3690	4.1733	6.5335
C6	12.0482	8.4750	7.0352	C7	11.6684	3.7078	5.3180
H7	11.1803	8.7873	7.6279	C6	10.1894	6.2561	6.9982
H8	12.2982	7.4412	7.2811	H7	10.0599	6.0044	8.0604
H9	12.8968	9.1215	7.2880	H8	10.3600	7.3397	6.9462
H10	10.9250	9.9045	2.7925	H9	11.7591	2.6850	4.9732
H11	11.2994	10.6442	5.4381	H10	11.0757	3.6355	7.4261
C12	11.3897	7.1768	2.3927	C11	12.9938	4.5961	3.4622
H13	12.0934	7.5426	1.6357	H12	12.8428	3.6117	2.9991
H14	11.6442	6.1474	2.6523	H13	12.8874	5.3458	2.6691
H15	10.3773	7.2034	1.9727	C14	14.3830	4.6951	4.0706
30				H15	15.1455	4.3985	3.3422
D5 dim				H16	14.5972	5.7231	4.3820
N1	12.7885	5.7226	6.3849	H17	14.4724	4.0428	4.9492
N2	12.3235	5.0242	4.2206	C18	8.9530	5.8742	6.2017
C3	12.2974	6.1321	5.1136	H19	8.0838	6.4612	6.5179
C4	13.1182	4.3576	6.2258	H20	9.1161	6.0530	5.1311
C7	12.8190	3.9476	4.9909	H21	8.7134	4.8119	6.3269
C6	13.8007	6.5729	6.9916	N22	12.3602	8.3292	5.6756
H7	14.1221	6.1293	7.9399	N23	12.2853	7.6082	3.4679
H8	14.6782	6.6960	6.3352	C24	12.1493	7.2014	4.8296
H9	13.3868	7.5672	7.1911	C25	12.7930	9.3532	4.8095
H10	12.8910	2.9598	4.5545	C26	12.7330	8.9532	3.5367
H11	13.5628	3.8099	7.0466	C27	13.0998	8.1645	6.9262
C12	11.1081	4.7766	3.4610	H28	13.1694	9.1638	7.3766
H13	11.2437	3.8807	2.8461	H29	12.5050	7.5442	7.6073
H14	10.2362	4.6329	4.1205	H30	12.9306	9.5084	2.6284
H15	10.8965	5.6259	2.8029	H31	13.1267	10.2978	5.2214

C32	11.0991	7.3901	2.6237	H10A	2.5906	1.5591	1.9563
H33	11.3921	7.6090	1.5871	H10B	4.1664	1.8716	1.2123
H34	10.8559	6.3200	2.6669	H10C	4.0773	0.9384	2.7028
C35	9.8957	8.2246	3.0289	C11	0.6290	-2.2449	-0.5560
H36	9.0149	7.9597	2.4337	H11A	0.3275	-3.0641	0.1115
H37	9.6553	8.0624	4.0875	H11B	0.6007	-2.6364	-1.5795
H38	10.0916	9.2947	2.8943	H11C	-0.1235	-1.4534	-0.4730
C39	14.4814	7.5541	6.7596	C12	5.6072	-2.9060	-0.5040
H40	15.0409	7.6048	7.7001	H12A	6.3113	-2.3627	-1.1475
H41	14.4078	6.5008	6.4686	H12B	5.3163	-3.8290	-1.0178
H42	15.0537	8.0860	5.9882	H12C	6.1536	-3.1691	0.4122
43				N2	5.8689	-0.4937	0.9496
M7				C44	6.4813	-1.3853	3.1801
C1	4.9328	1.2064	1.7246	C45	6.3223	-3.4564	4.1773
C2	4.9909	3.3957	0.8207	H46	7.1260	-4.0537	3.7220
H2A	5.2050	3.8017	-0.1751	H47	5.5823	-4.1487	4.5928
H2B	5.3277	4.1290	1.5729	C71	4.2707	-2.4125	3.4441
C3	3.5392	2.9989	1.0216	H72	3.9453	-1.4645	3.0037
H3A	2.9518	3.7442	1.5693	H73	4.0764	-2.3219	4.5332
H3B	3.0218	2.7868	0.0729	C74	3.4132	-3.5149	2.8811
C4	2.5713	1.0940	2.3629	C75	2.0307	-3.4383	3.0611
C5	1.9814	1.6358	3.5127	C35	1.1893	-4.4067	2.5312
C6	0.8700	0.9948	4.0560	H36	0.1136	-4.3310	2.6796
H6	0.4127	1.4046	4.9583	C37	1.7212	-5.4719	1.8081
C7	0.3419	-0.1621	3.4892	H38	1.0644	-6.2321	1.3907
C8	0.9605	-0.6841	2.3547	C39	3.0964	-5.5580	1.6286
H8	0.5673	-1.5973	1.9054	H40	3.5211	-6.3859	1.0642
C9	2.0689	-0.0731	1.7734	C41	3.9375	-4.5861	2.1628
C10	2.5403	2.8636	4.1690	H42	5.0128	-4.6386	2.0018
H10A	2.1310	3.7852	3.7314	N43	5.6963	-2.5830	3.1728
H10B	3.6311	2.9128	4.0659	C46	6.8996	-2.4692	5.1845
H10C	2.2915	2.8810	5.2353	H45	6.1294	-2.1247	5.8938
C11	-0.8696	-0.8204	4.0798	H48	7.7433	-2.8679	5.7557
H11A	-0.8725	-0.7494	5.1735	C47	8.7492	-1.1691	4.1969
H11B	-0.9229	-1.8803	3.8092	C48	9.3858	-0.5454	5.2895
H11C	-1.7942	-0.3470	3.7247	C49	10.7546	-0.3056	5.2406
C12	2.7208	-0.6596	0.5589	H50	11.2305	0.1978	6.0840
H12A	2.7699	0.0679	-0.2622	C51	11.5214	-0.6547	4.1322
H12B	2.1712	-1.5367	0.2028	C52	10.8757	-1.2807	3.0727
H12C	3.7552	-0.9504	0.7757	H53	11.4605	-1.5813	2.2013
C13	7.1197	2.0803	1.0820	C54	9.5072	-1.5625	3.0773
H13A	7.5269	2.4686	0.1364	C55	8.6080	-0.1138	6.4973
H13B	7.4042	1.0243	1.1603	H56	9.1765	0.6169	7.0817
C14	7.6763	2.8669	2.2453	H57	8.3735	-0.9554	7.1637
C15	8.6366	3.8572	2.0602	H58	7.6482	0.3368	6.2189
H15	9.0033	4.0662	1.0550	C59	12.9933	-0.3690	4.0962
C16	9.1270	4.5800	3.1448	H60	13.2059	0.6657	4.3930
H16	9.8759	5.3532	2.9873	H61	13.4090	-0.5234	3.0948
C17	8.6522	4.3207	4.4244	H62	13.5473	-1.0174	4.7871
H17	9.0292	4.8882	5.2724	C63	8.9457	-2.2801	1.8825
C18	7.6873	3.3344	4.6168	H64	8.2002	-3.0381	2.1454
H18	7.3121	3.1281	5.6170	H65	9.7545	-2.7806	1.3379
C19	7.2019	2.6132	3.5347	H66	8.4459	-1.5933	1.1869
H19	6.4442	1.8394	3.6703	N67	7.3383	-1.3737	4.3146
N1	5.6732	2.1092	1.0384	C68	7.0827	1.5068	1.0664
N2	3.6981	1.7499	1.7950	H69	8.0463	1.9409	0.7705
86				H70	6.3573	2.3390	1.1465
D7 dim				C72	6.9459	1.6244	3.5009
C1	6.4259	-0.4081	2.2442	H74	6.8832	0.9585	4.3718
C3	6.5862	0.4421	0.0936	H75	5.9655	2.1403	3.4244
H3A	5.9174	0.8494	-0.6738	C76	8.0253	2.6480	3.7248
H3B	7.4318	-0.0470	-0.4210	C77	7.7032	3.9008	4.2437
C4	4.5537	-0.9091	0.5977	C78	8.6974	4.8343	4.5168
C5	3.4294	-0.1439	0.9476	H77	8.4298	5.8079	4.9223
C6	2.1672	-0.5876	0.5557	C79	10.0292	4.5251	4.2647
H6	1.2906	-0.0020	0.8386	H79	10.8088	5.2542	4.4750
C7	1.9964	-1.7522	-0.1897	C80	10.3572	3.2800	3.7362
C8	3.1294	-2.4782	-0.5482	H81	11.3970	3.0283	3.5328
H8	3.0126	-3.4014	-1.1186	C82	9.3631	2.3485	3.4655
C9	4.4060	-2.0805	-0.1603	H83	9.6157	1.3720	3.0547
C10	3.5702	1.1183	1.7439	N84	7.1851	0.7941	2.3351

H85	1.6138	-2.5970	3.6165	H12B	4.8668	1.3146	4.1471
H86	6.6582	4.1455	4.4359	H12C	5.5096	-0.3127	3.8641
43				N2	7.8288	0.4587	1.5650
M8				C44	7.1389	-1.8035	2.3036
C1	3.3458	2.4352	10.8520	C45	6.3852	-3.5569	3.6287
C3	5.3624	3.4343	10.0921	H46	6.0935	-4.6125	3.5647
H3A	5.3930	3.5587	8.9969	H47	5.7775	-3.0862	4.4256
H3B	6.3960	3.3842	10.4563	C71	4.7766	-2.4980	2.1771
C4	4.5218	4.5009	10.7758	H72	4.7090	-1.6106	1.5354
H4A	4.9324	4.8013	11.7529	H73	4.3321	-2.1934	3.1446
H4B	4.3864	5.4057	10.1724	C74	3.9361	-3.5864	1.5646
C5	2.1481	4.4336	11.6045	C75	2.5526	-3.4485	1.4254
C6	1.4029	5.3751	10.8825	I35	1.5493	-1.7065	2.1483
C8	0.3489	6.0253	11.5222	C77	1.7776	-4.4130	0.7941
H8	-0.2422	6.7504	10.9598	H78	0.7043	-4.2697	0.6938
C9	0.0219	5.7620	12.8494	C79	2.3844	-5.5580	0.2894
C10	0.7746	4.8130	13.5381	H80	1.7792	-6.3135	-0.2069
H10	0.5261	4.5869	14.5764	C81	3.7566	-5.7231	0.4199
C11	1.8405	4.1436	12.9424	H82	4.2441	-6.6094	0.0192
C12	1.7067	5.6706	9.4436	C83	4.5174	-4.7467	1.0502
H12A	2.4992	6.4252	9.3381	H84	5.5987	-4.8550	1.1351
H12B	0.8218	6.0649	8.9326	N85	6.1725	-2.8647	2.3570
H12C	2.0424	4.7720	8.9138	C46	7.8533	-3.3543	3.9025
C13	-1.0992	6.4944	13.5260	H48	8.0863	-3.3592	4.9755
H13A	-0.7530	7.4427	13.9580	H49	8.4657	-4.1369	3.4214
H13B	-1.5291	5.9048	14.3433	C48	9.5140	-1.8142	3.0697
H13C	-1.9041	6.7360	12.8225	C49	10.2530	-1.2821	4.1412
C14	2.6260	3.1270	13.7148	C50	11.6226	-1.0841	3.9886
H14A	3.7038	3.3346	13.6776	H51	12.1870	-0.6533	4.8180
H14B	2.4930	2.1257	13.2892	C52	12.2835	-1.4088	2.8070
H14C	2.3183	3.1111	14.7654	C53	11.5392	-1.9765	1.7778
C15	5.0538	0.9349	9.9385	H54	12.0450	-2.2705	0.8557
H15A	4.4338	0.1799	10.4346	C55	10.1656	-2.1994	1.8833
H15B	6.0966	0.7631	10.2340	C56	9.5907	-0.9429	5.4423
C16	4.8848	0.8201	8.4407	H57	10.2081	-0.2490	6.0244
C17	5.9173	0.7098	7.5114	H58	9.4382	-1.8394	6.0602
C18	5.6594	0.6197	6.1442	H59	8.6067	-0.4888	5.2817
H18	6.4823	0.5319	5.4392	C60	13.7455	-1.1240	2.6347
C19	4.3491	0.6501	5.6875	H61	9.2473	2.4662	7.2416
H19	4.1509	0.5804	4.6201	H62	13.9038	-0.1139	2.2328
C20	3.3010	0.7769	6.5944	H63	14.2147	-1.8259	1.9359
H20	2.2720	0.8085	6.2434	C64	9.4622	-2.8440	0.7244
C21	3.5738	0.8622	7.9497	H65	8.6462	-3.5095	1.0299
H21	2.7693	0.9805	8.6776	H66	10.1729	-3.4293	0.1298
N1	3.2449	3.7795	10.9736	H67	9.0183	-2.0864	0.0632
N2	4.6133	2.2168	10.4400	N68	8.1206	-2.0468	3.2997
I1	7.9725	0.6980	8.0974	C69	6.8133	0.4409	-0.5266
86				H70	6.0226	0.8907	-1.1403
D8 dim				H71	7.6201	0.1063	-1.2068
C1	7.1016	-0.7549	1.4473	C72	6.2280	-1.9009	-0.5505
C3	7.3647	1.3748	0.5205	H74	6.3340	-2.7787	0.0990
H3A	8.1837	1.9993	0.1399	H75	7.0823	-1.9530	-1.2534
H3B	6.5823	2.0484	0.9125	C76	4.9421	-2.0370	-1.3214
C4	8.0136	1.1059	2.8281	C78	4.7400	-3.0865	-2.2217
C5	9.2697	1.6941	3.0618	I77	6.2819	-4.5162	-2.5977
C6	9.4887	2.3721	4.2574	C80	3.5326	-3.2635	-2.8851
H6	10.4726	2.8079	4.4424	H79	3.4053	-4.0970	-3.5719
C7	8.4937	2.4945	5.2246	C82	2.4894	-2.3717	-2.6603
C8	7.2445	1.9492	4.9482	H81	1.5425	-2.5101	-3.1775
H8	6.4400	2.0653	5.6773	C84	2.6653	-1.3172	-1.7748
C9	6.9738	1.2637	3.7622	H83	1.8524	-0.6203	-1.5818
C10	10.3624	1.6068	2.0392	C85	3.8781	-1.1568	-1.1169
H10A	11.3360	1.8336	2.4889	H85	4.0168	-0.3472	-0.4005
H10B	10.2062	2.3243	1.2206	N86	6.3090	-0.6907	0.2518
H10C	10.4064	0.6087	1.5895	47			
C11	8.7735	3.1640	6.5370	M9			
H11A	14.2841	-1.1801	3.5874	C20	4.2217	12.7097	15.0619
H11B	7.8541	3.5268	7.0101	H20A	4.9731	12.5174	15.8414
H11C	9.4551	4.0146	6.4207	H20B	3.2408	12.8096	15.5319
C12	5.5835	0.7379	3.5512	H20C	4.4730	13.6532	14.5555
H12A	5.2607	0.7735	2.5041	O1	4.1256	11.6412	14.1556

C1	3.5310	9.0633	9.9821	H46	5.8397	-4.1138	4.2567
C2	2.7484	11.1501	10.7828	H47	5.6773	-2.5432	5.0792
H2A	3.5864	11.8512	10.9409	C71	4.5909	-2.0531	2.8085
H2B	1.9198	11.4571	11.4316	H72	4.5071	-1.1475	2.1970
C3	2.3747	11.0322	9.3157	H73	4.1356	-1.7752	3.7789
H3A	1.2959	10.8708	9.1645	C74	3.7813	-3.1528	2.1736
H3B	2.6748	11.8970	8.7136	C75	2.4060	-2.9423	1.9724
C4	3.2628	9.4300	7.5756	C35	1.6207	-3.9093	1.3517
C5	2.4884	8.3837	7.0588	H36	0.5610	-3.7407	1.1843
C6	2.6432	8.0446	5.7171	C37	2.2061	-5.1013	0.9285
H6	2.0434	7.2310	5.3059	H38	1.5869	-5.8560	0.4469
C7	3.5408	8.7156	4.8877	C39	3.5610	-5.3245	1.1178
C8	4.3001	9.7475	5.4327	H40	4.0181	-6.2539	0.7840
H8	5.0168	10.2750	4.8009	C41	4.3393	-4.3450	1.7343
C9	4.1820	10.1159	6.7717	H42	5.4118	-4.4853	1.8632
C10	1.5312	7.6411	7.9396	N43	5.9876	-2.4206	2.9870
H10A	2.0612	7.1769	8.7796	C46	7.6992	-2.9795	4.4616
H10B	1.0025	6.8633	7.3796	H45	7.9902	-2.9758	5.5198
H10C	0.7820	8.3133	8.3780	H48	8.2251	-3.8165	3.9703
C11	3.7124	8.3102	3.4535	C47	9.4109	-1.5959	3.4790
H11A	4.4962	7.5489	3.3460	C48	10.2707	-1.1382	4.4945
H11B	4.0011	9.1612	2.8265	C49	11.6359	-1.0450	4.2381
H11C	2.7903	7.8841	3.0431	H50	12.2934	-0.6701	5.0250
C12	5.0390	11.2089	7.3381	C51	12.1784	-1.4032	3.0063
H12A	5.9385	11.3526	6.7307	C52	11.3161	-1.9033	2.0366
H12B	5.3461	10.9818	8.3667	H53	11.7249	-2.2273	1.0775
H12C	4.5128	12.1735	7.3647	C54	9.9418	-2.0222	2.2481
C13	3.7867	9.5039	12.3721	C55	9.7439	-0.7701	5.8483
H13A	3.9549	8.4224	12.4119	H56	10.4419	-0.1058	6.3705
H13B	3.0838	9.7731	13.1681	H57	9.6079	-1.6600	6.4793
C14	5.0956	10.2378	12.5449	H58	8.7714	-0.2709	5.7755
C15	6.1799	9.8877	11.7472	C59	13.6403	-1.2246	2.7265
H15	6.0395	9.0777	11.0299	H60	9.7092	2.5727	7.7351
C16	7.3952	10.5579	11.8392	H61	13.8551	-0.1980	2.4002
H16	8.2324	10.2658	11.2104	H62	13.9846	-1.8955	1.9315
C17	7.5246	11.6033	12.7434	C63	9.1056	-2.6157	1.1541
H17	8.4678	12.1383	12.8315	H64	8.2812	-3.2309	1.5329
C18	6.4511	11.9877	13.5411	H65	9.7267	-3.2400	0.5017
H18	6.5689	12.8168	14.2325	H66	8.6480	-1.8322	0.5356
C19	5.2348	11.3112	13.4367	N67	8.0246	-1.7119	3.8060
N1	3.1870	9.7773	11.0794	C68	6.7137	0.8935	0.0852
N2	3.1226	9.8157	8.9369	H69	5.9470	1.4206	-0.4964
94				H70	7.4549	0.4834	-0.6282
D9 dim				C72	5.8267	-1.3511	0.0747
C1	6.9803	-0.3466	2.0302	H74	5.9191	-2.2644	0.6731
C3	7.3967	1.7642	1.1075	H75	6.5771	-1.4706	-0.7308
H3A	8.2522	2.3122	0.6912	C76	4.4495	-1.3054	-0.5312
H3B	6.6991	2.5053	1.5359	C77	4.0424	-2.3804	-1.3410
C4	8.1263	1.4101	3.3803	C78	2.7632	-2.4157	-1.8877
C5	9.4341	1.9011	3.5439	H77	2.4439	-3.2539	-2.5002
C6	9.7751	2.5374	4.7341	C79	1.8785	-1.3693	-1.6321
H6	10.7978	2.8968	4.8640	H79	0.8802	-1.4000	-2.0648
C7	8.8533	2.7099	5.7645	C80	2.2638	-0.3018	-0.8361
C8	7.5524	2.2637	5.5571	H81	1.5710	0.5125	-0.6340
H8	6.8043	2.4238	6.3361	C82	3.5458	-0.2813	-0.2871
C9	7.1608	1.6262	4.3791	H83	3.8567	0.5296	0.3700
C10	10.4532	1.7577	2.4540	N84	6.1322	-0.1811	0.8862
H10A	11.4669	1.8903	2.8488	C85	0.6022	-1.4190	2.1215
H10B	10.3085	2.5111	1.6661	H86	0.4374	-0.4095	2.5055
H10C	10.3886	0.7733	1.9763	H87	0.4198	-1.4262	1.0363
C11	9.2626	3.3245	7.0694	H88	-0.1031	-2.1064	2.6115
H11A	14.2491	-1.4123	3.6184	C89	4.5885	-4.5153	-2.2035
H11B	8.4062	3.7570	7.5986	H90	5.4510	-5.1859	-2.1907
H11C	10.0101	4.1138	6.9288	H91	3.7418	-5.0079	-1.7019
C12	5.7244	1.2186	4.2359	H92	4.3169	-4.3041	-3.2483
H12A	5.3536	1.3141	3.2088	O93	4.9762	-3.3509	-1.5192
H12B	5.0916	1.8327	4.8868	O94	1.9372	-1.7459	2.4135
H12C	5.5798	0.1673	4.5183	49			
N2	7.8172	0.7993	2.1249	M10			
C44	6.9956	-1.4084	2.8712	N1	11.7899	8.4422	5.7197
C45	6.2093	-3.0813	4.2702	N2	11.4888	7.8293	3.5795

B3	11.5485	7.2124	4.9280	H23	15.4000	6.6198	11.7518
C4	12.0002	9.6619	4.9375	H24	15.5788	8.1897	10.9610
H5	13.0728	9.9212	4.8231	H25	16.8160	6.9475	10.7296
H6	11.5063	10.5399	5.3863	C26	11.4593	5.4715	8.7954
C7	11.4100	9.2933	3.5838	H27	11.1314	5.7405	9.8083
H8	10.3573	9.6322	3.5044	H28	11.2013	4.4160	8.6271
H9	11.9567	9.7665	2.7528	H29	10.8671	6.0582	8.0795
C10	12.0622	8.4338	7.0965	C30	12.4809	4.7427	2.5439
C11	11.0054	8.2230	8.0085	C31	13.6400	5.2078	1.8531
C12	11.2604	8.2488	9.3751	C32	13.7079	5.0926	0.4666
H13	10.4322	8.0846	10.0697	H33	14.6002	5.4758	-0.0405
C14	12.5400	8.4810	9.8829	C34	12.6955	4.5202	-0.2936
C15	13.5790	8.6499	8.9741	C35	11.5778	4.0260	0.3932
H16	14.5983	8.7839	9.3467	H36	10.7495	3.5882	-0.1741
C17	13.3646	8.6186	7.5948	C37	11.4578	4.1348	1.7743
C18	9.6393	7.9338	7.4753	C38	14.7862	5.7934	2.5932
H19	8.9305	7.7067	8.2817	H39	15.5975	6.0739	1.9062
H20	9.7134	7.0901	6.7675	H40	15.1943	5.0926	3.3383
H21	9.2523	8.7778	6.8879	H41	14.4637	6.6769	3.1843
C22	12.7799	8.5426	11.3639	C42	12.7545	4.4952	-1.7910
H23	13.8467	8.4466	11.6002	H43	12.3663	3.5508	-2.2052
H24	12.2476	7.7423	11.8947	H44	13.7843	4.6202	-2.1546
H25	12.4360	9.4923	11.7987	H45	12.1550	5.3053	-2.2385
C26	14.5230	8.6744	6.6468	C46	10.1786	3.6892	2.4214
H27	15.4597	8.4143	7.1548	H47	9.3487	3.7401	1.7034
H28	14.6582	9.6659	6.1929	H48	9.9425	4.3381	3.2752
H29	14.3512	7.9673	5.8223	H49	10.2259	2.6565	2.8012
C30	11.3578	7.1345	2.3688	N50	12.7564	8.8331	5.3666
C31	10.2110	6.3546	2.0983	N51	12.0893	8.3062	3.1592
C32	10.1086	5.6866	0.8826	B52	12.4217	7.6537	4.4836
H33	9.2153	5.0874	0.6862	C53	13.1862	9.9002	4.4822
C34	11.1015	5.7666	-0.0956	H54	14.2395	9.7693	4.1512
C35	12.2266	6.5339	0.1830	H55	13.0945	10.8916	4.9516
H36	13.0297	6.5947	-0.5564	C56	12.2759	9.7447	3.2747
C37	12.3736	7.2093	1.3954	H57	11.3058	10.2562	3.4553
C38	9.1324	6.2535	3.1284	H58	12.7165	10.2019	2.3727
H39	8.3440	5.5553	2.8195	C59	12.6652	9.0475	6.7393
H40	8.6759	7.2350	3.3213	C60	11.4579	8.7681	7.4456
H41	9.5906	5.9520	4.0882	C61	11.3994	8.9585	8.8216
C42	10.9677	5.0172	-1.3897	H62	10.4633	8.7186	9.3378
H43	11.6910	5.3713	-2.1344	C63	12.4799	9.4307	9.5623
H44	9.9639	5.1269	-1.8214	C64	13.6504	9.7366	8.8595
H45	11.1380	3.9393	-1.2591	H65	14.5295	10.0832	9.4127
C46	13.6380	7.9587	1.6881	C66	13.7627	9.5467	7.4873
H47	14.4686	7.5864	1.0759	C67	10.2591	8.2900	6.7093
H48	13.8927	7.8532	2.7511	H68	9.3971	8.1733	7.3814
H49	13.5478	9.0374	1.4932	H69	10.4757	7.3327	6.1937
98				H70	9.9831	8.9828	5.8983
D10A				C71	12.4319	9.5153	11.0574
N1	12.7732	5.4279	6.1924	H72	13.0053	10.3736	11.4408
N2	12.3328	4.8814	3.9326	H73	12.8561	8.6144	11.5327
B3	12.5552	6.0699	4.8397	H74	11.3998	9.6130	11.4220
C4	12.1097	4.1376	6.1427	C75	15.0995	9.7472	6.8364
H5	11.0116	4.2273	6.2916	H76	15.8783	9.8957	7.5962
H6	12.5028	3.4395	6.8973	H77	15.1342	10.6030	6.1472
C7	12.3700	3.6608	4.7228	H78	15.3642	8.8584	6.2473
H8	13.3674	3.1751	4.6565	C79	11.3181	7.8619	2.0732
H9	11.6268	2.9108	4.4049	C80	10.0252	7.2824	2.2457
C10	13.4978	5.7980	7.3229	C81	9.3241	6.8173	1.1361
C11	14.8609	6.2064	7.2189	H82	8.3434	6.3573	1.3002
C12	15.5597	6.5918	8.3566	C83	9.8165	6.9095	-0.1601
H13	16.6004	6.9137	8.2403	C84	11.0675	7.5176	-0.3295
C14	14.9898	6.5859	9.6282	H85	11.4980	7.5880	-1.3341
C15	13.6635	6.1531	9.7293	C86	11.8128	7.9788	0.7503
H16	13.1736	6.1591	10.7090	C87	9.4059	7.1851	3.5920
C17	12.9178	5.7755	8.6180	H88	8.4089	6.7248	3.5351
C18	15.5409	6.1925	5.8978	H89	9.3007	8.1728	4.0675
H19	16.5943	6.4936	5.9866	H90	10.0538	6.6071	4.2846
H20	15.0124	6.8520	5.1817	C91	9.0816	6.3160	-1.3238
H21	15.4970	5.1937	5.4351	H92	9.1743	6.9353	-2.2301
C22	15.7315	7.1064	10.8219	H93	8.0102	6.1996	-1.1072

H94 9.4668 5.3169 -1.5850
C95 13.1937 8.5092 0.4944
H96 13.6089 8.0667 -0.4215
H97 13.8544 8.2584 1.3346
H98 13.2188 9.6038 0.3749
98

D10B

N1 12.6375 5.3182 6.0470
N2 12.7849 5.0120 3.7684
B3 12.6168 6.0198 4.7894
C4 12.6205 3.8707 5.8027
H5 11.5882 3.4787 5.8688
H6 13.2272 3.3353 6.5444
C7 13.1563 3.7375 4.3912
H8 14.2565 3.6195 4.3816
H9 12.7322 2.8770 3.8588
C10 12.7603 5.7183 7.4045
C11 14.0422 6.0100 7.9052
C12 14.1904 6.3057 9.2561
H13 15.1855 6.5384 9.6403
C14 13.1018 6.3180 10.1262
C15 11.8464 6.0076 9.6137
H16 10.9826 6.0040 10.2818
C17 11.6559 5.6959 8.2673
C18 15.2343 5.9750 6.9974
H19 16.1449 6.2704 7.5307
H20 15.1077 6.6575 6.1465
H21 15.3939 4.9722 6.5758
C22 13.2725 6.7003 11.5661
H23 12.5443 6.1946 12.2101
H24 13.1308 7.7817 11.7041
H25 14.2770 6.4579 11.9321
C26 10.2881 5.3254 7.7773
H27 9.5115 5.8049 8.3843
H28 10.1210 4.2411 7.8353
H29 10.1386 5.6175 6.7324
C30 13.0040 5.0998 2.3703
C31 14.1075 5.8040 1.8587
C32 14.3224 5.8181 0.4809
H33 15.1794 6.3692 0.0878
C34 13.4767 5.1565 -0.4042
C35 12.4019 4.4482 0.1260
H36 11.7217 3.9248 -0.5500
C37 12.1537 4.3994 1.4950
C38 15.0457 6.5479 2.7606
H39 16.0009 6.7413 2.2609
H40 15.2473 6.0028 3.6923
H41 14.6224 7.5226 3.0499
C42 13.6875 5.2367 -1.8870
H43 13.4239 4.2963 -2.3846
H44 14.7285 5.4716 -2.1352
H45 13.0616 6.0219 -2.3346
C46 10.9843 3.6072 1.9997
H47 10.1506 3.6532 1.2891
H48 10.6284 3.9830 2.9647
H49 11.2422 2.5466 2.1282
N50 12.8402 8.8399 5.1498
N51 11.6397 8.1684 3.3038
B52 12.3588 7.6746 4.4553
C53 12.5955 10.0362 4.3353
H54 13.4947 10.2958 3.7453
H55 12.3480 10.9014 4.9638
C56 11.4592 9.6191 3.4248
H57 10.4728 9.8392 3.8754
H58 11.4921 10.1230 2.4516
C59 13.3761 9.0604 6.4447
C60 12.4915 9.1298 7.5360
C61 12.9937 9.4437 8.7947
H62 12.3056 9.4917 9.6408
C63 14.3489 9.6927 9.0014
C64 15.2038 9.6343 7.9053

H65 16.2683 9.8333 8.0464
C66 14.7400 9.3316 6.6252
C67 11.0238 8.9032 7.3354
H68 10.4830 8.9548 8.2871
H69 10.8283 7.9179 6.8921
H70 10.5891 9.6476 6.6529
C71 14.8801 9.9712 10.3758
H72 15.7565 10.6286 10.3469
H73 15.1897 9.0407 10.8729
H74 14.1231 10.4413 11.0143
C75 15.6950 9.3201 5.4695
H76 16.7179 9.1196 5.8083
H77 15.7088 10.2872 4.9484
H78 15.4234 8.5620 4.7268
C79 10.8178 7.5309 2.3406
C80 9.6879 6.7930 2.7337
C81 8.8644 6.2421 1.7525
H82 7.9888 5.6680 2.0628
C83 9.1281 6.3982 0.3952
C84 10.2443 7.1445 0.0285
H85 10.4765 7.2728 -1.0312
C86 11.0893 7.7200 0.9730
C87 9.3607 6.5741 4.1800
H88 8.3089 6.2967 4.3072
H89 9.5593 7.4650 4.7906
H90 9.9693 5.7582 4.6015
C91 8.2669 5.7440 -0.6435
H92 8.1796 6.3619 -1.5446
H93 7.2570 5.5476 -0.2671
H94 8.6906 4.7794 -0.9565
C95 12.2725 8.5165 0.5087
H96 12.6797 8.0985 -0.4197
H97 13.0714 8.5207 1.2568
H98 12.0012 9.5612 0.3016
98

D10C

N1 12.9305 5.4302 6.1921
N2 12.4048 4.9119 3.9670
B3 12.5771 6.0387 4.8857
C4 12.6520 4.0010 6.1078
H5 11.6101 3.7648 6.4239
H6 13.3291 3.4263 6.7620
C7 12.8060 3.6791 4.6326
H8 13.8604 3.4033 4.3900
H9 12.1772 2.8256 4.3206
C10 13.6434 5.9168 7.3150
C11 14.9133 6.5796 7.1145
C12 15.5996 7.0562 8.2408
H13 16.5588 7.5702 8.0812
C14 15.1261 6.9309 9.5361
C15 13.9113 6.2091 9.7345
H16 13.4489 6.1718 10.7272
C17 13.1736 5.7545 8.6188
C18 15.5140 6.6644 5.7575
H19 16.4995 7.1589 5.8013
H20 14.8752 7.2262 5.0318
H21 15.6522 5.6769 5.2753
C22 15.8986 7.4538 10.7024
H23 15.2877 8.1031 11.3536
H24 16.7581 8.0580 10.3700
H25 16.2884 6.6466 11.3778
C26 11.8184 5.1640 8.8832
H27 11.3511 5.7144 9.7182
H28 11.8541 4.0955 9.1671
H29 11.1697 5.2599 8.0057
C30 12.1807 4.8520 2.5530
C31 13.2920 4.9174 1.6312
C32 13.0986 4.6049 0.3012
H33 13.9468 4.7322 -0.3891
C34 11.8535 4.1842 -0.2295
C35 10.7975 3.9950 0.7226

H36	9.8019	3.6969	0.3632	C7	11.0754	8.2251	7.9733
C37	10.9552	4.3176	2.0553	C8	11.3275	8.2453	9.3453
C38	14.6387	5.3137	2.1402	H9	10.5027	8.0523	10.0363
H39	15.3426	5.4836	1.3089	C10	12.6036	8.4688	9.8531
H40	15.0882	4.5572	2.8155	C11	13.6451	8.6519	8.9434
H41	14.5898	6.2367	2.7445	H12	14.6605	8.8028	9.3187
C42	11.7011	3.7456	-1.6369	C13	13.4349	8.6533	7.5677
H43	11.7254	2.6311	-1.8336	C14	9.7207	7.8488	7.4539
H44	12.5013	4.1715	-2.2704	H15	9.0808	7.4725	8.2617
H45	10.7291	4.0705	-2.0820	H16	9.8299	7.0681	6.6871
C46	9.8161	4.1098	3.0109	H17	9.2014	8.6799	6.9608
H47	8.8663	3.9856	2.4673	C18	12.8559	8.5166	11.3322
H48	9.7181	4.9641	3.6955	H19	13.8252	8.0712	11.5905
H49	9.9602	3.2269	3.6637	H20	12.0805	7.9763	11.8893
N50	12.6504	8.8173	5.4829	H21	12.8682	9.5472	11.7155
N51	11.9988	8.3195	3.2912	C22	14.5696	8.8308	6.6105
B52	12.4075	7.6909	4.5488	H23	15.5348	8.8450	7.1324
C53	12.7244	10.0409	4.6931	H24	14.4646	9.7636	6.0405
H54	13.7638	10.2466	4.3496	H25	14.5290	8.0184	5.8634
H55	12.3950	10.9126	5.2822	C26	11.4158	7.1842	2.3478
C56	11.8560	9.7549	3.4830	C27	10.1649	6.6441	1.9989
H57	10.7933	10.0418	3.6711	C28	10.0214	5.9731	0.7836
H58	12.1824	10.3171	2.5908	H29	9.0450	5.5551	0.5249
C59	12.5749	8.9220	6.8919	C30	11.0911	5.7945	-0.0866
C60	11.4276	8.3733	7.5831	C31	12.3344	6.2993	0.2962
C61	11.3802	8.4862	8.9800	H32	13.1939	6.1577	-0.3643
H62	10.5150	8.0553	9.5039	C33	12.5184	6.9876	1.4909
C63	12.3691	9.0952	9.7329	C34	9.0242	6.7037	2.9703
C64	13.4581	9.7032	9.0401	H35	8.2166	6.0259	2.6669
H65	14.3092	10.1064	9.5994	H36	8.6031	7.7106	3.0842
C66	13.5628	9.5637	7.6386	H37	9.3809	6.4133	3.9691
C67	10.2864	7.8020	6.8230	C38	10.9321	5.0433	-1.3768
H68	9.4902	7.4714	7.5120	H39	11.4519	5.5438	-2.2046
H69	10.5796	6.9311	6.1855	H40	9.8762	4.9457	-1.6567
H70	9.8345	8.5150	6.1051	H41	11.3462	4.0272	-1.3134
C71	12.2695	9.1929	11.2200	C42	13.8636	7.5075	1.8855
H72	12.1659	10.2477	11.5883	H43	14.6418	7.1738	1.1875
H73	13.1663	8.7988	11.7299	H44	14.0793	7.1760	2.9169
H74	11.4096	8.6176	11.6002	H45	13.8694	8.6052	1.9200
C75	14.8220	10.0602	6.9874	H46	10.2185	9.5578	3.0611
H76	15.6704	9.8637	7.6656	H47	10.6028	10.2484	5.6486
H77	14.8079	11.1460	6.7765	94			
H78	15.0111	9.5349	6.0448	D11A			
C79	11.5496	7.7959	2.0373	N1	12.8371	5.3222	6.0605
C80	10.1931	7.4917	1.8220	N2	12.7911	4.9340	3.7969
C81	9.7143	7.1583	0.5595	B3	12.6560	6.0332	4.7911
H82	8.6809	6.8062	0.4499	C4	12.9933	3.9646	5.8219
C83	10.5864	7.1813	-0.5874	C5	12.9584	3.7261	4.4928
C84	11.8984	7.6107	-0.3764	C6	12.9446	5.7947	7.3948
H85	12.5762	7.6508	-1.2442	C7	14.2175	5.9434	7.9519
C86	12.4133	7.9294	0.8725	C8	14.3276	6.2500	9.3312
C87	9.2382	7.5339	2.9826	H9	15.3200	6.3511	9.7785
H88	8.3278	6.9539	2.7640	C10	13.1927	6.4420	10.1098
H89	8.9302	8.5642	3.2333	C11	11.9335	6.3083	9.5283
H90	9.7043	7.1226	3.8901	H12	11.0381	6.4341	10.1440
C91	10.0660	6.7946	-1.9172	C13	11.7846	5.9829	8.1615
H92	10.8636	6.7945	-2.6798	C14	15.4680	5.7458	7.1586
H93	9.2399	7.4500	-2.3072	H15	16.1600	6.5886	7.3330
H94	9.6186	5.7663	-1.9230	H16	15.2620	5.6819	6.0838
C95	13.7986	8.4646	1.0207	H17	16.0086	4.8298	7.4554
H96	14.3921	8.2614	0.1127	C18	13.3063	6.8380	11.5537
H97	14.3307	8.0214	1.8799	H19	13.1359	7.9194	11.6735
H98	13.8529	9.5656	1.2028	H20	14.3016	6.6122	11.9611
47				H21	12.5592	6.3250	12.1769
M11				C22	10.4114	5.7231	7.6237
N1	11.8911	8.4801	5.6873	H23	9.8145	6.6467	7.5370
N2	11.5721	7.9059	3.5446	H24	9.8474	5.0557	8.2990
B3	12.3582	7.4184	4.7317	H25	10.4472	5.2630	6.6310
C4	10.9893	9.3849	5.1128	C26	12.9341	4.9760	2.3951
C5	10.7971	9.0406	3.8224	C27	13.9433	5.7951	1.8101
C6	12.1302	8.4600	7.0731	C28	14.0090	5.8935	0.4243

H29	14.7699	6.5475	-0.0146	C4	12.9526	4.1790	5.9712
C30	13.1409	5.2042	-0.4196	C5	11.8718	3.9261	5.2102
C31	12.2018	4.3407	0.1749	C6	14.5274	5.9579	6.4381
H32	11.4941	3.8005	-0.4602	C7	15.6414	6.2971	5.6532
C33	12.0814	4.2277	1.5533	C8	16.8227	6.6692	6.2911
C34	14.9303	6.5244	2.6589	H9	17.6869	6.9379	5.6806
H35	15.7195	6.9749	2.0413	C10	16.9313	6.7098	7.6780
H36	15.4036	5.8624	3.4008	C11	15.8203	6.3435	8.4311
H37	14.4464	7.3191	3.2572	H12	15.8828	6.3691	9.5213
C38	13.2034	5.3516	-1.9091	C13	14.6181	5.9609	7.8395
H39	13.5412	4.4323	-2.4216	C14	15.5838	6.2620	4.1551
H40	13.8916	6.1574	-2.1990	H15	16.5917	6.2783	3.7268
H41	12.2170	5.5970	-2.3329	H16	15.0372	7.1277	3.7508
C42	10.9735	3.3908	2.1218	H17	15.0655	5.3649	3.7906
H43	10.2653	3.1087	1.3316	C18	18.1935	7.1776	8.3392
H44	10.4217	3.9624	2.8812	H19	18.1581	8.2578	8.5385
H45	11.3273	2.4735	2.6139	H20	19.0713	6.9953	7.7092
N46	12.6506	8.8410	5.2539	H21	18.3562	6.6795	9.3016
N47	11.6413	8.2221	3.2850	C22	13.4596	5.5700	8.7097
B52	12.3067	7.6413	4.4866	H23	12.5025	5.8530	8.2609
C49	12.2925	9.9743	4.5425	H24	13.5332	6.0598	9.6878
C50	11.7060	9.6219	3.3792	H25	13.4270	4.4868	8.8861
C51	13.1932	9.0055	6.5542	C26	10.5520	4.9363	3.4711
C52	12.3195	9.2608	7.6131	C27	11.0199	4.5778	2.1976
C53	12.8628	9.5995	8.8763	C28	10.0901	4.3310	1.1927
H54	12.1866	9.8197	9.7069	H29	10.4470	4.0580	0.1979
C55	14.2384	9.6323	9.0739	C30	8.7194	4.4244	1.4265
C56	15.0909	9.3575	8.0073	C31	8.2851	4.7590	2.7069
H57	16.1741	9.4115	8.1535	H32	7.2143	4.8144	2.9104
C58	14.5861	9.0417	6.7255	C33	9.1824	5.0121	3.7444
C59	10.8338	9.2391	7.4549	C34	12.4892	4.4305	1.9458
H60	10.3807	8.6296	8.2570	H35	12.6903	4.2099	0.8921
H61	10.5323	8.8204	6.4876	H36	12.9224	3.6292	2.5600
H62	10.3928	10.2481	7.5422	H37	13.0269	5.3509	2.2083
C63	14.8091	9.9128	10.4336	C38	7.7395	4.1989	0.3142
H64	15.0910	8.9748	10.9363	H39	6.7762	3.8355	0.6888
H65	14.0858	10.4300	11.0794	H40	8.1174	3.4722	-0.4145
H66	15.7169	10.5300	10.3751	H41	7.5444	5.1320	-0.2330
C67	15.5428	8.9034	5.5813	C42	8.6909	5.3449	5.1210
H68	16.1142	7.9607	5.6206	H43	7.6111	5.1833	5.2015
H69	16.2899	9.7163	5.6007	H44	8.8907	6.3944	5.3787
H70	15.0265	8.9322	4.6157	H45	9.1886	4.7386	5.8877
C71	10.8843	7.6233	2.2598	N46	12.7751	8.8840	5.2825
C72	9.8019	6.7566	2.5892	N47	12.2431	8.4285	3.1194
C73	9.1121	6.1199	1.5649	B52	12.4668	7.7748	4.3959
H74	8.2994	5.4367	1.8339	C49	12.7092	10.0802	4.5643
C75	9.4197	6.3105	0.2177	C50	12.3941	9.8096	3.2831
C76	10.4379	7.2311	-0.0897	C51	13.0322	8.9340	6.6795
H77	10.7091	7.3990	-1.1362	C52	12.0178	8.6031	7.5893
C78	11.1710	7.8742	0.8974	C53	12.2794	8.7306	8.9528
C79	9.3864	6.5474	4.0078	H54	11.4917	8.4765	9.6647
H80	8.4496	5.9753	4.0590	C55	13.5111	9.1682	9.4304
H81	9.2410	7.5016	4.5375	C56	14.4959	9.4968	8.5030
H82	10.1627	6.0171	4.5899	H57	15.4727	9.8323	8.8576
C83	8.6628	5.6109	-0.8694	C58	14.2797	9.3936	7.1318
H84	7.9117	6.2542	-1.3648	C59	10.6763	8.1235	7.1226
H85	8.1268	4.7348	-0.4793	H60	9.9349	8.1965	7.9252
H86	9.3359	5.2547	-1.6637	H61	10.7160	7.0708	6.8016
C87	12.3223	8.7477	0.4931	H62	10.3161	8.7024	6.2616
H88	12.5659	8.5894	-0.5657	C63	13.7903	9.2457	10.9019
H89	13.2131	8.4963	1.0856	H64	14.3096	8.3439	11.2558
H90	12.1347	9.8200	0.6479	H65	12.8669	9.3380	11.4841
H91	11.2511	10.2546	2.6276	H66	14.4323	10.0999	11.1463
H92	12.4895	10.9617	4.9454	C67	15.3730	9.7601	6.1721
H93	13.1191	3.2702	6.6460	H68	15.3973	9.0804	5.3129
H94	13.1024	2.7920	3.9634	H69	16.3501	9.7160	6.6668
94				H70	15.2483	10.7745	5.7733
D11B				C71	11.6858	7.9779	1.8862
N1	13.3326	5.5129	5.8078	C72	10.2882	7.9548	1.7597
N2	11.5058	5.0864	4.5198	C73	9.7348	7.5875	0.5376
B3	12.4281	6.1515	4.8626	H74	8.6483	7.5558	0.4381

C75	10.5326	7.2559	-0.5553
C76	11.9164	7.3196	-0.4099
H77	12.5543	7.0873	-1.2646
C78	12.5136	7.6850	0.7970
C79	9.4191	8.3424	2.9170
H80	8.3612	8.1682	2.6934
H81	9.5504	9.4008	3.1803
H82	9.6780	7.7628	3.8127
C83	9.9126	6.8069	-1.8447
H84	8.9390	7.2823	-2.0117
H85	9.7438	5.7207	-1.8409
H86	10.5534	7.0319	-2.7044
C87	14.0057	7.7687	0.9155
H88	14.4774	7.7492	-0.0724
H89	14.4142	6.9272	1.4929
H90	14.3222	8.6830	1.4318
H91	12.2503	10.4849	2.4481
H92	12.8870	11.0339	5.0434
H93	13.5086	3.5125	6.6178
H94	11.3177	3.0043	5.0797
94			
D11C			
N1	12.9749	5.3936	6.0633
N2	12.7481	4.9663	3.8148
B3	12.6720	6.0592	4.7997
C4	13.1916	4.0454	5.8195
C5	13.0639	3.7901	4.4970
C6	13.1247	5.8496	7.4171
C7	14.4223	6.0007	7.9205
C8	14.5824	6.2656	9.3299
H9	15.5853	6.4407	9.7355
C10	13.4659	6.4046	10.1445
C11	12.1814	6.2712	9.6239
H12	11.3069	6.4092	10.2713
C13	11.9865	6.0064	8.2199
C14	15.6527	5.8519	7.1006
H15	16.3263	6.7270	7.2496
H16	15.4332	5.7784	6.0278
H17	16.2666	4.9646	7.3884
C18	13.6310	6.7632	11.5955
H19	13.3060	7.8056	11.7611
H20	14.6759	6.6646	11.9305
H21	13.0019	6.1279	12.2463
C22	10.6095	5.6817	7.7517
H23	9.8947	6.4998	7.9831
H24	10.1894	4.7826	8.2624
H25	10.5700	5.5046	6.6696
C26	12.7123	4.9323	2.3830
C27	13.6931	5.6918	1.6607
C28	13.6775	5.5965	0.2519
H29	14.4298	6.1612	-0.3165
C30	12.7542	4.8389	-0.4401
C31	11.8331	4.0226	0.2847
H32	10.9368	3.6387	-0.2222
C33	11.7954	4.1394	1.7096
C34	14.7721	6.4361	2.3734
H35	15.4613	6.8973	1.6456
H36	15.3812	5.8009	3.0487
H37	14.3929	7.2483	3.0271
C38	12.7841	4.7754	-1.9336
H39	13.0711	3.7623	-2.3159
H40	13.4814	5.5179	-2.3543
H41	11.7922	4.9783	-2.3719
C42	10.6846	3.4217	2.4111
H43	9.7328	3.6978	1.9173
H44	10.6191	3.6708	3.4760
H45	10.7771	2.3209	2.3130
N46	12.5181	8.8373	5.3588
N47	11.6663	8.2463	3.3065
B52	12.2893	7.6716	4.5113
C49	12.0752	9.9711	4.6933
C50	11.5688	9.6277	3.4861
C51	13.0420	9.0101	6.6847
C52	12.1342	9.2308	7.7260
C53	12.6537	9.6062	9.0187
H54	11.9648	9.7301	9.8613
C55	14.0262	9.6992	9.2174
C56	14.9167	9.4515	8.1769
H57	15.9979	9.5070	8.3528
C58	14.4317	9.0716	6.8725
C59	10.6583	9.1459	7.5721
H60	10.2248	8.4892	8.3614
H61	10.3620	8.7375	6.5974
H62	10.1473	10.1297	7.7045
C63	14.5672	10.0113	10.5853
H64	15.0158	9.1044	11.0261
H65	13.7830	10.3801	11.2656
H66	15.3676	10.7728	10.5420
C67	15.4072	9.0241	5.7473
H68	16.2327	8.3093	5.9501
H69	15.9130	10.0055	5.5823
H70	14.9359	8.7246	4.8027
C71	11.0597	7.6646	2.1472
C72	9.9463	6.7761	2.3337
C73	9.3186	6.2707	1.1745
H74	8.4488	5.6106	1.3001
C75	9.7526	6.5593	-0.1044
C76	10.8199	7.4912	-0.2828
H77	11.3364	7.5501	-1.2506
C78	11.4931	7.9943	0.8719
C79	9.3781	6.5250	3.6905
H80	8.4933	5.8700	3.6173
H81	9.0632	7.4480	4.2194
H82	10.0914	6.0381	4.3865
C83	9.0320	6.0159	-1.2964
H84	8.4889	6.8096	-1.8720
H85	8.3075	5.2376	-1.0074
H86	9.7246	5.5588	-2.0228
C87	12.7134	8.8273	0.6288
H88	13.3700	8.2765	-0.0720
H89	13.2727	9.0434	1.5458
H90	12.4669	9.7886	0.1343
H91	11.1175	10.2491	2.7192
H92	12.1567	10.9466	5.1638
H93	13.4338	3.3761	6.6391
H94	13.1864	2.8591	3.9538
17			
M12			
N1	11.6931	8.4265	5.7343
N2	11.6413	7.8171	3.5578
B3	11.6371	7.1682	4.9141
C4	11.9170	9.6404	4.9812
H5	12.9884	9.9523	5.0077
H6	11.3452	10.5117	5.3649
C7	11.4990	9.2562	3.5668
H8	10.4466	9.5689	3.3657
H9	12.1128	9.7669	2.7950
C10	11.2806	7.1699	2.3461
H11	11.3350	6.0863	2.5162
H12	10.2459	7.4084	2.0047
H13	11.9436	7.4323	1.4923
C14	12.0314	8.4842	7.1125
H15	11.9109	7.4751	7.5287
H16	13.0828	8.8057	7.3000
H17	11.3966	9.1897	7.6928
34			
D12A			
N1	12.6111	5.4193	6.2586
N2	12.3769	4.8588	3.8956
B3	12.4917	6.0901	4.8519
C4	12.2194	4.0544	6.0627
H5	11.0986	3.9393	6.0398

H6	12.5900	3.3616	6.8534	H7	13.6324	10.5529	5.4811
C7	12.7603	3.7253	4.6844	H8	13.4147	10.1577	7.2871
H8	13.8803	3.6249	4.7578	C9	11.6370	9.8585	5.9549
H9	12.3840	2.7530	4.2896	H10	11.0294	10.2550	6.8416
N10	12.9372	8.8529	5.3519	H11	11.4666	10.5539	5.0426
N11	12.0306	8.3174	3.1510	C12	14.9457	8.4949	5.5646
B12	12.4863	7.6351	4.4816	C32	9.9478	8.4362	5.1330
C13	13.0474	9.9529	4.4395	N52	13.7411	5.2657	4.3695
H14	14.0204	9.9304	3.8713	N53	12.2471	6.0211	2.7258
H15	12.9940	10.9524	4.9302	B52	12.6461	6.3849	4.1701
C16	11.9163	9.7128	3.4580	C55	13.2768	4.1048	3.6463
H17	10.9432	9.9502	3.9746	H56	12.7117	3.3874	4.3502
H18	11.9663	10.3764	2.5638	H57	14.1471	3.5255	3.1730
C19	14.1474	8.6648	6.0851	C58	12.2587	4.5952	2.6112
H20	14.4141	9.5633	6.6935	H59	12.5107	4.2729	1.5410
H21	14.0564	7.8082	6.7671	H60	11.2278	4.1525	2.9034
H22	15.0156	8.4236	5.4134	C61	14.1302	4.8712	5.6746
C23	11.8447	6.0341	7.2945	C81	11.1762	6.5984	2.0050
H24	11.9743	5.5165	8.2762	H24	15.3049	7.4883	5.2908
H25	12.1415	7.0841	7.4227	H25	15.1074	9.1762	4.6497
H26	10.7479	6.0593	7.0502	H26	15.6161	8.8889	6.3936
C27	13.1485	4.9425	2.6974	H27	9.2015	8.8235	5.9067
H28	13.0190	4.0406	2.0506	H28	9.8057	9.0590	4.1752
H29	12.8567	5.8253	2.1121	H29	9.7118	7.3799	4.8617
H30	14.2452	5.0703	2.9084	H31	14.4279	5.7399	6.2863
C31	10.8206	7.8100	2.5884	H32	13.2764	4.3363	6.2338
H32	10.5481	8.3317	1.6387	H33	15.0088	4.1507	5.6447
H33	10.9164	6.7371	2.3714	H34	11.2671	6.3487	0.8937
H34	9.9547	7.8971	3.2999	H35	10.1484	6.2210	2.3602
34				H36	11.1764	7.7042	2.1550
D12B				15			
N1	12.4629	5.4960	5.9313	M13			
N2	12.6922	5.0421	3.6891	N1	11.8204	8.3958	5.7195
B3	12.4861	6.0958	4.6337	N2	11.4971	7.8130	3.5796
C4	12.4842	4.0462	5.8134	B3	12.2606	7.2947	4.7792
H5	11.4700	3.6342	5.9726	C4	10.9772	9.3315	5.1282
H6	13.1374	3.5804	6.5674	C5	10.7817	8.9790	3.8340
C7	12.9698	3.7945	4.3842	H6	10.1746	9.4870	3.0836
H8	14.0539	3.5757	4.3601	H7	10.5645	10.1894	5.6607
H9	12.4631	2.9379	3.9131	C8	12.2021	8.5163	7.0881
N10	12.9472	8.8293	4.9386	H9	12.8713	7.6730	7.3072
N11	11.4735	8.2958	3.2566	H10	12.7421	9.4579	7.3126
B12	12.3030	7.7368	4.2781	H11	11.3455	8.4724	7.7903
C13	12.6680	10.0729	4.2373	C12	11.4759	7.2084	2.2884
H14	13.5381	10.3688	3.6214	H13	12.1201	6.3201	2.3424
H15	12.4748	10.9044	4.9328	H14	10.4645	6.8859	1.9691
C16	11.4535	9.7464	3.3654	H15	11.8603	7.8702	1.4867
H17	10.5134	10.0818	3.8425	30			
H18	11.5004	10.2349	2.3798	D13A			
C19	10.4070	7.6792	2.5219	N1	13.0449	5.3781	6.0394
H20	10.4493	7.9300	1.4498	N2	13.6864	5.0043	3.7417
H21	10.4699	6.5899	2.6210	B3	13.0607	6.1132	4.6428
H22	9.4150	7.9957	2.8893	C4	13.3517	4.0424	5.7803
C23	14.0864	8.8120	5.8096	C5	13.7008	3.8326	4.4930
H24	13.9473	9.4737	6.6795	N6	12.1965	8.5982	5.6134
H25	14.2551	7.7942	6.1778	N7	12.1711	8.4720	3.1928
H26	15.0060	9.1427	5.2952	B8	12.5352	7.6011	4.4250
C27	11.9955	6.0450	7.1711	C9	11.6195	9.7333	5.0078
H28	12.6781	5.8146	8.0047	C10	11.5786	9.6283	3.6602
H29	11.9150	7.1343	7.0862	H11	11.0987	10.3229	2.9608
H30	11.0014	5.6528	7.4489	H12	11.2824	10.5907	5.6008
C31	13.0938	5.1283	2.3147	H13	13.3830	3.3029	6.5903
H32	12.5143	4.4437	1.6749	H14	14.0939	2.9026	4.0636
H33	12.9427	6.1496	1.9476	C15	13.2998	8.9059	6.4802
H34	14.1597	4.8742	2.1786	H16	14.1854	9.3324	5.9406
34				H17	13.0116	9.6334	7.2785
D12C				H18	13.6538	7.9856	6.9675
N3	13.5867	8.4674	5.9686	C23	13.3442	4.8948	2.3680
N4	11.2649	8.5132	5.6417	H24	13.8306	4.0177	1.8888
B3	12.4956	7.8072	5.0393	H25	12.2339	4.8320	2.1949
C6	13.1449	9.8186	6.2245	H26	13.6776	5.7965	1.8224

C27	11.8815	5.5816	6.8525	H30	14.0069	9.2121	2.5667
H28	11.9375	5.0055	7.8044	50			
H29	11.7729	6.6509	7.0891	M16			
H30	10.9409	5.2970	6.3196	N1	11.7505	8.5487	5.6132
H27	11.2608	8.6935	1.3148	N2	11.5820	8.0054	3.5753
H31	12.6412	7.5719	1.3500	C3	12.0782	7.5259	4.7476
H32	11.1141	7.0390	2.0767	C4	11.0351	9.6728	4.9981
C30	11.7766	7.9375	1.9410	C5	10.9203	9.2583	3.6767
30				C6	12.0673	8.4934	6.9884
D13B				C7	11.0359	8.5186	7.9419
N2	13.9238	5.7005	6.1333	C8	11.3672	8.4543	9.2937
N3	13.2658	4.8511	4.1367	H9	10.5623	8.4677	10.0328
B3	13.1587	6.0319	4.9570	C10	12.6873	8.3576	9.7257
C4	14.2128	6.5038	7.2940	C11	13.6902	8.3190	8.7612
C24	12.6728	4.5727	2.8533	H12	14.7330	8.2368	9.0767
N44	11.3585	8.1260	5.4495	C13	13.4048	8.3913	7.3982
N45	12.4330	8.2946	3.4604	C14	9.6050	8.5897	7.5118
B52	12.3347	7.4545	4.6277	H15	8.9281	8.4216	8.3581
C47	10.8128	7.7361	6.7248	H16	9.3961	7.8419	6.7346
C67	13.2807	8.1569	2.3036	H17	9.4120	9.5702	7.0561
C91	11.5701	9.3751	3.6064	C18	13.0154	8.3172	11.1903
C92	10.9278	9.2738	4.7933	H19	13.9994	7.8671	11.3693
C99	14.4346	4.4148	5.9929	H20	12.2734	7.7366	11.7534
C100	14.0409	3.9073	4.8017	H21	13.0343	9.3232	11.6328
H93	11.4844	10.1432	2.8467	C22	14.5133	8.3527	6.3908
H94	10.1868	9.9350	5.2269	H23	14.4252	7.4585	5.7626
H95	15.0493	3.9586	6.7601	H24	15.4933	8.3682	6.8835
H96	14.2508	2.9355	4.3704	H25	14.4482	9.2074	5.7050
H19	15.2944	6.5849	7.4672	C26	11.5191	7.1976	2.4166
H20	13.8182	7.5135	7.1411	C27	10.3498	6.4755	2.1541
H21	13.7560	6.0903	8.2047	C28	10.2842	5.7057	0.9936
H22	13.4319	4.4545	2.0667	H29	9.3759	5.1359	0.7851
H23	12.0703	3.6547	2.8798	C30	11.3545	5.6359	0.1044
H24	12.0149	5.4014	2.5730	C31	12.5159	6.3472	0.4041
H25	11.2016	6.7495	6.9965	H32	13.3686	6.2907	-0.2765
H26	11.0833	8.4439	7.5215	C33	12.6151	7.1327	1.5505
H27	9.7169	7.6716	6.6898	C34	9.2361	6.5070	3.1532
H28	13.9545	7.3067	2.4498	H35	8.3907	5.8849	2.8376
H29	12.7000	7.9817	1.3866	H36	8.8893	7.5333	3.3323
H30	13.8955	9.0535	2.1465	H37	9.6074	6.1505	4.1241
30				C38	11.2596	4.8272	-1.1571
D13C				H39	11.0230	5.4565	-2.0265
N2	13.9778	5.7216	6.1638	H40	10.4748	4.0648	-1.0869
N3	13.4802	4.8763	4.0927	H41	12.2051	4.3176	-1.3810
B3	13.2313	6.0267	4.9641	C42	13.8636	7.8875	1.8877
C4	14.3994	6.6433	7.1393	H43	14.6371	7.7451	1.1241
C24	12.6992	4.5059	2.9895	H44	14.2494	7.5620	2.8615
N44	11.2880	8.0988	5.4666	H45	13.6583	8.9611	1.9896
N45	12.2881	8.2999	3.4201	O46	10.6821	10.6937	5.6253
B52	12.2815	7.4698	4.6138	C47	10.3871	10.0231	2.5286
C47	10.6662	7.5305	6.5877	H48	11.1389	10.2510	1.7506
C67	13.2777	8.3476	2.4300	H49	9.5429	9.5391	2.0027
C91	11.4117	9.5712	3.5682	H50	10.0224	10.9793	2.9257
C92	10.5305	9.2511	4.7635	100			
C99	14.9298	4.5266	5.8965	D16 dimer			
C100	14.1954	3.7317	4.7943	N1	12.9962	5.5564	6.0029
H93	12.2299	10.3098	3.9365	N2	12.0746	5.0963	3.9808
H94	9.6169	8.6548	4.3661	C3	12.5821	6.1765	4.7845
H95	14.8915	3.8935	6.8529	C4	12.5491	4.1832	6.0268
H96	13.2554	3.1730	5.2638	C5	12.0655	3.8501	4.7780
H19	15.5272	6.7574	7.0811	C6	13.6515	6.0296	7.1489
H20	13.9069	7.6279	7.0404	C7	14.9864	6.4996	7.0906
H21	14.2231	6.3002	8.2034	C8	15.6546	6.8255	8.2624
H22	13.3064	3.9618	2.2137	H9	16.6828	7.1928	8.1917
H23	11.8864	3.7516	3.2526	C10	15.0663	6.6857	9.5189
H24	12.1862	5.3740	2.5336	C11	13.7709	6.1788	9.5676
H25	11.2577	6.6987	7.0189	H12	13.2763	6.0673	10.5376
H26	10.4532	8.2976	7.3852	C13	13.0466	5.8601	8.4225
H27	9.6330	7.1105	6.3618	C14	15.7111	6.5629	5.7867
H28	13.8911	7.4268	2.3918	H15	16.7167	6.9834	5.9187
H29	12.8440	8.5506	1.4096	H16	15.1748	7.1717	5.0463

H17	15.8071	5.5573	5.3523	H88	13.5586	8.6081	-0.2617
C18	15.7778	7.1351	10.7592	H89	14.0129	7.4156	0.9667
H19	15.4127	6.6081	11.6513	H90	13.6272	9.0547	1.4864
H20	15.6292	8.2124	10.9394	O91	13.2020	10.9598	5.0618
H21	16.8625	6.9713	10.6905	O92	11.7732	2.7545	4.2671
C22	11.6159	5.4426	8.5664	C93	11.6794	10.5335	2.4978
H23	10.9589	6.3005	8.3659	H94	10.5818	10.4119	2.5924
H24	11.4117	5.1031	9.5905	H95	11.9504	11.5207	2.8928
H25	11.3424	4.6620	7.8411	H96	11.8900	10.5249	1.4119
C26	12.2726	4.8621	2.5947	C97	13.2907	3.1803	6.8334
C27	13.5200	5.0641	1.9710	H98	14.3887	3.3012	6.7410
C28	13.6640	4.8019	0.6089	H99	13.0780	3.1885	7.9189
H29	14.6324	4.9982	0.1392	H100	13.0199	2.1934	6.4372
C30	12.6185	4.3088	-0.1602	52			
C31	11.4206	4.0261	0.4912	M17			
H32	10.5853	3.6146	-0.0833	N1	11.5913	8.6073	5.7087
C33	11.2268	4.2770	1.8459	N2	11.4789	8.0534	3.6551
C34	14.7066	5.5142	2.7562	C3	11.6104	7.5125	4.9185
H35	15.6176	5.4680	2.1449	C4	11.4967	9.8451	4.9954
H36	14.8459	4.8789	3.6409	C5	11.4142	9.4913	3.6199
H37	14.5878	6.5409	3.1297	C6	11.9324	8.4660	7.0853
C38	12.7437	4.1477	-1.6456	C7	10.9361	8.4585	8.0643
H39	12.1828	3.2762	-2.0110	C8	11.3085	8.2465	9.3883
H40	13.7910	4.0300	-1.9541	H9	10.5352	8.2370	10.1597
H41	12.3462	5.0282	-2.1740	C10	12.6402	8.0538	9.7537
C42	9.9010	3.9205	2.4471	C11	13.6093	8.0647	8.7554
H43	9.1059	3.9852	1.6910	H12	14.6566	7.9059	9.0216
H44	9.6520	4.5808	3.2820	C13	13.2744	8.2677	7.4171
H45	9.9485	2.9094	2.8638	C14	9.5143	8.7063	7.6783
N46	12.8997	8.6187	5.3490	H15	8.8322	8.5161	8.5153
N47	11.9757	8.1580	3.3278	H16	9.2216	8.0786	6.8270
C48	12.3913	7.5379	4.5460	H17	9.4160	9.7540	7.3645
C49	12.9082	9.8643	4.5515	C18	13.0180	7.8653	11.1943
C50	12.4229	9.5311	3.3033	H19	13.9861	7.3597	11.2926
C51	12.7038	8.8534	6.7354	H20	12.2704	7.2692	11.7325
C52	11.4563	8.6525	7.3600	H21	13.0992	8.8266	11.7203
C53	11.3144	8.9128	8.7229	C22	14.3175	8.2466	6.3420
H54	10.3462	8.7174	9.1936	H23	14.1582	7.3869	5.6776
C55	12.3621	9.4032	9.4912	H24	15.3265	8.1893	6.7665
C56	13.5592	9.6856	8.8384	H25	14.2493	9.1426	5.7109
H57	14.3957	10.0949	9.4125	C26	11.4321	7.2424	2.4949
C58	13.7510	9.4365	7.4833	C27	10.3752	6.3367	2.3309
C59	10.2683	8.2051	6.5752	C28	10.3570	5.5243	1.1984
H60	9.3578	8.2523	7.1873	H29	9.5367	4.8140	1.0732
H61	10.3853	7.1784	6.2009	C30	11.3506	5.5978	0.2261
H62	10.1295	8.8412	5.6909	C31	12.3908	6.5035	0.4164
C63	12.2402	9.5607	10.9773	H32	13.1903	6.5631	-0.3256
H64	12.8050	10.4292	11.3439	C33	12.4544	7.3270	1.5388
H65	12.6357	8.6773	11.5022	C34	9.2893	6.2368	3.3583
H66	11.1939	9.6813	11.2881	H35	8.5244	5.5128	3.0543
C67	15.0760	9.7921	6.8804	H36	8.8074	7.2105	3.5156
H68	15.8724	9.7258	7.6350	H37	9.7086	5.9459	4.3292
H69	15.0291	10.8038	6.4650	C38	11.2936	4.7417	-1.0058
H70	15.3229	9.1327	6.0443	H39	12.2951	4.4243	-1.3222
C71	11.3204	7.6837	2.1821	H40	10.8462	5.2792	-1.8534
C72	9.9861	7.2117	2.2413	H41	10.6915	3.8402	-0.8408
C73	9.3179	6.8836	1.0698	C42	13.6040	8.2653	1.7326
H74	8.2902	6.5149	1.1411	H43	14.4003	8.0698	1.0051
C75	9.9061	7.0229	-0.1870	H44	14.0190	8.1688	2.7447
C76	11.2007	7.5318	-0.2361	H45	13.2608	9.3024	1.6325
H77	11.6950	7.6427	-1.2063	O46	11.3023	10.1315	2.5666
C78	11.9246	7.8529	0.9083	C47	11.3590	11.1158	5.6150
C79	9.2625	7.1476	3.5459	C48	11.3698	12.3243	4.6972
H80	8.2571	6.7264	3.4148	O49	11.2475	11.2927	6.8392
H81	9.1664	8.1529	3.9810	H50	10.5779	13.0117	5.0162
H82	9.8001	6.5387	4.2854	H51	12.3229	12.8527	4.8297
C83	9.1964	6.5692	-1.4267	H52	11.2551	12.0547	3.6432
H84	9.5647	7.0912	-2.3204	104			
H85	8.1117	6.7353	-1.3615	D17 dimer			
H86	9.3436	5.4908	-1.6014	N1	12.3383	5.6180	6.1020
C87	13.3546	8.2717	0.7634	N2	11.8149	5.2499	3.9217

C3	12.0235	6.2752	4.8859	H74	7.2940	6.9767	2.0223
C4	12.2758	4.1840	5.9340	C75	8.5801	7.5868	0.4179
C5	11.9512	3.9562	4.5849	C76	9.8887	7.9385	0.1087
C6	12.3047	6.0326	7.4621	H77	10.1703	8.0885	-0.9371
C7	13.4617	5.8481	8.2363	C78	10.8788	8.0691	1.0787
C8	13.4059	6.0947	9.6057	C79	8.8219	7.0269	4.1404
H9	14.3167	5.9729	10.1973	H80	7.8129	6.5973	4.1079
C10	12.2324	6.4952	10.2346	H81	8.8043	7.8551	4.8557
C11	11.0813	6.6033	9.4616	H82	9.5026	6.2668	4.5397
H12	10.1369	6.8818	9.9368	C83	7.5518	7.3841	-0.6557
C13	11.0923	6.3580	8.0913	H84	7.6975	8.0793	-1.4927
C14	14.7349	5.3508	7.6270	H85	6.5336	7.5269	-0.2715
H15	15.5814	5.5091	8.3069	H86	7.5984	6.3665	-1.0718
H16	14.9532	5.8559	6.6800	C87	12.2746	8.3341	0.5980
H17	14.6403	4.2801	7.4092	H88	12.4424	7.8086	-0.3521
C18	12.2154	6.8383	11.6943	H89	13.0184	7.9755	1.3172
H19	12.9534	6.2537	12.2584	H90	12.4443	9.4063	0.4650
H20	11.2273	6.6616	12.1394	O91	11.7736	10.3434	2.4163
H21	12.4530	7.9014	11.8516	O92	12.4498	3.4235	6.8928
C22	9.8160	6.4021	7.3195	C93	13.6613	10.8823	4.7580
H23	8.9554	6.4953	7.9929	O94	14.3317	11.0804	5.7936
H24	9.6917	5.4960	6.7109	C95	11.8293	2.6786	3.9836
H25	9.8091	7.2551	6.6316	O96	11.5555	2.4384	2.7926
C26	12.4978	5.3241	2.6561	C97	13.6851	11.9749	3.6918
C27	13.8765	5.6032	2.5774	H98	14.4291	12.7156	4.0065
C28	14.5110	5.5503	1.3366	H99	13.9240	11.5867	2.6960
H29	15.5807	5.7690	1.2857	H100	12.7025	12.4515	3.5870
C30	13.8258	5.2208	0.1718	C101	12.0897	1.4804	4.8948
C31	12.4706	4.9391	0.2744	H102	12.0417	0.5857	4.2638
H32	11.9055	4.6762	-0.6240	H103	13.0594	1.5432	5.4001
C33	11.7912	4.9762	1.4924	H104	11.3503	1.4135	5.7012
C34	14.6999	5.9282	3.7883				
H35	15.7679	5.9032	3.5387				
H36	14.5203	5.2028	4.5917				
H37	14.4644	6.9230	4.1932				
C38	14.5180	5.2196	-1.1597				
H39	14.1015	4.4567	-1.8302				
H40	15.5935	5.0269	-1.0550				
H41	14.4116	6.1888	-1.6688				
C42	10.3387	4.6125	1.4824				
H43	9.8044	5.2091	0.7303				
H44	9.8755	4.7684	2.4586				
H45	10.2423	3.5484	1.2490				
N46	12.6357	8.6481	5.4131				
N47	11.5307	8.2295	3.4049				
C48	11.9310	7.6299	4.6595				
C49	12.8728	9.7454	4.4780				
C50	12.0681	9.5673	3.3316				
C51	12.1586	9.1343	6.6868				
C52	10.8971	9.7315	6.8592				
C53	10.4980	10.1214	8.1395				
H54	9.4975	10.5439	8.2706				
C55	11.3419	10.0103	9.2388				
C56	12.6334	9.5412	9.0222				
H57	13.3336	9.4874	9.8602				
C58	13.0616	9.1120	7.7682				
C59	9.9908	10.0679	5.7114				
H60	8.9461	10.1325	6.0460				
H61	10.0486	9.3576	4.8866				
H62	10.2715	11.0398	5.2836				
C63	10.8629	10.3462	10.6203				
H64	11.6809	10.7171	11.2516				
H65	10.4462	9.4603	11.1252				
H66	10.0738	11.1094	10.6033				
C67	14.4896	8.7145	7.5778				
H68	14.9376	8.3650	8.5171				
H69	15.0237	9.5915	7.1927				
H70	14.5845	7.9430	6.8102				
C71	10.5498	7.8807	2.4456				
C72	9.2450	7.4400	2.7647				
C73	8.2958	7.3188	1.7501				