

-Supplementary information-

Incorporation of *N*7-Platinated Guanine into *Thermus Aquaticus* DNA Polymerase (Taq DNA Polymerase): Atomistic Insights from Molecular Dynamics Simulations

Federica De Castro ¹, Giada Ciardullo ², Francesco Paolo Fanizzi ¹, Mario Prejanò ², Michele Benedetti ^{1,*} and Tiziana Marino ^{2,*}

¹ Dipartimento di Scienze e Tecnologie Biologiche ed Ambientali, Università del Salento, Prov.le Lecce-Monteroni, Centro Ecotekne, I-73100 Lecce, Italy

² Dipartimento di Chimica e Tecnologie Chimiche, Laboratorio PROMOCS cubo 14C, Università della Calabria, I-87036 RENDE (CS), Italy

* Correspondence: michele.benedetti@unisalento.it; tiziana.marino65@unical.it

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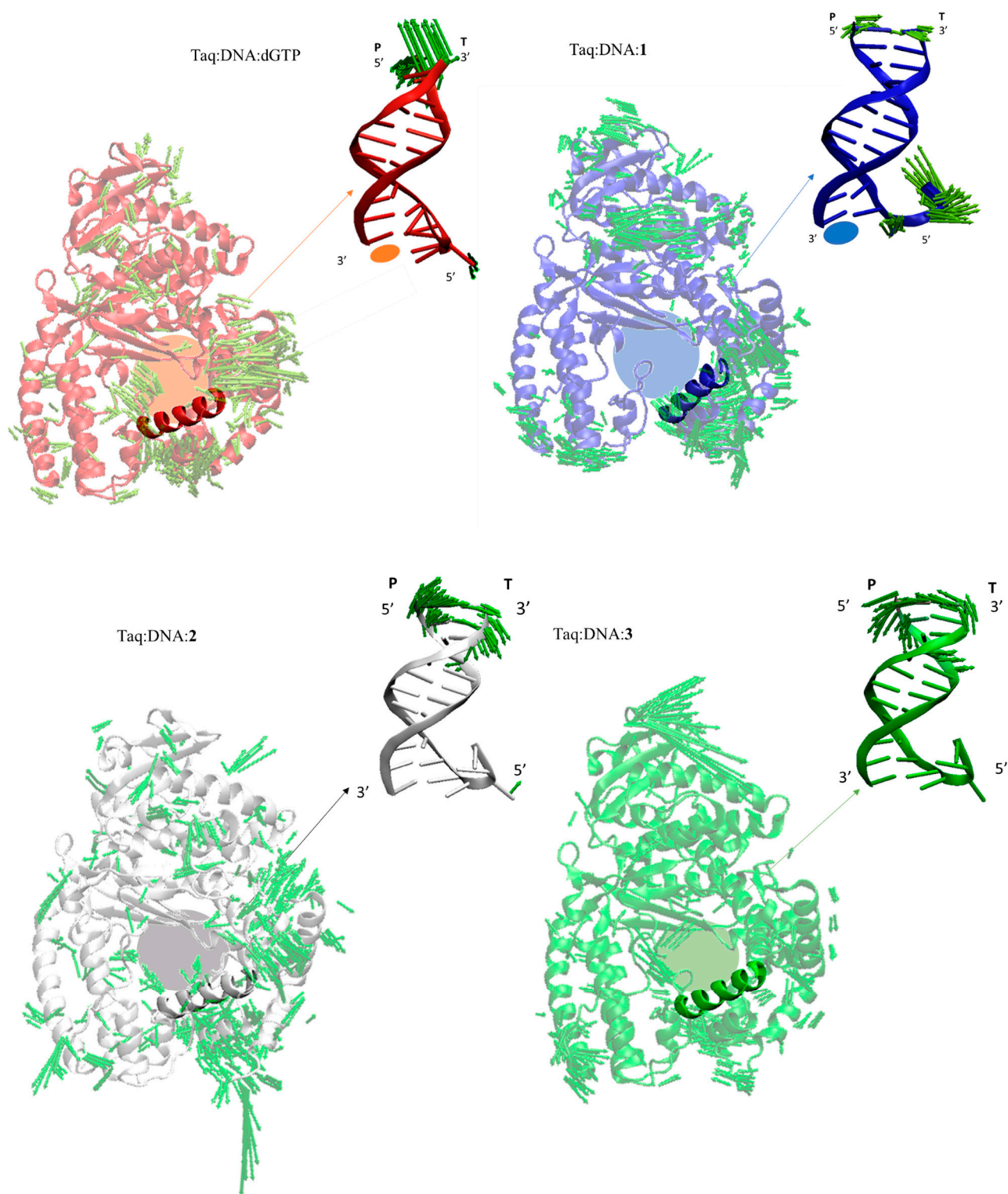


Figure S1. Principal component analysis (PCA) calculated for the investigated systems. Major fluctuations are represented as green arrows.

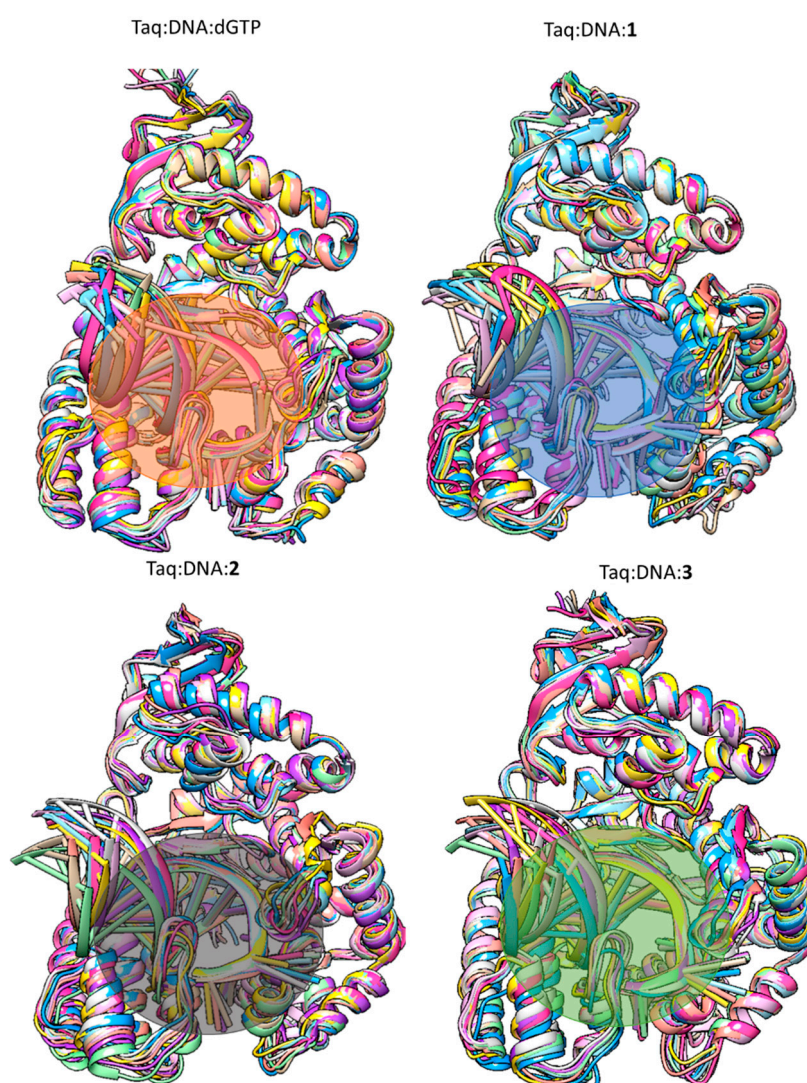
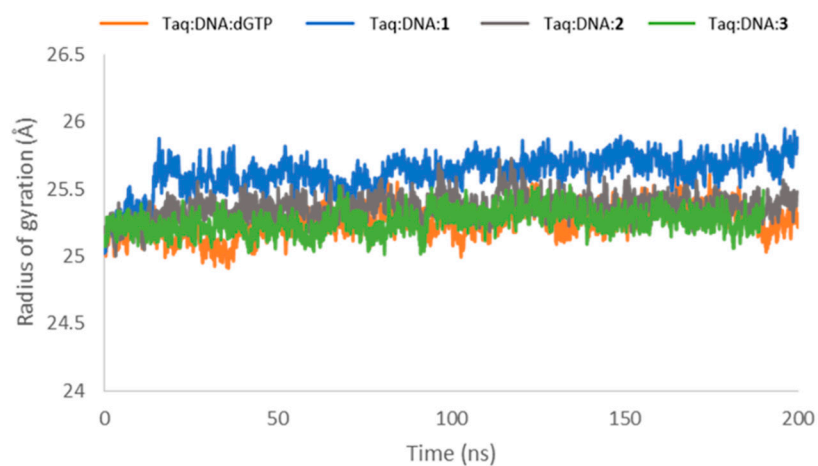
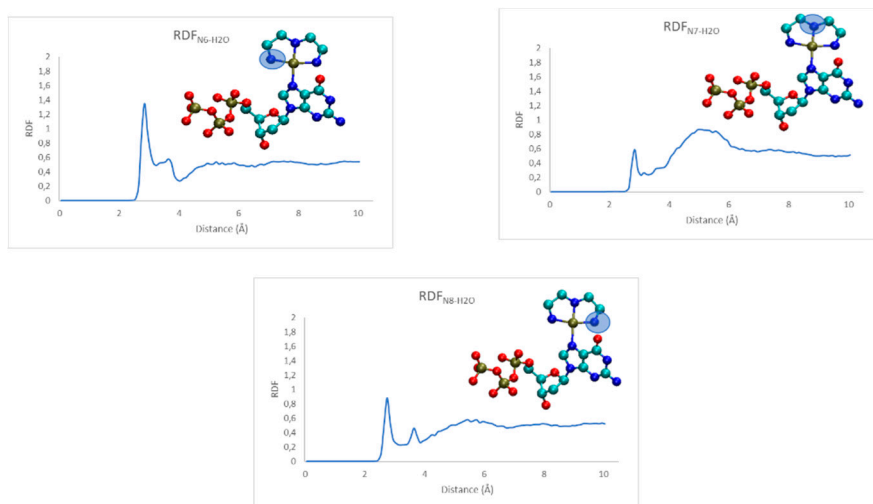
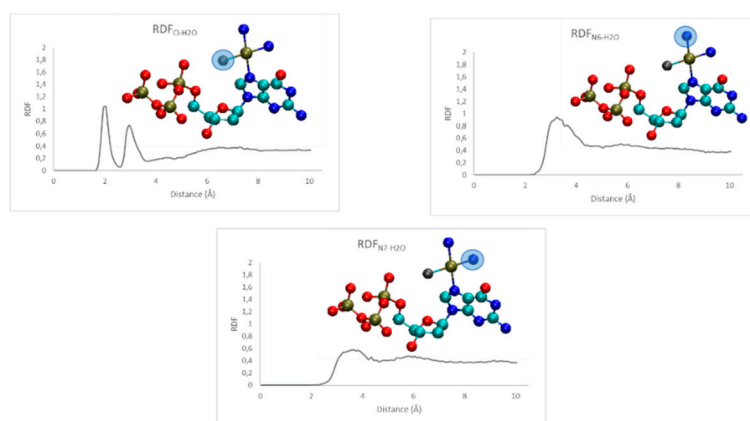


Figure S2. On the top, radius of gyration and, on the bottom, superposition of ten clustered geometries calculated for the investigated systems.

(a)



(b)



(c)

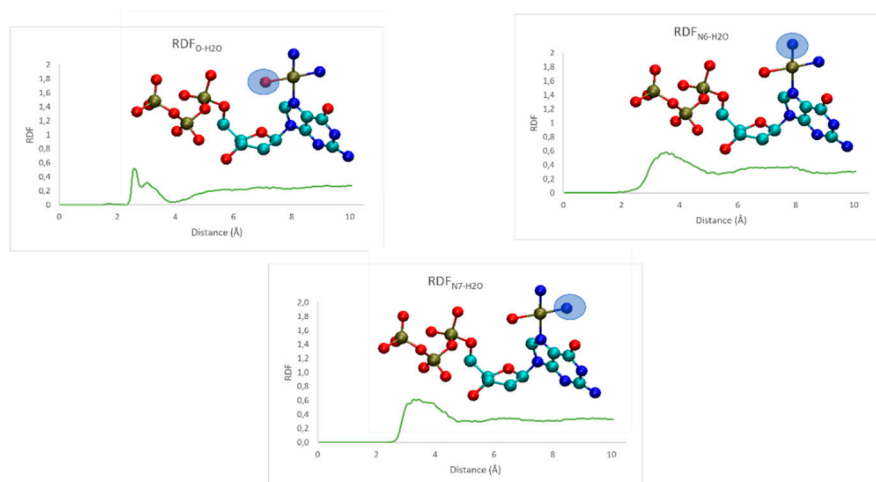


Figure S3. Radial distribution function (RDFs) of $N6-H_2O$, $C1-H_2O$ and $N7-H_2O$ pairs, concerning the water molecule and the ligand to the Pt atoms (circled in the images), calculated for (a) Taq:DNA:1, (b) Taq:DNA:2, and (c) Taq:DNA:3 complexes.

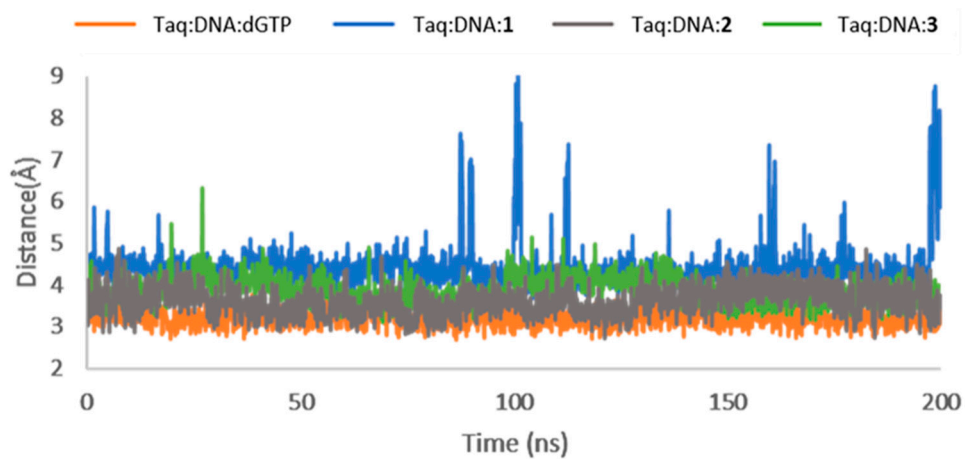
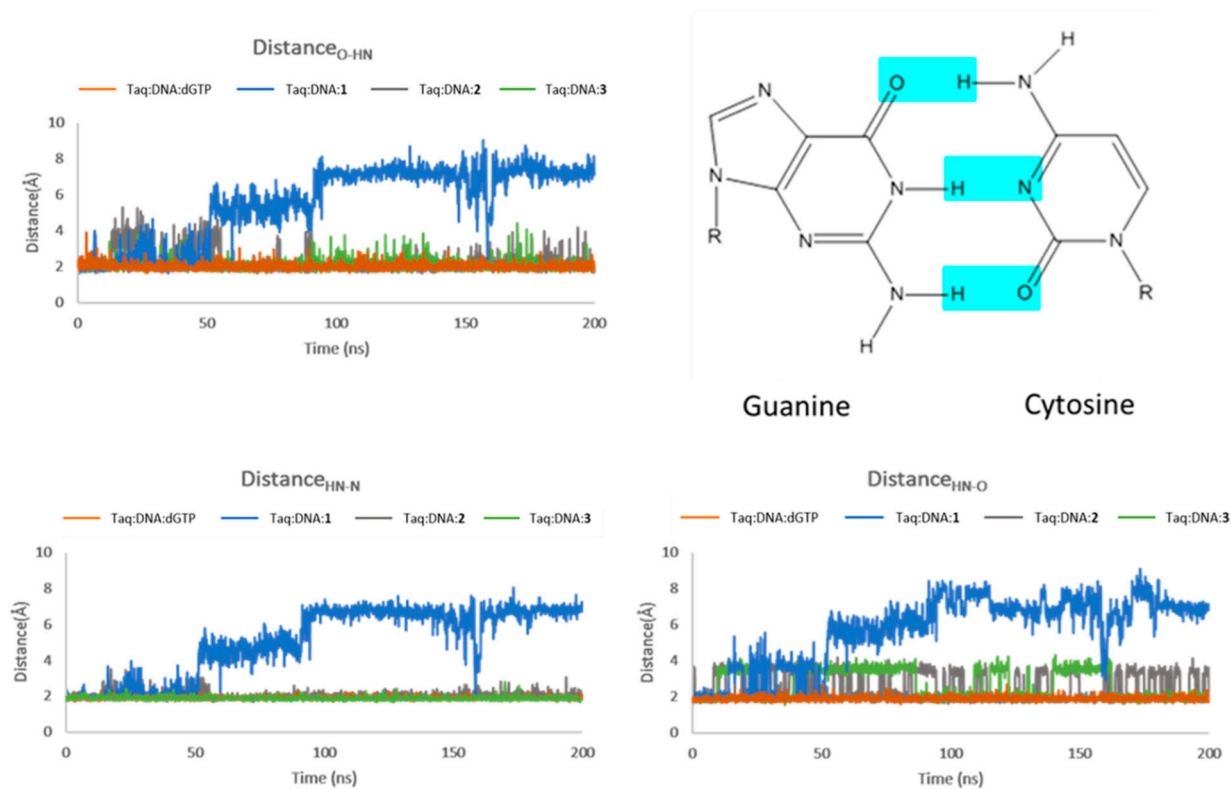


Figure S4. $O_{dNTP}-N_{K663}$ distance obtained from the molecular dynamics simulations of the four investigated systems.



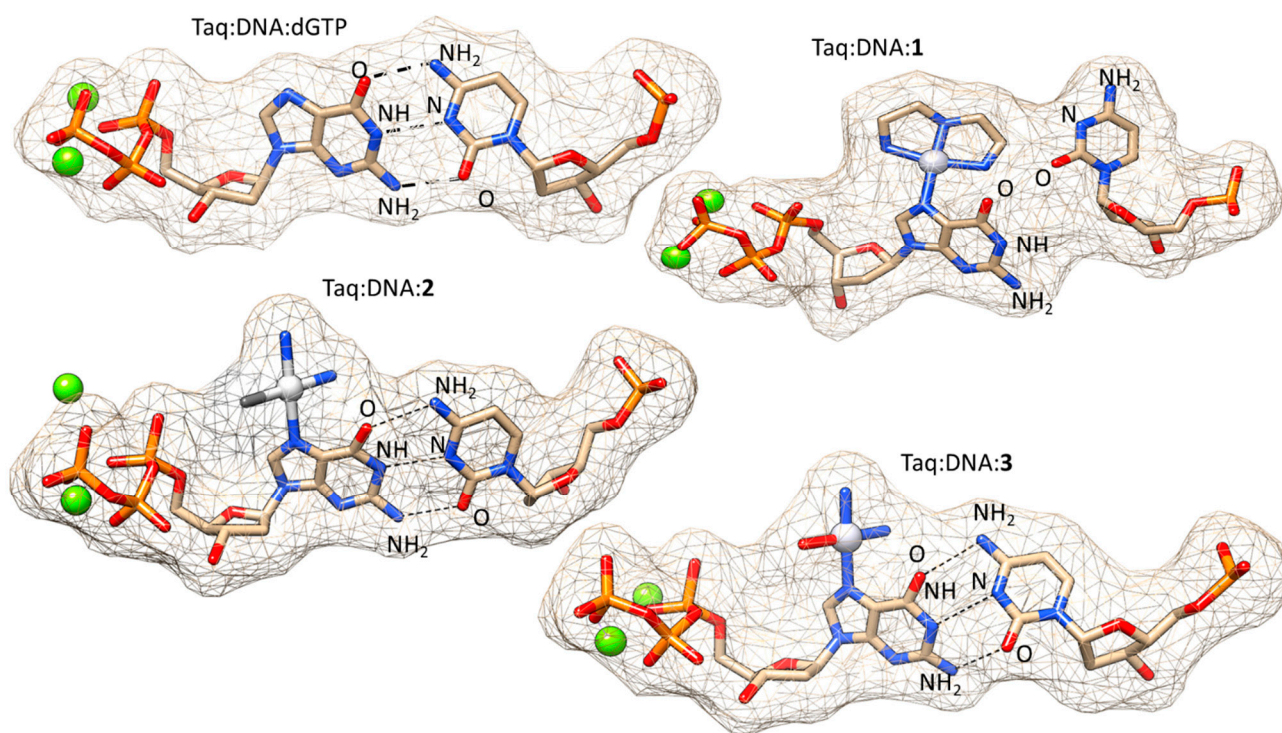


Figure S5. On the top, pairing of guanosine_{dNTP}-cytosine investigated in terms of hydrogen-bond parameters, obtained from the molecular dynamics simulations of the investigated ternary complexes. On the bottom, focus on the paired guanosine_{dNTP}-cytosine obtained from most representative clustered geometry.

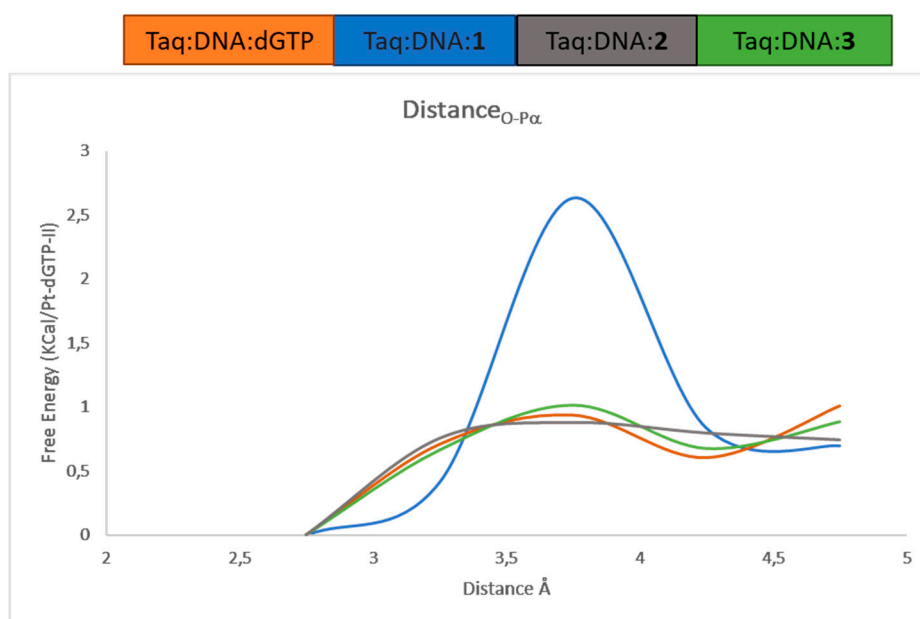


Figure S6. Potential of Mean Force (PMF) calculated for the four different ternary complexes, obtained from umbrella sampling simulations performed on the O_{dC551}-P_α_{dNTP} distance.

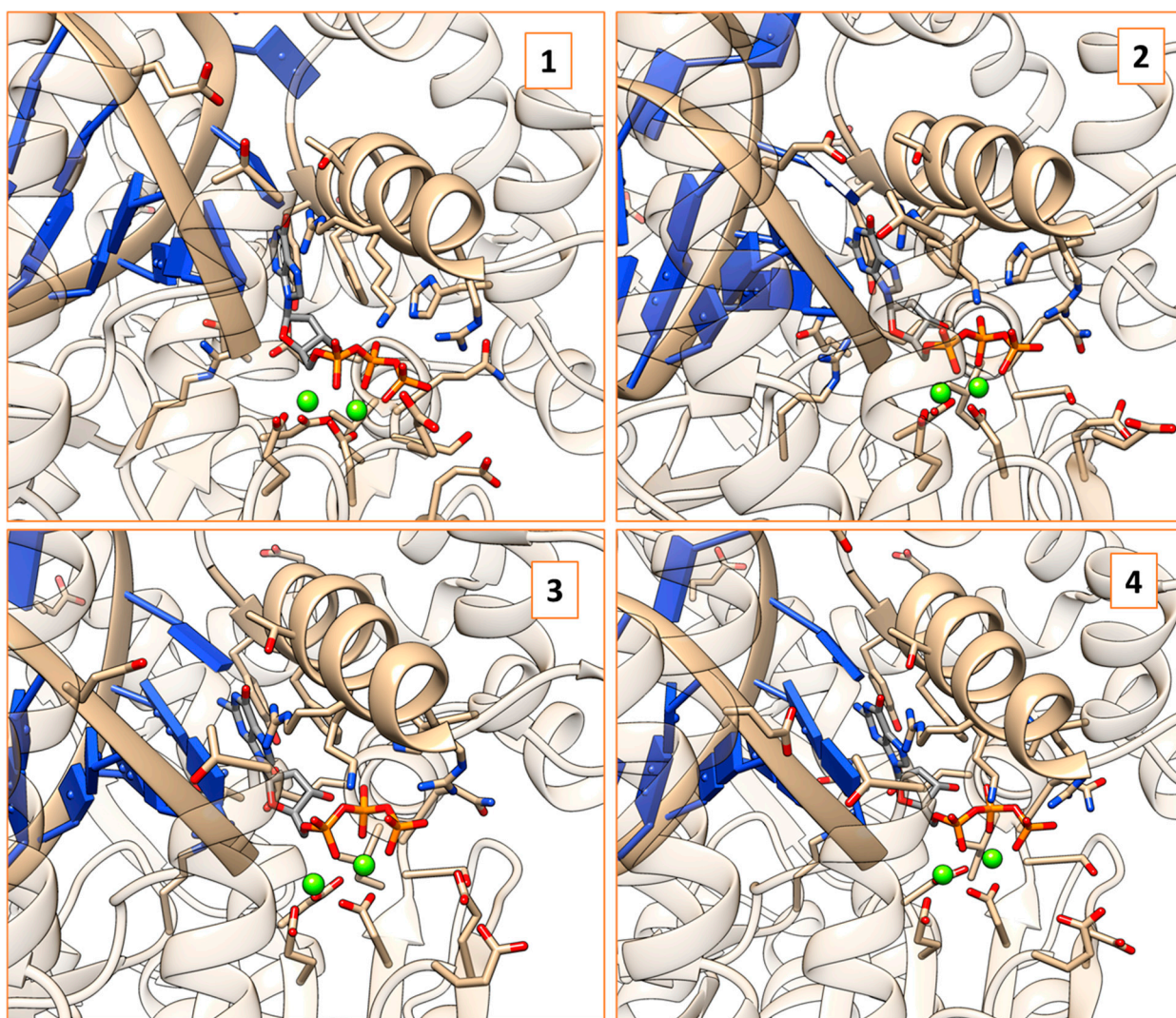


Figure S7. Snapshots of molecular dynamics trajectories, extrapolated at 50, 100, 150 and 200 ns, obtained for the Taq:DNA:dGTP system.

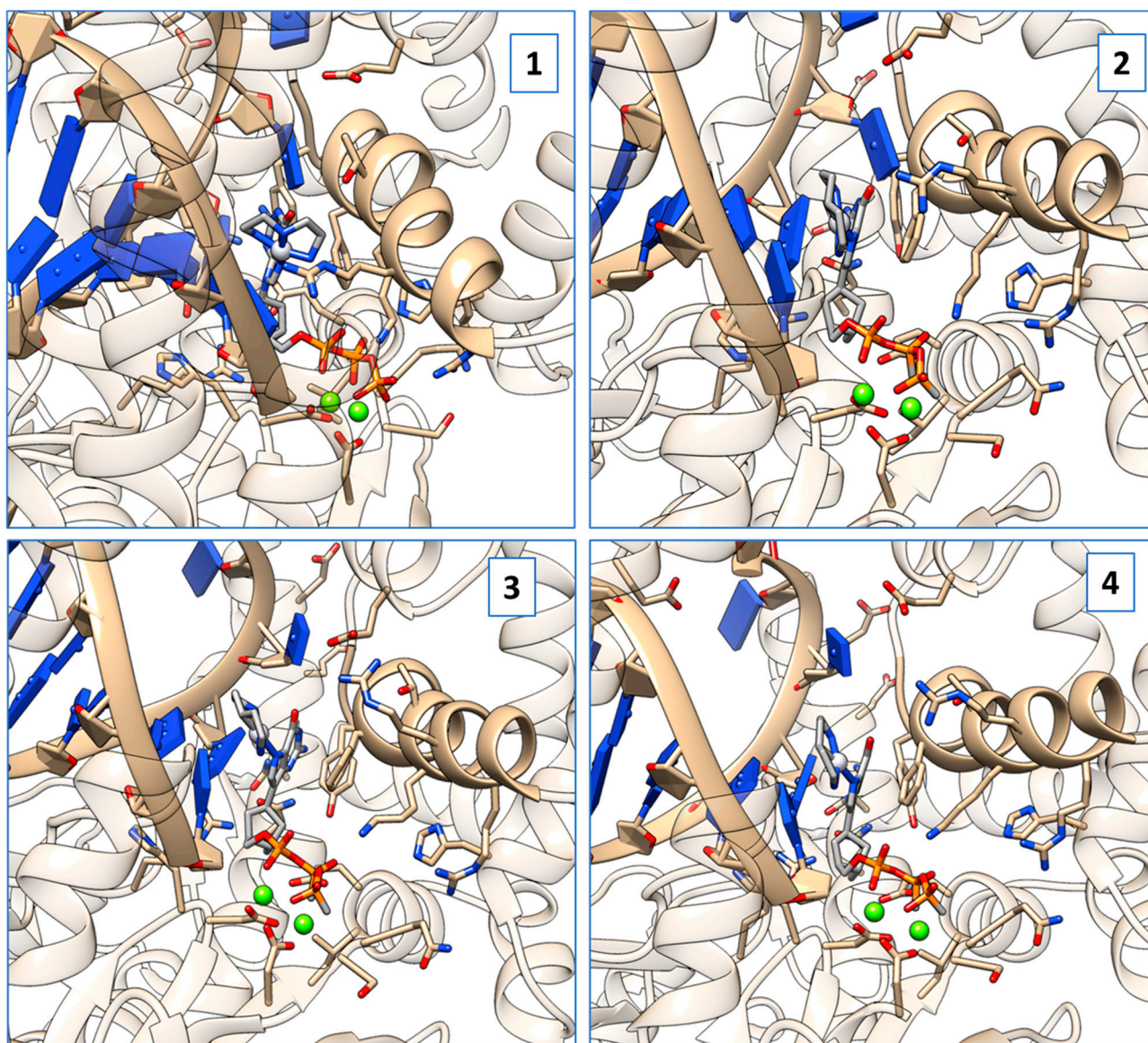


Figure S8. Snapshots of molecular dynamics trajectories, extrapolated at 50, 100, 150 and 200 ns, obtained for the Taq:DNA:1 system.

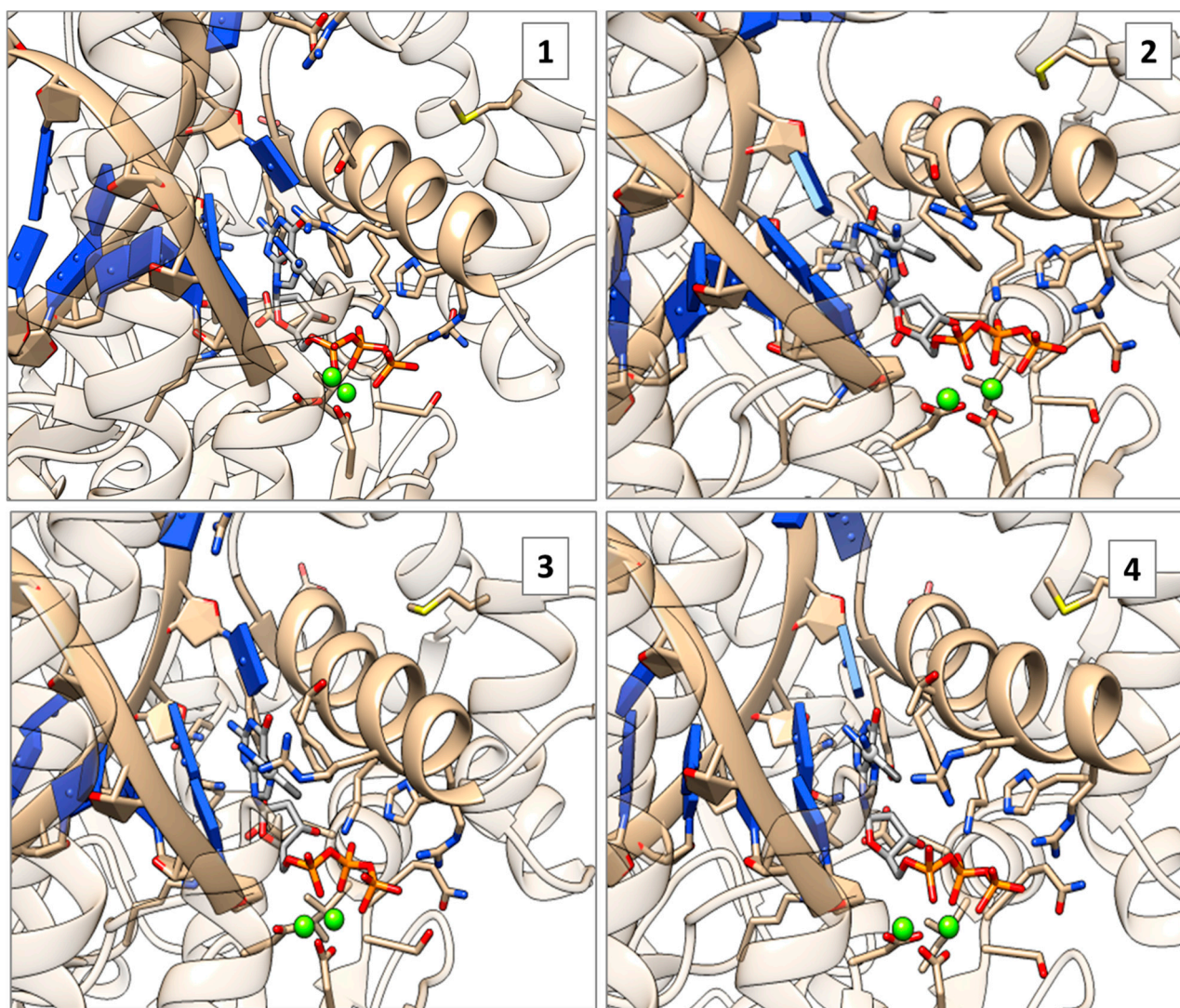


Figure S9. Snapshots of molecular dynamics trajectories, extrapolated at 50, 100, 150 and 200 ns, obtained for the Taq:DNA:2 system.

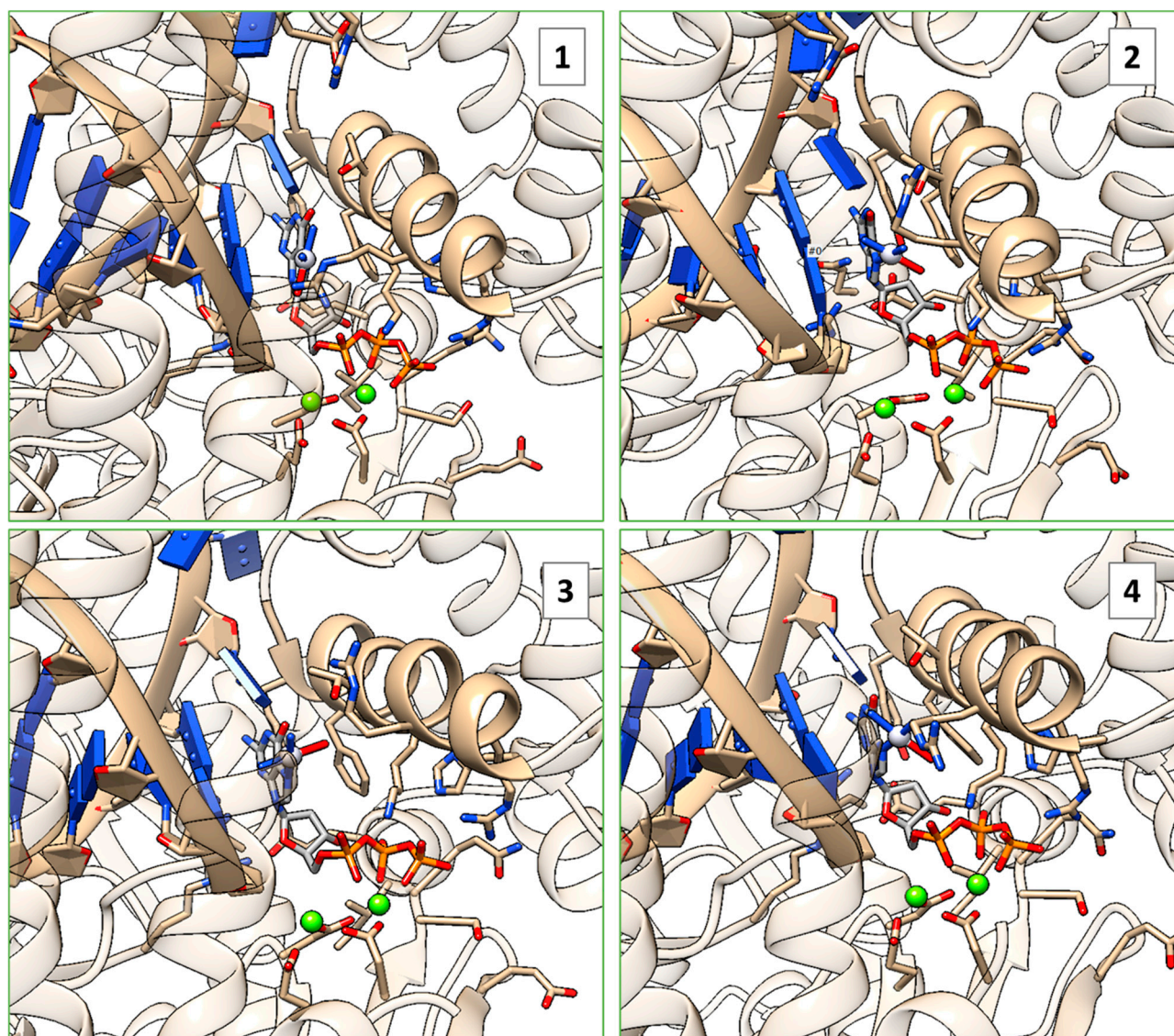


Figure S10. Snapshots of molecular dynamics trajectories, extrapolated at 50, 100, 150 and 200 ns, obtained for the Taq:DNA:3 system.

Replicas of Taq:DNA:ligand complexes

A control on the reproducibility of the conformational behavior observed for all considered complexes was preliminary performed. In detail, additional 100 ns of molecular dynamics were carried out starting from input geometry obtained from the heating phase of the simulation and selecting the same procedure (conditions, algorithms, and software) discussed in the main text (see Methods section). At this stage of the investigation, the attention was mainly focused on the structural parameters considered in the initial part of the study, like RMSD, RMSF, radius of gyration and visual inspection of representative structures obtained from the hierarchical clustering procedure.

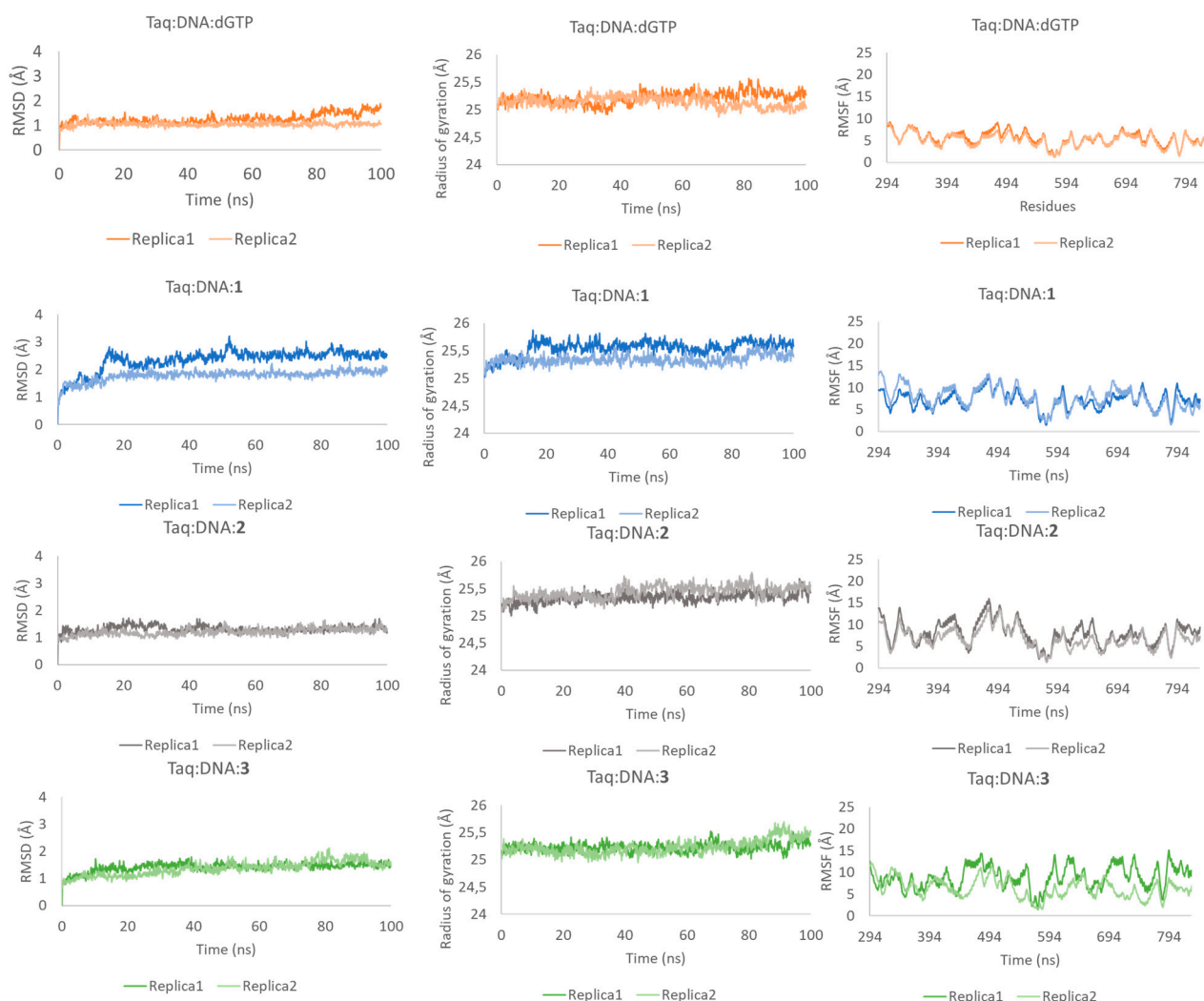


Figure S11. Structural parameters, considered for the analysis of the replicas carried out for the considered systems.

The comparison of each replica at 100 ns showed a reproducible RMSD trend for all the considered complexes, with higher values (>2 Å) obtained for Taq:DNA:1, as reported in **Figure S11**. Overall, the structural reproducibility was evinced by the analysis of RMSF and radius of gyration trends. In the case of the former, a good and superimposable trend between the replicas was obtained for Taq:DNA:dGTP and Taq:DNA:1 in particular, in proximity of residues of the O-helix deputed to the recognition of the ligands (from residue 656 to residue 672, see **Figure S11**). Also the superposition of the most populated cluster geometries (see **Figure S12**) further did not highlight any relevant structural variation of the secondary structures.

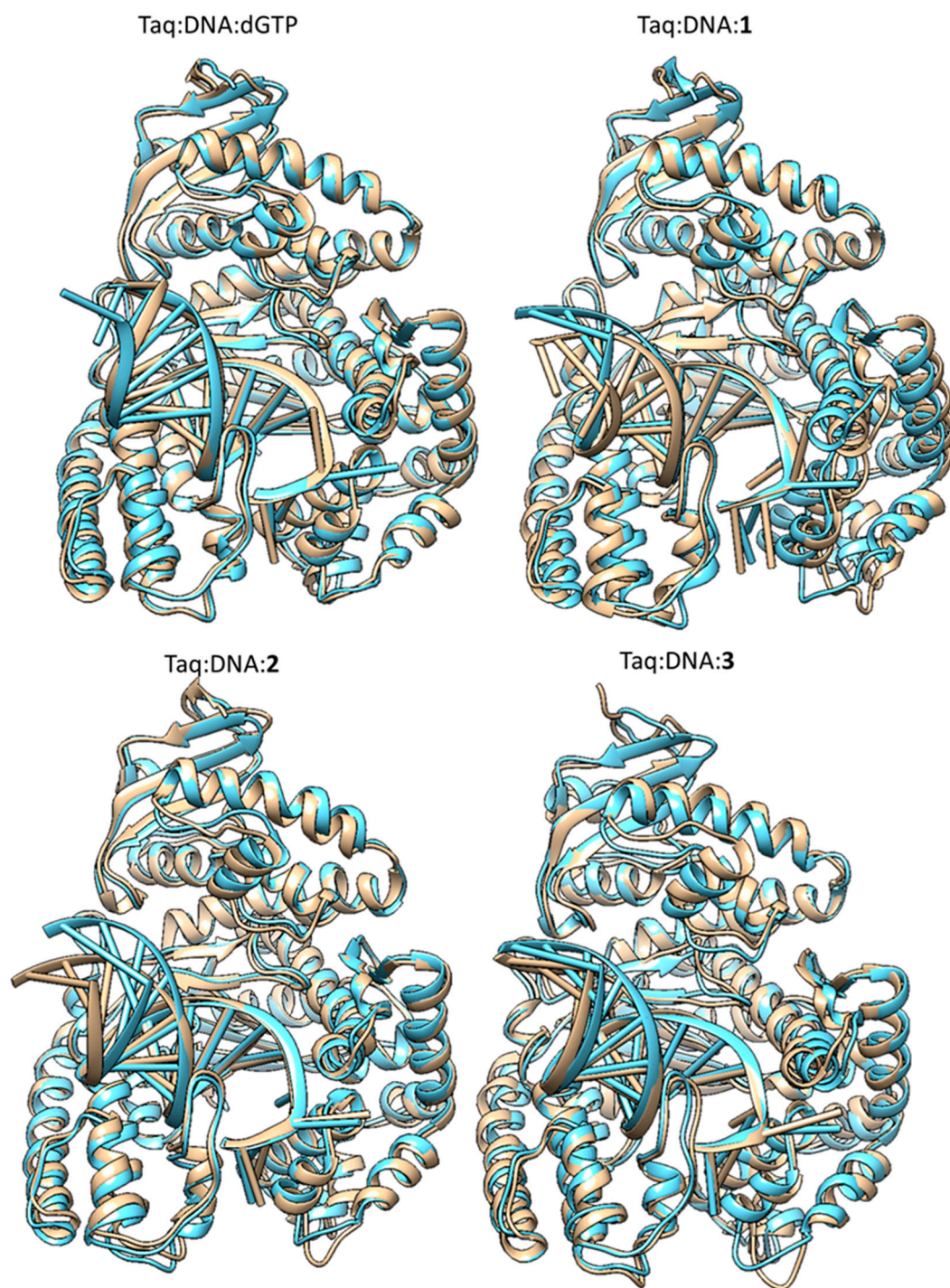


Figure S12. Superimposition of the most representative geometries obtained from hierarchical clustering procedure of replicas of the considered systems.

Table S1. Details of the ten clustered structures obtained from 200 ns of molecular dynamics simulation carried out for the investigated systems.

Taq:DNA:dGTP							Taq:DNA:1						
Cluster	Frames	Frac	AvgDist	Stdev	Centroid	AvgCDist	Cluster	Frames	Frac	AvgDist	Stdev	Centroid	AvgCDist
0	7429	0.371	1.126	0.147	3491	1.57	0	9164	0.458	1.187	0.155	15067	1.786
1	5397	0.27	1.124	0.153	16140	1.539	1	5018	0.251	1.164	0.161	7874	1.677
2	2502	0.125	1.077	0.157	18983	1.478	2	2869	0.143	1.188	0.169	2943	1.633
3	1499	0.075	1.059	0.129	8313	1.436	3	1088	0.054	1.122	0.148	364	1.831
4	1405	0.07	1.045	0.135	10719	1.472	4	706	0.035	1.095	0.172	1747	1.65
5	1086	0.054	1.063	0.166	11386	1.482	5	454	0.023	1.01	0.132	5336	1.815
6	365	0.018	1.045	0.163	9408	1.503	6	334	0.017	1.02	0.225	10851	2.028
7	181	0.009	0.82	0.101	5221	1.692	7	180	0.009	1.015	0.143	1320	1.826
8	116	0.006	0.922	0.063	7834	1.545	8	124	0.006	0.973	0.104	46	2.283
9	20	0.001	0	0	12954	1.784	9	63	0.003	0.965	0.258	2085	1.766

Taq:DNA:2							Taq:DNA:3						
Cluster	Frames	Frac	AvgDist	Stdev	Centroid	AvgCDist	Cluster	Frames	Frac	AvgDist	Stdev	Centroid	AvgCDist
0	13410	0.67	1.087	0.127	17621	1.315	0	6143	0.323	1.067	0.131	8386	1.293
1	3968	0.198	1.124	0.139	3198	1.346	1	5327	0.28	1.019	0.109	15418	1.295
2	1268	0.063	1.043	0.128	441	1.504	2	4185	0.22	1.066	0.119	3451	1.313
3	609	0.03	0.924	0.103	11763	1.5	3	1292	0.068	1.011	0.12	1623	1.279
4	283	0.014	0.956	0.154	17076	1.426	4	897	0.047	0.972	0.127	478	1.35
5	207	0.01	0.971	0.147	6090	1.462	5	867	0.046	1.023	0.153	9172	1.38
6	153	0.008	0.901	0.129	4309	1.442	6	193	0.01	0.977	0.047	11185	1.434
7	46	0.002	1.032	0.173	8834	1.431	7	54	0.003	0.829	0.088	897	1.355
8	41	0.002	0.83	0.033	15191	1.499	8	29	0.002	0	0	16953	1.599
9	15	0.001	0	0	4749	1.485	9	14	0.001	0	0	8580	1.398

Table S2. Results of MMPBSA calculation and, in the next page, Per-residue decomposition analysis carried out for all investigated systems.

Taq:DNA:dGTP				Taq:DNA:1			
Energy Component	Average (kcal/mol)	Std. Dev.	Std. Err. Of Mean	Energy Component	Average (kcal/mol)	Std. Dev.	Std. Err. Of Mean
VDWAALS	-30,5895	6,4703	0,647	VDWAALS	-16,8036	6,8919	0,6892
EEL	50,7178	72,0463	7,2046	EEL	-464,87	58,0147	5,8015
EPB	-217,0776	59,6789	5,9679	EPB	376,4979	49,8608	4,9861
ENPOLAR	-3,8233	0,0552	0,0055	ENPOLAR	-5,64	0,1595	0,016
EDISPER	0	0	0	EDISPER	0	0	0
D G gas	20,1766	70,3387	7,0339	D G gas	-479,1343	58,2689	5,8269
D G solv	-220,9009	59,6974	5,9697	D G solv	370,8578	49,846	4,9846
D TOTAL	-200,7243	14,1667	1,4167	D TOTAL	-108,2765	24,2667	2,4267

Taq:DNA:2				Taq:DNA:3			
Energy Component	Average (kcal/mol)	Std. Dev.	Std. Err. Of Mean	Energy Component	Average (kcal/mol)	Std. Dev.	Std. Err. Of Mean
VDWAALS	-41,0381	5,8275	0,5828	VDWAALS	-40,2914	7,2343	0,7234
EEL	141,7018	43,4112	4,3411	EEL	-79,3406	31,1848	3,1185
EPB	-202,5309	38,0495	3,8049	EPB	29,1732	29,7848	2,9785
ENPOLAR	-4,8065	0,0971	0,0097	ENPOLAR	-4,8069	0,121	0,0121
EDISPER	0	0	0	EDISPER	0	0	0
D G gas	100,6637	42,079	4,2079	D G gas	-119,632	31,2792	3,1279
D G solv	-207,3374	38,0494	3,8049	D G solv	24,3662	29,7683	2,9768
D TOTAL	-106,6737	9,0736	0,9074	D TOTAL	-95,2657	9,3905	0,939

Taq:DNA:dGTP

Resid Number	ΔG internal (kcal/mol)	ΔG vdw (kcal/mol)	ΔG eel (kcal/mol)	ΔG pol (kcal/mol)	ΔG TOT (kcal/mol)
R573	0	-0,482	-62,407	57,618	-5,271
D610	0	-0,951	145,217	-132,452	11,814
Y611	0	-0,722	19,885	-14,951	4,212
S612	0	-1,199	-14	11,632	-3,567
N613	0	-1,933	-19,824	14,504	-7,253
I614	0	-1,529	-8,493	6,414	-3,608
E615	0	-1,504	50,798	-41,045	8,249
H639	0	-1,498	-19,261	14,951	-5,808
R659	0	0,151	-127,343	108,912	-18,28
R660	0	-0,108	-81,505	73,89	-7,723
K663	0	-1,006	-158,545	140,663	-18,888
F667	0	-3,143	-5,433	5,047	-3,529
Y671	0	-0,756	-1,533	0,804	-1,485
Q754	0	-0,669	-0,542	-0,223	-1,434
D785	0	0,593	147,559	-128,492	19,66

Taq:DNA:1

Resid Number	ΔG internal (kcal/mol)	ΔG vdw (kcal/mol)	ΔG eel (kcal/mol)	ΔG pol (kcal/mol)	ΔG TOT (kcal/mol)
R573	0	-0,723	-35,935	31,866	-4,069
D610	0	0,402	127,891	-117,256	10,635
E615	0	0,236	44,809	-36,41	8,399
K663	0	-0,993	-81,605	81,77	0,165
F667	0	-3,886	-1,435	2,759	1,324
Y671	0	-0,817	-1,536	1,373	-0,163
N750	0	-0,271	0,445	-0,865	-0,42
Q754	0	-1,316	-3,169	1,478	-1,691
H784	0	-0,977	-0,325	1,241	0,916
D785	0	-1,206	127,557	-113,404	14,153

Taq:DNA:2

Resid Number	ΔG internal (kcal/mol)	ΔG vdw (kcal/mol)	ΔG eel (kcal/mol)	ΔG pol (kcal/mol)	ΔG TOT (kcal/mol)
R573	0	-1,034	-50,563	49,916	-1,681
D610	0	-0,988	112,165	-106,401	4,776
Y611	0	-0,423	15,648	-12,508	2,717
S612	0	-1,4	-12,007	11,01	-2,397
N613	0	-0,996	-14,656	13,088	-2,564
I614	0	-1,939	-5,967	5,158	-2,748
E615	0	-1,708	43,88	-37,804	4,368
H639	0	-1,4	-15,216	12,318	-4,298
R659	0	-0,036	-97,172	86,508	-10,7
R660	0	-0,213	-47,285	45,826	-1,672
K663	0	-0,992	-115,588	107,621	-8,959
T664	0	-0,615	0,281	1,179	0,845
F667	0	-3,712	-4,612	4,288	-4,036
Y671	0	-0,912	-0,288	0,158	-1,042
Q754	0	-0,728	-2,868	0,479	-3,117
D785	0	-1,045	105,871	-97,417	7,409

Taq:DNA:3

Resid Number	ΔG internal (kcal/mol)	ΔG vdw (kcal/mol)	ΔG eel (kcal/mol)	ΔG pol (kcal/mol)	ΔG TOT (kcal/mol)
R573	0	-0,579	-31,388	27,441	-4,526
D610	0	-0,599	93,879	-91,475	1,805
Y611	0	-0,844	12,952	-11,749	0,359
S612	0	-1,128	-10,15	9,32	-1,958
N613	0	-2,065	-12,516	11,309	-3,272
I614	0	-2,001	-6,058	5,243	-2,816
E615	0	-2,437	28,489	-18,935	7,117
H639	0	-1,208	-11,017	10,735	-1,49
R659	0	-0,05	-69,949	63,986	-6,013
R660	0	-0,368	-30,894	30,703	-0,559
K663	0	-0,075	-94,831	84,106	-10,8
F667	0	-3,815	-3,196	2,591	-4,42
Y671	0	-0,825	-0,666	0,181	-1,31
Q754	0	-1,048	-0,768	-0,363	-2,179
D785	0	-0,164	89,443	-86,925	2,354

Parameters in Amber16 format

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Parameters complex 1

4 O3 20.1600 6.7270 0.5290 o 1
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File mol2:

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MOL

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SMALL

bcc

@<TRIPOS>ATOM

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2 O1 20.0490 8.6620 2.2080 o 1

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5 O4 21.4390 6.7790 2.5730 os 1
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6 P2 22.9680 6.3840 2.8980 p5 1
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7 O5 22.8700 5.1690 3.7780 o 1
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8 O6 23.8440 6.3660 1.7030 o 1
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9 O7 23.3530 7.5350 3.9730 os 1
MOL -0.817800

10 P3 24.3560 8.7590 3.6350 p5 1
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MOL	-0.926500				MOL	0.545800			
12 O9	24.4740	8.9970	2.1750 o	1	29 N4	27.9970	3.8790	11.4040 nh	1
MOL	-0.926500				MOL	-0.862800			
13 O10	25.7280	8.1780	4.2120 os	1	30 N5	27.5110	5.1290	9.5380 nc	1
MOL	-0.603200				MOL	-0.596100			
14 C1	26.5990	7.3450	3.4680 c3	1	31 C10	26.8720	6.1500	8.9400 cc	1
MOL	0.194400				MOL	0.016800			
15 C2	27.1220	6.2410	4.3650 c3	1	32 C11	23.9650	9.4640	11.6690 c3	1
MOL	0.093100				MOL	0.144300			
16 O11	27.8070	6.7860	5.5060 os	1	33 C12	24.0130	11.4100	7.2970 c3	1
MOL	-0.453600				MOL	0.144300			
17 C3	26.0340	5.3420	4.9500 c3	1	34 N6	25.1670	10.9030	7.9230 Y2	1
MOL	0.148100				MOL	-0.931800			
18 O12	25.6780	4.2710	4.0760 oh	1	35 N7	23.6430	8.2890	10.9800 Y3	1
MOL	-0.686800				MOL	-0.931800			
19 C4	26.7350	4.8090	6.1630 c3	1	36 C13	23.0130	10.4220	11.1140 c3	1
MOL	-0.113400				MOL	0.171300			
20 C5	27.6250	5.9470	6.6250 c3	1	37 C14	23.2640	11.5480	8.4720 c3	1
MOL	0.250800				MOL	0.171300			
21 N1	26.8590	6.5940	7.7210 na	1	38 N8	22.8980	10.8800	9.6470 Y4	1
MOL	-0.167900				MOL	-0.861200			
22 C6	26.0230	7.6410	7.6390 cc	1	39 H1	27.4350	7.9350	3.0920 h1	1
MOL	0.482400				MOL	0.028200			
23 N2	25.4980	7.9260	8.8270 Y1	1	40 H2	26.0560	6.9070	2.6300 h1	1
MOL	-0.647000				MOL	0.028200			
24 C7	25.9650	7.0220	9.6940 cd	1	41 H3	27.8190	5.6260	3.7960 h1	1
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25 C8	25.8140	6.7290	11.1130 c	1	42 H4	25.1560	5.9240	5.2310 h1	1
MOL	0.641000				MOL	0.180700			
26 O13	25.0580	7.4180	11.8400 o	1	43 H5	27.3320	3.9330	5.9080 hc	1
MOL	-0.667500				MOL	0.064700			
27 N3	26.5120	5.6870	11.6080 n	1	44 H6	26.0100	4.5590	6.9380 hc	1
MOL	-0.491400				MOL	0.064700			

45 H7	28.5810	5.5700	6.9880 h2	1	62 H24	23.5770	10.6990	6.6270 h1	1
MOL	0.052700				MOL	0.034950			
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MOL	0.172100				MOL	0.372800			
47 H9	26.4210	5.4700	12.5900 hn	1	64 PT	24.2490	9.4200	9.2380 M1	1
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50 H12	24.1960	12.3540	6.8270 h1	1	4	1	5	1	
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51 H13	23.1760	11.3390	11.6810 h1	1	6	6	7	1	
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52 H14	22.0230	10.0640	11.3970 h1	1	8	6	9	1	
MOL	-0.005050				9	9	10	1	
53 H15	22.3050	11.8120	8.0770 h1	1	10	10	11	1	
MOL	-0.005050				11	10	12	1	
54 H16	22.0440	10.4230	9.3600 hn	1	12	10	13	1	
MOL	0.454800				13	13	14	1	
55 H17	28.6020	3.3070	10.8330 hn	1	14	14	15	1	
MOL	0.366800				15	14	39	1	
56 H18	27.8910	3.6760	12.3880 hn	1	16	14	40	1	
MOL	0.366800				17	15	16	1	
57 H19	22.9190	7.7940	11.4800 hn	1	18	15	17	1	
MOL	0.372800				19	15	41	1	
58 H20	23.3140	8.5210	10.0540 hn	1	20	16	20	1	
MOL	0.372800				21	17	18	1	
59 H21	24.7370	4.3100	3.8880 ho	1	22	17	19	1	
MOL	0.493000								
60 H22	25.9540	11.0230	7.3180 hn	1					
MOL	0.372800								
61 H23	23.6650	12.4580	8.8670 h1	1					
MOL	-0.005050								

23 17 42 1
 24 18 59 1
 25 19 20 1
 26 19 43 1
 27 19 44 1
 28 20 21 1
 29 20 45 1
 30 21 22 1
 31 21 31 1
 32 22 23 2
 33 22 46 1
 34 23 24 1
 35 24 25 1
 36 24 31 2
 37 25 26 2
 38 25 27 1
 39 27 28 1
 40 27 47 1
 41 28 29 1
 42 28 30 2
 43 29 55 1
 44 29 56 1
 45 30 31 1
 46 32 35 1
 47 32 36 1
 48 32 48 1
 49 32 49 1
 50 33 34 1

51 33 37 1
 52 33 50 1
 53 33 62 1
 54 34 60 1
 55 34 63 1
 56 35 57 1
 57 35 58 1
 58 36 38 1
 59 36 51 1
 60 36 52 1
 61 37 38 1
 62 37 53 1
 63 37 61 1
 64 38 54 1

@<TRIPOS>SUBSTRUCTURE

1 MOL 1 TEMP 0 ***** 0 ROOT

File frcmod:

remark goes here

MASS

p5 30.970 1.538

o 16.000 0.434

os 16.000 0.465

c3 12.010 0.878

oh 16.000 0.465

na 14.010 0.530

cc 12.010 0.360

cd 12.010 0.360

c	12.010	0.616		c3-na	262.85	1.463	
n	14.010	0.530		c3-h2	377.33	1.096	
nh	14.010	0.530		na-cc	354.49	1.380	
nc	14.010	0.530		cc-h5	403.49	1.082	
h1	1.008	0.135		cd-c	295.35	1.468	
hc	1.008	0.135		cd-cc	416.13	1.373	
h2	1.008	0.135		c-o	652.57	1.218	
h5	1.008	0.135		c-n	356.21	1.379	
hn	1.008	0.161		n-cd	353.83	1.381	
ho	1.008	0.135		n-hn	527.31	1.013	
M1	195.08			cd-nh	363.47	1.373	
Y1	14.01	0.530	Sp2 N in non-pure aromatic systems, identical to nc	cd-nc	450.71	1.317	
Y2	14.01	0.530	Sp3 N with three connected atoms	nh-hn	529.46	1.012	
Y3	14.01	0.530	Sp3 N with three connected atoms	nc-cc	369.10	1.369	
Y4	14.01	0.530	Sp3 N with three connected atoms	Y1-M1	80.2	2.1471	Created by Seminario method using MCPB.py
				na-M1	80.2	2.1471	Created by Seminario method using MCPB.py
				o-M1	80.2	2.1471	Created by Seminario method using MCPB.py
BOND				Y2-M1	107.7	2.1129	Created by Seminario method using MCPB.py
p5-o	529.55	1.487		os-M1	80.2	2.1471	Created by Seminario method using MCPB.py
p5-os	346.25	1.615		Y3-M1	120.0	2.0740	Created by Seminario method using MCPB.py
os-c3	284.76	1.432		Y4-M1	166.0	2.0051	Created by Seminario method using MCPB.py
c3-c3	232.52	1.538		Y1-cd	441.1	1.3694	SOURCE1_SOURCE5
c3-h1	375.92	1.097			2269	0.0086	
c3-oh	293.40	1.423					
oh-ho	563.51	0.973					
c3-hc	375.92	1.097					

Y2-hn	392.4	1.0190	SOURCE3_SOURCE5	os-c3-Y1	109.296	109.030
5944	0.0012			os-c3-h2	62.442	109.580
Y3-hn	392.4	1.0190	SOURCE3_SOURCE5	c3-oh-ho	49.027	107.260
5944	0.0012			c3-c3-hc	46.816	109.800
Y4-hn	392.4	1.0190	SOURCE3_SOURCE5	oh-c3-h1	62.540	110.260
5944	0.0012			c3-c3-na	82.668	112.880
c3-Y2	325.9	1.4647	SOURCE1_SOURCE5	c3-c3-h2	46.730	110.220
15206	0.0039			c3-na-cc	63.695	126.460
c3-Y3	325.9	1.4647	SOURCE1_SOURCE5	c3-Y1-cc	63.695	126.460
15206	0.0039			na-c3-h2	61.963	107.310
c3-Y4	325.9	1.4647	SOURCE1_SOURCE5	na-cc-h5	61.226	121.550
15206	0.0039			na-cc-cd	92.653	106.990
cc-Y1	525.4	1.3172	SOURCE3_SOURCE5	Y1-cc-cd	92.653	106.990
4612	0.0083			na-cc-nc	108.812	121.950
c3-Y1	525.4	1.3172	SOURCE3_SOURCE5	Y1-cc-nc	108.812	121.950
4612	0.0083			cc-na-cc	70.492	109.900
ANGLE				cd-c -o	86.736	123.930
p5-os-p5	106.787	126.250		cd-c -n	87.145	112.700
o -p5-o	85.510	115.800		cd-cc-nc	91.057	111.650
o -p5-os	81.819	115.460		c -cd-cc	67.187	121.350
os-p5-os	83.949	101.840		c -n -cd	66.579	123.270
p5-os-c3	83.251	119.540		c -n -hn	48.691	117.550
os-c3-c3	85.306	107.970		o -c -n	113.811	123.050
p5-o-c3	83.251	119.540		n -cd-nh	110.935	116.940
os-c3-h1	62.377	109.780		n -cd-nc	110.304	123.000
c3-c3-c3	64.888	111.510		cd-n -hn	48.285	119.260
c3-c3-h1	46.868	109.560		cd-nh-hn	49.267	115.630
c3-os-c3	66.293	112.480		cd-nc-cc	73.871	105.490
c3-c3-oh	84.642	110.190				
os-c3-na	109.296	109.030				

nh-cd-nc	111.697	120.650			
h1-c3-h1	38.802	108.460			
hc-c3-hc	38.960	107.580			
hn-nh-hn	39.519	115.120			
M1-Y1-cd	183.02	128.65	Created by Seminario method using MCPB.py		
M1-Y2-hn	65.00	112.03	Created by Seminario method using MCPB.py		
M1-Y3-hn	61.05	107.39	Created by Seminario method using MCPB.py		
M1-Y4-hn	6.91	75.07	Created by Seminario method using MCPB.py		
Y1-M1-Y2	182.93	101.20	Created by Seminario method using MCPB.py		
Y1-M1-Y3	183.35	92.13	Created by Seminario method using MCPB.py		
Y1-M1-Y4	159.20	174.42	Created by Seminario method using MCPB.py		
Y2-M1-Y3	150.16	166.59	Created by Seminario method using MCPB.py		
Y2-M1-Y4	141.81	82.68	Created by Seminario method using MCPB.py		
Y3-M1-Y4	143.98	83.92	Created by Seminario method using MCPB.py		
c3-Y2-M1	159.25	107.57	Created by Seminario method using MCPB.py		
na-M1-Y1	182.93	101.20	Created by Seminario method using MCPB.py		
na-M1-Y2	183.35	92.13	Created by Seminario method using MCPB.py		
na-M1-Y3	159.20	174.42	Created by Seminario method using MCPB.py		
na-M1-Y4	150.16	166.59	Created by Seminario method using MCPB.py		
c3-Y3-M1	135.52	107.32	Created by Seminario method using MCPB.py		
c3-Y4-M1	149.84	109.73	Created by Seminario method using MCPB.py		
cc-Y1-M1	152.18	123.14	Created by Seminario method using MCPB.py		
cc-na-M1	152.18	123.14	Created by Seminario method using MCPB.py		
c3-na-M1	152.18	123.14	Created by Seminario method using MCPB.py		
c3-Y1-M1	152.18	123.14	Created by Seminario method using MCPB.py		
Y1-cd-c	66.22	123.32		CORR_SOURCE5	
27	2.2025				
Y1-cd-cc	72.17	111.65		CORR_SOURCE5	
1656	1.8430				
c3-Y2-hn	47.42	109.29		SOURCE3_SOURCE5	
6742	0.6614				
c3-Y3-hn	47.42	109.29		SOURCE3_SOURCE5	
6742	0.6614				
c3-Y4-c3	63.82	112.35		SOURCE3_SOURCE5	
10425	1.3688				
c3-Y4-hn	47.42	109.29		SOURCE3_SOURCE5	
6742	0.6614				
c3-c3-Y2	66.02	111.04		SOURCE3_SOURCE5	
12086	1.5519				
c3-c3-Y3	66.02	111.04		SOURCE3_SOURCE5	
12086	1.5519				
c3-c3-Y4	66.02	111.04		SOURCE3_SOURCE5	
12086	1.5519				
c3-c3-Y1	66.02	111.04		SOURCE3_SOURCE5	
12086	1.5519				

cc-Y1-cd	71.76	105.49	CORR_SOURCE5		o -p5-os-p5	1	0.550	0.000	3.000
1810	1.9032				os-c3-c3-os	1	0.000	0.000	-3.000
h1-c3-Y2	49.53	109.88	SOURCE3_SOURCE5		os-c3-c3-os	1	0.000	180.000	-2.000
20428	1.2681				os-c3-c3-os	1	0.170	180.000	1.000
h1-c3-Y3	49.53	109.88	SOURCE3_SOURCE5		os-c3-c3-c3	1	0.156	0.000	3.000
20428	1.2681				os-c3-c3-h1	1	0.000	0.000	-3.000
h1-c3-Y4	49.53	109.88	SOURCE3_SOURCE5		os-c3-c3-h1	1	0.250	0.000	1.000
20428	1.2681				c3-c3-os-c3	1	0.240	0.000	-3.000
h5-cc-Y1	50.58	125.52	SOURCE3_SOURCE5		c3-c3-os-c3	1	0.160	0.000	2.000
1309	0.7276				c3-c3-c3-oh	1	0.210	0.000	3.000
hn-Y2-hn	41.40	106.40	SOURCE3_SOURCE5		c3-c3-c3-c3	1	0.130	0.000	-3.000
2019	0.9777				c3-c3-c3-c3	1	0.290	180.000	-2.000
hn-Y3-hn	41.40	106.40	SOURCE3_SOURCE5		c3-c3-c3-c3	1	0.110	0.000	1.000
2019	0.9777				c3-c3-c3-h1	1	0.156	0.000	3.000
na-cc-Y1	74.90	112.22	SOURCE3_SOURCE5		c3-os-c3-na	1	0.383	0.000	-3.000
2726	1.5103				c3-os-c3-na	1	0.650	0.000	2.000
cc-Y1-cc	71.76	105.49	CORR_SOURCE5		c3-os-c3-h2	1	0.383	0.000	3.000
1810	1.9032				c3-c3-oh-ho	1	0.000	0.000	3.000
Y1-c3-h2	61.963	107.310			c3-c3-c3-hc	1	0.080	0.000	3.000
DIHE					os-c3-c3-oh	1	1.010	0.000	-3.000
p5-os-c3-c3	1	0.383	0.000	-3.000	os-c3-c3-oh	1	0.000	0.000	-2.000
p5-os-p5-o	1	0.800	0.000	2.000	os-c3-c3-oh	1	0.020	180.000	1.000
p5-os-p5-os	1	0.800	0.000	2.000	os-c3-c3-hc	1	0.000	0.000	-3.000
os-p5-os-c3	1	0.000	0.000	-3.000	os-c3-c3-hc	1	0.250	0.000	1.000
os-p5-os-c3	1	2.610	0.000	2.000	os-c3-na-cc	1	0.000	0.000	2.000
p5-os-c3-c3	1	3.950	180.000	1.000	os-c3-Y1-cc	1	0.000	0.000	2.000
p5-os-c3-h1	1	0.217	0.000	3.000	os-c3-Y1-M1	1	0.000	0.000	2.000
o -p5-os-c3	1	0.800	0.000	-2.000	c3-c3-c3-na	1	0.156	0.000	3.000
o -p5-os-c3	1	0.800	0.000	-2.000					
o -p5-os-p5	1	0.550	0.000	3.000					

c3-c3-c3-h2	1	0.156	0.000	3.000	n -c -cd-cc	1	2.875	180.000	2.000	
oh-c3-c3-h1	1	0.000	0.000	-3.000	n -cd-nh-hn	1	1.050	180.000	2.000	
oh-c3-c3-h1	1	0.250	0.000	1.000	n -cd-nc-cc	1	4.750	180.000	2.000	
oh-c3-c3-hc	1	0.180	0.000	-3.000	nh-cd-n -hn	1	1.650	180.000	2.000	
oh-c3-c3-hc	1	0.510	0.000	1.000	nh-cd-nc-cc	1	4.750	180.000	2.000	
c3-c3-na-cc	1	0.000	0.000	2.000	nc-cd-n -hn	1	1.650	180.000	2.000	
c3-os-c3-h1	1	0.337	0.000	3.000	nc-cd-nh-hn	1	1.050	180.000	2.000	
#c3-na-cc-nd	1	1.700	180.000	2.000	cc-na-cc-h5	1	1.700	180.000	2.000	
c3-na-cc-h5	1	1.700	180.000	2.000	cc-Y1-c3-h2	1	1.700	180.000	2.000	
c3-na-cc-cd	1	1.700	180.000	2.000	h1-c3-c3-h1	1	0.156	0.000	3.000	
c3-na-cc-nc	1	1.700	180.000	2.000	h1-c3-oh-ho	1	0.113	0.000	3.000	
na-c3-c3-hc	1	0.156	0.000	3.000	h1-c3-c3-hc	1	0.156	0.000	3.000	
na-cc-cd-c	1	4.000	180.000	2.000	hc-c3-c3-h2	1	0.156	0.000	3.000	
na-cc-nc-cd	1	4.750	180.000	2.000	X -Y1-cd-X	2	9.5	180.0	2.0	statistiv value from parm94
cc-na-c3-h2	1	0.000	0.000	2.000	X -c3-Y2-X	6	1.8	0.0	3.0	Junmei et al, 1999
cc-na-cc-cd	1	1.700	180.000	2.000	X -c3-Y3-X	6	1.8	0.0	3.0	Junmei et al, 1999
cc-na-cc-nc	1	1.700	180.000	2.000	X -c3-Y4-X	6	1.8	0.0	3.0	Junmei et al, 1999
nd-cd-cc-nc	1	4.000	180.000	2.000	X -cc-Y1-X	2	9.5	180.0	2.0	statistic value from parm94
cd-c -n -cd	1	2.500	180.000	2.000	M1-Y1-cd-c	3	0.00	0.00	3.0	Treat as zero by MCPB.py
cd-c -n -hn	1	2.500	180.000	2.000	M1-Y1-cd-cc	3	0.00	0.00	3.0	Treat as zero by MCPB.py
cd-cc-nc-cd	1	4.750	180.000	2.000	Y1-M1-Y2-c3	3	0.00	0.00	3.0	Treat as zero by MCPB.py
c -cd-cc-nc	1	4.000	180.000	2.000	Y1-M1-Y2-hn	3	0.00	0.00	3.0	Treat as zero by MCPB.py
c -n -cd-nh	1	1.650	180.000	2.000						
c -n -cd-nc	1	1.650	180.000	2.000						
o -c -cd-cc	1	2.875	180.000	2.000						
o -c -n -cd	1	2.500	180.000	2.000						
o -c -n -hn	1	2.500	180.000	-2.000						
o -c -n -hn	1	2.000	0.000	1.000						

Y1-M1-Y3-c3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-na-M1-Y4 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y1-M1-Y3-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-na-M1-Y2 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y1-M1-Y4-c3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-Y1-M1-Y3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y1-M1-Y4-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-Y2-M1-Y4 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y2-M1-Y1-cd by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-Y3-M1-Y2 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y2-M1-Y3-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-Y3-M1-Y4 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y2-M1-Y4-c3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y2-M1 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y2-M1-Y4-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y3-M1 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y3-M1-Y1-cd by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y4-M1 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y3-M1-Y2-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y1-M1 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y3-M1-Y4-c3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y4-c3	1	0.3	0.0	-3.0	Junmei et al, 1999
Y3-M1-Y4-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y4-c3	1	0.48	180.0	2.0	Junmei et al, 1999
Y4-M1-Y1-cd by MCPB.py	3	0.00	0.00	3.0	Treat as zero	c3-c3-Y1-cc	1	0.48	180.0	2.0	Junmei et al, 1999
Y4-M1-Y2-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	cc-Y1-M1-Y2 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
Y4-M1-Y3-hn by MCPB.py	3	0.00	0.00	3.0	Treat as zero	cc-Y1-M1-Y3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
c3-Y2-M1-Y3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero	cc-Y1-M1-Y4 by MCPB.py	3	0.00	0.00	3.0	Treat as zero
c3-na-M1-Y3 by MCPB.py	3	0.00	0.00	3.0	Treat as zero	h1-c3-Y2-M1 by MCPB.py	3	0.00	0.00	3.0	Treat as zero

h1-c3-Y3-M1	3	0.00	0.00	3.0	Treat as zero	cd-n -c -o	10.5	180.0	2.0	General
by MCPB.py						improper torsional angle (2 general atom types)				
h1-c3-Y4-M1	3	0.00	0.00	3.0	Treat as zero	c -cd-n -hn	1.1	180.0	2.0	General
by MCPB.py						improper torsional angle (2 general atom types)				
h2-c3-Y1-M1	3	0.00	0.00	3.0	Treat as zero	n -nc-cd-nh	1.1	180.0	2.0	Using
by MCPB.py						default value				
h5-cc-Y1-M1	3	0.00	0.00	3.0	Treat as zero	cd-hn-nh-hn	1.1	180.0	2.0	Using
by MCPB.py						default value				
na-cc-Y1-M1	3	0.00	0.00	3.0	Treat as zero	cd-na-cc-nc	1.1	180.0	2.0	Using
by MCPB.py						default value				
na-cc-cd-Y1	3	0.00	0.00	3.0		Y1-h5-cc-na	1.1	180.0	2.0	Using
Y1-cc-na-cc	3	0.00	0.00	3.0		default value				
Y1-cd-c-o	3	0.00	0.00	3.0		Y1-c -cd-cc	1.1	180.0	2.0	Using
Y1-cd-c-n	3	0.00	0.00	3.0		default value				
Y1-cd-cc-nc	3	0.00	0.00	3.0						
Y2-c3-c3-Y4	3	0.00	0.00	3.0		NONBON				
Y2-c3-c3-h1	3	0.00	0.00	3.0		p5	2.0732	0.2295		
Y2-c3-c3-h1	3	0.00	0.00	3.0		o	1.7107	0.1463		
Y2-c3-c3-h1	3	0.00	0.00	3.0		os	1.7713	0.0726		
Y3-c3-c3-Y4	3	0.00	0.00	3.0		c3	1.9069	0.1078		
Y3-c3-c3-h1	3	0.00	0.00	3.0		oh	1.8200	0.0930		
Y3-c3-c3-h1	3	0.00	0.00	3.0		na	1.7992	0.2042		
Y4-c3-c3-h1	3	0.00	0.00	3.0		cc	1.8606	0.0988		
Y4-c3-c3-h1	3	0.00	0.00	3.0		cd	1.8606	0.0988		
Y4-c3-c3-h1	3	0.00	0.00	3.0		c	1.8606	0.0988		
Y4-c3-c3-h1	3	0.00	0.00	3.0		n	1.7852	0.1636		
c3-na-cc-Y1	3	0.00	0.00	3.0		nh	1.7903	0.2150		
						nc	1.8993	0.0941		
IMPROPER						h1	1.3593	0.0208		
c3-cc-na-cc	1.1	180.0	2.0	Using		hc	1.4593	0.0208		
default value										

h2	1.2593	0.0208	
h5	1.3735	0.0161	
hn	0.6210	0.0100	
ho	0.3019	0.0047	
M1	1.2660	0.0030764200	CM set for Pt2+ ion in TIP3P water from Li et al. JCTC, 2013, 9, 2733
Y1	1.8240	0.1700	OPLS
Y2	1.8240	0.1700	OPLS
Y3	1.8240	0.1700	OPLS
Y4	1.8240	0.1700	OPLS

Parameters complex 2

File off:

!!index array str

"MOL"

!entry.MOL.unit.atoms table str name str type int typex
int resx int flags int seq int elmnt dbl chg

"P1" "p5" 0 1 131072 1 15 1.125285

"O1" "o" 0 1 131072 2 8 -0.754291

"O2" "oh" 0 1 131072 3 8 -0.604601

"O3" "o" 0 1 131072 4 8 -0.754291

"O4" "os" 0 1 131072 5 8 -0.418199

"P2" "p5" 0 1 131072 6 15 1.098280

"O5" "o" 0 1 131072 7 8 -0.716928

"O6" "o" 0 1 131072 8 8 -0.716928

"O7" "os" 0 1 131072 9 8 -0.469240

"P3" "p5" 0 1 131072 10 15 1.333113

"O8" "oh" 0 1 131072 11 8 -0.813652

"O9" "o" 0 1 131072 12 8 -0.743769

"O10" "os" 0 1 131072 13 8 -0.393679

"C1" "c3" 0 1 131072 14 6 -0.177885

"C2" "c3" 0 1 131072 15 6 0.367282

"O11" "os" 0 1 131072 16 8 -0.527228

"C3" "c3" 0 1 131072 17 6 0.531252

"O12" "oh" 0 1 131072 18 8 -0.773339

"C4" "c3" 0 1 131072 19 6 -0.304723

"C5" "c3" 0 1 131072 20 6 0.244592

"N1" "na" 0 1 131072 21 7 -0.241174

"C6" "c2" 0 1 131072 22 6 0.328615

"N2" "Y1" 0 1 131072 23 7 -0.363914

"C7" "c2" 0 1 131072 24 6 -0.322051

"C8" "c2" 0 1 131072 25 6 0.877511

"O13" "o" 0 1 131072 26 8 -0.665214

"N3" "ns" 0 1 131072 27 7 -0.828823

"C9" "c2" 0 1 131072 28 6 0.997780

"N4" "nv" 0 1 131072 29 7 -0.993401

"N5" "n2" 0 1 131072 30 7 -0.797741

"C10" "c2" 0 1 131072 31 6 0.581560

"N6" "Y2" 0 1 131072 32 7 -0.944276

"N7" "Y3" 0 1 131072 33 7 -1.215885

"H1" "h1" 0 1 131072 34 1 0.107442

"H2" "h1" 0 1 131072 35 1 0.107442

"H3" "h1" 0 1 131072 36 1 -0.013563

"H4" "h1" 0 1 131072 37 1 -0.057286

"H5" "hc" 0 1 131072 38 1 0.095972

"H6" "hc" 0 1 131072 39 1 0.095972

"H7" "h2" 0 1 131072 40 1 0.092949

"H8" "h5" 0 1 131072 41 1 0.192431	"O9" "o" 0 -1 0.0
"H9" "hn" 0 1 131072 42 1 0.417336	"O10" "os" 0 -1 0.0
"H10" "ho" 0 1 131072 43 1 0.342319	"C1" "c3" 0 -1 0.0
"H11" "hn" 0 1 131072 44 1 0.399656	"C2" "c3" 0 -1 0.0
"H12" "hn" 0 1 131072 45 1 0.399656	"O11" "os" 0 -1 0.0
"H13" "hn" 0 1 131072 46 1 0.392561	"C3" "c3" 0 -1 0.0
"H14" "hn" 0 1 131072 47 1 0.392561	"O12" "oh" 0 -1 0.0
"H15" "ho" 0 1 131072 48 1 0.436381	"C4" "c3" 0 -1 0.0
"PT" "M1" 0 1 131072 49 -1 0.531600	"C5" "c3" 0 -1 0.0
"H16" "hn" 0 1 131072 50 1 0.392561	"N1" "na" 0 -1 0.0
"H17" "hn" 0 1 131072 51 1 0.363593	"C6" "c2" 0 -1 0.0
"H18" "hn" 0 1 131072 52 1 0.363593	"N2" "Y2" 0 -1 0.0
"O14" "Y4" 0 1 131072 53 8 -1.039336	"C7" "c2" 0 -1 0.0
"H19" "ho" 0 1 131072 54 1 0.472268	"C8" "c2" 0 -1 0.0
"H20" "ho" 0 1 131072 55 1 0.569852	"O13" "o" 0 -1 0.0
!entry.MOL.unit.atomsptinfo table str pname str	"N3" "ns" 0 -1 0.0
pctype int ptypex int pelmnt dbl pchg	"C9" "c2" 0 -1 0.0
"P1" "p5" 0 -1 0.0	"N4" "nv" 0 -1 0.0
"O1" "o" 0 -1 0.0	"N5" "n2" 0 -1 0.0
"O2" "oh" 0 -1 0.0	"C10" "c2" 0 -1 0.0
"O3" "o" 0 -1 0.0	"N6" "Y2" 0 -1 0.0
"O4" "os" 0 -1 0.0	"N7" "Y3" 0 -1 0.0
"P2" "p5" 0 -1 0.0	"H1" "h1" 0 -1 0.0
"O5" "o" 0 -1 0.0	"H2" "h1" 0 -1 0.0
"O6" "o" 0 -1 0.0	"H3" "h1" 0 -1 0.0
"O7" "os" 0 -1 0.0	"H4" "h1" 0 -1 0.0
"P3" "p5" 0 -1 0.0	"H5" "hc" 0 -1 0.0
"O8" "oh" 0 -1 0.0	"H6" "hc" 0 -1 0.0

"H7" "h2" 0 -1 0.0	!entry.MOL.unit.connectivity table int atom1x int
"H8" "h5" 0 -1 0.0	atom2x int flags
"H9" "hn" 0 -1 0.0	1 2 1
"H10" "ho" 0 -1 0.0	1 3 1
"H11" "hn" 0 -1 0.0	1 4 1
"H12" "hn" 0 -1 0.0	1 5 1
"H13" "hn" 0 -1 0.0	5 6 1
"H14" "hn" 0 -1 0.0	6 7 1
"H15" "ho" 0 -1 0.0	6 8 1
"PT" "M1" 0 -1 0.0	6 9 1
"H16" "hn" 0 -1 0.0	9 10 1
"H17" "hn" 0 -1 0.0	10 11 1
"H18" "hn" 0 -1 0.0	10 12 1
"O14" "Y4" 0 -1 0.0	10 13 1
"H19" "ho" 0 -1 0.0	13 14 1
"H20" "ho" 0 -1 0.0	14 15 1
!entry.MOL.unit.boundbox array dbl	14 34 1
-1.000000	14 35 1
0.0	15 16 1
0.0	15 17 1
0.0	15 36 1
0.0	16 20 1
!entry.MOL.unit.childsequence single int	17 18 1
2	17 19 1
!entry.MOL.unit.connect array int	17 37 1
0	18 48 1
0	19 20 1
	19 38 1

19 39 1	49 53 1
20 21 1	53 54 1
20 40 1	53 55 1
21 22 1	!entry.MOL.unit.hierarchy table str abovetype int
21 31 1	abovex str belowtype int belowx
22 23 1	"U" 0 "R" 1
22 41 1	"R" 1 "A" 1
23 24 1	"R" 1 "A" 2
23 49 1	"R" 1 "A" 3
24 25 1	"R" 1 "A" 4
24 31 1	"R" 1 "A" 5
25 26 1	"R" 1 "A" 6
25 27 1	"R" 1 "A" 7
27 28 1	"R" 1 "A" 8
27 42 1	"R" 1 "A" 9
28 29 1	"R" 1 "A" 10
28 30 1	"R" 1 "A" 11
29 44 1	"R" 1 "A" 12
29 45 1	"R" 1 "A" 13
30 31 1	"R" 1 "A" 14
32 46 1	"R" 1 "A" 15
32 47 1	"R" 1 "A" 16
32 50 1	"R" 1 "A" 17
32 49 1	"R" 1 "A" 18
33 51 1	"R" 1 "A" 19
33 52 1	"R" 1 "A" 20
33 43 1	"R" 1 "A" 21
33 49 1	"R" 1 "A" 22

"R" 1 "A" 23	"R" 1 "A" 51
"R" 1 "A" 24	"R" 1 "A" 52
"R" 1 "A" 25	"R" 1 "A" 53
"R" 1 "A" 26	"R" 1 "A" 54
"R" 1 "A" 27	"R" 1 "A" 55
"R" 1 "A" 28	!entry.MOL.unit.name single str
"R" 1 "A" 29	"MOL"
"R" 1 "A" 30	!entry.MOL.unit.positions table dbl x dbl y dbl z
"R" 1 "A" 31	0.636000 -0.359000 2.908000
"R" 1 "A" 32	-0.713000 -0.039000 2.264000
"R" 1 "A" 33	1.141000 -1.828000 2.340000
"R" 1 "A" 34	0.783000 -0.328000 4.394000
"R" 1 "A" 35	1.714000 0.674000 2.140000
"R" 1 "A" 36	3.256000 0.511000 1.609000
"R" 1 "A" 37	3.611000 1.842000 0.971000
"R" 1 "A" 38	4.165000 -0.191000 2.552000
"R" 1 "A" 39	2.913000 -0.581000 0.313000
"R" 1 "A" 40	3.154000 -0.406000 -1.269000
"R" 1 "A" 41	2.505000 -1.690000 -1.894000
"R" 1 "A" 42	4.540000 -0.100000 -1.705000
"R" 1 "A" 43	2.090000 0.784000 -1.710000
"R" 1 "A" 44	2.580000 2.085000 -2.029000
"R" 1 "A" 45	1.574000 3.147000 -1.630000
"R" 1 "A" 46	0.317000 2.995000 -2.364000
"R" 1 "A" 47	1.182000 3.197000 -0.140000
"R" 1 "A" 48	2.091000 3.923000 0.634000
"R" 1 "A" 49	-0.199000 3.857000 -0.225000
"R" 1 "A" 50	-0.766000 3.239000 -1.506000

-1.459000 1.949000 -1.224000	-0.909000 -2.200000 -0.671000
-0.957000 0.693000 -1.440000	-2.772000 -2.296000 1.167000
-1.747000 -0.246000 -0.949000	0.410000 -3.927000 0.567000
-2.819000 0.420000 -0.384000	0.783000 -4.012000 -0.974000
-3.998000 -0.044000 0.287000	0.112000 -1.963000 -2.462000
-4.421000 -1.172000 0.527000	0.080000 -2.824000 -2.899000
-4.785000 1.072000 0.715000	1.456000 -1.750000 -2.108000
-4.486000 2.395000 0.534000	!entry.MOL.unit.residueconnect table int c1x int c2x
-5.409000 3.314000 1.038000	int c3x int c4x int c5x int c6x
-3.432000 2.813000 -0.110000	0 0 0 0 0 0
-2.638000 1.796000 -0.549000	!entry.MOL.unit.residues table str name int seq int
-1.753000 -2.311000 1.216000	childseq int startatomx str restype int imagingx
-0.033000 -4.003000 -0.355000	"MOL" 1 56 1 "?" 0
2.778000 2.141000 -3.109000	!entry.MOL.unit.residuesPdbSequenceNumber array int
3.508000 2.288000 -1.487000	0
2.003000 4.126000 -1.897000	!entry.MOL.unit.solventcap array dbl
1.073000 2.175000 0.241000	-1.000000
-0.076000 4.938000 -0.367000	0.0
-0.827000 3.682000 0.650000	0.0
-1.506000 3.871000 -2.012000	0.0
-0.004000 0.519000 -1.914000	0.0
-5.644000 0.802000 1.176000	!entry.MOL.unit.velocities table dbl x dbl y dbl z
1.464000 -1.735000 1.427000	0.0 0.0 0.0
-5.010000 4.245000 0.991000	0.0 0.0 0.0
-5.711000 3.103000 1.983000	0.0 0.0 0.0
-1.442000 -3.150000 1.699000	0.0 0.0 0.0
-1.392000 -1.458000 1.735000	0.0 0.0 0.0
2.743000 3.224000 0.931000	0.0 0.0 0.0

[illegible]

0.0 0.0 0.0
0.0 0.0 0.0
0.0 0.0 0.0
0.0 0.0 0.0
0.0 0.0 0.0
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0.0 0.0 0.0
0.0 0.0 0.0
0.0 0.0 0.0
File frcmoc
remark gY4
MASS
p5 30.970
oh 16.000
os 16.000
c3 12.010
na 14.010

c2 12.010	0.360		c3-h2 377.33	1.096	
n2 14.010	0.530		na-c2 327.66	1.401	
ns 14.010	0.530		c2-n2 518.67	1.282	
nv 14.010	0.530		c2-h5 386.13	1.091	
n9 14.010	0.530		c2-c2 481.83	1.334	
h1 1.008	0.135		c2-Y4 635.23	1.225	
hc 1.008	0.135		c2-ns 330.19	1.399	same as c2-n
h2 1.008	0.135		ns-hn 527.31	1.013	same as hn-n
h5 1.008	0.135		c2-nv 345.39	1.387	same as c2-nh
hn 1.008	0.161		nv-hn 529.46	1.012	same as hn-nh
ho 1.008	0.135		n9-hn 511.28	1.019	same as hn-n3
M1 195.08		Pt iY4n	n2-hn 501.09	1.023	
Y1 14.010	0.530		Y4 -ho 540.28	0.981	
Y2 14.010	0.530		Y1-M1 80.3	2.1471	Created by Seminar iY4 method using MCPB.py
Y3 14.010	0.530		Y2-M1 80.3	2.1471	Created by Seminar iY4 method using MCPB.py
Y4 16.000	0.434		Y3-M1 80.0	2.1471	Created by Seminar iY4 method using MCPB.py
BOND			Y4-M1 78.6	2.3471	Created by Seminar iY4 method using MCPB.py
p5-Y4 529.55	1.487		Y4 -M1 78.6	2.6178	Created by Seminario method using MCPB.py
p5-oh 346.03	1.615		Y2-hn 511.28	1.019	same as hn-n3
p5-os 346.25	1.615		Y3-hn 501.09	1.023	
oh-ho 563.51	0.973		Y3-ho 501.09	1.023	
os-c3 284.76	1.432		Y4-ho 540.28	0.981	
c3-c3 232.52	1.538		c2-Y1 518.67	1.282	
c3-h1 375.92	1.097		oh-M1 81.6	2.6178	
c3-oh 293.40	1.423				
c3-hc 375.92	1.097				
c3-na 262.85	1.463				

ANGLE			na-c2-h5	59.208	126.390	
p5-oh-ho	58.997	110.080	na-c2-c2	87.232	121.940	
p5-oh-M1	58.997	110.080	c2-na-c2	69.273	110.370	
p5-Y4 -M1	58.997	110.080	c2-n2-c2	73.202	118.180	
p5-os-p5	106.787	126.250	n2-c2-h5	64.543	121.700	
Y4 -p5-oh	81.901	115.210	n2-c2-c2	89.778	126.010	
Y4 -p5-o	85.510	115.800	c2-c2-Y4	89.793	130.890	
Y4 -p5-os	81.819	115.460	c2-c2-ns	86.693	123.670	same as c2-c2-n
oh-p5-os	83.902	101.940	c2-c2-c2	71.539	121.810	
os-p5-os	83.949	101.840	c2-ns-c2	67.455	116.750	same as c2-n -c2
p5-os-c3	83.251	119.540	c2-ns-hn	47.988	117.900	same as c2-n -hn
os-c3-c3	85.306	107.970	Y4 -c2-ns	75.107	117.460	Calculated with empirical approach
os-c3-h1	62.377	109.780	ns-c2-nv	113.384	109.350	same as n -c2-nh
c3-c3-c3	64.888	111.510	ns-c2-n2	110.772	122.820	same as n -c2-n2
c3-c3-h1	46.868	109.560	c2-nv-hn	48.954	115.090	same as c2-nh-hn
c3-os-c3	66.293	112.480	nv-c2-n2	110.708	124.270	same as n2-c2-nh
c3-c3-oh	84.642	110.190	h1-c3-h1	38.802	108.460	
os-c3-na	109.296	109.030	hc-c3-hc	38.960	107.580	
o -c2-ns	109.296	109.030	hn-nv-hn	39.519	115.120	same as hn-nh-hn
os-c3-h2	62.442	109.580	hn-n9-hn	40.828	106.400	same as hn-n3-hn
c3-oh-ho	49.027	107.260	hn-n2-hn	38.294	120.000	
c3-c3-hc	46.816	109.800	M1-Y2-hn	67.55	104.19	Created by SeminariY4 method using MCPB.py
oh-c3-h1	62.540	110.260	M1-Y3-hn	67.39	104.19	Created by SeminariY4 method using MCPB.py
c3-c3-na	82.668	112.880	M1-Y3-ho	67.39	104.19	
c3-c3-h2	46.730	110.220	M1-Y4-ho	50.01	91.15	Created by Seminario method using MCPB.py
c3-na-c2	65.723	117.200				
na-c3-h2	61.963	107.310				
na-c2-n2	110.312	123.620				

Y1-M1-Y2 182.00 90.94 Created by SeminariY4
method using MCPB.py

Y1-M1-Y3 182.48 180.46 Created by SeminariY4
method using MCPB.py

Y1-M1-Y4 182.50 89.61 Created by SeminariY4
method using MCPB.py

Y2-M1-Y3 182.00 87.62 Created by SeminariY4
method using MCPB.py

Y2-M1-Y4 182.43 183.80 Created by SeminariY4
method using MCPB.py

Y3-M1-Y4 182.71 96.43 Created by SeminariY4
method using MCPB.py

oh-M1-Y2 180.43 83.80 Created by Seminario
method using MCPB.py

oh-M1-Y3 182.43 83.80 Created by Seminario
method using MCPB.py

oh-M1-Y1 182.43 93.80 Created by Seminario
method using MCPB.py

oh-M1-Y4 182.71 86.43 Created by Seminario
method using MCPB.py

ho -oh-M1 72.71 86.43 Created by Seminario
method using MCPB.py

Y4 -M1-oh 72.71 76.43 Created by Seminario
method using MCPB.py

c2-Y1-M1 11.11 120.68 Created by SeminariY4
method using MCPB.py

Y1-c2-c2 89.778 126.010

c2-Y1-c2 73.202 118.180

h5-c2-Y1 52.26 121.70 SOURCE4_SOURCE5
71 2.1538

hn-Y2-hn 40.828 106.400 same as hn-n3-hn

hn-Y3-hn 38.294 120.000

ho-Y3-hn 38.294 120.000

na-c2-Y1 110.312 123.620

ho-Y4-ho 38.294 120.000

DIHE

p5-os-p5-o 1 0.800 0.000 2.000

p5-os-p5-os 1 0.800 0.000 2.000

Y4 -p5-oh-ho 1 0.367 0.000 3.000

oh-p5-os-p5 1 0.800 0.000 2.000

os-p5-oh-ho 1 0.533 0.000 3.000

os-p5-os-c3 1 0.000 0.000 -3.000

os-p5-os-c3 1 2.610 0.000 2.000

p5-os-c3-c3 1 0.383 0.000 -3.000

p5-os-c3-c3 1 3.950 180.000 1.000

p5-os-c3-h1 1 0.217 0.000 3.000

oh-p5-os-c3 1 0.250 0.000 -3.000

oh-p5-os-c3 1 1.200 0.000 2.000

Y4 -p5-os-c3 1 0.800 0.000 -2.000

Y4 -p5-os-c3 1 0.550 0.000 3.000

os-c3-c3-os 1 0.000 0.000 -3.000

os-c3-c3-os 1 0.000 180.000 -2.000

os-c3-c3-os 1 0.170 180.000 1.000

os-c3-c3-c3 1 0.156 0.000 3.000

os-c3-c3-h1 1 0.000 0.000 -3.000

os-c3-c3-h1 1 0.250 0.000 1.000

c3-c3-os-c3 1 0.240 0.000 -3.000

c3-c3-os-c3 1 0.160 0.000 2.000

c3-c3-c3-oh 1 0.210 0.000 3.000

c3-c3-c3-c3 1 0.130 0.000 -3.000

c3-c3-c3-c3	1	0.290	180.000	-2.000	na-c2-c2-n2	1	6.650	180.000	2.000	
c3-c3-c3-c3	1	0.110	0.000	1.000	na-c2-c2-c2	1	6.650	180.000	2.000	
c3-c3-c3-h1	1	0.156	0.000	3.000	c2-na-c3-h2	1	0.000	0.000	2.000	
c3-os-c3-na	1	0.383	0.000	-3.000	c2-na-c2-c2	1	0.625	180.000	2.000	
c3-os-c3-na	1	0.650	0.000	2.000	c2-na-c2-n2	1	0.625	180.000	2.000	
c3-os-c3-h2	1	0.383	0.000	3.000	c2-n2-c2-c2	1	4.150	180.000	2.000	
c3-c3-oh-ho	1	0.000	0.000	3.000	n2-c2-c2-Y4	1	6.650	180.000	2.000	
c3-c3-c3-hc	1	0.080	0.000	3.000	n2-c2-c2-ns	1	6.650	180.000	2.000	
os-c3-c3-oh	1	1.010	0.000	-3.000	n2-c2-c2-n2	1	6.650	180.000	2.000	
os-c3-c3-oh	1	0.000	0.000	-2.000	c2-n2-c2-h5	1	4.150	180.000	2.000	
os-c3-c3-oh	1	0.020	180.000	1.000	c2-c2-ns-c2	1	0.650	180.000	2.000	same as X -c2-n -X
os-c3-c3-hc	1	0.000	0.000	-3.000	c2-c2-ns-hn	1	0.650	180.000	2.000	same as X -c2-n -X
os-c3-c3-hc	1	0.250	0.000	1.000	c2-c2-c2-n2	1	6.650	180.000	2.000	
os-c3-na-c2	1	0.000	0.000	-2.000	c2-ns-c2-nv	1	0.650	180.000	2.000	same as X -c2-n -X
os-c3-na-c2	1	2.500	0.000	1.000	c2-ns-c2-n2	1	0.650	180.000	2.000	same as X -c2-n -X
c3-c3-c3-na	1	0.156	0.000	3.000	Y4 -c2-c2-c2	1	6.650	180.000	2.000	
c3-c3-c3-h2	1	0.156	0.000	3.000	Y4 -c2-ns-c2	1	0.650	180.000	2.000	same as X -c2-n -X
oh-c3-c3-h1	1	0.000	0.000	-3.000	Y4 -c2-ns-hn	1	0.650	180.000	2.000	same as X -c2-n -X
oh-c3-c3-h1	1	0.250	0.000	1.000	ns-c2-c2-c2	1	6.650	180.000	2.000	
oh-c3-c3-hc	1	0.180	0.000	-3.000	ns-c2-nv-hn	1	0.675	180.000	2.000	same as X -c2-nh-X
oh-c3-c3-h	1	0.510	0.000	1.000	ns-c2-n2-c2	1	4.150	180.000	2.000	
c3-c3-na-c2	1	0.000	0.000	2.000	nv-c2-ns-hn	1	0.650	180.000	2.000	same as X -c2-n -X
c3-os-c3-h1	1	0.337	0.000	3.000	nv-c2-n2-c2	1	4.150	180.000	2.000	
c3-na-c2-n2	1	0.625	180.000	2.000						
c3-na-c2-h5	1	0.625	180.000	2.000						
c3-na-c2-c2	1	0.625	180.000	2.000						
na-c3-c3-hc	1	0.156	0.000	3.000						
na-c2-n2-c2	1	4.150	180.000	2.000						

n2-c2-ns-hn	1	0.650	180.000	2.000	same	Y2-M1-Y4-ho	3	0.00	0.00	3.0	Treat as zero by MCPB.py
as X -c2-n -X											
n2-c2-nv-hn	1	0.675	180.000	2.000	same	Y3-M1-Y1-c2	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
as X -c2-nh-X											
c2-na-c2-h5	1	0.625	180.000	2.000		Y3-M1-Y2-hn	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
h1-c3-c3-h1	1	0.156	0.000	3.000		Y3-M1-Y4-ho	3	0.00	0.00	3.0	Treat as zero by MCPB.py
h1-c3-oh-ho	1	0.113	0.000	3.000		Y4-M1-Y1-c2	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
h1-c3-c3-hc	1	0.156	0.000	3.000		Y4-M1-Y2-hn	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
hc-c3-c3-h2	1	0.156	0.000	3.000		Y4-M1-Y3-hn	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
M1-Y1-c2-c2	3	0.00	0.00	3.0	Treat as	c2-Y1-M1-Y2	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
zerY4 by MCPB.py						c2-Y1-M1-Y3	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
Y1-M1-Y2-hn	3	0.00	0.00	3.0	Treat as	c2-Y1-M1-Y4	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
zerY4 by MCPB.py						c2-Y1-c2-c2	1	4.15	180.0	2.0	
oh-M1-Y2-hn	3	0.00	0.00	3.0	Treat as	c2-na-c2-Y1	1	0.625	180.0	2.0	
zero by MCPB.py						c3-na-c2-Y1	1	0.625	180.0	2.0	
oh-M1-Y3-hn	3	0.00	0.00	3.0	Treat as	h5-c2-Y1-M1	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
zero by MCPB.py						h5-c2-Y1-c2	1	4.15	180.0	2.0	
oh-M1-Y1-c2	3	0.00	0.00	3.0	Treat as	na-c2-Y1-M1	3	0.00	0.00	3.0	Treat as zerY4 by MCPB.py
zero by MCPB.py						na-c2-Y1-c2	1	4.15	180.0	2.0	
oh-M1-Y4-ho	3	0.00	0.00	3.0	Treat as	na-c2-c2-Y1	1	6.65	180.0	2.0	
zero by MCPB.py						ho-Y3-M1-Y4	3	0.00	0.00	3.0	
Y1-M1-Y3-hn	3	0.00	0.00	3.0	Treat as	Y2-M1-Y3-ho	3	0.00	0.00	3.0	
zerY4 by MCPB.py											
Y1-M1-Y4-ho	3	0.00	0.00	3.0	Treat as						
zero by MCPB.py											
Y1-c2-c2-n2	1	6.65	180.0	2.0							
Y1-c2-c2-ns	1	6.65	180.0	2.0							
Y1-c2-c2-Y4	1	6.65	180.0	2.0							
Y2-M1-Y1-c2	3	0.00	0.00	3.0	Treat as						
zerY4 by MCPB.py											
Y2-M1-Y3-hn	3	0.00	0.00	3.0	Treat as						
zerY4 by MCPB.py											

Y1-M1-Y3-ho	3	0.00	0.00	3.0		na	1.7992	0.2042	
o -c2-ns-c2	3	0.00	0.00	3.0		c2	1.8606	0.0988	
o -c2-ns-hn	3	0.00	0.00	3.0		n2	1.8993	0.0941	
						ns	1.8352	0.1174	
IMPROPER						nv	1.8903	0.1120	
c2-c2-na-c3		1.1	180.0	2.0		n9	2.2700	0.0095	
h5-n2-c2-na		1.1	180.0	2.0	Using	h1	1.3593	0.0208	
default value						hc	1.4593	0.0208	
c2-c2-c2-n2		1.1	180.0	2.0	Using	h2	1.2593	0.0208	
default value						h5	1.3735	0.0161	
c2-ns-c2-Y4		1.1	180.0	2.0	Using	hn	0.6210	0.0100	
default value						ho	0.3019	0.0047	
c2-c2-ns-hn		1.1	180.0	2.0	Using	M1	1.2660	0.0030764200	CM set fY4r Pt2+
default value						ion in TIP3P water from Li et al. JCTC, 2013, 9, 2733			
n2-ns-c2-nv		1.1	180.0	2.0	Using	Y1	1.8993	0.0941	
default value						Y2	2.2700	0.0095	
c2-hn-nv-hn		1.1	180.0	2.0	Using	Y3	1.8993	0.0941	
default value						Parameters complex 3			
c2-n2-c2-na		1.1	180.0	2.0	Using	<u>File off:</u>			
default value						!!index array str			
Y1-c2-c2-c2		1.1	180.0	2.0	Using	"MOL"			
default value						!entry.MOL.unit.atoms table str name str type int typex			
Y1-h5-c2-na		1.1	180.0	2.0	Using	int resx int flags int seq int elmnt dbl chg			
default value						"P1" "p5" 0 1 131072 1 15 1.342234			
NONBON						"O1" "o" 0 1 131072 2 8 -0.739517			
p5		2.0732	0.2295			"O2" "oh" 0 1 131072 3 8 -0.845220			
Y4		1.7107	0.1463			"O3" "o" 0 1 131072 4 8 -0.845220			
oh		1.8200	0.0930			"O4" "os" 0 1 131072 5 8 -0.592939			
os		1.7713	0.0726			"P2" "p5" 0 1 131072 6 15 1.252046			
c3		1.9069	0.1078						

"O5" "o" 0 1 131072 7 8 -0.795894	"H2" "h1" 0 1 131072 35 1 0.067284
"O6" "o" 0 1 131072 8 8 -0.795894	"H3" "h1" 0 1 131072 36 1 -0.012167
"O7" "os" 0 1 131072 9 8 -0.476918	"H4" "h1" 0 1 131072 37 1 0.098376
"P3" "p5" 0 1 131072 10 15 1.117531	"H5" "hc" 0 1 131072 38 1 0.118713
"O8" "oh" 0 1 131072 11 8 -0.750455	"H6" "hc" 0 1 131072 39 1 0.118713
"O9" "o" 0 1 131072 12 8 -0.750455	"H7" "h2" 0 1 131072 40 1 0.076893
"O10" "os" 0 1 131072 13 8 -0.367908	"H8" "h5" 0 1 131072 41 1 0.178019
"C1" "c3" 0 1 131072 14 6 -0.063722	"H9" "hn" 0 1 131072 42 1 0.418070
"C2" "c3" 0 1 131072 15 6 0.200863	"H10" "hn" 0 1 131072 43 1 0.292595
"O11" "os" 0 1 131072 16 8 -0.513040	"H11" "hn" 0 1 131072 44 1 0.373510
"C3" "c3" 0 1 131072 17 6 0.741353	"H12" "hn" 0 1 131072 45 1 0.373510
"O12" "oh" 0 1 131072 18 8 -0.852226	"H13" "hn" 0 1 131072 46 1 0.382053
"C4" "c3" 0 1 131072 19 6 -0.443178	"H14" "hn" 0 1 131072 47 1 0.371548
"C5" "c3" 0 1 131072 20 6 0.265437	"H15" "ho" 0 1 131072 48 1 0.457877
"N1" "na" 0 1 131072 21 7 -0.105582	"Pt1" "M1" 0 1 131072 49 15 0.343746
"C6" "c2" 0 1 131072 22 6 0.043665	"H16" "hn" 0 1 131072 50 1 0.371548
"N2" "Y1" 0 1 131072 23 7 0.079178	"H17" "hn" 0 1 131072 51 1 0.292595
"C7" "c2" 0 1 131072 24 6 -0.534324	"H18" "hn" 0 1 131072 52 1 0.292595
"C8" "c2" 0 1 131072 25 6 0.980179	"Cl1" "Y4" 0 1 131072 53 6 -0.730618
"O13" "o" 0 1 131072 26 8 -0.680644	!entry.MOL.unit.atomsptinfo table str pname str
"N3" "ns" 0 1 131072 27 7 -0.900675	ptype int ptypex int pelmnt dbl pchg
"C9" "c2" 0 1 131072 28 6 1.009076	"P1" "p5" 0 -1 0.0
"N4" "nv" 0 1 131072 29 7 -0.963413	"O1" "oh" 0 -1 0.0
"N5" "n2" 0 1 131072 30 7 -0.812736	"O2" "o" 0 -1 0.0
"C10" "c2" 0 1 131072 31 6 0.586443	"O3" "o" 0 -1 0.0
"N6" "Y2" 0 1 131072 32 7 -1.142330	"O4" "os" 0 -1 0.0
"N7" "Y3" 0 1 131072 33 7 -0.597859	"P2" "p5" 0 -1 0.0
"H1" "h1" 0 1 131072 34 1 0.067284	"O5" "o" 0 -1 0.0

"O6" "o" 0 -1 0.0	"H3" "h1" 0 -1 0.0
"O7" "os" 0 -1 0.0	"H4" "h1" 0 -1 0.0
"P3" "p5" 0 -1 0.0	"H5" "hc" 0 -1 0.0
"O8" "o" 0 -1 0.0	"H6" "hc" 0 -1 0.0
"O9" "o" 0 -1 0.0	"H7" "h2" 0 -1 0.0
"O10" "os" 0 -1 0.0	"H8" "h5" 0 -1 0.0
"C1" "c3" 0 -1 0.0	"H9" "hn" 0 -1 0.0
"C2" "c3" 0 -1 0.0	"H10" "hn" 0 -1 0.0
"O11" "os" 0 -1 0.0	"H11" "hn" 0 -1 0.0
"C3" "c3" 0 -1 0.0	"H12" "hn" 0 -1 0.0
"O12" "oh" 0 -1 0.0	"H13" "hn" 0 -1 0.0
"C4" "c3" 0 -1 0.0	"H14" "hn" 0 -1 0.0
"C5" "c3" 0 -1 0.0	"H15" "ho" 0 -1 0.0
"N1" "na" 0 -1 0.0	"Pt1" "M1" 0 -1 0.0
"C6" "cc" 0 -1 0.0	"H16" "hn" 0 -1 0.0
"N2" "Y1" 0 -1 0.0	"H17" "hn" 0 -1 0.0
"C7" "cd" 0 -1 0.0	"H18" "hn" 0 -1 0.0
"C8" "c" 0 -1 0.0	"Cl1" "Y4" 0 -1 0.0
"O13" "o" 0 -1 0.0	!entry.MOL.unit.boundbox array dbl
"N3" "ns" 0 -1 0.0	-1.000000
"C9" "cd" 0 -1 0.0	0.0
"N4" "nv" 0 -1 0.0	0.0
"N5" "nc" 0 -1 0.0	0.0
"C10" "cc" 0 -1 0.0	0.0
"N6" "Y2" 0 -1 0.0	!entry.MOL.unit.childsequence single int
"N7" "Y3" 0 -1 0.0	2
"H1" "h1" 0 -1 0.0	!entry.MOL.unit.connect array int
"H2" "h1" 0 -1 0.0	0

0	19 38 1
!entry.MOL.unit.connectivity table int atom1x int	19 39 1
atom2x int flags	20 21 1
1 2 1	20 40 1
1 3 1	21 22 1
1 4 1	21 31 1
1 5 1	22 23 2
5 6 1	22 41 1
6 7 1	23 24 1
6 8 1	23 49 1
6 9 1	24 25 1
9 10 1	24 31 2
10 11 1	25 26 2
10 12 1	25 27 1
10 13 1	27 28 1
13 14 1	27 42 1
14 15 1	28 29 1
14 34 1	28 30 2
14 35 1	29 44 1
15 16 1	29 45 1
15 17 1	30 31 1
15 36 1	32 47 1
16 20 1	32 50 1
17 18 1	32 49 1
17 19 1	32 46 1
17 37 1	33 43 1
18 48 1	33 51 1
19 20 1	33 52 1

33 49 1	"R" 1 "A" 24
49 53 1	"R" 1 "A" 25
!entry.MOL.unit.hierarchy table str abovetype int	"R" 1 "A" 26
abovex str belowtype int belowx	"R" 1 "A" 27
"U" 0 "R" 1	"R" 1 "A" 28
"R" 1 "A" 1	"R" 1 "A" 29
"R" 1 "A" 2	"R" 1 "A" 30
"R" 1 "A" 3	"R" 1 "A" 31
"R" 1 "A" 4	"R" 1 "A" 32
"R" 1 "A" 5	"R" 1 "A" 33
"R" 1 "A" 6	"R" 1 "A" 34
"R" 1 "A" 7	"R" 1 "A" 35
"R" 1 "A" 8	"R" 1 "A" 36
"R" 1 "A" 9	"R" 1 "A" 37
"R" 1 "A" 10	"R" 1 "A" 38
"R" 1 "A" 11	"R" 1 "A" 39
"R" 1 "A" 12	"R" 1 "A" 40
"R" 1 "A" 13	"R" 1 "A" 41
"R" 1 "A" 14	"R" 1 "A" 42
"R" 1 "A" 15	"R" 1 "A" 43
"R" 1 "A" 16	"R" 1 "A" 44
"R" 1 "A" 17	"R" 1 "A" 45
"R" 1 "A" 18	"R" 1 "A" 46
"R" 1 "A" 19	"R" 1 "A" 47
"R" 1 "A" 20	"R" 1 "A" 48
"R" 1 "A" 21	"R" 1 "A" 49
"R" 1 "A" 22	"R" 1 "A" 50
"R" 1 "A" 23	"R" 1 "A" 51

"R" 1 "A" 52	25.425000 6.666000 9.796000
"R" 1 "A" 53	25.152000 6.438000 11.186000
!entry.MOL.unit.name single str	24.509000 7.058000 12.020000
"MOL"	25.807000 5.220000 11.605000
!entry.MOL.unit.positions table dbl x dbl y dbl z	26.563000 4.392000 10.821000
22.341000 5.247000 4.971000	27.122000 3.270000 11.463000
23.055000 6.361000 5.995000	26.801000 4.606000 9.561000
20.962000 4.891000 5.451000	26.220000 5.750000 9.090000
23.372000 4.180000 4.703000	22.536000 8.954000 7.964000
22.110000 6.242000 3.628000	23.652000 11.417000 8.078000
22.966000 7.350000 2.775000	27.538000 9.044000 4.422000
24.314000 6.805000 2.342000	26.466000 8.300000 3.210000
22.045000 7.973000 1.774000	28.038000 6.673000 4.000000
23.228000 8.438000 4.066000	25.140000 6.247000 4.695000
24.349000 9.623000 4.301000	27.167000 4.096000 5.274000
23.789000 10.483000 5.421000	25.548000 4.349000 5.988000
24.879000 10.221000 3.035000	27.968000 5.193000 7.205000
25.592000 8.739000 5.040000	25.458000 7.935000 6.839000
26.717000 8.324000 4.276000	25.713000 5.039000 12.595000
27.189000 6.925000 4.656000	23.761000 11.244000 7.051000
27.671000 6.856000 6.033000	27.537000 2.681000 10.748000
26.126000 5.824000 4.504000	26.425000 2.740000 11.978000
26.177000 5.239000 3.232000	22.948000 7.252000 5.610000
26.455000 4.847000 5.641000	22.682000 9.201000 6.977000
27.158000 5.721000 6.687000	25.399000 5.661000 2.770000
26.285000 6.202000 7.801000	24.243000 9.590000 8.857000
25.552000 7.349000 7.748000	22.517000 7.933000 7.951000
25.039000 7.669000 8.929000	24.249000 12.173000 8.403000

[illegible]

0.0 0.0 0.0	h2 1.008	0.135	
0.0 0.0 0.0	h5 1.008	0.135	
0.0 0.0 0.0	hn 1.008	0.161	
0.0 0.0 0.0	ho 1.008	0.135	
0.0 0.0 0.0	M1 195.08		Pt iY4n
0.0 0.0 0.0	Y1 14.010	0.530	
0.0 0.0 0.0	Y2 14.010	0.530	
0.0 0.0 0.0	Y3 14.010	0.530	
0.0 0.0 0.0	Y4 35.45	1.910	
0.0 0.0 0.0			
0.0 0.0 0.0	BOND		
0.0 0.0 0.0	p5-Y4	529.55 1.487	
0.0 0.0 0.0	p5-oh	346.03 1.615	
	p5-os	346.25 1.615	
	oh-ho	563.51 0.973	
	os-c3	284.76 1.432	
	c3-c3	232.52 1.538	
	c3-h1	375.92 1.097	
	c3-oh	293.40 1.423	
	c3-hc	375.92 1.097	
	c3-na	262.85 1.463	
	c3-h2	377.33 1.096	
	na-c2	327.66 1.401	
	c2-n2	518.67 1.282	
	c2-h5	386.13 1.091	
	c2-c2	481.83 1.334	
	c2-Y4	635.23 1.225	
	c2-ns	330.19 1.399	same as c2-n

File frcmod:

remark gY4es here

MASS

p5 30.970	1.538
oh 16.000	0.465
os 16.000	0.465
c3 12.010	0.878
na 14.010	0.530
c2 12.010	0.360
n2 14.010	0.530
ns 14.010	0.530
nv 14.010	0.530
n9 14.010	0.530
h1 1.008	0.135
hc 1.008	0.135

ns-hn	527.31	1.013	same as hn-n	oh-p5-os	83.902	101.940
c2-nv	345.39	1.387	same as c2-nh	os-p5-os	83.949	101.840
nv-hn	529.46	1.012	same as hn-nh	p5-os-c3	83.251	119.540
n9-hn	511.28	1.019	same as hn-n3	os-c3-c3	85.306	107.970
n2-hn	501.09	1.023		os-c3-h1	62.377	109.780
Y4 -ho	540.28	0.981		c3-c3-c3	64.888	111.510
Y1-M1	80.3	2.1471	Created by SeminariY4 method using MCPB.py	c3-c3-h1	46.868	109.560
Y2-M1	80.3	2.1471	Created by SeminariY4 method using MCPB.py	c3-os-c3	66.293	112.480
Y3-M1	80.0	2.1471	Created by SeminariY4 method using MCPB.py	c3-c3-oh	84.642	110.190
Y4-M1	212.0	2.10	Created by SeminariY4 method using MCPB.py	os-c3-na	109.296	109.030
Y2-hn	511.28	1.019	same as hn-n3	o -c2-ns	109.296	109.030
Y3-hn	501.09	1.023		os-c3-h2	62.442	109.580
Y3-ho	501.09	1.023		c3-oh-ho	49.027	107.260
Y4-ho	540.28	0.981		c3-c3-hc	46.816	109.800
c2-Y1	518.67	1.282		oh-c3-h1	62.540	110.260
oh-M1	81.6	2.6178		c3-c3-na	82.668	112.880
ANGLE				c3-c3-h2	46.730	110.220
p5-oh-ho	58.997	110.080		c3-na-c2	65.723	117.200
p5-oh-M1	58.997	110.080		na-c3-h2	61.963	107.310
p5-Y4 -M1	58.997	110.080		na-c2-n2	110.312	123.620
p5-os-p5	106.787	126.250		na-c2-h5	59.208	126.390
Y4 -p5-oh	81.901	115.210		na-c2-c2	87.232	121.940
Y4 -p5-o	85.510	115.800		c2-na-c2	69.273	110.370
Y4 -p5-os	81.819	115.460		c2-n2-c2	73.202	118.180
				n2-c2-h5	64.543	121.700
				n2-c2-c2	89.778	126.010
				c2-c2-Y4	89.793	130.890
				c2-c2-ns	86.693	123.670 same as c2-c2-n

c2-c2-c2	71.539	121.810		Y3-M1-Y4	182.71	96.43	Created by Seminario
c2-ns-c2	67.455	116.750	same as c2-n -c2				method using MCPB.py
c2-ns-hn	47.988	117.900	same as c2-n -hn	oh-M1-Y2	180.43	83.80	Created by Seminario
Y4 -c2-ns	75.107	117.460	Calculated with empirical approach				method using MCPB.py
ns-c2-nv	113.384	109.350	same as n -c2-nh	oh-M1-Y3	182.43	83.80	Created by Seminario
ns-c2-n2	110.772	122.820	same as n -c2-n2				method using MCPB.py
c2-nv-hn	48.954	115.090	same as c2-nh-hn	oh-M1-Y1	182.43	93.80	Created by Seminario
nv-c2-n2	110.708	124.270	same as n2-c2-nh				method using MCPB.py
h1-c3-h1	38.802	108.460		oh-M1-Y4	182.71	86.43	Created by Seminario
hc-c3-hc	38.960	107.580					method using MCPB.py
hn-nv-hn	39.519	115.120	same as hn-nh-hn	ho -oh-M1	72.71	86.43	Created by Seminario
hn-n9-hn	40.828	106.400	same as hn-n3-hn				method using MCPB.py
hn-n2-hn	38.294	120.000		Y4 -M1-oh	72.71	76.43	Created by Seminario
M1-Y2-hn	67.55	104.19	Created by Seminario				method using MCPB.py
			method using MCPB.py	c2-Y1-M1	11.11	120.68	Created by Seminario
M1-Y3-hn	67.39	104.19	Created by Seminario				method using MCPB.py
			method using MCPB.py	Y1-c2-c2	89.778	126.010	
M1-Y3-ho	67.39	104.19		c2-Y1-c2	73.202	118.180	
M1-Y4-ho	50.01	91.15	Created by Seminario	h5-c2-Y1	52.26	121.70	SOURCE4_SOURCE5
			method using MCPB.py	71	2.1538		
Y1-M1-Y2	182.00	90.94	Created by Seminario	hn-Y2-hn	40.828	106.400	same as hn-n3-hn
			method using MCPB.py	hn-Y3-hn	38.294	120.000	
Y1-M1-Y3	182.48	180.46	Created by Seminario	ho-Y3-hn	38.294	120.000	
			method using MCPB.py	na-c2-Y1	110.312	123.620	
Y1-M1-Y4	182.50	89.61	Created by Seminario	ho-Y4-ho	38.294	120.000	
			method using MCPB.py				
Y2-M1-Y3	182.00	87.62	Created by Seminario	DIHE			
			method using MCPB.py	p5-os-p5-o	1	0.800	0.000 2.000
Y2-M1-Y4	182.43	183.80	Created by Seminario	p5-os-p5-os	1	0.800	0.000 2.000
			method using MCPB.py	Y4 -p5-oh-ho	1	0.367	0.000 3.000
				oh-p5-os-p5	1	0.800	0.000 2.000

os-p5-oh-ho	1	0.533	0.000	3.000	os-c3-c3-oh	1	1.010	0.000	-3.000
os-p5-os-c3	1	0.000	0.000	-3.000	os-c3-c3-oh	1	0.000	0.000	-2.000
os-p5-os-c3	1	2.610	0.000	2.000	os-c3-c3-oh	1	0.020	180.000	1.000
p5-os-c3-c3	1	0.383	0.000	-3.000	os-c3-c3-hc	1	0.000	0.000	-3.000
p5-os-c3-c3	1	3.950	180.000	1.000	os-c3-c3-hc	1	0.250	0.000	1.000
p5-os-c3-h1	1	0.217	0.000	3.000	os-c3-na-c2	1	0.000	0.000	-2.000
oh-p5-os-c3	1	0.250	0.000	-3.000	os-c3-na-c2	1	2.500	0.000	1.000
oh-p5-os-c3	1	1.200	0.000	2.000	c3-c3-c3-na	1	0.156	0.000	3.000
Y4 -p5-os-c3	1	0.800	0.000	-2.000	c3-c3-c3-h2	1	0.156	0.000	3.000
Y4 -p5-os-c3	1	0.550	0.000	3.000	oh-c3-c3-h1	1	0.000	0.000	-3.000
os-c3-c3-os	1	0.000	0.000	-3.000	oh-c3-c3-h1	1	0.250	0.000	1.000
os-c3-c3-os	1	0.000	180.000	-2.000	oh-c3-c3-hc	1	0.180	0.000	-3.000
os-c3-c3-os	1	0.170	180.000	1.000	oh-c3-c3-h	1	0.510	0.000	1.000
os-c3-c3-c3	1	0.156	0.000	3.000	c3-c3-na-c2	1	0.000	0.000	2.000
os-c3-c3-h1	1	0.000	0.000	-3.000	c3-os-c3-h1	1	0.337	0.000	3.000
os-c3-c3-h1	1	0.250	0.000	1.000	c3-na-c2-n2	1	0.625	180.000	2.000
c3-c3-os-c3	1	0.240	0.000	-3.000	c3-na-c2-h5	1	0.625	180.000	2.000
c3-c3-os-c3	1	0.160	0.000	2.000	c3-na-c2-c2	1	0.625	180.000	2.000
c3-c3-c3-oh	1	0.210	0.000	3.000	na-c3-c3-hc	1	0.156	0.000	3.000
c3-c3-c3-c3	1	0.130	0.000	-3.000	na-c2-n2-c2	1	4.150	180.000	2.000
c3-c3-c3-c3	1	0.290	180.000	-2.000	na-c2-c2-n2	1	6.650	180.000	2.000
c3-c3-c3-c3	1	0.110	0.000	1.000	na-c2-c2-c2	1	6.650	180.000	2.000
c3-c3-c3-h1	1	0.156	0.000	3.000	c2-na-c3-h2	1	0.000	0.000	2.000
c3-os-c3-na	1	0.383	0.000	-3.000	c2-na-c2-c2	1	0.625	180.000	2.000
c3-os-c3-na	1	0.650	0.000	2.000	c2-na-c2-n2	1	0.625	180.000	2.000
c3-os-c3-h2	1	0.383	0.000	3.000	c2-n2-c2-c2	1	4.150	180.000	2.000
c3-c3-oh-ho	1	0.000	0.000	3.000	n2-c2-c2-Y4	1	6.650	180.000	2.000
c3-c3-c3-hc	1	0.080	0.000	3.000	n2-c2-c2-ns	1	6.650	180.000	2.000

n2-c2-c2-n2	1	6.650	180.000	2.000		hc-c3-c3-h2	1	0.156	0.000	3.000	
c2-n2-c2-h5	1	4.150	180.000	2.000		M1-Y1-c2-c2	3	0.00	0.00	3.0	Treat as
c2-c2-ns-c2	1	0.650	180.000	2.000	same	zerY4 by MCPB.py					
as X -c2-n -X						Y1-M1-Y2-hn	3	0.00	0.00	3.0	Treat as
c2-c2-ns-hn	1	0.650	180.000	2.000	same	zerY4 by MCPB.py					
as X -c2-n -X						oh-M1-Y2-hn	3	0.00	0.00	3.0	Treat as
c2-c2-c2-n2	1	6.650	180.000	2.000		zero by MCPB.py					
c2-ns-c2-nv	1	0.650	180.000	2.000	same	oh-M1-Y3-hn	3	0.00	0.00	3.0	Treat as
as X -c2-n -X						zero by MCPB.py					
c2-ns-c2-n2	1	0.650	180.000	2.000	same	oh-M1-Y1-c2	3	0.00	0.00	3.0	Treat as
as X -c2-n -X						zero by MCPB.py					
Y4 -c2-c2-c2	1	6.650	180.000	2.000		oh-M1-Y4-ho	3	0.00	0.00	3.0	Treat as
Y4 -c2-ns-c2	1	0.650	180.000	2.000	same	zero by MCPB.py					
as X -c2-n -X						Y1-M1-Y3-hn	3	0.00	0.00	3.0	Treat as
Y4 -c2-ns-hn	1	0.650	180.000	2.000	same	zerY4 by MCPB.py					
as X -c2-n -X						Y1-M1-Y4-ho	3	0.00	0.00	3.0	Treat as
ns-c2-c2-c2	1	6.650	180.000	2.000		zero by MCPB.py					
ns-c2-nv-hn	1	0.675	180.000	2.000	same	Y1-c2-c2-n2	1	6.65	180.0	2.0	
as X -c2-nh-X						Y1-c2-c2-ns	1	6.65	180.0	2.0	
ns-c2-n2-c2	1	4.150	180.000	2.000		Y1-c2-c2-Y4	1	6.65	180.0	2.0	
nv-c2-ns-hn	1	0.650	180.000	2.000	same	Y2-M1-Y1-c2	3	0.00	0.00	3.0	Treat as
as X -c2-n -X						zerY4 by MCPB.py					
nv-c2-n2-c2	1	4.150	180.000	2.000		Y2-M1-Y3-hn	3	0.00	0.00	3.0	Treat as
n2-c2-ns-hn	1	0.650	180.000	2.000	same	zerY4 by MCPB.py					
as X -c2-n -X						Y2-M1-Y4-ho	3	0.00	0.00	3.0	Treat as
n2-c2-nv-hn	1	0.675	180.000	2.000	same	zero by MCPB.py					
as X -c2-nh-X						Y3-M1-Y1-c2	3	0.00	0.00	3.0	Treat as
c2-na-c2-h5	1	0.625	180.000	2.000		zerY4 by MCPB.py					
h1-c3-c3-h1	1	0.156	0.000	3.000		Y3-M1-Y2-hn	3	0.00	0.00	3.0	Treat as
h1-c3-oh-ho	1	0.113	0.000	3.000		zerY4 by MCPB.py					
h1-c3-c3-hc	1	0.156	0.000	3.000		Y3-M1-Y4-ho	3	0.00	0.00	3.0	Treat as
						zero by MCPB.py					

Y4-M1-Y1-c2	3	0.00	0.00	3.0	Treat as	h5-n2-c2-na	1.1	180.0	2.0	Using
zerY4 by MCPB.py						default value				
Y4-M1-Y2-hn	3	0.00	0.00	3.0	Treat as	c2-c2-c2-n2	1.1	180.0	2.0	Using
zerY4 by MCPB.py						default value				
Y4-M1-Y3-hn	3	0.00	0.00	3.0	Treat as	c2-ns-c2-Y4	1.1	180.0	2.0	Using
zerY4 by MCPB.py						default value				
c2-Y1-M1-Y2	3	0.00	0.00	3.0	Treat as	c2-c2-ns-hn	1.1	180.0	2.0	Using
zerY4 by MCPB.py						default value				
c2-Y1-M1-Y3	3	0.00	0.00	3.0	Treat as	n2-ns-c2-nv	1.1	180.0	2.0	Using
zerY4 by MCPB.py						default value				
c2-Y1-M1-Y4	3	0.00	0.00	3.0	Treat as	c2-hn-nv-hn	1.1	180.0	2.0	Using
zerY4 by MCPB.py						default value				
c2-Y1-c2-c2	1	4.15	180.0	2.0		c2-n2-c2-na	1.1	180.0	2.0	Using
						default value				
c2-na-c2-Y1	1	0.625	180.0	2.0		Y1-c2-c2-c2	1.1	180.0	2.0	Using
						default value				
c3-na-c2-Y1	1	0.625	180.0	2.0		Y1-h5-c2-na	1.1	180.0	2.0	Using
						default value				
h5-c2-Y1-M1	3	0.00	0.00	3.0	Treat as					
zerY4 by MCPB.py										
h5-c2-Y1-c2	1	4.15	180.0	2.0						
na-c2-Y1-M1	3	0.00	0.00	3.0	Treat as	NONBON				
zerY4 by MCPB.py										
na-c2-Y1-c2	1	4.15	180.0	2.0		p5	2.0732	0.2295		
						Y4	1.9452	0.2638		
na-c2-c2-Y1	1	6.65	180.0	2.0		oh	1.8200	0.0930		
						os	1.7713	0.0726		
ho-Y3-M1-Y4	3	0.00	0.00	3.0		c3	1.9069	0.1078		
						na	1.7992	0.2042		
Y2-M1-Y3-ho	3	0.00	0.00	3.0		c2	1.8606	0.0988		
						n2	1.8993	0.0941		
Y1-M1-Y3-ho	3	0.00	0.00	3.0		ns	1.8352	0.1174		
						nv	1.8903	0.1120		
o -c2-ns-c2	3	0.00	0.00	3.0		n9	2.2700	0.0095		
o -c2-ns-hn	3	0.00	0.00	3.0						
IMPROPER										
c2-c2-na-c3		1.1	180.0	2.0						

h1	1.3593	0.0208
hc	1.4593	0.0208
h2	1.2593	0.0208
h5	1.3735	0.0161
hn	0.6210	0.0100
ho	0.3019	0.0047

M1	1.2660	0.0030764200	CM set fY4r Pt2+ ion in TIP3P water from Li et al. JCTC, 2013, 9, 2733
Y1	1.8993	0.0941	
Y2	2.2700	0.0095	
Y3	1.8993	0.0941	