

## SUPPORTING INFORMATION

# “Rational design of novel fluorescent analogues of cholesterol: a “step-by-step” computational study”

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## 1. Optical properties

### 1.1. CASPT2 energies for probes **8** and **10**

In order to check the presence that no dark states are below the bright excited state (characterized by a HOMO → LUMO transition), we performed a CASPT2 excited state analysis based on a multireference CASSCF(8, 8)-SA11 wavefunction using the ANO-L-VDZP basis-set. These computations were performed with MOLCAS 8.0 quantum chemistry software<sup>1</sup>. As we can see, no dark excited states are present below the bright excited state ( $S_1$ ) for both probe **8** and, more importantly, for probe **10**. As in TD-DFT computations, presented in the main article, the bright state  $S_1$  is described by a single excitation HOMO → LUMO transition. The dominant CI-coefficient, for this HOMO → LUMO configuration, are -0.91155 and 0.74507 for probes **8** and **10**, respectively.

Table 1: CASPT2 energies for the ground state and the first 10 singlet excited state

Probe	State	CASPT2 energy [Hartree]	$\Delta E$ [eV]	$\Delta E$ [nm]	$f(S_0 \rightarrow S_n)$
<b>8</b>	$S_0$	-1124.84427339	/	/	/
	$S_1$	-1124.72750955	3.177	390	0.394
	$S_2$	-1124.69752380	3.993	311	0.000
	$S_3$	-1124.67543105	4.594	270	0.037
	$S_4$	-1124.66983915	4.747	261	0.002
	$S_5$	-1124.65621984	5.117	242	0.001
	$S_6$	-1124.63694614	5.642	220	0.006
	$S_7$	-1124.63500171	5.695	218	0.014
	$S_8$	-1124.61830540	6.149	202	0.018
	$S_9$	-1124.61011564	6.372	195	0.002
<b>10</b>	$S_{10}$	-1124.52684766	8.638	144	0.001
	$S_0$	-1124.84783247	/	/	/
	$S_1$	-1124.69332332	4.204	295	0.668
	$S_2$	-1124.68748496	4.363	284	0.001
	$S_3$	-1124.67038366	4.829	257	0.010
	$S_4$	-1124.65748752	5.180	239	0.032
	$S_5$	-1124.65584240	5.224	237	0.340
	$S_6$	-1124.62510039	6.061	205	0.012
	$S_7$	-1124.61558977	6.320	196	0.230
	$S_8$	-1124.60706099	6.552	189	0.005
	$S_9$	-1124.58373118	7.187	173	0.001
	$S_{10}$	-1124.57949729	7.302	170	0.044

## 2. Molecular dynamics in POPC

### 2.1. GAFF force field for cholesterol, probe **8** and **10**

In the following tables (Table 2, Table 3 and Table 4) we list GAFF atoms type and RESP charges used in the molecular dynamics simulation performed in this work together with their ground state geometries (obtained at B3LYP/6-31+G(d) level of theory in vacuum).

Table 2: Ground state geometry, atom names, GAFF atom-types and RESP charges and isotropic polarizability  $\alpha_{iso}$  for cholesterol.

Cholesterol							
Atom number	Name	X [Å]	Y [Å]	Z [Å]	GAFF atom-type	RESP Charge	$\alpha_{iso}$
1	C3	6.879	-0.082	-0.801	c3	0.378828	7.52532704
2	O1	8.294	-0.065	-0.568	oh	-0.751689	5.78273641
3	HO1	8.718	-0.682	-1.185	ho	0.423029	1.45927443
4	H31	6.675	0.260	-1.830	h1	-0.019871	1.85606657
5	C4	6.254	0.909	0.180	c3	-0.008496	8.28625582
6	H41	6.591	0.624	1.188	hc	0.072237	1.85981222
7	H42	6.651	1.913	-0.008	hc	0.072237	1.90550381
8	C5	4.738	0.922	0.100	c2	-0.410084	9.38013989
9	C6	4.079	2.079	-0.056	c2	-0.181935	9.48966611
10	H61	4.656	3.000	-0.150	ha	0.118276	2.13067117
11	C7	2.584	2.222	-0.114	c3	-0.008375	8.01595289
12	H71	2.274	3.084	0.494	hc	0.026749	2.00261433
13	H72	2.283	2.471	-1.146	hc	0.026749	1.87565226
14	C8	1.840	0.959	0.350	c3	0.055053	8.30976134
15	H81	1.896	0.925	1.448	hc	0.002266	1.64573912
16	C14	0.367	0.998	-0.079	c3	0.018813	8.42204347
17	H141	0.379	1.013	-1.184	hc	-0.016960	1.71879767
18	C15	-0.500	2.186	0.360	c3	-0.119530	7.74240985
19	H151	-0.352	2.401	1.426	hc	0.018976	1.74644610
20	H152	-0.256	3.107	-0.182	hc	0.018976	1.94576940
21	C16	-1.949	1.704	0.078	c3	-0.052708	7.57859951
22	H161	-2.593	1.868	0.952	hc	0.017247	1.83108783
23	H162	-2.399	2.269	-0.745	hc	0.017247	1.89258326
24	C17	-1.872	0.176	-0.268	c3	-0.032017	8.75505836
25	H171	-1.796	0.080	-1.364	hc	-0.037496	1.76402732
26	C13	-0.475	-0.252	0.304	c3	0.460490	8.56285985
27	C18	-0.563	-0.454	1.834	c3	-0.453169	6.99109867
28	H181	0.423	-0.511	2.302	hc	0.087135	1.90436703
29	H182	-1.107	0.360	2.325	hc	0.087135	1.72393117
30	H183	-1.089	-1.385	2.073	hc	0.087135	1.87649897
31	C12	0.209	-1.463	-0.366	c3	-0.056081	7.73584487
32	H121	-0.296	-2.401	-0.114	hc	-0.010733	1.94597956
33	H122	0.134	-1.352	-1.459	hc	-0.010733	1.74600306
34	C11	1.693	-1.581	0.040	c3	-0.112061	7.54481247
35	H111	1.741	-1.831	1.105	hc	0.018586	1.60138308
36	H112	2.132	-2.434	-0.492	hc	0.018586	1.93903499
37	C9	2.526	-0.306	-0.231	c3	-0.036960	8.42607955
38	H91	2.542	-0.159	-1.325	hc	-0.016648	1.68257052
39	C10	4.027	-0.436	0.207	c3	0.570986	8.82640231
40	C19	4.150	-0.949	1.667	c3	-0.209529	7.19731538
41	H191	3.844	-1.997	1.752	hc	0.031785	1.84347048
42	H192	5.179	-0.882	2.032	hc	0.031785	1.92703743
43	H193	3.529	-0.353	2.345	hc	0.031785	1.74234186
44	C1	4.744	-1.431	-0.757	c3	-0.098195	7.73358052
45	H11	4.342	-2.441	-0.613	hc	0.004995	1.87933166
46	H12	4.501	-1.145	-1.791	hc	0.004995	1.72468700
47	C2	6.273	-1.473	-0.607	c3	-0.127133	7.82100959

48	H21	6.693	-2.175	-1.342	hc	0.030841	2.19436286
49	H22	6.560	-1.844	0.386	hc	0.030841	1.79025242
50	C20	-3.150	-0.592	0.158	c3	0.187074	8.78398973
51	H201	-3.283	-0.452	1.243	hc	-0.031548	1.78696205
52	C21	-3.076	-2.104	-0.116	c3	-0.298829	7.28927144
53	H211	-2.860	-2.304	-1.175	hc	0.059599	1.74095443
54	H212	-4.022	-2.598	0.129	hc	0.059599	2.11061672
55	H213	-2.301	-2.594	0.479	hc	0.059599	1.86141947
56	C22	-4.392	0.022	-0.536	c3	-0.039615	8.56967625
57	H221	-4.379	1.110	-0.401	hc	0.003047	1.69440175
58	H222	-4.312	-0.150	-1.622	hc	0.003047	1.82432582
59	C23	-5.755	-0.492	-0.043	c3	0.048794	8.37314366
60	H231	-5.875	-1.558	-0.279	hc	-0.002549	1.74572690
61	H232	-5.790	-0.411	1.053	hc	-0.002549	1.74419898
62	C24	-6.928	0.282	-0.665	c3	-0.183603	8.45937794
63	H241	-6.863	1.342	-0.372	hc	0.027806	1.80320613
64	H242	-6.816	0.264	-1.759	hc	0.027806	1.84661334
65	C25	-8.334	-0.246	-0.312	c3	0.440534	8.42696324
66	H251	-8.368	-1.312	-0.587	hc	-0.075146	1.94260581
67	C26	-8.644	-0.136	1.190	c3	-0.313515	7.48903306
68	H261	-9.655	-0.502	1.409	hc	0.058435	2.12100669
69	H262	-7.944	-0.719	1.799	hc	0.058435	1.86713877
70	H263	-8.588	0.909	1.523	hc	0.058435	1.77742806
71	C27	-9.405	0.483	-1.138	c3	-0.313515	7.77477783
72	H271	-9.219	0.380	-2.215	hc	0.058435	1.88779141
73	H272	-10.407	0.086	-0.932	hc	0.058435	2.12417547
74	H273	-9.420	1.557	-0.904	hc	0.058435	1.82240413

Table 3: Ground state geometry, atom names, GAFF atom-types and RESP charges and isotropic polarizability  $\alpha_{iso}$  for probe 8.

**8**

Atom number	Name	X [Å]	Y [Å]	Z [Å]	GAFF atom-type	RESP Charge	$\alpha_{iso}$
1	C3	-6.821	-0.372	0.302	c3	0.379537	7.83287132
2	O1	-8.207	-0.458	-0.049	oh	-0.751834	5.97916166
3	HO1	-8.640	-1.094	0.543	ho	0.426420	1.48724839
4	H31	-6.727	-0.015	1.341	h1	-0.017557	1.85515560
5	C4	-6.191	0.653	-0.640	c3	-0.042345	8.80726285
6	H41	-6.424	0.321	-1.665	hc	0.086903	2.06119467
7	H42	-6.677	1.627	-0.513	hc	0.086903	1.98966366
8	C5	-4.696	0.787	-0.459	c2	-0.372427	10.74453013
9	C6	-4.082	1.986	-0.584	ce	-0.104696	10.85790610
10	H61	-4.682	2.877	-0.765	ha	0.120410	2.18964225
11	C7	-2.640	2.132	-0.552	ce	-0.160831	11.28198516
12	H71	-2.205	3.079	-0.865	ha	0.122079	1.79207853
13	C8	-1.838	1.128	-0.113	cf	-0.053637	13.38385678
14	C14	-0.409	1.279	0.108	cf	-0.240229	13.66594762
15	C15	0.293	2.375	0.511	ce	-0.098201	10.39188990
16	H152	-0.113	3.371	0.660	ha	0.121469	1.84788643
17	C16	1.664	1.990	0.825	ce	-0.200848	9.45199949
18	H162	2.424	2.690	1.163	ha	0.138680	2.19139163
19	C17	1.830	0.653	0.658	c2	-0.290057	11.96376277
20	C13	0.521	0.075	0.123	c3	0.611216	11.27203078
21	C18	0.761	-0.454	-1.316	c3	-0.340933	6.88150629
22	H181	-0.172	-0.761	-1.797	hc	0.066290	1.68130482
23	H182	1.214	0.327	-1.936	hc	0.066290	1.67345670
24	H183	1.436	-1.319	-1.303	hc	0.066290	1.87687509
25	C12	-0.140	-0.998	1.023	c3	-0.020602	8.76388288
26	H121	0.479	-1.906	1.053	hc	-0.005185	1.98576558
27	H122	-0.192	-0.611	2.050	hc	-0.005185	1.66070977
28	C11	-1.551	-1.359	0.526	c3	-0.086898	8.96664695
29	H111	-1.464	-1.862	-0.443	hc	-0.000889	1.54280150
30	H112	-1.985	-2.095	1.213	hc	-0.000889	1.87630289
31	C9	-2.510	-0.148	0.408	c3	0.161217	11.08238011
32	H91	-2.807	0.116	1.442	hc	-0.008512	1.70117090
33	C10	-3.852	-0.488	-0.329	c3	0.392685	11.05577120
34	C19	-3.606	-1.018	-1.769	c3	-0.131381	6.90892338
35	H191	-3.124	-2.002	-1.762	hc	0.025392	1.70777342
36	H192	-4.548	-1.124	-2.316	hc	0.025392	1.90294260
37	H193	-2.971	-0.327	-2.334	hc	0.025392	1.61651702
38	C1	-4.607	-1.563	0.506	c3	-0.048603	8.20283188
39	H11	-4.111	-2.534	0.384	hc	-0.002412	1.77831532
40	H12	-4.524	-1.302	1.571	hc	-0.002412	1.71409144
41	C2	-6.097	-1.711	0.165	c3	-0.142672	8.31944624
42	H21	-6.549	-2.458	0.834	hc	0.036590	2.21969606
43	H22	-6.238	-2.080	-0.860	hc	0.036590	1.79780880
44	C20	3.078	-0.168	0.894	c3	0.292584	10.03691504
45	H201	2.945	-1.136	0.389	hc	-0.022635	1.69452968
46	C21	3.262	-0.466	2.401	c3	-0.333432	7.97522051
47	H211	3.470	0.458	2.955	hc	0.074820	1.78140752
48	H212	4.091	-1.162	2.573	hc	0.074820	2.31056817
49	H213	2.358	-0.913	2.829	hc	0.074820	1.86735417
50	C22	4.333	0.496	0.277	c3	-0.101012	9.01100958
51	H221	4.108	0.749	-0.769	hc	0.034646	1.77569958
52	H222	4.525	1.451	0.787	hc	0.034646	1.74074811
53	C23	5.607	-0.362	0.314	c3	0.042340	8.70607620
54	H231	5.914	-0.537	1.355	hc	-0.005945	1.81779238
55	H232	5.384	-1.349	-0.115	hc	-0.005945	1.72475514
56	C24	6.773	0.291	-0.445	c3	-0.148286	8.62563957
57	H241	6.494	0.412	-1.504	hc	0.017384	1.79406452
58	H242	6.918	1.309	-0.052	hc	0.017384	1.84783087
59	C25	8.119	-0.457	-0.369	c3	0.444907	8.54011875
60	H251	8.368	-0.590	0.696	hc	-0.079539	1.96685106
61	C26	8.057	-1.850	-1.016	c3	-0.307153	7.50560426
62	H261	9.032	-2.349	-0.965	hc	0.056040	2.14834221

63	H262	7.327	-2.503	-0.523	hc	0.056040	1.87954772
64	H263	7.777	-1.774	-2.076	hc	0.056040	1.77273807
65	C27	9.237	0.382	-1.008	c3	-0.307153	7.84044954
66	H271	9.328	1.364	-0.525	hc	0.056040	1.89460056
67	H272	10.209	-0.121	-0.930	hc	0.056040	2.15627136
68	H273	9.036	0.554	-2.075	hc	0.056040	1.81391535

Table 4: Ground state geometry, atom names, GAFF atom-types and RESP charges and isotropic polarizability  $\alpha_{iso}$  for probe 10.

Atom number	Name	X [Å]	Y [Å]	Z [Å]	GAFF atom-type	RESP Charge	$\alpha_{iso}$
1	C3	6.934	-0.131	-0.665	c3	0.462102	8.57667181
2	O1	8.221	0.196	-0.108	oh	-0.703286	6.36886752
3	HO1	8.845	-0.509	-0.347	ho	0.391200	1.71731622
4	H31	7.061	-0.283	-1.753	h1	-0.050194	2.19094507
5	C4	6.043	1.059	-0.463	c2	-0.254458	11.67352752
6	H42	6.502	2.026	-0.667	ha	0.146240	2.16403734
7	C5	4.737	0.998	-0.122	ce	-0.106399	12.17971468
8	C6	3.933	2.212	-0.036	ce	-0.181129	12.33636836
9	H61	4.460	3.164	-0.056	ha	0.138004	2.14704981
10	C7	2.581	2.193	0.067	cf	-0.214416	12.39709056
11	H71	2.046	3.134	0.172	ha	0.136980	1.82637149
12	C8	1.818	0.958	0.014	cf	0.023657	14.23670776
13	C14	0.475	0.912	0.228	ce	-0.026294	14.38557838
14	C15	-0.452	2.022	0.416	ce	-0.150062	11.07901447
15	H152	-0.167	3.031	0.697	ha	0.133945	1.82405626
16	C16	-1.708	1.633	0.123	c2	-0.317713	10.90757318
17	H162	-2.567	2.295	0.142	ha	0.160393	2.33860489
18	C17	-1.769	0.182	-0.334	c3	0.070291	10.40472434
19	H171	-1.726	0.186	-1.439	hc	0.021437	1.79648409
20	C13	-0.372	-0.366	0.176	c3	0.266833	10.57410738
21	C18	-0.501	-0.910	1.624	c3	-0.331857	7.15351214
22	H181	0.484	-1.102	2.059	hc	0.076482	1.66934731
23	H182	-1.003	-0.182	2.270	hc	0.076482	1.71084648
24	H183	-1.070	-1.845	1.657	hc	0.076482	2.02612741
25	C12	0.322	-1.390	-0.742	c3	-0.151221	9.24886275
26	H121	-0.184	-2.361	-0.705	hc	0.032766	1.94793234
27	H122	0.258	-1.039	-1.783	hc	0.032766	1.68721894
28	C11	1.802	-1.577	-0.362	c3	0.003416	9.18656977
29	H111	1.864	-2.000	0.647	hc	0.018739	1.54105062
30	H112	2.249	-2.322	-1.031	hc	0.018739	1.85092184
31	C9	2.605	-0.266	-0.450	c3	-0.101945	11.28437242
32	H91	2.800	-0.086	-1.523	hc	0.042366	1.79639515
33	C10	4.025	-0.320	0.211	c3	0.416652	11.42862728
34	C19	3.911	-0.442	1.753	c3	-0.149815	6.84847595
35	H191	3.445	-1.392	2.042	hc	0.030785	1.69613391
36	H192	4.898	-0.397	2.224	hc	0.030785	1.84016311
37	H193	3.310	0.371	2.172	hc	0.030785	1.62739712
38	C1	4.838	-1.507	-0.358	c3	-0.110359	8.88077671
39	H11	4.448	-2.452	0.040	hc	0.000629	1.74631583
40	H12	4.700	-1.546	-1.449	hc	0.000629	1.72906295
41	C2	6.339	-1.400	-0.049	c3	-0.033784	8.94577250
42	H21	6.867	-2.286	-0.428	hc	0.006495	2.27185228
43	H22	6.509	-1.366	1.034	hc	0.006495	1.70632121
44	C20	-3.068	-0.582	0.044	c3	0.155761	9.35465557
45	H201	-3.152	-0.602	1.140	hc	0.002741	1.81626404
46	C21	-3.048	-2.035	-0.467	c3	-0.318300	7.39193533
47	H211	-2.959	-2.060	-1.562	hc	0.068930	1.74719391
48	H212	-3.967	-2.565	-0.197	hc	0.068930	2.19576233
49	H213	-2.215	-2.609	-0.054	hc	0.068930	1.82397620
50	C22	-4.316	0.157	-0.501	c3	-0.042846	9.00419394
51	H221	-4.288	1.205	-0.181	hc	0.005802	1.73910424
52	H222	-4.269	0.174	-1.601	hc	0.005802	1.84508832
53	C23	-5.671	-0.422	-0.059	c3	0.071084	8.74013463
54	H231	-5.813	-1.428	-0.475	hc	-0.007237	1.77704838
55	H232	-5.669	-0.536	1.034	hc	-0.007237	1.76963068
56	C24	-6.852	0.460	-0.494	c3	-0.188363	8.72734175
57	H241	-6.765	1.449	-0.016	hc	0.026465	1.82722806
58	H242	-6.773	0.640	-1.577	hc	0.026465	1.86614238
59	C25	-8.252	-0.115	-0.197	c3	0.452439	8.60395444
60	H251	-8.307	-1.112	-0.663	hc	-0.082185	1.95936142
61	C26	-8.514	-0.281	1.308	c3	-0.301590	7.57294713
62	H261	-9.522	-0.675	1.489	hc	0.054393	2.17001359
63	H262	-7.803	-0.969	1.779	hc	0.054393	1.88913793
64	H263	-8.439	0.685	1.826	hc	0.054393	1.788091237
65	C27	-9.340	0.761	-0.839	c3	-0.301590	7.88091443
66	H271	-9.192	0.855	-1.922	hc	0.054393	1.89129709

67	H272	-10.340	0.341	-0.672	hc	0.054393	2.16941131
68	H273	-9.331	1.774	-0.413	hc	0.054393	1.82702304

## 2.2. Atom numbering for POPC

In Figure 1 we presents the atom numbering convention used for the analysis of the order parameter for the lipid 1-palmitoyl-2-oleoyl-phosphatidylcholine (POPC).

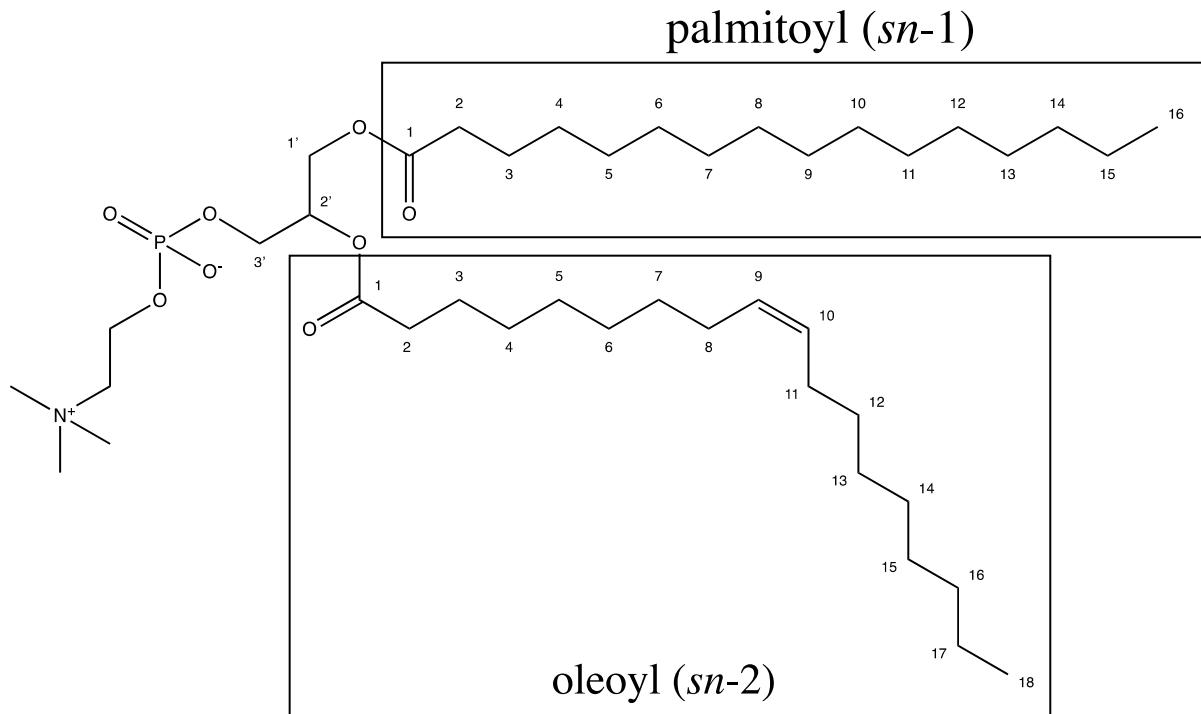


Figure 1: Chemical structure of of 1-palmitoyl-2-oleoyl-phosphatidylcholine (POPC) with the atom numbering of the *sn*-1, *sn*-2 used in this study. Atoms labelled with 1', 2' and 3' are those used to extract the rotational autocorrelation function.

## 2.3. POPC rotational correlation time

The estimation of POPC rotational correlation time was based on a previous study (see Supporting Information of ref. 2). The autocorrelation function  $C_f(t)$  is defined by<sup>3</sup>:

$$C_f(t) = \langle f(\xi)f(\xi + t) \rangle_{\xi},$$

where the average is over the time origin  $\xi$ . The integral over the time define the correlation time  $\tau_f$ :

$$\tau_f = \int_0^\infty C_f(t) dt$$

We performed the fitting of the rotational correlation function  $C_f(t)$  using a double exponential function defined by:

$$y(t) = a_1 e^{-(t/\tau_1)} + a_2 e^{-(t/\tau_2)}$$

The mean correlation time is the integral of the fitting function  $y$  from time origin 0 to infinity (last frame), this equals to the sum of correlation time  $\tau_1$  and  $\tau_2$  each one multiplied by their prefactors  $a_1$  and  $a_2$ :

$$\tau_f = \int_0^\infty a_1 e^{-(t/\tau_1)} + a_2 e^{-(t/\tau_2)} dt$$

In Figure 2 we presents the rotational correlation function  $C_f(t)$  for each simulated sterol (cholesterol, probe **8** and probe **10**, continuous lines) together with their double exponential fitting function.

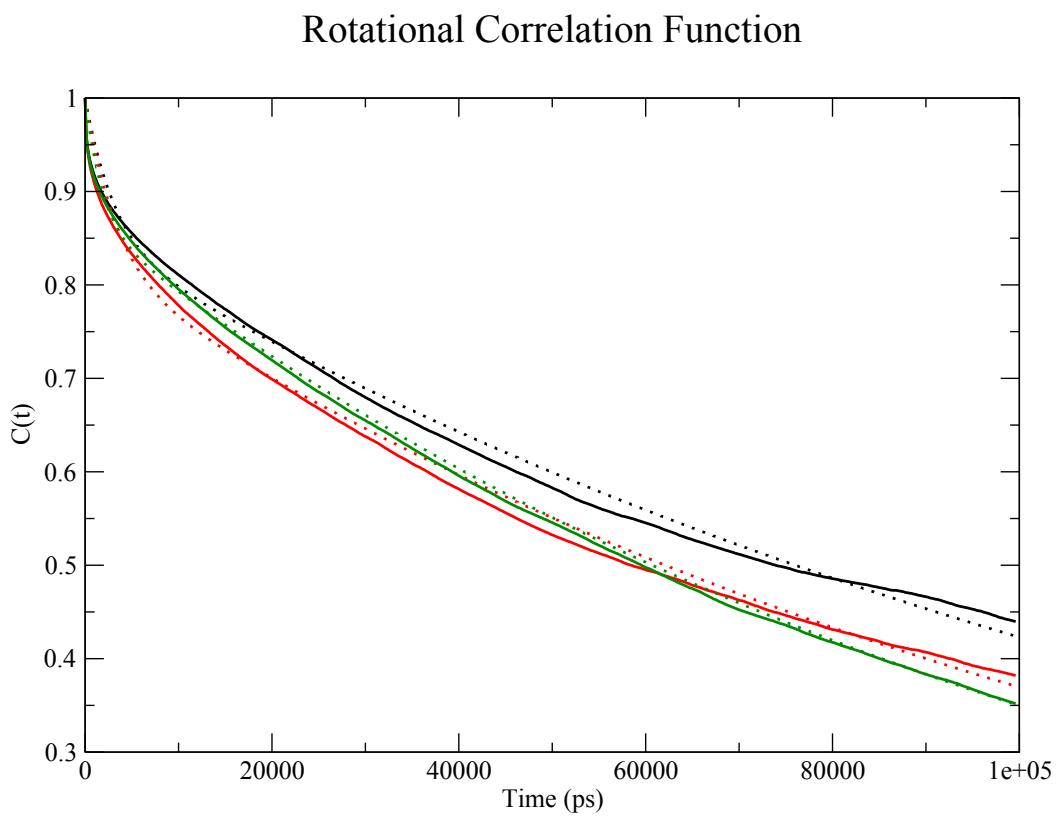


Figure 2: Rotational correlation function (straight lines) and fitting (dotted lines) for: cholesterol (black), probe 8 (red) and probe (10).

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