

Supplemental Information for

Orientation of nitro-group governs the fluorescence lifetime of nitrobenzoxadiazole (NBD)-labeled lipids in lipid bilayers

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Table S1: average lifetimes of C₆NBD-PC and C₁₂NBD-PC in DOPC and POPC LUVs

Average fluorescence lifetime, $\langle\tau\rangle$, of C₆NBD-PC or C₁₂NBD-PC in DOPC or POPC LUVs of different diameter, in H₂O buffer. Error are deviations from at least 3 independent samples.

$\langle\tau\rangle$ (ns)	POPC	DOPC
C₆NBD-PC	7.16 ± 0.18	6.35 ± 0.13
C₁₂NBD-PC	6.21 ± 0.04	5.76 ± 0.05

Topology file for C₆NBD-PC (Gromos 53a6 force field)

[moleculetype]

; Name nrexcl

NBD 3

[atoms]

; nr	type	resnr	residu	atom	cgnr	charge	mass
1	CH3L	1	NBD	C1	1	0.4000	15.0350
2	CH3L	1	NBD	C2	1	0.4000	15.0350
3	CH3L	1	NBD	C3	1	0.4000	15.0350
4	NL	1	NBD	N4	1	-0.5000	14.0067
5	CH2	1	NBD	C5	1	0.3000	14.0270
6	CH2	1	NBD	C6	2	0.4000	14.0270
7	OA	1	NBD	O7	2	-0.8000	15.9994
8	P	1	NBD	P8	2	1.7000	30.9738
9	OML	1	NBD	O9	2	-0.8000	15.9994
10	OML	1	NBD	O10	2	-0.8000	15.9994
11	OA	1	NBD	O11	2	-0.7000	15.9994
12	CH2	1	NBD	C12	3	0.4000	14.0270
13	CH1	1	NBD	C13	3	0.3000	13.0190
14	OE	1	NBD	O14	3	-0.7000	15.9994
15	C	1	NBD	C15	3	0.7000	12.0110
16	O	1	NBD	O16	3	-0.7000	15.9994
17	CH2	1	NBD	C17	4	0.0000	14.0270
18	CH2	1	NBD	C18	4	0.0000	14.0270
19	CH2	1	NBD	C19	5	0.0000	14.0270
20	CH2	1	NBD	C20	5	0.0000	14.0270
21	CH2	1	NBD	C21	6	0.25	14.0270
22	NE	1	NBD	N22	6	-0.34	14.0067
23	C	1	NBD	C23	6	-0.01	12.0110
24	CR1	1	NBD	C24	6	-0.12	13.0190
25	CR1	1	NBD	C25	6	0.19	13.0190
26	C	1	NBD	C26	6	-0.37	12.0110
27	NR	1	NBD	N27	6	0.86	14.0067
28	O	1	NBD	O28	6	-0.51	15.9994
29	O	1	NBD	O29	6	-0.46	15.9994
30	NR	1	NBD	N30	6	-0.30	14.0067
31	C	1	NBD	C31	6	0.41	12.0110
32	C	1	NBD	C32	6	0.41	12.0110
33	OE	1	NBD	O33	6	0.02	15.9994
34	NR	1	NBD	N34	6	-0.32	14.0067
35	H	1	NBD	H35	6	0.29	1.008
36	CH2	1	NBD	C36	7	0.5000	14.0270
37	OE	1	NBD	O37	7	-0.7000	15.9994
38	C	1	NBD	C38	7	0.8000	12.0110
39	O	1	NBD	O39	7	-0.6000	15.9994
40	CH2	1	NBD	C40	8	0.0000	14.0270
41	CH2	1	NBD	C41	8	0.0000	14.0270
42	CH2	1	NBD	C42	9	0.0000	14.0270
43	CH2	1	NBD	C43	9	0.0000	14.0270
44	CH2	1	NBD	C44	10	0.0000	14.0270
45	CH2	1	NBD	C45	10	0.0000	14.0270
46	CH2	1	NBD	C46	11	0.0000	14.0270
47	CH2	1	NBD	C47	11	0.0000	14.0270

48	CH2	1	NBD	C48	12	0.0000	14.0270
49	CH2	1	NBD	C49	12	0.0000	14.0270
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51	CH2	1	NBD	C51	13	0.0000	14.0270
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53	CH2	1	NBD	C53	14	0.0000	14.0270
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[bonds]

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12		13	2	gb_27	
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13		36	2	gb_27	
14		15	2	gb_10	
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23		24	2	0.13940E+00	1.0800e+07
23		32	2	0.14440E+00	1.0800e+07
24		25	2	0.14140E+00	1.0800e+07
25		26	2	0.13800E+00	1.0800e+07
26		27	2	0.14450E+00	8.5400e+06
26		31	2	0.14310E+00	1.0800e+07
27		28	2	0.12370E+00	1.6600e+07
27		29	2	0.12320E+00	1.6600e+07
30		31	2	0.13240E+00	1.1800e+07
30		33	2	0.13780E+00	1.1000e+07
31		32	2	0.14370E+00	1.0800e+07
32		34	2	0.13160E+00	1.1800e+07
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21		22	2	0.14570E+00	8.7100e+06
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[ angles ]
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   12     13     14     2 ga_13
   12     13     36     2 ga_13
   13     14     15     2 ga_22
   13     36     37     2 ga_13
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   14     15     16     2 ga_33
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25	26	31	2	117.70	560.00
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26	31	30	2	133.60	560.00
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31	30	33	2	104.70	465.00
31	32	34	2	109.40	465.00
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21	22	23	2	124.00	730.00
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37	38	40	2	ga_16	
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	6	7	8	11	1	gd_20		
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	7	8	11	12	1	gd_20		
	7	8	11	12	1	gd_27		
	8	11	12	13	1	gd_29		
	11	12	13	36	1	gd_34		

12	13	14	15	1	gd_29			
12	13	36	37	1	gd_34			
13	14	15	17	1	gd_13			
13	36	37	38	1	gd_29			
14	15	17	18	1	gd_40			
15	17	18	19	1	gd_34			
17	18	19	20	1	gd_34			
18	19	20	21	1	gd_34			
19	20	21	22	1	gd_34			
20	21	22	35	1	gd_41			
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31	26	27	29	1	0.0	33.50	2	
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37	38	40	41	1	gd_40			
38	40	41	42	1	gd_34			
40	41	42	43	1	gd_34			
41	42	43	44	1	gd_34			
42	43	44	45	1	gd_34			
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45	46	47	48	1	gd_34			
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47	48	49	50	1	gd_34			
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[dihedrals]

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	26	25	24	23	2	0.00000E+00	0.16740E+03	
	25	24	23	32	2	0.00000E+00	0.16740E+03	
	24	23	32	31	2	0.00000E+00	0.16740E+03	
	29	27	26	31	2	0.00000E+00	0.16740E+03	
	25	26	27	28	2	0.00000E+00	0.16740E+03	

Topology file for C₁₂NBD-PC (Gromos 53a6 force field)

[moleculetype]

; Name nrexcl

NBD 3

[atoms]

; nr	type	resnr	residu	atom	cgnr	charge	mass
1	CH3L	1	NBD	C1	1	0.4000	15.0350
2	CH3L	1	NBD	C2	1	0.4000	15.0350
3	CH3L	1	NBD	C3	1	0.4000	15.0350
4	NL	1	NBD	N4	1	-0.5000	14.0067
5	CH2	1	NBD	C5	1	0.3000	14.0270
6	CH2	1	NBD	C6	2	0.4000	14.0270
7	OA	1	NBD	O7	2	-0.8000	15.9994
8	P	1	NBD	P8	2	1.7000	30.9738
9	OML	1	NBD	O9	2	-0.8000	15.9994
10	OML	1	NBD	O10	2	-0.8000	15.9994
11	OA	1	NBD	O11	2	-0.7000	15.9994
12	CH2	1	NBD	C12	3	0.4000	14.0270
13	CH1	1	NBD	C13	3	0.3000	13.0190
14	OE	1	NBD	O14	3	-0.7000	15.9994
15	C	1	NBD	C15	3	0.7000	12.0110
16	O	1	NBD	O16	3	-0.7000	15.9994
17	CH2	1	NBD	C17	4	0.0000	14.0270
18	CH2	1	NBD	C18	4	0.0000	14.0270
19	CH2	1	NBD	C19	5	0.0000	14.0270
20	CH2	1	NBD	C20	5	0.0000	14.0270
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25	CH2	1	NBD	C25	8	0.0000	14.0270
26	CH2	1	NBD	C26	8	0.0000	14.0270
27	CH2	1	NBD	C27	9	0.25	14.0270
28	NE	1	NBD	N28	9	-0.34	14.0067
29	C	1	NBD	C29	9	-0.01	12.0110
30	CR1	1	NBD	C30	9	-0.12	13.0190
31	CR1	1	NBD	C31	9	0.19	13.0190
32	C	1	NBD	C32	9	-0.37	12.0110
33	NR	1	NBD	N33	9	0.86	14.0067
34	O	1	NBD	O34	9	-0.51	15.9994
35	O	1	NBD	O35	9	-0.46	15.9994
36	NR	1	NBD	N36	9	-0.30	14.0067
37	C	1	NBD	C37	9	0.41	12.0110
38	C	1	NBD	C38	9	0.41	12.0110
39	OE	1	NBD	O39	9	0.02	15.9994
40	NR	1	NBD	N40	9	-0.32	14.0067
41	H	1	NBD	H41	9	0.29	1.008
42	CH2	1	NBD	C42	10	0.5000	14.0270
43	OE	1	NBD	O43	10	-0.7000	15.9994
44	C	1	NBD	C44	10	0.8000	12.0110
45	O	1	NBD	O45	10	-0.6000	15.9994
46	CH2	1	NBD	C46	11	0.0000	14.0270
47	CH2	1	NBD	C47	11	0.0000	14.0270

48	CH2	1	NBD	C48	12	0.0000	14.0270
49	CH2	1	NBD	C49	12	0.0000	14.0270
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51	CH2	1	NBD	C51	13	0.0000	14.0270
52	CH2	1	NBD	C52	14	0.0000	14.0270
53	CH2	1	NBD	C53	14	0.0000	14.0270
54	CH2	1	NBD	C54	15	0.0000	14.0270
55	CH2	1	NBD	C55	15	0.0000	14.0270
56	CH2	1	NBD	C56	16	0.0000	14.0270
57	CH2	1	NBD	C57	16	0.0000	14.0270
58	CH2	1	NBD	C58	17	0.0000	14.0270
59	CH2	1	NBD	C59	17	0.0000	14.0270
60	CH3	1	NBD	C60	18	0.0000	15.0350

[bonds]

```

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2 4 2 gb_21
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4 5 2 gb_21
5 6 2 gb_27
6 7 2 gb_18
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8 10 2 gb_24
8 11 2 gb_28
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13 42 2 gb_27
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24 25 2 gb_27
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26 27 2 gb_27
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28 41 2 0.10120E+00 1.8700e+07
29 30 2 0.13940E+00 1.0800e+07
29 38 2 0.14440E+00 1.0800e+07
30 31 2 0.14140E+00 1.0800e+07
31 32 2 0.13800E+00 1.0800e+07
32 33 2 0.14450E+00 8.5400e+06
32 37 2 0.14310E+00 1.0800e+07
33 34 2 0.12370E+00 1.6600e+07
33 35 2 0.12320E+00 1.6600e+07
36 37 2 0.13240E+00 1.1800e+07
36 39 2 0.13780E+00 1.1000e+07
37 38 2 0.14370E+00 1.0800e+07

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38	40	2	0.13160E+00	1.1800e+07
39	40	2	0.13600E+00	1.1000e+07
27	28	2	0.14570E+00	8.7100e+06
42	43	2	gb_18	
43	44	2	gb_10	
44	45	2	gb_5	
44	46	2	gb_23	
46	47	2	gb_27	
47	48	2	gb_27	
48	49	2	gb_27	
49	50	2	gb_27	
50	51	2	gb_27	
51	52	2	gb_27	
52	53	2	gb_27	
53	54	2	gb_27	
54	55	2	gb_27	
55	56	2	gb_27	
56	57	2	gb_27	
57	58	2	gb_27	
58	59	2	gb_27	
59	60	2	gb_27	

[pairs]

;	ai	aj	funct
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	2	6	1
	3	6	1
	4	7	1
	5	8	1
	6	9	1
	6	10	1
	6	11	1
	7	12	1
	8	13	1
	9	12	1
	10	12	1
	11	14	1
	11	42	1
	12	15	1
	12	43	1
	13	16	1
	13	17	1
	13	44	1
	14	18	1
	14	43	1
	15	19	1
	15	42	1
	16	18	1
	17	20	1
	18	21	1
	19	22	1
	20	23	1
	21	24	1
	22	25	1
	23	26	1
	24	27	1

28	31	1
28	40	1
28	37	1
29	32	1
29	36	1
29	39	1
30	33	1
30	37	1
30	40	1
30	41	1
31	34	1
31	35	1
31	36	1
31	38	1
32	39	1
32	40	1
33	36	1
33	38	1
34	37	1
35	37	1
38	41	1
25	28	1
27	30	1
27	38	1
26	29	1
26	41	1
42	45	1
42	46	1
43	47	1
44	48	1
45	47	1
46	49	1
47	50	1
48	51	1
49	52	1
50	53	1
51	54	1
52	55	1
53	56	1
54	57	1
55	58	1
56	59	1
57	60	1

```
[ angles ]
; ai    aj    ak funct
      1     4     2     2 ga_13
      1     4     3     2 ga_13
      1     4     5     2 ga_13
      2     4     3     2 ga_13
      2     4     5     2 ga_13
      3     4     5     2 ga_13
      4     5     6     2 ga_15
      5     6     7     2 ga_15
      6     7     8     2 ga_26
      7     8     9     2 ga_14
```

7	8	10	2	ga_14	
7	8	11	2	ga_5	
8	11	12	2	ga_26	
9	8	10	2	ga_29	
9	8	11	2	ga_14	
10	8	11	2	ga_14	
11	12	13	2	ga_15	
12	13	14	2	ga_13	
12	13	42	2	ga_13	
13	14	15	2	ga_22	
13	42	43	2	ga_13	
14	13	42	2	ga_13	
14	15	16	2	ga_33	
14	15	17	2	ga_16	
15	17	18	2	ga_15	
16	15	17	2	ga_30	
17	18	19	2	ga_15	
18	19	20	2	ga_15	
19	20	21	2	ga_15	
20	21	22	2	ga_15	
21	22	23	2	ga_15	
22	23	24	2	ga_15	
23	24	25	2	ga_15	
24	25	26	2	ga_15	
25	26	27	2	ga_15	
28	29	30	2	125.60	750.00
28	29	38	2	119.00	750.00
29	28	41	2	117.20	390.00
29	30	31	2	121.40	560.00
29	38	37	2	123.30	560.00
29	38	40	2	127.30	560.00
30	29	38	2	115.50	560.00
30	31	32	2	123.90	560.00
31	32	33	2	120.10	560.00
31	32	37	2	117.70	560.00
32	33	34	2	118.00	750.00
32	33	35	2	117.30	750.00
32	37	36	2	133.60	560.00
32	37	38	2	118.30	560.00
33	32	37	2	122.20	560.00
34	33	35	2	124.80	750.00
36	37	38	2	108.10	465.00
36	39	40	2	113.10	450.00
37	36	39	2	104.70	465.00
37	38	40	2	109.40	465.00
38	40	39	2	104.70	465.00
27	28	41	2	118.60	460.00
27	28	29	2	124.00	730.00
26	27	28	2	111.10	530.00
42	43	44	2	ga_22	
43	44	45	2	ga_33	
43	44	46	2	ga_16	
44	46	47	2	ga_15	
45	44	46	2	ga_30	
46	47	48	2	ga_15	
47	48	49	2	ga_15	

48	49	50	2	ga_15
49	50	51	2	ga_15
50	51	52	2	ga_15
51	52	53	2	ga_15
52	53	54	2	ga_15
53	54	55	2	ga_15
54	55	56	2	ga_15
55	56	57	2	ga_15
56	57	58	2	ga_15
57	58	59	2	ga_15
58	59	60	2	ga_15

[dihedrals]

;	ai	aj	ak	al	funct	phi0	cp	mult
	1	4	5	6	1	gd_29		
	4	5	6	7	1	gd_4		
	4	5	6	7	1	gd_36		
	5	6	7	8	1	gd_29		
	6	7	8	11	1	gd_20		
	6	7	8	11	1	gd_27		
	7	8	11	12	1	gd_20		
	7	8	11	12	1	gd_27		
	8	11	12	13	1	gd_29		
	11	12	13	42	1	gd_34		
	12	13	14	15	1	gd_29		
	12	13	42	43	1	gd_34		
	13	14	15	17	1	gd_13		
	13	42	43	44	1	gd_29		
	14	15	17	18	1	gd_40		
	15	17	18	19	1	gd_34		
	17	18	19	20	1	gd_34		
	18	19	20	21	1	gd_34		
	19	20	21	22	1	gd_34		
	20	21	22	23	1	gd_34		
	21	22	23	24	1	gd_34		
	22	23	24	25	1	gd_34		
	23	24	25	26	1	gd_34		
	24	25	26	27	1	gd_34		
	25	26	27	28	1	gd_34		
	26	27	28	41	1	gd_41		
	27	28	29	30	1	0.0	33.50	2
	37	32	33	35	1	0.0	33.50	2
	42	43	44	46	1	gd_13		
	43	44	46	47	1	gd_40		
	44	46	47	48	1	gd_34		
	46	47	48	49	1	gd_34		
	47	48	49	50	1	gd_34		
	48	49	50	51	1	gd_34		
	49	50	51	52	1	gd_34		
	50	51	52	53	1	gd_34		
	51	52	53	54	1	gd_34		
	52	53	54	55	1	gd_34		
	53	54	55	56	1	gd_34		
	54	55	56	57	1	gd_34		
	55	56	57	58	1	gd_34		
	56	57	58	59	1	gd_34		

57 58 59 60 1 gd_34

[dihedrals]

```
; ai    aj    ak    al funct
    13    14    42    12      2    gi_2
    15    14    16    17      2    gi_1
    44    43    45    46      2    gi_1
    28    27    41    29      2  0.00000E+00 0.16740E+03
    29    28    38    30      2  0.00000E+00 0.16740E+03
    38    29    37    40      2  0.00000E+00 0.16740E+03
    37    38    32    36      2  0.00000E+00 0.16740E+03
    32    37    33    31      2  0.00000E+00 0.16740E+03
    33    32    35    34      2  0.00000E+00 0.16740E+03
    38    40    39    36      2  0.00000E+00 0.16740E+03
    40    39    36    37      2  0.00000E+00 0.16740E+03
    39    36    37    38      2  0.00000E+00 0.16740E+03
    36    37    38    40      2  0.00000E+00 0.16740E+03
    37    38    40    39      2  0.00000E+00 0.16740E+03
    29    38    37    32      2  0.00000E+00 0.16740E+03
    38    37    32    31      2  0.00000E+00 0.16740E+03
    37    32    31    30      2  0.00000E+00 0.16740E+03
    32    31    30    29      2  0.00000E+00 0.16740E+03
    31    30    29    38      2  0.00000E+00 0.16740E+03
    30    29    38    37      2  0.00000E+00 0.16740E+03
    35    33    32    37      2  0.00000E+00 0.16740E+03
    31    32    33    34      2  0.00000E+00 0.16740E+03
```


Topology file for C₄NBD-PC (Martini 2.0 force field)

```
[moleculetype]
; molname      nrexcl
  NC04         1

[atoms]
; id type  resnr      residu    atom  cgnr  charge
  1 Q0     1    NC04  NC3     1    1.0
  2 Qa     1    NC04  PO4     2   -1.0
  3 Na     1    NC04  GL1     3    0
  4 Na     1    NC04  GL2     4    0
  5 C2     1    NC04  C1A     5    0
  6 SNda   1    NC04  NHC     4    0
  7 C1     1    NC04  C1B     7    0
  8 C1     1    NC04  C2B     8    0
  9 C1     1    NC04  C3B     9    0
 10 C1     1    NC04  C4B    10    0
 11 SP3    1    NC04  NO2    11    0
 12 SC4    1    NC04  CC1    12    0
 13 SNa    1    NC04  NON    13    0
 14 SC4    1    NC04  CC2    14    0

[bonds]
; i  j      funct      length      force.c.
  1  2      1         0.47 1250
  2  3      1         0.47 1250
  3  4      1         0.37 1250
  3  5      1         0.47 1250
  5  6      1         0.34 1250
  4  7      1         0.47 1250
  7  8      1         0.47 1250
  8  9      1         0.47 1250
  9 10      1         0.47 1250

[constraints]
; i  j      funct      length
 11 12      1         0.317
 11 14      1         0.311
 12 13      1         0.153
  6 12      1         0.251
  6 14      1         0.247

[angles]
; i  j  k      funct      angle      force.c.
  2  3  4      2         120.0     25.0
  2  3  5      2         180.0     25.0
  3  5  6      2         180.0     25.0
  4  7  8      2         180.0     25.0
  7  8  9      2         180.0     25.0
  8  9 10      2         180.0     25.0
 11 12 13      2         116.0    1800.0
 11 14  6      2         130.0    2400.0
 12  6 14      2           58.0    6000.0
```

13	12	6	2	117.5	1900.0
14	6	5	2	135.0	25.0

[dihedrals]

;	i	j	k	l	funct	angle	force.c.
	11	12	6	14	2	0.0	25.0
	12	6	14	11	2	0.0	25.0
	6	14	11	12	2	0.0	25.0
	14	11	12	6	2	0.0	25.0
	12	11	13	6	2	0.0	200.0

[exclusions]

11	12	13	6	14
12	13	6	14	
13	6	14		
6	14			
12	14			

Topology file for C₁₂NBD-PC (Martini 2.0 force field)

```
[moleculetype]
; molname      nrexcl
  NC12         1

[atoms]
; id type  resnr      residu    atom  cgnr  charge
  1 Q0     1    NC12  NC3     1    1.0
  2 Qa     1    NC12  PO4     2   -1.0
  3 Na     1    NC12  GL1     3    0
  4 Na     1    NC12  GL2     4    0
  5 C1     1    NC12  C1A     5    0
  6 C1     1    NC12  C2A     6    0
  7 C2     1    NC12  C3A     7    0
  8 SNda   1    NC12  NHC     8    0
  9 C1     1    NC12  C1B     9    0
 10 C1     1    NC12  C2B    10    0
 11 C1     1    NC12  C3B    11    0
 12 C1     1    NC12  C4B    12    0
 13 SP3    1    NC12  NO2    13    0
 14 SC4    1    NC12  CC1    14    0
 15 SNa    1    NC12  NON    15    0
 16 SC4    1    NC12  CC2    16    0

[bonds]
; i  j      funct      length      force.c.
  1  2      1         0.47 1250
  2  3      1         0.47 1250
  3  4      1         0.37 1250
  3  5      1         0.47 1250
  5  6      1         0.47 1250
  6  7      1         0.47 1250
  7  8      1         0.34 1250
  4  9      1         0.47 1250
  9 10      1         0.47 1250
 10 11      1         0.47 1250
 11 12      1         0.47 1250

[constraints]
; i  j      funct      length
 13 14      1         0.317
 13 16      1         0.311
 14 15      1         0.153
  8 14      1         0.251
  8 16      1         0.247

[angles]
; i  j  k      funct      angle      force.c.
  2  3  4      2         120.0     25.0
  2  3  5      2         180.0     25.0
  3  5  6      2         180.0     25.0
  5  6  7      2         180.0     25.0
  6  7  8      2         180.0     25.0
```

4	9	10	2	180.0	25.0
9	10	11	2	180.0	25.0
10	11	12	2	180.0	25.0
13	14	15	2	116.0	1800.0
13	16	8	2	130.0	2400.0
14	8	16	2	58.0	6000.0
15	14	8	2	117.5	1900.0
16	8	7	2	135.0	25.0

[dihedrals]

;	i	j	k	l	funct	angle	force.c.
	13	14	8	16	2	0.0	25.0
	14	8	16	13	2	0.0	25.0
	8	16	13	14	2	0.0	25.0
	16	13	14	8	2	0.0	25.0
	14	13	15	8	2	0.0	200.0

[exclusions]

13	14	15	8	16
14	15	8	16	
15	8	16		
8	16			
14	16			

Description of the Martini parameterization of the NBD fluorophore

The number and distribution of the Martini beads, i.e. mapping, was chosen in order to preserve the geometrical structure of the NBD, which could be done with a five bead configuration, as shown in Fig. S1. Then, the attributed bonded and non-bonded (i.e. bead types) parameters, defined based on the mapping process, were tested and validated comparing the coarse grained (CG) model to atomistic data for the probes composed by the NBD group attached to an alkyl chains of different sizes, NBD-C₄ and NBD-C₈.^{1, 2} The bonded parameters were validated comparing 10 ns simulations of the molecule NBD-C₈ in water at both resolution levels. As shown in Fig. S2 to S4, the bond, angle and dihedral distribution of the CG model matched very well the atomistic data. Then, the non-bonded parameters were validated against our previous reported potential of mean force (PMF) profiles for the NBD-C_n homologous series,² in this case for both NBD-C₄ and NBD-C₈. The sampling at each umbrella window was done during 200 ns, discarding the first 50 ns as equilibration time. A time step of 20 fs was used in all simulations. As shown in Fig. S5, the PMFs obtained with the CG parameterizations are similar to those obtained at atomistic level, validating the type of Martini beads used.

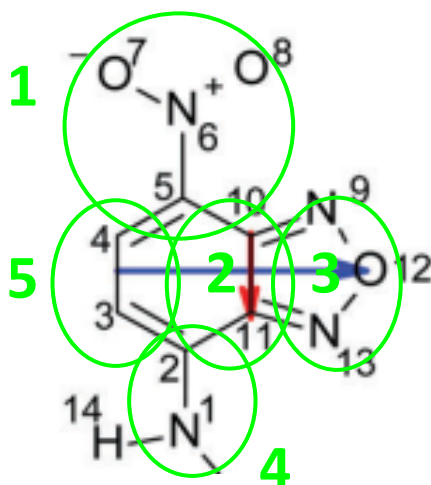


Fig. S1 – Coarse grained mapping of the NBD fluorophore for the Martini parameterization regarding its atomistic structure. The groups of atoms (5, 6, 7, 8), (10, 11), (9, 12, 13), (1,2) and (3, 4) were defined as SP3, SC4, SNa, SNda, and SC4 Martini beads, respectively. Bead numbers are indicated in green.

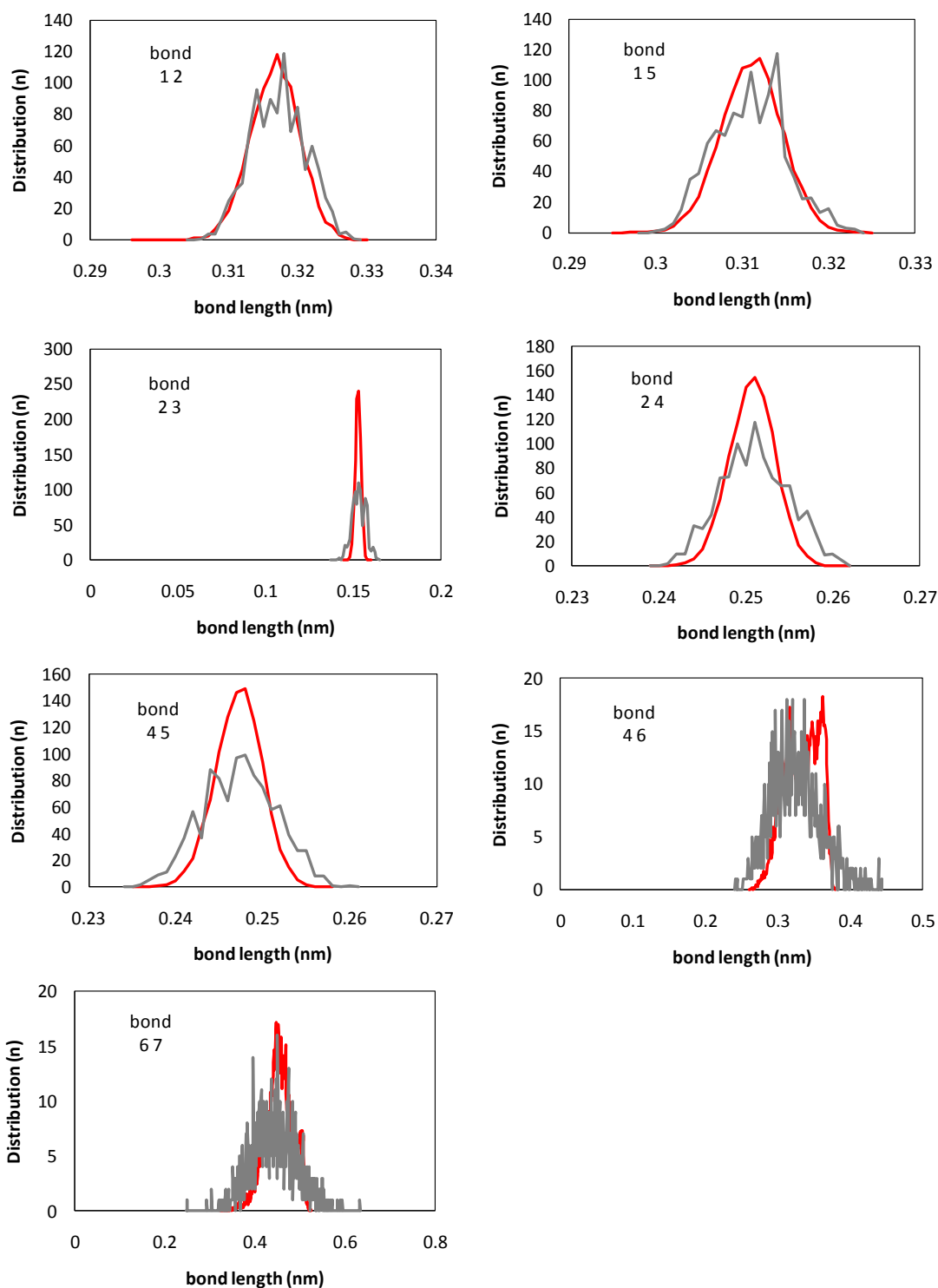


Fig. S2 – Comparison of bond length distributions for the martini (gray) and mapped G53A6FF (red) parameterization of NBD-C₈ as used in reference ¹. For bond identification, please refer to bead number in Figure S1. Bonds (1-2), (1-5), (2-3), (2-4) and (4-5) were set to constraints and bonds (4-6) and (6-7) were set as regular bonds.

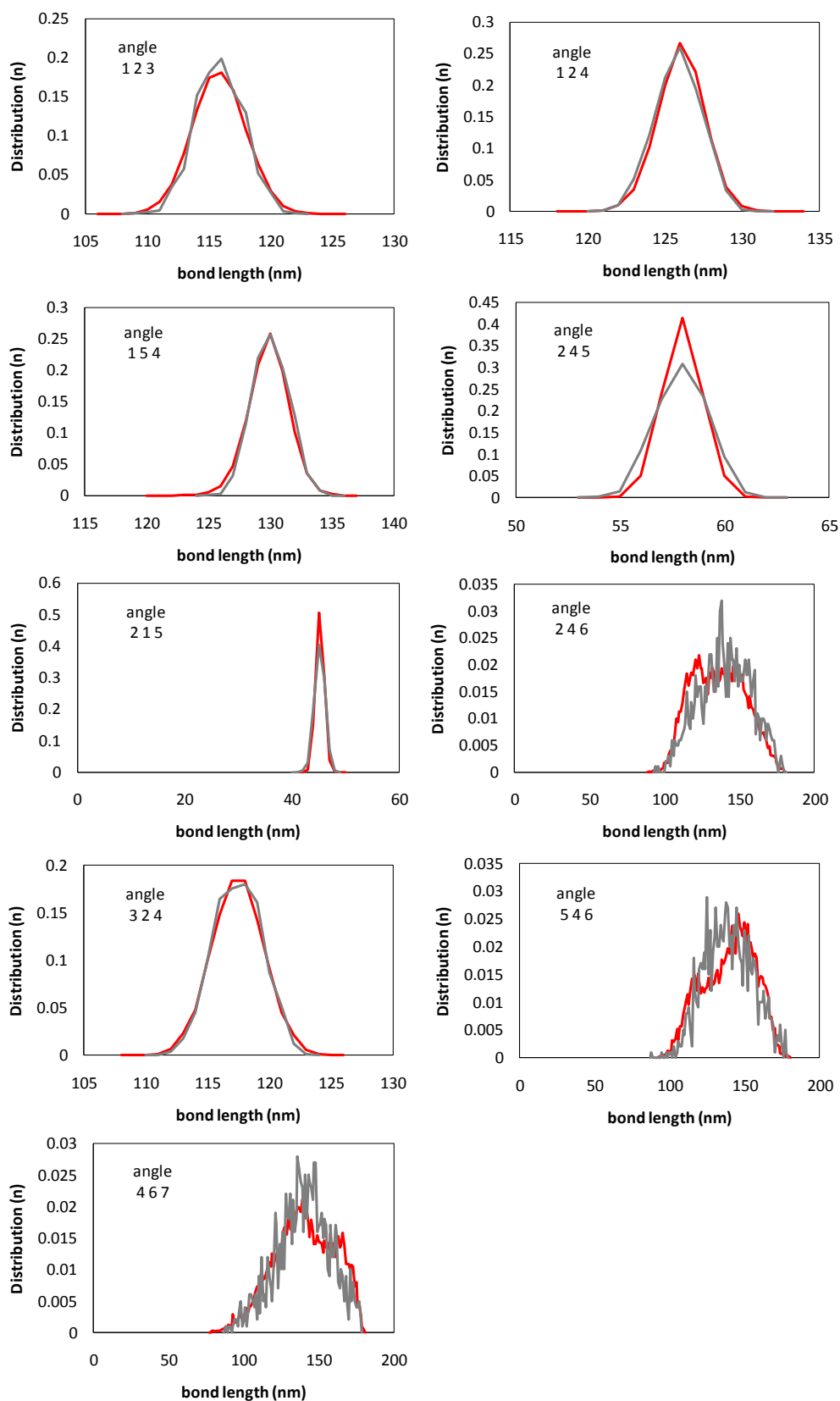


Fig. S3 – Comparison of angle distributions for the martini (gray) and mapped G53A6FF (red) parameterization of NBD-C₈ as used in reference ¹. For angle identification, please refer to bead number in Figure S1. Note that angles (1-2-4), (2-1-5) and (2-4-6) are actually not defined in the parametrization file.

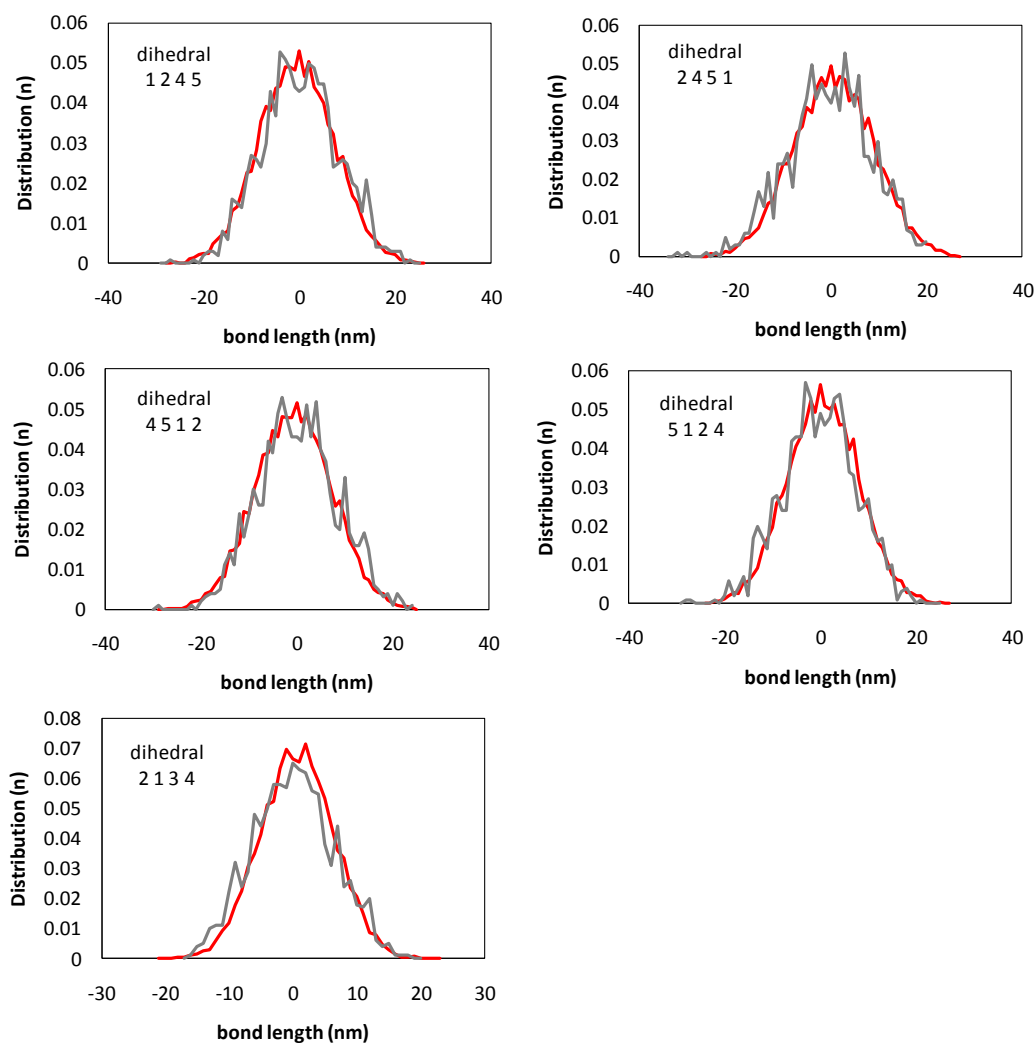


Fig. S4 – Comparison of bond dihedral distributions for the martini (gray) and mapped G53A6FF (red) parameterization of NBD-C₈ as used in reference ¹. For dihedral identification, please refer to bead number in Figure S1.

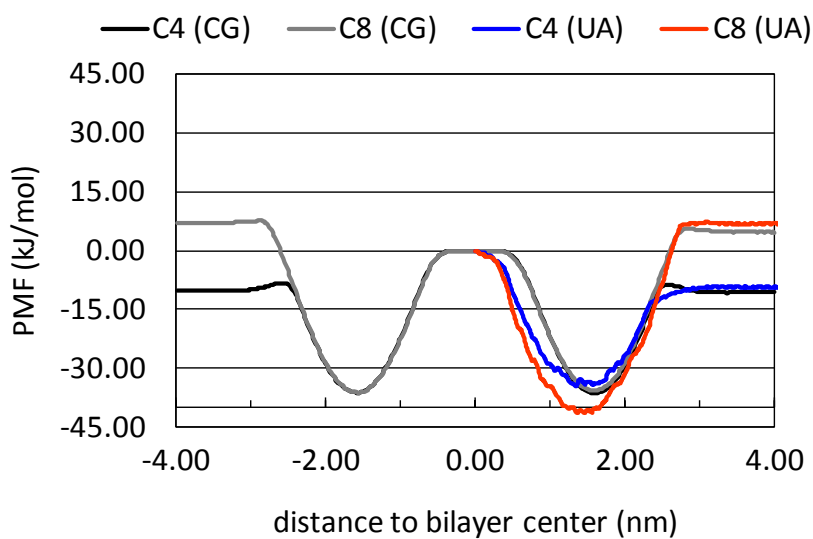


Fig. S5 – Potential of Mean Force (PMF) profiles for the interaction of NBD-C₄ and NBD-C₈ with a POPC bilayer at atomistic ² and coarse grained (this work) level.

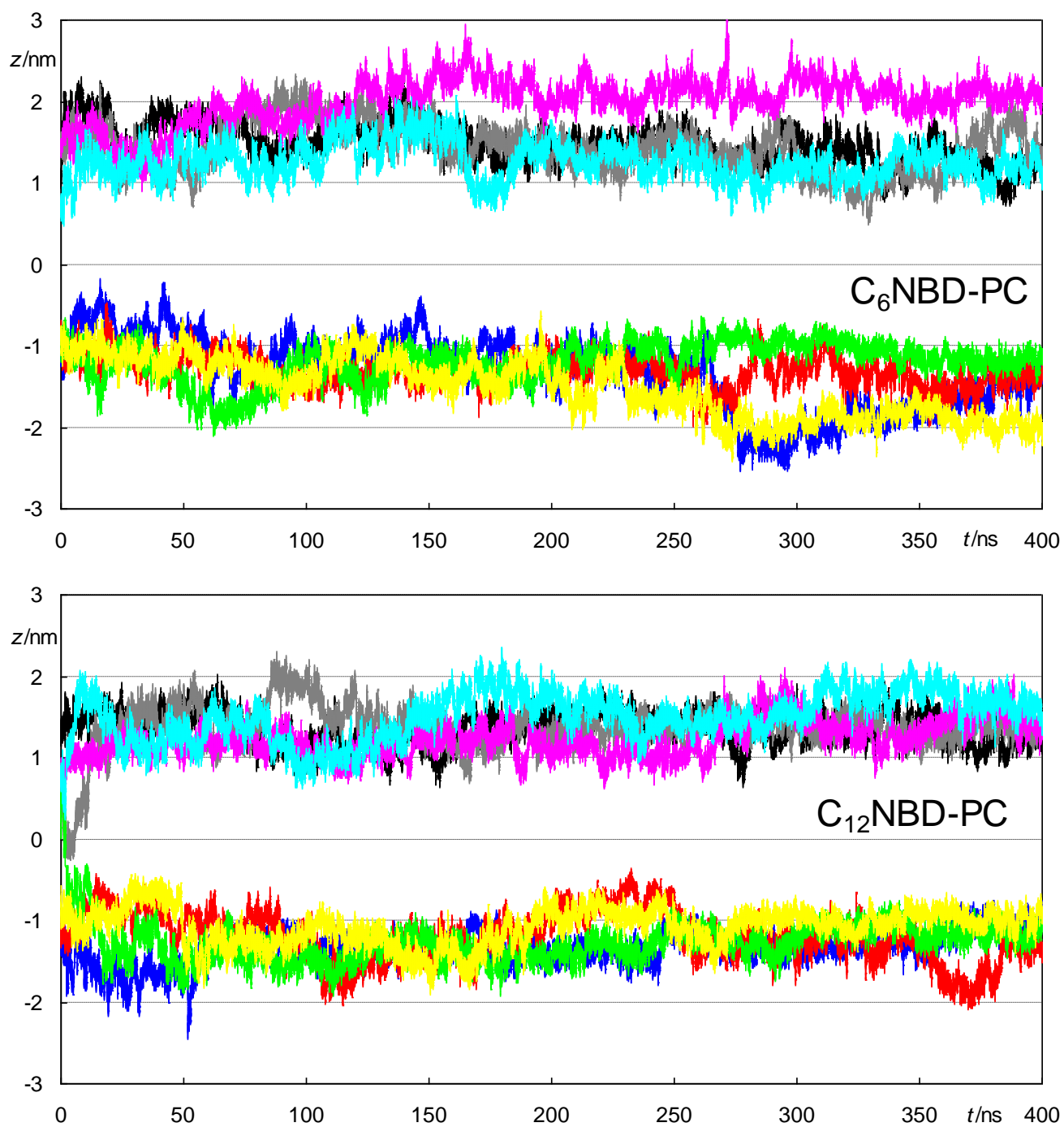


Fig. S6 - Time evolution of the center of mass of the fluorophores of the eight NBD-PC molecules considered in the atomistic simulations, each of them depicted in a distinct color. Top: C_6 NBD-PC; bottom: C_{12} NBD-PC.

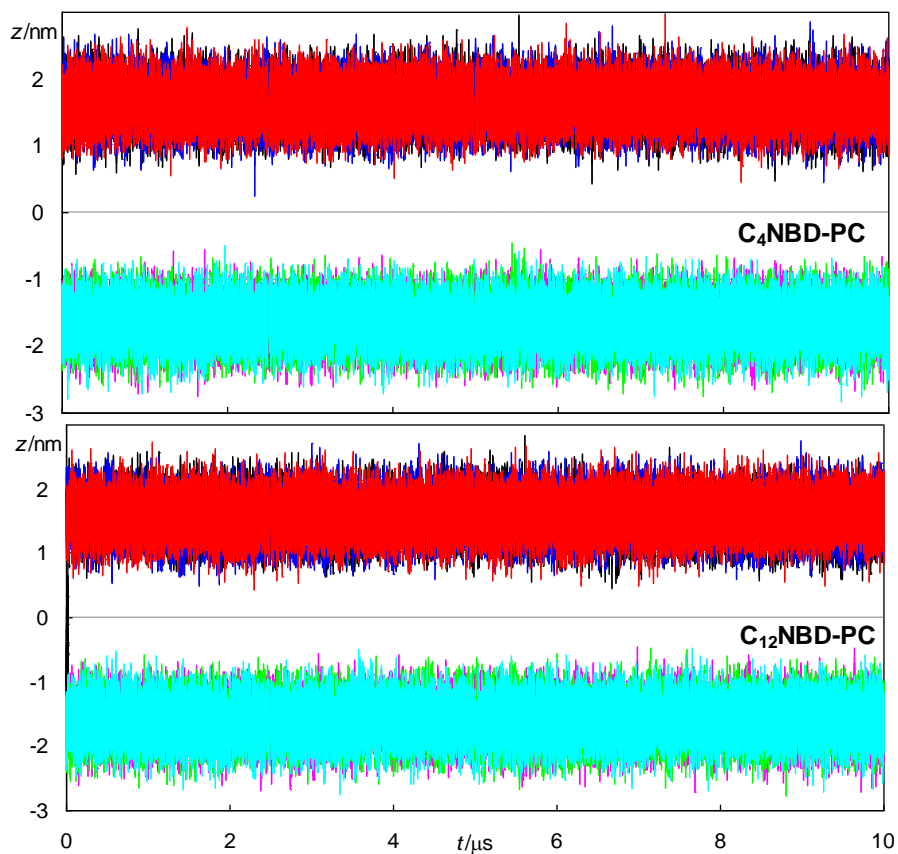


Fig. S7 - Time evolution of the transverse location z of the center of mass of the NBD fluorophore of each individual probe (black, blue and red for the top leaflet molecules; pink, green and cyan for the bottom leaflet molecules) in the CG simulations of C₄NBD-PC (top) and C₁₂NBD-PC (bottom).

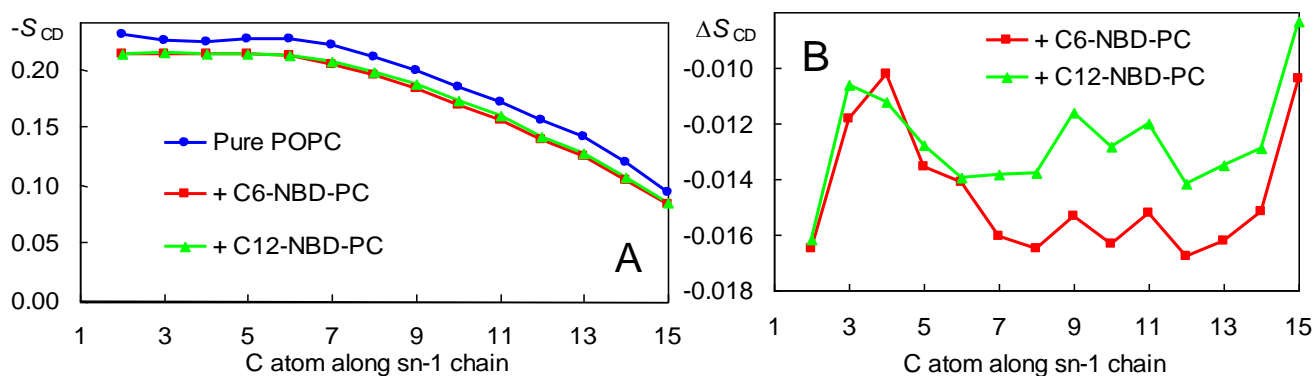


Fig. S8 - (A) Deuterium order parameters (S_{CD}) for the POPC *sn*-1 atoms in the absence and in the presence of C₆- or C₁₂NBD-PC, calculated from the atomistic simulations. (B) Difference order parameter plots (ΔS_{CD}), obtained by subtraction of the pure POPC order parameter profile from those calculated in the presence of C₆- or C₁₂NBD-PC.

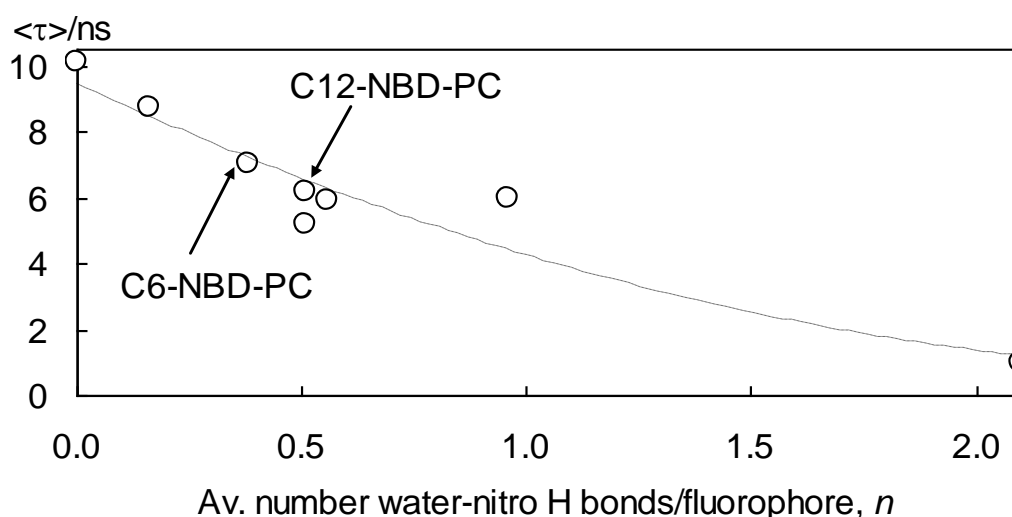


Fig. S9 - Intensity-averaged lifetimes $\langle \tau \rangle$ obtained for different NBD probes in fluid PC bilayers,³⁻⁷ represented as a function of their average number of water-NBD NO₂ hydrogen bonds determined by MD simulation.^{3, 8-10} The point (0, 10.1 ns) corresponds to polar aprotic solvents such as acetone, tetrahydrofuran or dioxane¹¹. For water, $\langle \tau \rangle = 1.0$ ns,¹¹ and $n = 2.1$ was calculated from a simulation of NBD-C₄ in water. The points corresponding to the probes addressed in this work are highlighted with arrows. For more details, see Table S2.

Table S2 - MD simulation data on orientation and H-bonding, and experimental values of fluorescence lifetime for different NBD lipid probes. Unless stated otherwise, the data were obtained at room temperature/298 K.

Probe	Orientation and H-bonding from MD				Fluorescence lifetime		
	Short axis tilt/ °	Average number of water-NBD NO ₂ H-bonds/ fluorophore	System	Ref.	Intensity-averaged τ /ns	System	Ref.
NBD-C _n (n ≥ 8)	130	0.51	POPC	⁸	5.2	POPC	⁴
22-NBD-Chol	130	0.96	POPC	⁹	6.0 ⁵ , 6.2 ⁶	DMPC, 40 °C ⁵ or 37 °C ⁶	^{5,6}
C ₁₂ -NBD-PC	130	0.51	POPC	This work	6.2	POPC	This work
NBD-PSH	100	0.56	DOPC	³	5.9	DOPC	³
C ₆ -NBD-PC	100	0.38	POPC	This work	7.0	POPC	This work
N-NBD-PE	70	0.16	POPC	¹⁰	8.7	Egg-PC	⁷

NBD-C_n: *N*-[(7-nitro-2-1,3-benzoxadiazol-4-yl)amino]alkyl; 22-NBD-Chol: 22-(*N*-(7-nitrobenz-2-1,3-benzoxadiazol-4-yl)amino)-23,24-bisnor-5-cholen-3 β -ol; C₁₂-NBD-PC: 1-palmitoyl-2-[12-[(7-nitro-2-1,3-benzoxadiazol-4-yl)amino]dodecanoyl]-*sn*-glycero-3-phosphocholine; NBD-PSH: 1,2-dioleoyl-*sn*-glycero-3-phospho-L-serine-*N*-(7-nitro-2-1,3-benzoxadiazol-4-yl); C₆-NBD-PC: 1-palmitoyl-2-[6-[(7-nitro-2-1,3-benzoxadiazol-4-yl)amino]hexanoyl]-*sn*-glycero-3-phosphocholine; *N*-NBD-PE: 1,2-dipalmitoyl-*sn*-glycero-3-phosphoethanolamine-*N*-(7-nitro-2-1,3-benzoxadiazol-4-yl); POPC: 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine; DOPC: 1,2-dioleoyl-*sn*-glycero-3-phosphocholine; DMPC: 1,2-dimyristoyl-*sn*-glycero-3-phosphocholine; Egg-PC: L- α -phosphatidylcholine from chicken egg.

References

1. H. A. L. Filipe, D. Bowman, T. Palmeira, R. M. S. Cardoso, L. M. S. Loura and M. J. Moreno, *Phys. Chem. Chem. Phys.*, 2015, **17**, 27534-27547.
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