

Supporting information for:

Limited Perturbation of a DPPC Bilayer by Fluorescent Lipid Probes: A Molecular Dynamics Study

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S1 DiI Topology Files

The topology files for DiI-C18:0, DiI-C18:2 and DiI-C12:0 are listed below:

S1.1 DiI-C18:0

```
[ moleculetype ]
; Name nrexcl
DII      3

[ atoms ]
;  nr      type  resnr  resid  atom  cgnr  charge  mass
  1      LP3     1    DII     C1     1    0.000  15.0350
  2      LP2     1    DII     C2     2    0.000  14.0270
  3      LP2     1    DII     C3     3    0.000  14.0270
  4      LP2     1    DII     C4     4    0.000  14.0270
  5      LP2     1    DII     C5     5    0.000  14.0270
  6      LP2     1    DII     C6     6    0.000  14.0270
  7      LP2     1    DII     C7     7    0.000  14.0270
  8      LP2     1    DII     C8     8    0.000  14.0270
  9      LP2     1    DII     C9     9    0.000  14.0270
 10      LP2     1    DII    C10    10    0.000  14.0270
 11      LP2     1    DII    C11    11    0.000  14.0270
 12      LP2     1    DII    C12    12    0.000  14.0270
 13      LP2     1    DII    C13    13    0.000  14.0270
 14      LP2     1    DII    C14    14    0.000  14.0270
 15      LP2     1    DII    C15    15    0.000  14.0270
 16      LP2     1    DII    C16    16    0.000  14.0270
 17      LP2     1    DII    C17    17    0.000  14.0270
 18      LP2     1    DII    C18    18    0.150  14.0270
 19      NR      1    DII   N19    19   -0.010  14.0067
 20      C       1    DII   C20    20    0.100  12.0110
 21      CR1     1    DII   C21    21   -0.040  13.0190
 22      CR1     1    DII   C22    22    0.040  13.0190
 23      CR1     1    DII   C23    23    0.010  13.0190
 24      CR1     1    DII   C24    24    0.010  13.0190
 25      C       1    DII   C25    25   -0.100  12.0110
 26      CH1     1    DII   C26    26    0.400  12.0110
 27      CH3     1    DII   C27    27   -0.060  15.0350
 28      CH3     1    DII   C28    28   -0.070  15.0350
 29      C       1    DII   C29    29    0.300  12.0110
 30      CR1     1    DII   C30    30   -0.430  13.0190
 31      CR1     1    DII   C31    31    0.370  13.0190
 32      CR1     1    DII   C32    32   -0.340  13.0190
 33      C       1    DII   C33    33    0.410  12.0110
 34      CH1     1    DII   C34    34    0.140  12.0110
 35      CH3     1    DII   C35    35   -0.030  15.0350
 36      CH3     1    DII   C36    36   -0.040  15.0350
 37      C       1    DII   C37    37    0.030  12.0110
 38      CR1     1    DII   C38    38   -0.020  13.0190
 39      CR1     1    DII   C39    39    0.010  13.0190
 40      CR1     1    DII   C40    40    0.040  13.0190
 41      CR1     1    DII   C41    41   -0.060  13.0190
 42      C       1    DII   C42    42    0.140  12.0110
 43      N       1    DII  N43    43   -0.100  14.0067
 44      LP2     1    DII   C44    44    0.150  14.0270
 45      LP2     1    DII   C45    45    0.000  14.0270
 46      LP2     1    DII   C46    46    0.000  14.0270
 47      LP2     1    DII   C47    47    0.000  14.0270
 48      LP2     1    DII   C48    48    0.000  14.0270
 49      LP2     1    DII   C49    49    0.000  14.0270
 50      LP2     1    DII   C50    50    0.000  14.0270
 51      LP2     1    DII   C51    51    0.000  14.0270
 52      LP2     1    DII   C52    52    0.000  14.0270
 53      LP2     1    DII   C53    53    0.000  14.0270
 54      LP2     1    DII   C54    54    0.000  14.0270
 55      LP2     1    DII   C55    55    0.000  14.0270
 56      LP2     1    DII   C56    56    0.000  14.0270
```

57	LP2	1	DII	C57	57	0.000	14.0270
58	LP2	1	DII	C58	58	0.000	14.0270
59	LP2	1	DII	C59	59	0.000	14.0270
60	LP2	1	DII	C60	60	0.000	14.0270
61	LP3	1	DII	C61	61	0.000	15.0350

[bonds]

```
; ai  aj  fu  c0, c1, ...
1  2  1  0.153  334720.0  0.153  334720.0 ; CAA CAR
2  3  1  0.153  334720.0  0.153  334720.0 ; CAR CAT
3  4  1  0.153  334720.0  0.153  334720.0 ; CAT CAV
4  5  1  0.153  334720.0  0.153  334720.0 ; CAV CAX
5  6  1  0.153  334720.0  0.153  334720.0 ; CAX CAZ
6  7  1  0.153  334720.0  0.153  334720.0 ; CAZ CBB
7  8  1  0.153  334720.0  0.153  334720.0 ; CBB CBD
8  9  1  0.153  334720.0  0.153  334720.0 ; CBD CBF
9 10  1  0.153  334720.0  0.153  334720.0 ; CBF CBH
10 11  1  0.153  334720.0  0.153  334720.0 ; CBH CBJ
11 12  1  0.153  334720.0  0.153  334720.0 ; CBJ CBL
12 13  1  0.153  334720.0  0.153  334720.0 ; CBL CBN
13 14  1  0.153  334720.0  0.153  334720.0 ; CBN CBP
14 15  1  0.153  334720.0  0.153  334720.0 ; CBP CBR
15 16  1  0.153  334720.0  0.153  334720.0 ; CBR CBT
16 17  1  0.153  334720.0  0.153  334720.0 ; CBT CBV
17 18  1  0.153  334720.0  0.153  334720.0 ; CBV CBX
18 19  1  0.147  376560.0  0.147  376560.0 ; CBX NCF
19 20  1  0.141  418400.0  0.133  418400.0 ; NCF CCB
19 29  1  0.133  418400.0  0.133  418400.0 ; NCF CBZ
20 21  1  0.139  418400.0  0.139  418400.0 ; CCB CAN
20 25  1  0.139  418400.0  0.139  418400.0 ; CCB CCD
21 22  1  0.139  418400.0  0.139  418400.0 ; CAN CAJ
22 23  1  0.139  418400.0  0.139  418400.0 ; CAJ CAL
23 24  1  0.139  418400.0  0.139  418400.0 ; CAL CAP
24 25  1  0.139  418400.0  0.139  418400.0 ; CAP CCD
25 26  1  0.153  334720.0  0.153  334720.0 ; CCD CCH
26 27  1  0.153  334720.0  0.153  334720.0 ; CCH CAC
26 28  1  0.153  334720.0  0.153  334720.0 ; CCH CAD
26 29  1  0.153  334720.0  0.153  334720.0 ; CCH CBZ
29 30  1  0.139  418400.0  0.139  418400.0 ; CBZ CAH
30 31  1  0.139  418400.0  0.139  418400.0 ; CAH CAG
31 32  1  0.139  418400.0  0.139  418400.0 ; CAG CAI
32 33  1  0.139  418400.0  0.139  418400.0 ; CAI CCA
33 34  1  0.153  334720.0  0.153  334720.0 ; CCA CCI
33 43  1  0.133  418400.0  0.133  418400.0 ; CCA NCG
34 35  1  0.153  334720.0  0.153  334720.0 ; CCI CAE
34 36  1  0.153  334720.0  0.153  334720.0 ; CCI CAF
34 37  1  0.153  334720.0  0.153  334720.0 ; CCI CCE
37 38  1  0.139  418400.0  0.139  418400.0 ; CCE CAQ
37 42  1  0.139  418400.0  0.139  418400.0 ; CCE CCC
38 39  1  0.139  418400.0  0.139  418400.0 ; CAQ CAM
39 40  1  0.139  418400.0  0.139  418400.0 ; CAM CAK
40 41  1  0.139  418400.0  0.139  418400.0 ; CAK CAO
41 42  1  0.139  418400.0  0.139  418400.0 ; CAO CCC
42 43  1  0.141  418400.0  0.133  418400.0 ; CCC NCG
43 44  1  0.147  376560.0  0.147  376560.0 ; NCG CBY
44 45  1  0.153  334720.0  0.153  334720.0 ; CBY CBW
45 46  1  0.153  334720.0  0.153  334720.0 ; CBW CBU
46 47  1  0.153  334720.0  0.153  334720.0 ; CBU CBS
47 48  1  0.153  334720.0  0.153  334720.0 ; CBS CBQ
48 49  1  0.153  334720.0  0.153  334720.0 ; CBQ CBO
49 50  1  0.153  334720.0  0.153  334720.0 ; CBO CBM
50 51  1  0.153  334720.0  0.153  334720.0 ; CBM CBK
51 52  1  0.153  334720.0  0.153  334720.0 ; CBK CBI
52 53  1  0.153  334720.0  0.153  334720.0 ; CBI CBG
53 54  1  0.153  334720.0  0.153  334720.0 ; CBG CBE
54 55  1  0.153  334720.0  0.153  334720.0 ; CBE CBC
55 56  1  0.153  334720.0  0.153  334720.0 ; CBC CBA
56 57  1  0.153  334720.0  0.153  334720.0 ; CBA CAY
57 58  1  0.153  334720.0  0.153  334720.0 ; CAY CAW
58 59  1  0.153  334720.0  0.153  334720.0 ; CAW CAU
59 60  1  0.153  334720.0  0.153  334720.0 ; CAU CAS
```

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60 61 1 0.153 334720.0 0.153 334720.0 ; CAS CAB

[ pairs ]
; ai aj fu c0, c1, ...
19 22 1 ; NCF CAJ
19 24 1 ; NCF CAP
19 27 1 ; NCF CAC
19 28 1 ; NCF CAD
19 31 1 ; NCF CAG
20 23 1 ; CCB CAL
20 27 1 ; CCB CAC
20 28 1 ; CCB CAD
20 30 1 ; CCB CAH
21 24 1 ; CAN CAP
21 26 1 ; CAN CCH
21 29 1 ; CAN CBZ
22 25 1 ; CAJ CCD
23 26 1 ; CAL CCH
24 27 1 ; CAP CAC
24 28 1 ; CAP CAD
24 29 1 ; CAP CBZ
25 30 1 ; CCD CAH
26 31 1 ; CCH CAG
27 30 1 ; CAC CAH
28 30 1 ; CAD CAH
29 32 1 ; CBZ CAI
30 33 1 ; CAH CCA
31 34 1 ; CAG CCI
31 43 1 ; CAG NCG
32 35 1 ; CAI CAE
32 36 1 ; CAI CAF
32 37 1 ; CAI CCE
32 42 1 ; CAI CCC
33 38 1 ; CCA CAQ
33 41 1 ; CCA CAO
34 39 1 ; CCI CAM
34 41 1 ; CCI CAO
35 38 1 ; CAE CAQ
35 42 1 ; CAE CCC
35 43 1 ; CAE NCG
36 38 1 ; CAF CAQ
36 42 1 ; CAF CCC
36 43 1 ; CAF NCG
37 40 1 ; CCE CAK
38 41 1 ; CAQ CAO
38 43 1 ; CAQ NCG
39 42 1 ; CAM CCC
40 43 1 ; CAK NCG

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[ angles ]
; ai aj ak fu c0, c1, ...
1 2 3 1 111.0 460.2 111.0 460.2 ; CAA CAR CAT
2 3 4 1 111.0 460.2 111.0 460.2 ; CAR CAT CAV
3 4 5 1 111.0 460.2 111.0 460.2 ; CAT CAV CAX
4 5 6 1 111.0 460.2 111.0 460.2 ; CAV CAX CAZ
5 6 7 1 111.0 460.2 111.0 460.2 ; CAX CAZ CBB
6 7 8 1 111.0 460.2 111.0 460.2 ; CAZ CBB CBD
7 8 9 1 111.0 460.2 111.0 460.2 ; CBB CBD CBF
8 9 10 1 111.0 460.2 111.0 460.2 ; CBD CBF CBH
9 10 11 1 111.0 460.2 111.0 460.2 ; CBF CBH CBJ
10 11 12 1 111.0 460.2 111.0 460.2 ; CBH CBJ CBL
11 12 13 1 111.0 460.2 111.0 460.2 ; CBJ CBL CBN
12 13 14 1 111.0 460.2 111.0 460.2 ; CBL CBN CBP
13 14 15 1 111.0 460.2 111.0 460.2 ; CBN CBP CBR
14 15 16 1 111.0 460.2 111.0 460.2 ; CBP CBR CBT
15 16 17 1 111.0 460.2 111.0 460.2 ; CBR CBT CBV
16 17 18 1 111.0 460.2 111.0 460.2 ; CBT CBV CBX
17 18 19 1 109.5 460.2 109.5 460.2 ; CBV CBX NCF
18 19 20 1 120.0 418.4 120.0 418.4 ; CBX NCF CCB

```

18	19	29	1	120.0	418.4	120.0	418.4 ;	CBX	NCF	CBZ
20	19	29	1	108.0	418.4	108.0	418.4 ;	CCB	NCF	CBZ
19	20	21	1	120.0	418.4	120.0	418.4 ;	NCF	CCB	CAN
19	20	25	1	108.0	418.4	108.0	418.4 ;	NCF	CCB	CCD
21	20	25	1	120.0	418.4	120.0	418.4 ;	CAN	CCB	CCD
20	21	22	1	120.0	418.4	120.0	418.4 ;	CCB	CAN	CAJ
21	22	23	1	120.0	418.4	120.0	418.4 ;	CAN	CAJ	CAL
22	23	24	1	120.0	418.4	120.0	418.4 ;	CAJ	CAL	CAP
23	24	25	1	120.0	418.4	120.0	418.4 ;	CAL	CAP	CCD
20	25	24	1	120.0	418.4	120.0	418.4 ;	CCB	CCD	CAP
20	25	26	1	108.0	418.4	108.0	418.4 ;	CCB	CCD	CCH
24	25	26	1	132.0	418.4	132.0	418.4 ;	CAP	CCD	CCH
25	26	27	1	109.5	397.5	109.5	397.5 ;	CCD	CCH	CAC
25	26	28	1	109.5	397.5	109.5	397.5 ;	CCD	CCH	CAD
25	26	29	1	104.0	460.2	104.0	460.2 ;	CCD	CCH	CBZ
27	26	28	1	111.0	460.2	111.0	460.2 ;	CAC	CCH	CAD
27	26	29	1	109.5	397.5	109.5	397.5 ;	CAC	CCH	CBZ
28	26	29	1	109.5	397.5	109.5	397.5 ;	CAD	CCH	CBZ
19	29	26	1	108.0	418.4	108.0	418.4 ;	NCF	CBZ	CCH
19	29	30	1	132.0	418.4	132.0	418.4 ;	NCF	CBZ	CAH
26	29	30	1	132.0	418.4	132.0	418.4 ;	CCH	CBZ	CAH
29	30	31	1	120.0	418.4	120.0	418.4 ;	CBZ	CAH	CAG
30	31	32	1	135.0	418.4	135.0	418.4 ;	CAH	CAG	CAI
31	32	33	1	125.0	418.4	125.0	418.4 ;	CAG	CAI	CCA
32	33	34	1	120.0	418.4	120.0	418.4 ;	CAI	CCA	CCI
32	33	43	1	132.0	418.4	132.0	418.4 ;	CAI	CCA	NCG
34	33	43	1	108.0	418.4	108.0	418.4 ;	CCI	CCA	NCG
33	34	35	1	109.5	397.5	109.5	397.5 ;	CCA	CCI	CAE
33	34	36	1	109.5	397.5	109.5	397.5 ;	CCA	CCI	CAF
33	34	37	1	104.0	460.2	104.0	460.2 ;	CCA	CCI	CCE
35	34	36	1	111.0	460.2	111.0	460.2 ;	CAE	CCI	CAF
35	34	37	1	109.5	397.5	109.5	397.5 ;	CAE	CCI	CCE
36	34	37	1	109.5	397.5	109.5	397.5 ;	CAF	CCI	CCE
34	37	38	1	132.0	418.4	132.0	418.4 ;	CCI	CCE	CAQ
34	37	42	1	108.0	418.4	108.0	418.4 ;	CCI	CCE	CCC
38	37	42	1	120.0	418.4	120.0	418.4 ;	CAQ	CCE	CCC
37	38	39	1	120.0	418.4	120.0	418.4 ;	CCE	CAQ	CAM
38	39	40	1	120.0	418.4	120.0	418.4 ;	CAQ	CAM	CAK
39	40	41	1	120.0	418.4	120.0	418.4 ;	CAM	CAK	CAO
40	41	42	1	120.0	418.4	120.0	418.4 ;	CAK	CAO	CCC
37	42	41	1	120.0	418.4	120.0	418.4 ;	CCE	CCC	CAO
37	42	43	1	108.0	418.4	108.0	418.4 ;	CCE	CCC	NCG
41	42	43	1	132.0	418.4	132.0	418.4 ;	CAO	CCC	NCG
33	43	42	1	108.0	418.4	108.0	418.4 ;	CCA	NCG	CCC
33	43	44	1	120.0	418.4	120.0	418.4 ;	CCA	NCG	CBY
42	43	44	1	120.0	418.4	120.0	418.4 ;	CCC	NCG	CBY
43	44	45	1	109.5	460.2	109.5	460.2 ;	NCG	CBY	CBW
44	45	46	1	111.0	460.2	111.0	460.2 ;	CBY	CBW	CBU
45	46	47	1	111.0	460.2	111.0	460.2 ;	CBW	CBU	CBS
46	47	48	1	111.0	460.2	111.0	460.2 ;	CBU	CBS	CBQ
47	48	49	1	111.0	460.2	111.0	460.2 ;	CBS	CBQ	CBO
48	49	50	1	111.0	460.2	111.0	460.2 ;	CBQ	CBO	CBM
49	50	51	1	111.0	460.2	111.0	460.2 ;	CBO	CBM	CBK
50	51	52	1	111.0	460.2	111.0	460.2 ;	CBM	CBK	CBI
51	52	53	1	111.0	460.2	111.0	460.2 ;	CBK	CBI	CBG
52	53	54	1	111.0	460.2	111.0	460.2 ;	CBI	CBG	CBE
53	54	55	1	111.0	460.2	111.0	460.2 ;	CBG	CBE	CBC
54	55	56	1	111.0	460.2	111.0	460.2 ;	CBE	CBC	CBA
55	56	57	1	111.0	460.2	111.0	460.2 ;	CBC	CBA	CAY
56	57	58	1	111.0	460.2	111.0	460.2 ;	CBA	CAY	CAW
57	58	59	1	111.0	460.2	111.0	460.2 ;	CAY	CAW	CAU
58	59	60	1	111.0	460.2	111.0	460.2 ;	CAW	CAU	CAS
59	60	61	1	111.0	460.2	111.0	460.2 ;	CAU	CAS	CAB

[dihedrals]

; ai	aj	ak	al	fu	c0,	c1,	m,	...				
19	18	20	29	2	0.0	1673.6			0.0	1673.6	; imp	NCF CBX CCB CBZ
20	25	21	19	2	0.0	1673.6			0.0	1673.6	; imp	CCB CCD CAN NCF
25	20	26	24	2	0.0	1673.6			0.0	1673.6	; imp	CCD CCB CCH CAP
29	30	26	19	2	0.0	1673.6			0.0	1673.6	; imp	CBZ CAH CCH NCF
33	32	34	43	2	0.0	1673.6			0.0	1673.6	; imp	CCA CAI CCI NCG

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37 34 42 38 2      0.0 1673.6      0.0 1673.6 ; imp CCE CCI CCC CAQ
42 37 41 43 2      0.0 1673.6      0.0 1673.6 ; imp CCC CCE CAO NCG
43 33 42 44 2      0.0 1673.6      0.0 1673.6 ; imp NCG CCA CCC CBY
26 25 28 27 2     35.3 836.8     35.3 836.8 ; imp CCH CCD CAD CAC
34 33 35 36 2     35.3 836.8     35.3 836.8 ; imp CCI CCA CAE CAF
37 38 39 40 2      0.0 1673.6      0.0 1673.6 ; imp CCE CAQ CAM CAK
38 39 40 41 2      0.0 1673.6      0.0 1673.6 ; imp CAQ CAM CAK CAO
39 40 41 42 2      0.0 1673.6      0.0 1673.6 ; imp CAM CAK CAO CCC
40 41 42 37 2      0.0 1673.6      0.0 1673.6 ; imp CAK CAO CCC CCE
41 42 37 38 2      0.0 1673.6      0.0 1673.6 ; imp CAO CCC CCE CAQ
42 37 38 39 2      0.0 1673.6      0.0 1673.6 ; imp CCC CCE CAQ CAM
20 21 22 23 2      0.0 1673.6      0.0 1673.6 ; imp CCB CAN CAJ CAL
21 22 23 24 2      0.0 1673.6      0.0 1673.6 ; imp CAN CAJ CAL CAP
22 23 24 25 2      0.0 1673.6      0.0 1673.6 ; imp CAJ CAL CAP CCD
23 24 25 20 2      0.0 1673.6      0.0 1673.6 ; imp CAL CAP CCD CCB
24 25 20 21 2      0.0 1673.6      0.0 1673.6 ; imp CAP CCD CCB CAN
25 20 21 22 2      0.0 1673.6      0.0 1673.6 ; imp CCD CCB CAN CAJ
4 3 2 1 3
5 4 3 2 3
6 5 4 3 3
7 6 5 4 3
8 7 6 5 3
9 8 7 6 3
10 9 8 7 3
11 10 9 8 3
12 11 10 9 3
13 12 11 10 3
14 13 12 11 3
15 14 13 12 3
16 15 14 13 3
17 16 15 14 3
18 17 16 15 3
19 18 17 16 1      0.0 5.9 3      0.0 5.9 3 ; dih NCF CBX CBV CBT
17 18 19 29 1      0.0 3.8 6      0.0 3.8 6 ; dih CBV CBX NCF CBZ
18 19 20 25 1     180.0 33.5 2     180.0 33.5 2 ; dih CBX NCF CCB CCD
18 19 29 30 1     180.0 33.5 2     180.0 33.5 2 ; dih CBX NCF CBZ CAH
20 25 26 29 1      0.0 0.4 6      0.0 0.4 6 ; dih CCB CCD CCH CBZ
25 26 29 30 1      0.0 0.4 6      0.0 0.4 6 ; dih CCD CCH CBZ CAH
19 29 30 31 1     180.0 41.8 2     180.0 41.8 2 ; dih NCF CBZ CAH CAG
29 30 31 32 1     180.0 41.8 2     180.0 41.8 2 ; dih CBZ CAH CAG CAI
30 31 32 33 1     180.0 41.8 2     180.0 41.8 2 ; dih CAH CAG CAI CCA
43 33 32 31 1     180.0 41.8 2     180.0 41.8 2 ; dih NCG CCA CAI CAG
32 33 34 37 1      0.0 0.4 6      0.0 0.4 6 ; dih CAI CCA CCI CCE
32 33 43 44 1     180.0 33.5 2     180.0 33.5 2 ; dih CAI CCA NCG CBY
33 34 37 42 1      0.0 0.4 6      0.0 0.4 6 ; dih CCA CCI CCE CCC
37 42 43 44 1     180.0 33.5 2     180.0 33.5 2 ; dih CCE CCC NCG CBY
33 43 44 45 1      0.0 3.8 6      0.0 3.8 6 ; dih CCA NCG CBY CBW
46 45 44 43 1      0.0 5.9 3      0.0 5.9 3 ; dih CBU CBW CBY NCG
47 46 45 44 3
48 47 46 45 3
49 48 47 46 3
50 49 48 47 3
51 50 49 48 3
52 51 50 49 3
53 52 51 50 3
54 53 52 51 3
55 54 53 52 3
56 55 54 53 3
57 56 55 54 3
58 57 56 55 3
59 58 57 56 3
60 59 58 57 3
61 60 59 58 3

; Strong position restraints for InflateGRO
#ifdef STRONG_POSRES
#include "strong_posre.itp"
#endif

```

S1.2 DiI-C18:2

```
[ moleculetype ]
; Name nrexcl
DII      3

[ atoms ]
;  nr      type  resnr  resid  atom  cgnr  charge  mass
  1      LP3     1    DII     C1     1    0.000  15.0350
  2      LP2     1    DII     C2     2    0.000  14.0270
  3      LP2     1    DII     C3     3    0.000  14.0270
  4      LP2     1    DII     C4     4    0.000  14.0270
  5      LP2     1    DII     C5     5    0.000  14.0270
  6      LH1     1    DII     C6     6    0.000  13.0190
  7      LH1     1    DII     C7     7    0.000  13.0190
  8      LP2     1    DII     C8     8    0.000  14.0270
  9      LH1     1    DII     C9     9    0.000  13.0190
 10      LH1     1    DII    C10    10    0.000  13.0190
 11      LP2     1    DII    C11    11    0.000  14.0270
 12      LP2     1    DII    C12    12    0.000  14.0270
 13      LP2     1    DII    C13    13    0.000  14.0270
 14      LP2     1    DII    C14    14    0.000  14.0270
 15      LP2     1    DII    C15    15    0.000  14.0270
 16      LP2     1    DII    C16    16    0.000  14.0270
 17      LP2     1    DII    C17    17    0.000  14.0270
 18      LP2     1    DII    C18    18    0.150  14.0270
 19      NR      1    DII   N19    19   -0.010  14.0067
 20      C       1    DII   C20    20    0.100  12.0110
 21      CR1     1    DII   C21    21   -0.040  13.0190
 22      CR1     1    DII   C22    22    0.040  13.0190
 23      CR1     1    DII   C23    23    0.010  13.0190
 24      CR1     1    DII   C24    24    0.010  13.0190
 25      C       1    DII   C25    25   -0.100  12.0110
 26      CH1     1    DII   C26    26    0.400  12.0110
 27      CH3     1    DII   C27    27   -0.060  15.0350
 28      CH3     1    DII   C28    28   -0.070  15.0350
 29      C       1    DII   C29    29    0.300  12.0110
 30      CR1     1    DII   C30    30   -0.430  13.0190
 31      CR1     1    DII   C31    31    0.370  13.0190
 32      CR1     1    DII   C32    32   -0.340  13.0190
 33      C       1    DII   C33    33    0.410  12.0110
 34      CH1     1    DII   C34    34    0.140  12.0110
 35      CH3     1    DII   C35    35   -0.030  15.0350
 36      CH3     1    DII   C36    36   -0.040  15.0350
 37      C       1    DII   C37    37    0.030  12.0110
 38      CR1     1    DII   C38    38   -0.020  13.0190
 39      CR1     1    DII   C39    39    0.010  13.0190
 40      CR1     1    DII   C40    40    0.040  13.0190
 41      CR1     1    DII   C41    41   -0.060  13.0190
 42      C       1    DII   C42    42    0.140  12.0110
 43      N       1    DII  N43    43   -0.100  14.0067
 44      LP2     1    DII   C44    44    0.150  14.0270
 45      LP2     1    DII   C45    45    0.000  14.0270
 46      LP2     1    DII   C46    46    0.000  14.0270
 47      LP2     1    DII   C47    47    0.000  14.0270
 48      LP2     1    DII   C48    48    0.000  14.0270
 49      LP2     1    DII   C49    49    0.000  14.0270
 50      LP2     1    DII   C50    50    0.000  14.0270
 51      LP2     1    DII   C51    51    0.000  14.0270
 52      LH1     1    DII   C52    52    0.000  13.0190
 53      LH1     1    DII   C53    53    0.000  13.0190
 54      LP2     1    DII   C54    54    0.000  14.0270
 55      LH1     1    DII   C55    55    0.000  13.0190
 56      LH1     1    DII   C56    56    0.000  13.0190
 57      LP2     1    DII   C57    57    0.000  14.0270
 58      LP2     1    DII   C58    58    0.000  14.0270
 59      LP2     1    DII   C59    59    0.000  14.0270
 60      LP2     1    DII   C60    60    0.000  14.0270
 61      LP3     1    DII   C61    61    0.000  15.0350
```

```

[ bonds ]
; ai aj fu c0, c1, ...
  1  2  1  0.153 334720.0 0.153 334720.0 ; CAA CAR
  2  3  1  0.153 334720.0 0.153 334720.0 ; CAR CAT
  3  4  1  0.153 334720.0 0.153 334720.0 ; CAT CAV
  4  5  1  0.153 334720.0 0.153 334720.0 ; CAV CAX
  5  6  1  0.153 334720.0 0.153 334720.0 ; CAX CAZ

  6  7  1  0.139 418400.0 0.139 418400.0 ; CAZ CBB

  7  8  1  0.153 334720.0 0.153 334720.0 ; CBB CBD
  8  9  1  0.153 334720.0 0.153 334720.0 ; CBD CBF

  9 10  1  0.139 418400.0 0.139 418400.0 ; CBF CBH

10 11  1  0.153 334720.0 0.153 334720.0 ; CBH CBJ
11 12  1  0.153 334720.0 0.153 334720.0 ; CBJ CBL
12 13  1  0.153 334720.0 0.153 334720.0 ; CBL CBN
13 14  1  0.153 334720.0 0.153 334720.0 ; CBN CBP
14 15  1  0.153 334720.0 0.153 334720.0 ; CBP CBR
15 16  1  0.153 334720.0 0.153 334720.0 ; CBR CBT
16 17  1  0.153 334720.0 0.153 334720.0 ; CBT CBV
17 18  1  0.153 334720.0 0.153 334720.0 ; CBV CBX
18 19  1  0.147 376560.0 0.147 376560.0 ; CBX NCF
19 20  1  0.141 418400.0 0.133 418400.0 ; NCF CCB
19 29  1  0.133 418400.0 0.133 418400.0 ; NCF CBZ
20 21  1  0.139 418400.0 0.139 418400.0 ; CCB CAN
20 25  1  0.139 418400.0 0.139 418400.0 ; CCB CCD
21 22  1  0.139 418400.0 0.139 418400.0 ; CAN CAJ
22 23  1  0.139 418400.0 0.139 418400.0 ; CAJ CAL
23 24  1  0.139 418400.0 0.139 418400.0 ; CAL CAP
24 25  1  0.139 418400.0 0.139 418400.0 ; CAP CCD
25 26  1  0.153 334720.0 0.153 334720.0 ; CCD CCH
26 27  1  0.153 334720.0 0.153 334720.0 ; CCH CAC
26 28  1  0.153 334720.0 0.153 334720.0 ; CCH CAD
26 29  1  0.153 334720.0 0.153 334720.0 ; CCH CBZ
29 30  1  0.139 418400.0 0.139 418400.0 ; CBZ CAH
30 31  1  0.139 418400.0 0.139 418400.0 ; CAH CAG
31 32  1  0.139 418400.0 0.139 418400.0 ; CAG CAI
32 33  1  0.139 418400.0 0.139 418400.0 ; CAI CCA
33 34  1  0.153 334720.0 0.153 334720.0 ; CCA CCI
33 43  1  0.133 418400.0 0.133 418400.0 ; CCA NCG
34 35  1  0.153 334720.0 0.153 334720.0 ; CCI CAE
34 36  1  0.153 334720.0 0.153 334720.0 ; CCI CAF
34 37  1  0.153 334720.0 0.153 334720.0 ; CCI CCE
37 38  1  0.139 418400.0 0.139 418400.0 ; CCE CAQ
37 42  1  0.139 418400.0 0.139 418400.0 ; CCE CCC
38 39  1  0.139 418400.0 0.139 418400.0 ; CAQ CAM
39 40  1  0.139 418400.0 0.139 418400.0 ; CAM CAK
40 41  1  0.139 418400.0 0.139 418400.0 ; CAK CAO
41 42  1  0.139 418400.0 0.139 418400.0 ; CAO CCC
42 43  1  0.141 418400.0 0.133 418400.0 ; CCC NCG
43 44  1  0.147 376560.0 0.147 376560.0 ; NCG CBY
44 45  1  0.153 334720.0 0.153 334720.0 ; CBY CBW
45 46  1  0.153 334720.0 0.153 334720.0 ; CBW CBU
46 47  1  0.153 334720.0 0.153 334720.0 ; CBU CBS
47 48  1  0.153 334720.0 0.153 334720.0 ; CBS CBQ
48 49  1  0.153 334720.0 0.153 334720.0 ; CBQ CBO
49 50  1  0.153 334720.0 0.153 334720.0 ; CBO CBM
50 51  1  0.153 334720.0 0.153 334720.0 ; CBM CBK
51 52  1  0.153 334720.0 0.153 334720.0 ; CBK CBI

52 53  1  0.139 418400.0 0.139 418400.0 ; CBI CBG

53 54  1  0.153 334720.0 0.153 334720.0 ; CBG CBE
54 55  1  0.153 334720.0 0.153 334720.0 ; CBE CBC

55 56  1  0.139 418400.0 0.139 418400.0 ; CBC CBA

56 57  1  0.153 334720.0 0.153 334720.0 ; CBA CAY
57 58  1  0.153 334720.0 0.153 334720.0 ; CAY CAW

```



```

58 59 1 0.153 334720.0 0.153 334720.0 ; CAW CAU
59 60 1 0.153 334720.0 0.153 334720.0 ; CAU CAS
60 61 1 0.153 334720.0 0.153 334720.0 ; CAS CAB

[ pairs ]
; ai aj fu c0, c1, ...
19 22 1 ; NCF CAJ
19 24 1 ; NCF CAP
19 27 1 ; NCF CAC
19 28 1 ; NCF CAD
19 31 1 ; NCF CAG
20 23 1 ; CCB CAL
20 27 1 ; CCB CAC
20 28 1 ; CCB CAD
20 30 1 ; CCB CAH
21 24 1 ; CAN CAP
21 26 1 ; CAN CCH
21 29 1 ; CAN CBZ
22 25 1 ; CAJ CCD
23 26 1 ; CAL CCH
24 27 1 ; CAP CAC
24 28 1 ; CAP CAD
24 29 1 ; CAP CBZ
25 30 1 ; CCD CAH
26 31 1 ; CCH CAG
27 30 1 ; CAC CAH
28 30 1 ; CAD CAH
29 32 1 ; CBZ CAI
30 33 1 ; CAH CCA
31 34 1 ; CAG CCI
31 43 1 ; CAG NCG
32 35 1 ; CAI CAE
32 36 1 ; CAI CAF
32 37 1 ; CAI CCE
32 42 1 ; CAI CCC
33 38 1 ; CCA CAQ
33 41 1 ; CCA CAO
34 39 1 ; CCI CAM
34 41 1 ; CCI CAO
35 38 1 ; CAE CAQ
35 42 1 ; CAE CCC
35 43 1 ; CAE NCG
36 38 1 ; CAF CAQ
36 42 1 ; CAF CCC
36 43 1 ; CAF NCG
37 40 1 ; CCE CAK
38 41 1 ; CAQ CAO
38 43 1 ; CAQ NCG
39 42 1 ; CAM CCC
40 43 1 ; CAK NCG

[ angles ]
; ai aj ak fu c0, c1, ...
1 2 3 1 111.0 460.2 111.0 460.2 ; CAA CAR CAT
2 3 4 1 111.0 460.2 111.0 460.2 ; CAR CAT CAV
3 4 5 1 111.0 460.2 111.0 460.2 ; CAT CAV CAX
4 5 6 1 111.0 460.2 111.0 460.2 ; CAV CAX CAZ

5 6 7 1 120.0 502.08 120.0 502.08 ; CAX CAZ CBB
6 7 8 1 120.0 502.08 120.0 502.08 ; CAZ CBB CBD

7 8 9 1 111.0 460.2 111.0 460.2 ; CBB CBD CBF

8 9 10 1 120.0 502.08 120.0 502.08 ; CBD CBF CBH
9 10 11 1 120.0 502.08 120.0 502.08 ; CBF CBH CBJ

10 11 12 1 111.0 460.2 111.0 460.2 ; CBH CBJ CBL
11 12 13 1 111.0 460.2 111.0 460.2 ; CBJ CBL CBN
12 13 14 1 111.0 460.2 111.0 460.2 ; CBL CBN CBP

```

13	14	15	1	111.0	460.2	111.0	460.2 ;	CBN	CBP	CBR
14	15	16	1	111.0	460.2	111.0	460.2 ;	CBP	CBR	CBT
15	16	17	1	111.0	460.2	111.0	460.2 ;	CBR	CBT	CBV
16	17	18	1	111.0	460.2	111.0	460.2 ;	CBT	CBV	CBX
17	18	19	1	109.5	460.2	109.5	460.2 ;	CBV	CBX	NCF
18	19	20	1	120.0	418.4	120.0	418.4 ;	CBX	NCF	CCB
18	19	29	1	120.0	418.4	120.0	418.4 ;	CBX	NCF	CBZ
20	19	29	1	108.0	418.4	108.0	418.4 ;	CCB	NCF	CBZ
19	20	21	1	120.0	418.4	120.0	418.4 ;	NCF	CCB	CAN
19	20	25	1	108.0	418.4	108.0	418.4 ;	NCF	CCB	CCD
21	20	25	1	120.0	418.4	120.0	418.4 ;	CAN	CCB	CCD
20	21	22	1	120.0	418.4	120.0	418.4 ;	CCB	CAN	CAJ
21	22	23	1	120.0	418.4	120.0	418.4 ;	CAN	CAJ	CAL
22	23	24	1	120.0	418.4	120.0	418.4 ;	CAJ	CAL	CAP
23	24	25	1	120.0	418.4	120.0	418.4 ;	CAL	CAP	CCD
20	25	24	1	120.0	418.4	120.0	418.4 ;	CCB	CCD	CAP
20	25	26	1	108.0	418.4	108.0	418.4 ;	CCB	CCD	CCH
24	25	26	1	132.0	418.4	132.0	418.4 ;	CAP	CCD	CCH
25	26	27	1	109.5	397.5	109.5	397.5 ;	CCD	CCH	CAC
25	26	28	1	109.5	397.5	109.5	397.5 ;	CCD	CCH	CAD
25	26	29	1	104.0	460.2	104.0	460.2 ;	CCD	CCH	CBZ
27	26	28	1	111.0	460.2	111.0	460.2 ;	CAC	CCH	CAD
27	26	29	1	109.5	397.5	109.5	397.5 ;	CAC	CCH	CBZ
28	26	29	1	109.5	397.5	109.5	397.5 ;	CAD	CCH	CBZ
19	29	26	1	108.0	418.4	108.0	418.4 ;	NCF	CBZ	CCH
19	29	30	1	132.0	418.4	132.0	418.4 ;	NCF	CBZ	CAH
26	29	30	1	132.0	418.4	132.0	418.4 ;	CCH	CBZ	CAH
29	30	31	1	120.0	418.4	120.0	418.4 ;	CBZ	CAH	CAG
30	31	32	1	135.0	418.4	135.0	418.4 ;	CAH	CAG	CAI
31	32	33	1	125.0	418.4	125.0	418.4 ;	CAG	CAI	CCA
32	33	34	1	120.0	418.4	120.0	418.4 ;	CAI	CCA	CCI
32	33	43	1	132.0	418.4	132.0	418.4 ;	CAI	CCA	NCG
34	33	43	1	108.0	418.4	108.0	418.4 ;	CCI	CCA	NCG
33	34	35	1	109.5	397.5	109.5	397.5 ;	CCA	CCI	CAE
33	34	36	1	109.5	397.5	109.5	397.5 ;	CCA	CCI	CAF
33	34	37	1	104.0	460.2	104.0	460.2 ;	CCA	CCI	CCE
35	34	36	1	111.0	460.2	111.0	460.2 ;	CAE	CCI	CAF
35	34	37	1	109.5	397.5	109.5	397.5 ;	CAE	CCI	CCE
36	34	37	1	109.5	397.5	109.5	397.5 ;	CAF	CCI	CCE
34	37	38	1	132.0	418.4	132.0	418.4 ;	CCI	CCE	CAQ
34	37	42	1	108.0	418.4	108.0	418.4 ;	CCI	CCE	CCC
38	37	42	1	120.0	418.4	120.0	418.4 ;	CAQ	CCE	CCC
37	38	39	1	120.0	418.4	120.0	418.4 ;	CCE	CAQ	CAM
38	39	40	1	120.0	418.4	120.0	418.4 ;	CAQ	CAM	CAK
39	40	41	1	120.0	418.4	120.0	418.4 ;	CAM	CAK	CAO
40	41	42	1	120.0	418.4	120.0	418.4 ;	CAK	CAO	CCC
37	42	41	1	120.0	418.4	120.0	418.4 ;	CCE	CCC	CAO
37	42	43	1	108.0	418.4	108.0	418.4 ;	CCE	CCC	NCG
41	42	43	1	132.0	418.4	132.0	418.4 ;	CAO	CCC	NCG
33	43	42	1	108.0	418.4	108.0	418.4 ;	CCA	NCG	CCC
33	43	44	1	120.0	418.4	120.0	418.4 ;	CCA	NCG	CBY
42	43	44	1	120.0	418.4	120.0	418.4 ;	CCC	NCG	CBY
43	44	45	1	109.5	460.2	109.5	460.2 ;	NCG	CBY	CBW
44	45	46	1	111.0	460.2	111.0	460.2 ;	CBY	CBW	CBU
45	46	47	1	111.0	460.2	111.0	460.2 ;	CBW	CBU	CBS
46	47	48	1	111.0	460.2	111.0	460.2 ;	CBU	CBS	CBQ
47	48	49	1	111.0	460.2	111.0	460.2 ;	CBS	CBQ	CBO
48	49	50	1	111.0	460.2	111.0	460.2 ;	CBQ	CBO	CBM
49	50	51	1	111.0	460.2	111.0	460.2 ;	CBO	CBM	CBK
50	51	52	1	111.0	460.2	111.0	460.2 ;	CBM	CBK	CBI
51	52	53	1	111.0	460.2	111.0	460.2 ;	CBK	CBI	CBG
52	53	54	1	111.0	460.2	111.0	460.2 ;	CBI	CBG	CBE
53	54	55	1	111.0	460.2	111.0	460.2 ;	CBG	CBE	CBC
54	55	56	1	120.0	502.08	120.0	502.08 ;	CBE	CBC	CBA
55	56	57	1	120.0	502.08	120.0	502.08 ;	CBC	CBA	CAY
56	57	58	1	111.0	460.2	111.0	460.2 ;	CBA	CAY	CAW
57	58	59	1	111.0	460.2	111.0	460.2 ;	CAY	CAW	CAU
58	59	60	1	111.0	460.2	111.0	460.2 ;	CAW	CAU	CAS

```

59 60 61 1 111.0 460.2 111.0 460.2 ; CAU CAS CAB

[ dihedrals ]
; ai aj ak al fu c0, c1, m, ...
19 18 20 29 2 0.0 1673.6 0.0 1673.6 ; imp NCF CBX CCB CBZ
20 25 21 19 2 0.0 1673.6 0.0 1673.6 ; imp CCB CCD CAN NCF
25 20 26 24 2 0.0 1673.6 0.0 1673.6 ; imp CCD CCB CCH CAP
29 30 26 19 2 0.0 1673.6 0.0 1673.6 ; imp CBZ CAH CCH NCF
33 32 34 43 2 0.0 1673.6 0.0 1673.6 ; imp CCA CAI CCI NCG
37 34 42 38 2 0.0 1673.6 0.0 1673.6 ; imp CCE CCI CCC CAQ
42 37 41 43 2 0.0 1673.6 0.0 1673.6 ; imp CCC CCE CAO NCG
43 33 42 44 2 0.0 1673.6 0.0 1673.6 ; imp NCG CCA CCC CBY
26 25 28 27 2 35.3 836.8 35.3 836.8 ; imp CCH CCD CAD CAC
34 33 35 36 2 35.3 836.8 35.3 836.8 ; imp CCI CCA CAE CAF
37 38 39 40 2 0.0 1673.6 0.0 1673.6 ; imp CCE CAQ CAM CAK
38 39 40 41 2 0.0 1673.6 0.0 1673.6 ; imp CAQ CAM CAK CAO
39 40 41 42 2 0.0 1673.6 0.0 1673.6 ; imp CAM CAK CAO CCC
40 41 42 37 2 0.0 1673.6 0.0 1673.6 ; imp CAK CAO CCC CCE
41 42 37 38 2 0.0 1673.6 0.0 1673.6 ; imp CAO CCC CCE CAQ
42 37 38 39 2 0.0 1673.6 0.0 1673.6 ; imp CCC CCE CAQ CAM
20 21 22 23 2 0.0 1673.6 0.0 1673.6 ; imp CCB CAN CAJ CAL
21 22 23 24 2 0.0 1673.6 0.0 1673.6 ; imp CAN CAJ CAL CAP
22 23 24 25 2 0.0 1673.6 0.0 1673.6 ; imp CAJ CAL CAP CCD
23 24 25 20 2 0.0 1673.6 0.0 1673.6 ; imp CAL CAP CCD CCB
24 25 20 21 2 0.0 1673.6 0.0 1673.6 ; imp CAP CCD CCB CAN
25 20 21 22 2 0.0 1673.6 0.0 1673.6 ; imp CCD CCB CAN CAJ
4 3 2 1 3
5 4 3 2 3
6 5 4 3 3
7 6 5 4 1 0.0 5.858 3 0.0 5.858 3
; 8 7 6 5 3
9 8 7 6 1 0.0 5.858 3 0.0 5.858 3
10 9 8 7 1 0.0 5.858 3 0.0 5.858 3
; 11 10 9 8 3
12 11 10 9 1 0.0 5.858 3 0.0 5.858 3
13 12 11 10 3
14 13 12 11 3
15 14 13 12 3
16 15 14 13 3
17 16 15 14 3
18 17 16 15 3
19 18 17 16 1 0.0 5.9 3 0.0 5.9 3 ; dih NCF CBX CBV CBT
17 18 19 29 1 0.0 3.8 6 0.0 3.8 6 ; dih CBV CBX NCF CBZ
18 19 20 25 1 180.0 33.5 2 180.0 33.5 2 ; dih CBX NCF CCB CCD
18 19 29 30 1 180.0 33.5 2 180.0 33.5 2 ; dih CBX NCF CBZ CAH
20 25 26 29 1 0.0 0.4 6 0.0 0.4 6 ; dih CCB CCD CCH CBZ
25 26 29 30 1 0.0 0.4 6 0.0 0.4 6 ; dih CCD CCH CBZ CAH
19 29 30 31 1 180.0 41.8 2 180.0 41.8 2 ; dih NCF CBZ CAH CAG
29 30 31 32 1 180.0 41.8 2 180.0 41.8 2 ; dih CBZ CAH CAG CAI
30 31 32 33 1 180.0 41.8 2 180.0 41.8 2 ; dih CAH CAG CAI CCA
43 33 32 31 1 180.0 41.8 2 180.0 41.8 2 ; dih NCG CCA CAI CAG
32 33 34 37 1 0.0 0.4 6 0.0 0.4 6 ; dih CAI CCA CCI CCE
32 33 43 44 1 180.0 33.5 2 180.0 33.5 2 ; dih CAI CCA NCG CBY
33 34 37 42 1 0.0 0.4 6 0.0 0.4 6 ; dih CCA CCI CCE CCC
37 42 43 44 1 180.0 33.5 2 180.0 33.5 2 ; dih CCE CCC NCG CBY
33 43 44 45 1 0.0 3.8 6 0.0 3.8 6 ; dih CCA NCG CBY CBW
46 45 44 43 1 0.0 5.9 3 0.0 5.9 3 ; dih CBU CBW CBY NCG
47 46 45 44 3
48 47 46 45 3
49 48 47 46 3
50 49 48 47 3
51 50 49 48 3
52 51 50 49 3
53 52 51 50 1 0.0 5.858 3 0.0 5.858 3
; 54 53 52 51 3
55 54 53 52 1 0.0 5.858 3 0.0 5.858 3
56 55 54 53 1 0.0 5.858 3 0.0 5.858 3
; 57 56 55 54 3
58 57 56 55 1 0.0 5.858 3 0.0 5.858 3
59 58 57 56 3
60 59 58 57 3

```

```

61 60 59 58 3

[ dihedrals ]
; ai    aj    ak    al funct
    11    10     9     8      2 0.000      167.360
    8     7     6     5      2 0.000      167.360
    54    53    52    51      2 0.000      167.360
    57    56    55    54      2 0.000      167.360

; Strong position restraints for InflateGRO
#ifdef STRONG_POSRES
#include "strong_posre.itp"
#endif

```

S1.3 DiI-C12:0

```
[ moleculetype ]
; Name nrexcl
DII      3

[ atoms ]
;  nr      type  resnr  resid  atom  cgndr  charge  mass
  1      LP3    1   DII    C1     1     0.000  15.0350
  2      LP2    1   DII    C2     2     0.000  14.0270
  3      LP2    1   DII    C3     3     0.000  14.0270
  4      LP2    1   DII    C4     4     0.000  14.0270
  5      LP2    1   DII    C5     5     0.000  14.0270
  6      LP2    1   DII    C6     6     0.000  14.0270
  7      LP2    1   DII    C7     7     0.000  14.0270
  8      LP2    1   DII    C8     8     0.000  14.0270
  9      LP2    1   DII    C9     9     0.000  14.0270
 10      LP2    1   DII   C10    10     0.000  14.0270
 11      LP2    1   DII   C11    11     0.000  14.0270
 12      LP2    1   DII   C12    12     0.150  14.0270
 13      NR     1   DII   N13    13    -0.010  14.0067
 14      C      1   DII   C14    14     0.100  12.0110
 15      CR1    1   DII   C15    15    -0.040  13.0190
 16      CR1    1   DII   C16    16     0.040  13.0190
 17      CR1    1   DII   C17    17     0.010  13.0190
 18      CR1    1   DII   C18    18     0.010  13.0190
 19      C      1   DII   C19    19    -0.100  12.0110
 20      CH1    1   DII   C20    20     0.400  12.0110
 21      CH3    1   DII   C21    21    -0.060  15.0350
 22      CH3    1   DII   C22    22    -0.070  15.0350
 23      C      1   DII   C23    23     0.300  12.0110
 24      CR1    1   DII   C24    24    -0.430  13.0190
 25      CR1    1   DII   C25    25     0.370  13.0190
 26      CR1    1   DII   C26    26    -0.340  13.0190
 27      C      1   DII   C27    27     0.410  12.0110
 28      CH1    1   DII   C28    28     0.140  12.0110
 29      CH3    1   DII   C29    29    -0.030  15.0350
 30      CH3    1   DII   C30    30    -0.040  15.0350
 31      C      1   DII   C31    31     0.030  12.0110
 32      CR1    1   DII   C32    32    -0.020  13.0190
 33      CR1    1   DII   C33    33     0.010  13.0190
 34      CR1    1   DII   C34    34     0.040  13.0190
 35      CR1    1   DII   C35    35    -0.060  13.0190
 36      C      1   DII   C36    36     0.140  12.0110
 37      N      1   DII   N37    37    -0.100  14.0067
 38      LP2    1   DII   C38    38     0.150  14.0270
 39      LP2    1   DII   C39    39     0.000  14.0270
 40      LP2    1   DII   C40    40     0.000  14.0270
 41      LP2    1   DII   C41    41     0.000  14.0270
 42      LP2    1   DII   C42    42     0.000  14.0270
 43      LP2    1   DII   C43    43     0.000  14.0270
 44      LP2    1   DII   C44    44     0.000  14.0270
 45      LP2    1   DII   C45    45     0.000  14.0270
 46      LP2    1   DII   C46    46     0.000  14.0270
 47      LP2    1   DII   C47    47     0.000  14.0270
 48      LP2    1   DII   C48    48     0.000  14.0270
 49      LP3    1   DII   C49    49     0.000  15.0350

[ bonds ]
; ai  aj  fu  c0, c1, ...
  1  2  1  0.153  334720.0  0.153  334720.0 ; CBB CBD
  2  3  1  0.153  334720.0  0.153  334720.0 ; CBD CBF
  3  4  1  0.153  334720.0  0.153  334720.0 ; CBF CBH
  4  5  1  0.153  334720.0  0.153  334720.0 ; CBH CBJ
  5  6  1  0.153  334720.0  0.153  334720.0 ; CBJ CBL
  6  7  1  0.153  334720.0  0.153  334720.0 ; CBL CBN
  7  8  1  0.153  334720.0  0.153  334720.0 ; CBN CBP
  8  9  1  0.153  334720.0  0.153  334720.0 ; CBP CBR
  9 10  1  0.153  334720.0  0.153  334720.0 ; CBR CBT
 10 11  1  0.153  334720.0  0.153  334720.0 ; CBT CBV
```

11	12	1	0.153	334720.0	0.153	334720.0 ;	CBV	CBX
12	13	1	0.147	376560.0	0.147	376560.0 ;	CBX	NCF
13	14	1	0.141	418400.0	0.133	418400.0 ;	NCF	CCB
13	23	1	0.133	418400.0	0.133	418400.0 ;	NCF	CBZ
14	15	1	0.139	418400.0	0.139	418400.0 ;	CCB	CAN
14	19	1	0.139	418400.0	0.139	418400.0 ;	CCB	CCD
15	16	1	0.139	418400.0	0.139	418400.0 ;	CAN	CAJ
16	17	1	0.139	418400.0	0.139	418400.0 ;	CAJ	CAL
17	18	1	0.139	418400.0	0.139	418400.0 ;	CAL	CAP
18	19	1	0.139	418400.0	0.139	418400.0 ;	CAP	CCD
19	20	1	0.153	334720.0	0.153	334720.0 ;	CCD	CCH
20	21	1	0.153	334720.0	0.153	334720.0 ;	CCH	CAC
20	22	1	0.153	334720.0	0.153	334720.0 ;	CCH	CAD
20	23	1	0.153	334720.0	0.153	334720.0 ;	CCH	CBZ
23	24	1	0.139	418400.0	0.139	418400.0 ;	CBZ	CAH
24	25	1	0.139	418400.0	0.139	418400.0 ;	CAH	CAG
25	26	1	0.139	418400.0	0.139	418400.0 ;	CAG	CAI
26	27	1	0.139	418400.0	0.139	418400.0 ;	CAI	CCA
27	28	1	0.153	334720.0	0.153	334720.0 ;	CCA	CCI
27	37	1	0.133	418400.0	0.133	418400.0 ;	CCA	NCG
28	29	1	0.153	334720.0	0.153	334720.0 ;	CCI	CAE
28	30	1	0.153	334720.0	0.153	334720.0 ;	CCI	CAF
28	31	1	0.153	334720.0	0.153	334720.0 ;	CCI	CCE
31	32	1	0.139	418400.0	0.139	418400.0 ;	CCE	CAQ
31	36	1	0.139	418400.0	0.139	418400.0 ;	CCE	CCC
32	33	1	0.139	418400.0	0.139	418400.0 ;	CAQ	CAM
33	34	1	0.139	418400.0	0.139	418400.0 ;	CAM	CAK
34	35	1	0.139	418400.0	0.139	418400.0 ;	CAK	CAO
35	36	1	0.139	418400.0	0.139	418400.0 ;	CAO	CCC
36	37	1	0.141	418400.0	0.133	418400.0 ;	CCC	NCG
37	38	1	0.147	376560.0	0.147	376560.0 ;	NCG	CBY
38	39	1	0.153	334720.0	0.153	334720.0 ;	CBY	CBW
39	40	1	0.153	334720.0	0.153	334720.0 ;	CBW	CBU
40	41	1	0.153	334720.0	0.153	334720.0 ;	CBU	CBS
41	42	1	0.153	334720.0	0.153	334720.0 ;	CBS	CBQ
42	43	1	0.153	334720.0	0.153	334720.0 ;	CBQ	CBO
43	44	1	0.153	334720.0	0.153	334720.0 ;	CBO	CBM
44	45	1	0.153	334720.0	0.153	334720.0 ;	CBM	CBK
45	46	1	0.153	334720.0	0.153	334720.0 ;	CBK	CBI
46	47	1	0.153	334720.0	0.153	334720.0 ;	CBI	CBG
47	48	1	0.153	334720.0	0.153	334720.0 ;	CBG	CBE
48	49	1	0.153	334720.0	0.153	334720.0 ;	CBE	CBC

```
[ pairs ]
; ai aj fu c0, c1, ...
13 16 1 ; NCF CAJ
13 18 1 ; NCF CAP
13 21 1 ; NCF CAC
13 22 1 ; NCF CAD
13 25 1 ; NCF CAG
14 17 1 ; CCB CAL
14 21 1 ; CCB CAC
14 22 1 ; CCB CAD
14 24 1 ; CCB CAH
15 18 1 ; CAN CAP
15 20 1 ; CAN CCH
15 23 1 ; CAN CBZ
16 19 1 ; CAJ CCD
17 20 1 ; CAL CCH
18 21 1 ; CAP CAC
18 22 1 ; CAP CAD
18 23 1 ; CAP CBZ
19 24 1 ; CCD CAH
20 25 1 ; CCH CAG
21 24 1 ; CAC CAH
22 24 1 ; CAD CAH
23 26 1 ; CBZ CAI
24 27 1 ; CAH CCA
25 28 1 ; CAG CCI
25 37 1 ; CAG NCG
26 29 1 ; CAI CAE
```

```

26 30 1
26 31 1
26 36 1
27 32 1
27 35 1
28 33 1
28 35 1
29 32 1
29 36 1
29 37 1
30 32 1
30 36 1
30 37 1
31 34 1
32 35 1
32 37 1
33 36 1
34 37 1

```

```

; CAI CAF
; CAI CCE
; CAI CCC
; CCA CAQ
; CCA CAO
; CCI CAM
; CCI CAO
; CAE CAQ
; CAE CCC
; CAE NCG
; CAF CAQ
; CAF CCC
; CAF NCG
; CCE CAK
; CAQ CAO
; CAQ NCG
; CAM CCC
; CAK NCG

```

```

[ angles ]
; ai aj ak fu c0, c1, ...
1 2 3 1 111.0 460.2 111.0 460.2 ; CBB CBD CBF
2 3 4 1 111.0 460.2 111.0 460.2 ; CBD CBF CBH
3 4 5 1 111.0 460.2 111.0 460.2 ; CBF CBH CBJ
4 5 6 1 111.0 460.2 111.0 460.2 ; CBH CBJ CBL
5 6 7 1 111.0 460.2 111.0 460.2 ; CBJ CBL CBN
6 7 8 1 111.0 460.2 111.0 460.2 ; CBL CBN CBP
7 8 9 1 111.0 460.2 111.0 460.2 ; CBN CBP CBR
8 9 10 1 111.0 460.2 111.0 460.2 ; CBP CBR CBT
9 10 11 1 111.0 460.2 111.0 460.2 ; CBR CBT CBV
10 11 12 1 111.0 460.2 111.0 460.2 ; CBT CBV CBX
11 12 13 1 109.5 460.2 109.5 460.2 ; CBV CBX NCF
12 13 14 1 120.0 418.4 120.0 418.4 ; CBX NCF CCB
12 13 23 1 120.0 418.4 120.0 418.4 ; CBX NCF CBZ
14 13 23 1 108.0 418.4 108.0 418.4 ; CCB NCF CBZ
13 14 15 1 120.0 418.4 120.0 418.4 ; NCF CCB CAN
13 14 19 1 108.0 418.4 108.0 418.4 ; NCF CCB CCD
15 14 19 1 120.0 418.4 120.0 418.4 ; CAN CCB CCD
14 15 16 1 120.0 418.4 120.0 418.4 ; CCB CAN CAJ
15 16 17 1 120.0 418.4 120.0 418.4 ; CAN CAJ CAL
16 17 18 1 120.0 418.4 120.0 418.4 ; CAJ CAL CAP
17 18 19 1 120.0 418.4 120.0 418.4 ; CAL CAP CCD
14 19 18 1 120.0 418.4 120.0 418.4 ; CCB CCD CAP
14 19 20 1 108.0 418.4 108.0 418.4 ; CCB CCD CCH
18 19 20 1 132.0 418.4 132.0 418.4 ; CAP CCD CCH
19 20 21 1 109.5 397.5 109.5 397.5 ; CCD CCH CAC
19 20 22 1 109.5 397.5 109.5 397.5 ; CCD CCH CAD
19 20 23 1 104.0 460.2 104.0 460.2 ; CCD CCH CBZ
21 20 22 1 111.0 460.2 111.0 460.2 ; CAC CCH CAD
21 20 23 1 109.5 397.5 109.5 397.5 ; CAC CCH CBZ
22 20 23 1 109.5 397.5 109.5 397.5 ; CAD CCH CBZ
13 23 20 1 108.0 418.4 108.0 418.4 ; NCF CBZ CCH
13 23 24 1 132.0 418.4 132.0 418.4 ; NCF CBZ CAH
20 23 24 1 132.0 418.4 132.0 418.4 ; CCH CBZ CAH
23 24 25 1 120.0 418.4 120.0 418.4 ; CBZ CAH CAG
24 25 26 1 135.0 418.4 135.0 418.4 ; CAH CAG CAI
25 26 27 1 125.0 418.4 125.0 418.4 ; CAG CAI CCA
26 27 28 1 120.0 418.4 120.0 418.4 ; CAI CCA CCI
26 27 37 1 132.0 418.4 132.0 418.4 ; CAI CCA NCG
28 27 37 1 108.0 418.4 108.0 418.4 ; CCI CCA NCG
27 28 29 1 109.5 397.5 109.5 397.5 ; CCA CCI CAE
27 28 30 1 109.5 397.5 109.5 397.5 ; CCA CCI CAF
27 28 31 1 104.0 460.2 104.0 460.2 ; CCA CCI CCE
29 28 30 1 111.0 460.2 111.0 460.2 ; CAE CCI CAF
29 28 31 1 109.5 397.5 109.5 397.5 ; CAE CCI CCE
30 28 31 1 109.5 397.5 109.5 397.5 ; CAF CCI CCE
28 31 32 1 132.0 418.4 132.0 418.4 ; CCI CCE CAQ
28 31 36 1 108.0 418.4 108.0 418.4 ; CCI CCE CCC
32 31 36 1 120.0 418.4 120.0 418.4 ; CAQ CCE CCC

```

31	32	33	1	120.0	418.4	120.0	418.4 ;	CCE	CAQ	CAM
32	33	34	1	120.0	418.4	120.0	418.4 ;	CAQ	CAM	CAK
33	34	35	1	120.0	418.4	120.0	418.4 ;	CAM	CAK	CAO
34	35	36	1	120.0	418.4	120.0	418.4 ;	CAK	CAO	CCC
31	36	35	1	120.0	418.4	120.0	418.4 ;	CCE	CCC	CAO
31	36	37	1	108.0	418.4	108.0	418.4 ;	CCE	CCC	NCG
35	36	37	1	132.0	418.4	132.0	418.4 ;	CAO	CCC	NCG
27	37	36	1	108.0	418.4	108.0	418.4 ;	CCA	NCG	CCC
27	37	38	1	120.0	418.4	120.0	418.4 ;	CCA	NCG	CBY
36	37	38	1	120.0	418.4	120.0	418.4 ;	CCC	NCG	CBY
37	38	39	1	109.5	460.2	109.5	460.2 ;	NCG	CBY	CBW
38	39	40	1	111.0	460.2	111.0	460.2 ;	CBY	CBW	CBU
39	40	41	1	111.0	460.2	111.0	460.2 ;	CBW	CBU	CBS
40	41	42	1	111.0	460.2	111.0	460.2 ;	CBU	CBS	CBQ
41	42	43	1	111.0	460.2	111.0	460.2 ;	CBS	CBQ	CBO
42	43	44	1	111.0	460.2	111.0	460.2 ;	CBQ	CBO	CBM
43	44	45	1	111.0	460.2	111.0	460.2 ;	CBO	CBM	CBK
44	45	46	1	111.0	460.2	111.0	460.2 ;	CBM	CBK	CBI
45	46	47	1	111.0	460.2	111.0	460.2 ;	CBK	CBI	CBG
46	47	48	1	111.0	460.2	111.0	460.2 ;	CBI	CBG	CBE
47	48	49	1	111.0	460.2	111.0	460.2 ;	CBG	CBE	CBC

[dihedrals]

; ai aj ak al fu c0, c1, m, ... FIXED

13	12	14	23	2	0.0	1673.6	0.0	1673.6 ;	imp	NCF	CBX	CCB	CBZ
14	19	15	13	2	0.0	1673.6	0.0	1673.6 ;	imp	CCB	CCD	CAN	NCF
19	14	20	18	2	0.0	1673.6	0.0	1673.6 ;	imp	CCD	CCB	CCH	CAP
23	24	20	13	2	0.0	1673.6	0.0	1673.6 ;	imp	CBZ	CAH	CCH	NCF
27	26	28	37	2	0.0	1673.6	0.0	1673.6 ;	imp	CCA	CAI	CCI	NCG
31	28	36	32	2	0.0	1673.6	0.0	1673.6 ;	imp	CCE	CCI	CCC	CAQ
36	31	35	37	2	0.0	1673.6	0.0	1673.6 ;	imp	CCC	CCE	CAO	NCG
37	27	36	38	2	0.0	1673.6	0.0	1673.6 ;	imp	NCG	CCA	CCC	CBY
20	19	22	21	2	35.3	836.8	35.3	836.8 ;	imp	CCH	CCD	CAD	CAC
28	27	29	30	2	35.3	836.8	35.3	836.8 ;	imp	CCI	CCA	CAE	CAF
31	32	33	34	2	0.0	1673.6	0.0	1673.6 ;	imp	CCE	CAQ	CAM	CAK
32	33	34	35	2	0.0	1673.6	0.0	1673.6 ;	imp	CAQ	CAM	CAK	CAO
33	34	35	36	2	0.0	1673.6	0.0	1673.6 ;	imp	CAM	CAK	CAO	CCC
34	35	36	31	2	0.0	1673.6	0.0	1673.6 ;	imp	CAK	CAO	CCC	CCE
35	36	31	32	2	0.0	1673.6	0.0	1673.6 ;	imp	CAO	CCC	CCE	CAQ
36	31	32	33	2	0.0	1673.6	0.0	1673.6 ;	imp	CCC	CCE	CAQ	CAM
14	15	16	17	2	0.0	1673.6	0.0	1673.6 ;	imp	CCB	CAN	CAJ	CAL
15	16	17	18	2	0.0	1673.6	0.0	1673.6 ;	imp	CAN	CAJ	CAL	CAP
16	17	18	19	2	0.0	1673.6	0.0	1673.6 ;	imp	CAJ	CAL	CAP	CCD
17	18	19	14	2	0.0	1673.6	0.0	1673.6 ;	imp	CAL	CAP	CCD	CCB
18	19	14	15	2	0.0	1673.6	0.0	1673.6 ;	imp	CAP	CCD	CCB	CAN
19	14	15	16	2	0.0	1673.6	0.0	1673.6 ;	imp	CCD	CCB	CAN	CAJ
4	3	2	1	3									
5	4	3	2	3									
6	5	4	3	3									
7	6	5	4	3									
8	7	6	5	3									
9	8	7	6	3									
10	9	8	7	3									
11	10	9	8	3									
12	11	10	9	3									
13	12	11	10	1	0.0	5.9 3	0.0	5.9 3 ;	dih	NCF	CBX	CBV	CBT
11	12	13	23	1	0.0	3.8 6	0.0	3.8 6 ;	dih	CBV	CBX	NCF	CBZ
12	13	14	19	1	180.0	33.5 2	180.0	33.5 2 ;	dih	CBX	NCF	CCB	CCD
12	13	23	24	1	180.0	33.5 2	180.0	33.5 2 ;	dih	CBX	NCF	CBZ	CAH
14	19	20	23	1	0.0	0.4 6	0.0	0.4 6 ;	dih	CCB	CCD	CCH	CBZ
19	20	23	24	1	0.0	0.4 6	0.0	0.4 6 ;	dih	CCD	CCH	CBZ	CAH
13	23	24	25	1	180.0	41.8 2	180.0	41.8 2 ;	dih	NCF	CBZ	CAH	CAG
23	24	25	26	1	180.0	41.8 2	180.0	41.8 2 ;	dih	CBZ	CAH	CAG	CAI
24	25	26	27	1	180.0	41.8 2	180.0	41.8 2 ;	dih	CAH	CAG	CAI	CCA
37	27	26	25	1	180.0	41.8 2	180.0	41.8 2 ;	dih	NCG	CCA	CAI	CAG
26	27	28	31	1	0.0	0.4 6	0.0	0.4 6 ;	dih	CAI	CCA	CCI	CCE
26	27	37	38	1	180.0	33.5 2	180.0	33.5 2 ;	dih	CAI	CCA	NCG	CBY
27	28	31	36	1	0.0	0.4 6	0.0	0.4 6 ;	dih	CCA	CCI	CCE	CCC
31	36	37	38	1	180.0	33.5 2	180.0	33.5 2 ;	dih	CCE	CCC	NCG	CBY
26	37	38	39	1	0.0	3.8 6	0.0	3.8 6 ;	dih	CCA	NCG	CBY	CBW
40	39	38	37	1	0.0	5.9 3	0.0	5.9 3 ;	dih	CBU	CBW	CBY	NCG


```
41 40 39 38 3
42 41 40 39 3
43 42 41 40 3
44 43 42 41 3
45 44 43 42 3
46 45 44 43 3
47 46 45 44 3
48 47 46 45 3
49 48 47 46 3

; Strong position restraints for InflateGRO
#ifdef STRONG_POSRES
#include "strong_posre.itp"
#endif
```

S2 Simulation Setup

Construction of the desired bilayers was a multi-step process. First, the PDB file of a template bilayer consisting of 128 DPPC molecules and water was obtained from the Tieleman website [<http://people.ucalgary.ca/~tieleman/download.html>]. Water was then removed and the system was tiled 2 times in both the x and y dimensions to create a bilayer with 512 DPPC molecules, with the z -axis being normal to the bilayer. One DPPC per leaflet was chosen so that their separation distance, including periodic copies, was maximized. These two DPPC molecules were then removed from the template bilayer and replaced with either two identical probe molecules (DiI-C18:0, DiI-C18:2, DiI-C12:0) or two DPPC (as a control). These inserted molecules were randomly rotated about their z -axis, and placed at the (x,y) position of the removed DPPC molecule. To minimize the possibility of atomic overlap, the inserted molecules were initially placed so that they were protruding from the bilayer, with only the ends of their hydrocarbon chains within the bilayer, and energy minimization was performed to reduce unfavorable interactions or overlaps. The inserted molecules were then moved 1 nm into the bilayer in 0.05 nm increments along the bilayer normal, with energy minimization performed at each step.

We tested a variety of simulation parameters, choosing particular values based on their ability to reproduce experimentally measured parameters. Importantly, we found that our results are robust even under different conditions including a lower hydration level or lower pressure (results not shown). We also saw that for the pure DPPC control simulations, insertion of the DPPC molecule did not cause bilayer perturbations.

Table S1. Number of lipids per shell after partitioning. The shell assignments described in Equation 3 partition the leaflets such that there are $\sim 6n$ lipids in shell n , for all systems studied. For shell $n > 7$ the shell diameters exceed the (x,y) box dimensions, leading to a decrease in the number of lipids per shell for the remaining shells.

inserted molecule	leaf	shell 0	shell 1	shell 2	shell 3	shell 4	shell 5	shell 6	shell 7	shell 8	shell 9	shell 10	shell 11
DPPC	same		6.01 $\pm$ 0.02	12.00 $\pm$ 0.03	18.04 $\pm$ 0.03	23.99 $\pm$ 0.03	30.00 $\pm$ 0.04	35.99 $\pm$ 0.04	42.00 $\pm$ 0.04	44.23 $\pm$ 0.05	24.86 $\pm$ 0.04	13.32 $\pm$ 0.03	4.56 $\pm$ 0.01
DPPC	opp.	1.000 $\pm$ 0.007	6.00 $\pm$ 0.02	12.00 $\pm$ 0.03	18.00 $\pm$ 0.04	23.99 $\pm$ 0.05	30.01 $\pm$ 0.03	35.99 $\pm$ 0.03	42.01 $\pm$ 0.05	44.23 $\pm$ 0.04	24.88 $\pm$ 0.04	13.32 $\pm$ 0.03	4.56 $\pm$ 0.01
DiI-C18:0	same		6.0 $\pm$ 0.1	12.1 $\pm$ 0.3	18.1 $\pm$ 0.4	24.0 $\pm$ 0.4	30.1 $\pm$ 0.6	36.0 $\pm$ 0.5	41.8 $\pm$ 0.5	44.3 $\pm$ 0.6	24.8 $\pm$ 0.5	13.3 $\pm$ 0.4	4.5 $\pm$ 0.1
DiI-C18:0	opp.	0.97 $\pm$ 0.09	5.8 $\pm$ 0.2	12.0 $\pm$ 0.3	18.2 $\pm$ 0.3	24.2 $\pm$ 0.3	30.0 $\pm$ 0.4	36.1 $\pm$ 0.5	41.9 $\pm$ 0.5	44.0 $\pm$ 0.6	24.6 $\pm$ 0.3	12.9 $\pm$ 0.3	4.4 $\pm$ 0.2
DiI-C18:2	same		5.9 $\pm$ 0.2	12.0 $\pm$ 0.3	18.1 $\pm$ 0.3	24.1 $\pm$ 0.4	30.1 $\pm$ 0.4	36.2 $\pm$ 0.4	41.7 $\pm$ 0.4	44.3 $\pm$ 0.4	24.7 $\pm$ 0.3	13.3 $\pm$ 0.2	4.6 $\pm$ 0.2
DiI-C18:2	opp.	0.97 $\pm$ 0.09	5.8 $\pm$ 0.1	12.0 $\pm$ 0.3	18.2 $\pm$ 0.3	24.0 $\pm$ 0.4	30.1 $\pm$ 0.6	36.1 $\pm$ 0.5	41.8 $\pm$ 0.4	44.1 $\pm$ 0.4	24.6 $\pm$ 0.3	12.9 $\pm$ 0.2	4.4 $\pm$ 0.2
DiI-C12:0	same		5.9 $\pm$ 0.2	12.2 $\pm$ 0.4	17.9 $\pm$ 0.4	24.0 $\pm$ 0.5	29.9 $\pm$ 0.4	36.4 $\pm$ 0.5	41.8 $\pm$ 0.5	44.3 $\pm$ 0.4	24.7 $\pm$ 0.5	13.2 $\pm$ 0.3	4.5 $\pm$ 0.1
DiI-C12:0	opp.	0.98 $\pm$ 0.09	5.9 $\pm$ 0.2	12.0 $\pm$ 0.2	17.9 $\pm$ 0.4	24.2 $\pm$ 0.4	30.1 $\pm$ 0.4	35.9 $\pm$ 0.5	42.0 $\pm$ 0.6	44.2 $\pm$ 0.6	24.5 $\pm$ 0.4	12.9 $\pm$ 0.3	4.4 $\pm$ 0.2

Table S2. DPPC chain order by shell. The average acyl chain order parameter S_{CD} (Eq. 4) for DPPC partitioned into shells, for each leaflet. Overall average for each system is shown in the last column.

inserted molecule	leaf	shell 0	shell 1	shell 2	shell 3	shell 4	shell 5	shell 6	shell 7	shell 8	shell 9	shell 10	shell 11	Avg.
DPPC	same		0.169 $\pm$ 0.001	0.166 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.166 $\pm$ 0.001	0.166 $\pm$ 0.001	0.165$\pm$0.001
DPPC	opp.	0.165 $\pm$ 0.002	0.166 $\pm$ 0.002	0.166 $\pm$ 0.002	0.166 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.165 $\pm$ 0.001	0.166 $\pm$ 0.001	0.165 $\pm$ 0.001	0.166 $\pm$ 0.001	
DiI-C18:0	same		0.15 $\pm$ 0.01	0.172 $\pm$ 0.009	0.171 $\pm$ 0.006	0.169 $\pm$ 0.004	0.167 $\pm$ 0.004	0.166 $\pm$ 0.003	0.165 $\pm$ 0.004	0.165 $\pm$ 0.004	0.167 $\pm$ 0.004	0.166 $\pm$ 0.006	0.16 $\pm$ 0.01	0.166$\pm$0.001
DiI-C18:0	opp.	0.15 $\pm$ 0.01	0.158 $\pm$ 0.008	0.166 $\pm$ 0.006	0.170 $\pm$ 0.005	0.170 $\pm$ 0.004	0.168 $\pm$ 0.004	0.166 $\pm$ 0.002	0.165 $\pm$ 0.002	0.165 $\pm$ 0.003	0.166 $\pm$ 0.004	0.165 $\pm$ 0.007	0.16 $\pm$ 0.01	
DiI-C18:2	same		0.148 $\pm$ 0.009	0.169 $\pm$ 0.005	0.170 $\pm$ 0.005	0.170 $\pm$ 0.005	0.169 $\pm$ 0.003	0.167 $\pm$ 0.004	0.165 $\pm$ 0.002	0.165 $\pm$ 0.003	0.165 $\pm$ 0.003	0.165 $\pm$ 0.005	0.16 $\pm$ 0.01	0.166$\pm$0.001
DiI-C18:2	opp.	0.15 $\pm$ 0.01	0.16 $\pm$ 0.01	0.164 $\pm$ 0.007	0.170 $\pm$ 0.004	0.170 $\pm$ 0.004	0.168 $\pm$ 0.003	0.167 $\pm$ 0.004	0.166 $\pm$ 0.003	0.165 $\pm$ 0.003	0.166 $\pm$ 0.005	0.166 $\pm$ 0.006	0.17 $\pm$ 0.01	
DiI-C12:0	same		0.152 $\pm$ 0.009	0.171 $\pm$ 0.006	0.169 $\pm$ 0.005	0.168 $\pm$ 0.004	0.167 $\pm$ 0.004	0.167 $\pm$ 0.003	0.166 $\pm$ 0.002	0.166 $\pm$ 0.003	0.166 $\pm$ 0.004	0.164 $\pm$ 0.006	0.16 $\pm$ 0.01	0.1664$\pm$0.0008
DiI-C12:0	opp.	0.16 $\pm$ 0.01	0.16 $\pm$ 0.01	0.164 $\pm$ 0.008	0.168 $\pm$ 0.004	0.168 $\pm$ 0.004	0.167 $\pm$ 0.004	0.167 $\pm$ 0.003	0.166 $\pm$ 0.003	0.166 $\pm$ 0.003	0.166 $\pm$ 0.004	0.164 $\pm$ 0.007	0.16 $\pm$ 0.01	

Table S3. P-N vector angle by shell with respect to the bilayer normal. The average P-N vector angle of all shells in each leaflet for each system is shown. Overall average for each system is shown in the last column.

inserted molecule	leaf	shell 0	shell 1	shell 2	shell 3	shell 4	shell 5	shell 6	shell 7	shell 8	shell 9	shell 10	shell 11	Avg
DPPC	same		79.6±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.6±0.2	79.5±0.2	79.5±0.2
DPPC	opp.	79.6±0.4	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	79.5±0.2	
DiI-C18:0	same		71±2	77±2	79±1	79±1	80±1	79.3±0.7	79.4±0.7	79.5±0.6	78.9±0.9	79±1	79±2	79.1±0.2
DiI-C18:0	opp.	79±4	79±2	79±1	79±1	79.4±0.9	79.4±0.7	79.3±0.8	79.2±0.6	79±1	78.4±0.8	77±1	77±3	
DiI-C18:2	same		71±3	78±2	79±1	79±1	79.2±0.9	79.5±0.7	79.4±0.8	79.5±0.8	79.3±0.9	79±1	79±2	79.0±0.2
DiI-C18:2	opp.	80±4	79±2	79±1	79±1	79.0±0.9	79.5±0.7	79.2±0.6	79.4±0.7	79.0±0.7	78.6±0.8	78±2	77±3	
DiI-C12:0	same		71±2	78±1	79±1	79±1	79.4±0.9	79.4±0.9	79.3±0.6	79.4±0.7	79.2±0.6	80±1	79±2	79.0±0.1
DiI-C12:0	opp.	80±5	79±1	79±2	79±1	79±1	79.5±0.7	79.4±0.6	79.3±0.7	79.0±0.8	79.5±0.9	78±1	77±4	

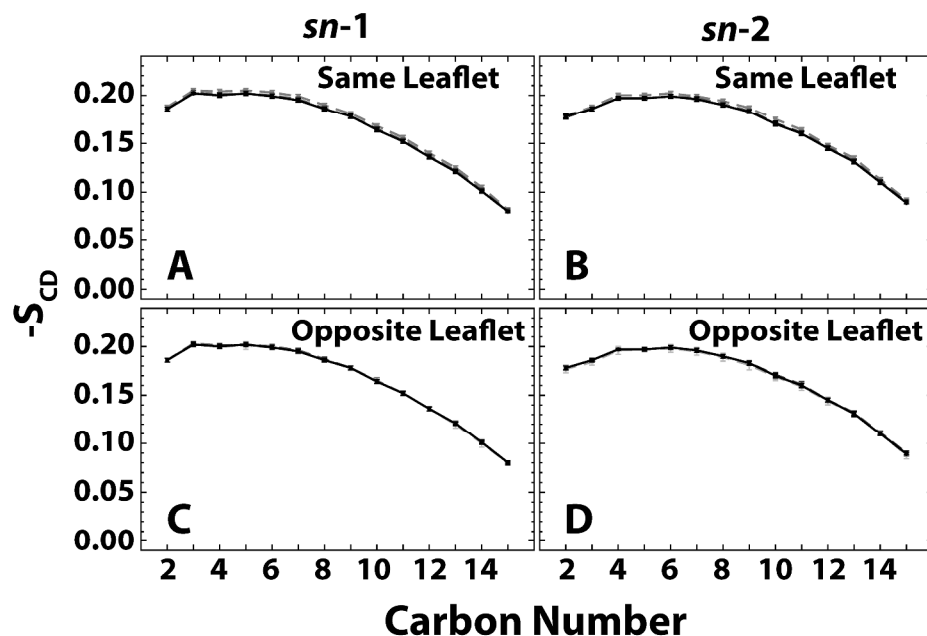


Figure S1. Order parameter by shell for a pure DPPC bilayer reveals a partitioning artifact. Order parameter of DPPC carbons shown for same leaflet lipids (*A,B*) and opposite leaflet lipids (*C,D*) for a pure DPPC bilayer. Shells $n \leq 3$ are shown (gray dashed) with darker grays indicating increasing shell number. Overall average order for the pure DPPC bilayers (black solid) is shown for comparison. For same leaflet lipids, first shell order is marginally increased due to the partitioning scheme, while opposite leaflet lipids are unaffected.

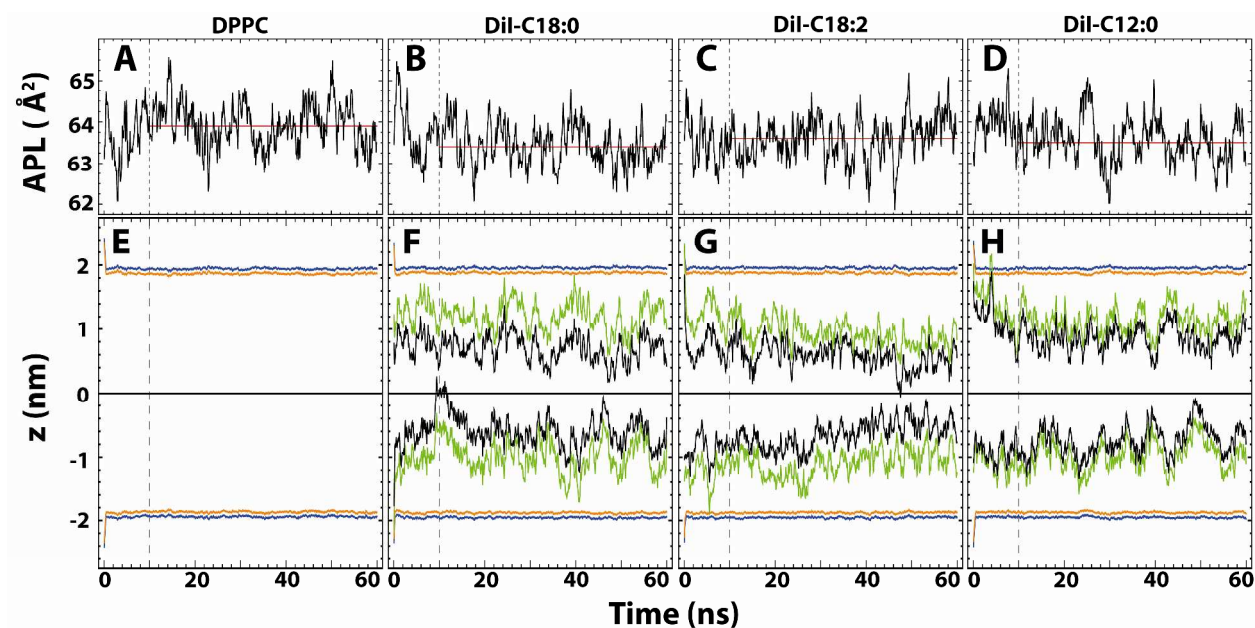


Figure S2. Representative data showing equilibration of pure DPPC and DiI-containing, simulations. *A-D*, APL during a representative 60 ns simulation of the systems studied. The average value over the final 50 ns is shown (red line). *E-H*, transverse bilayer position with respect to the bilayer midplane ($z = 0$ nm) of DPPC nitrogen (blue) and phosphate (yellow), and center-of-mass of the DiI chromophore (green) and entire DiI molecule (black).

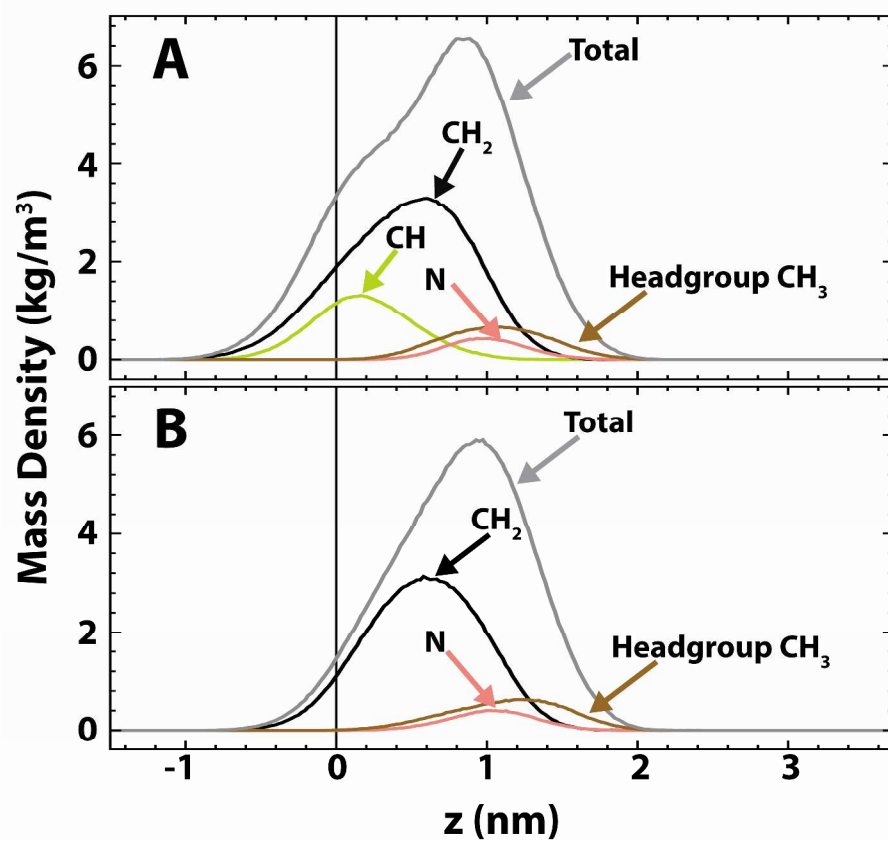


Figure S3. Average single-leaflet mass density profiles for simulations. *A*, average densities of DiI-C18:2 components. *B*, average densities for DiI-C12:0 components. Bilayer midplane at $z = 0$ nm.

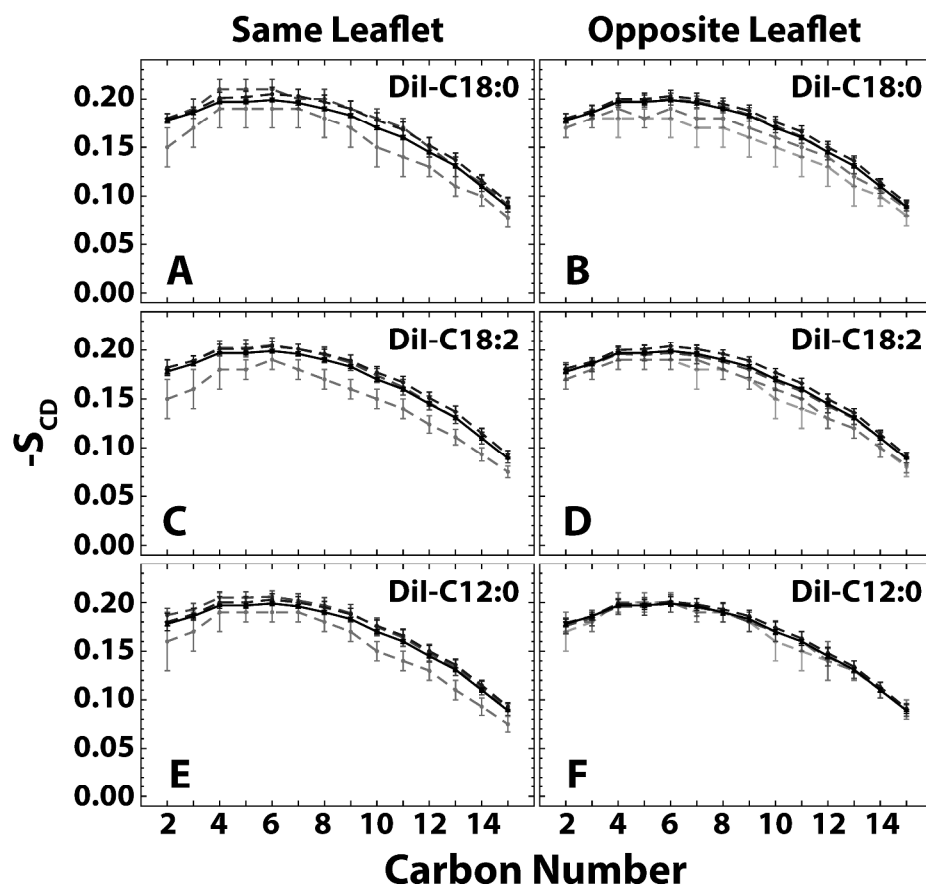


Figure S4. DPPC *sn*-2 carbon order perturbation depends on shell, not carbon number. Order parameter of DPPC *sn*-2 carbons shown for same leaflet lipids (*left*) and opposite leaflet lipids (*right*) for bilayers containing DiI-C18:0 (A,B), DiI-C18:2 (C,D) and DiI-C12:0 (E,F). Shells $n \leq 3$ are shown (gray dashed), with darker grays indicating increased shell number. Overall average *sn*-2 order for pure DPPC bilayers (black solid) is shown for comparison. Error bars indicate the standard deviation.

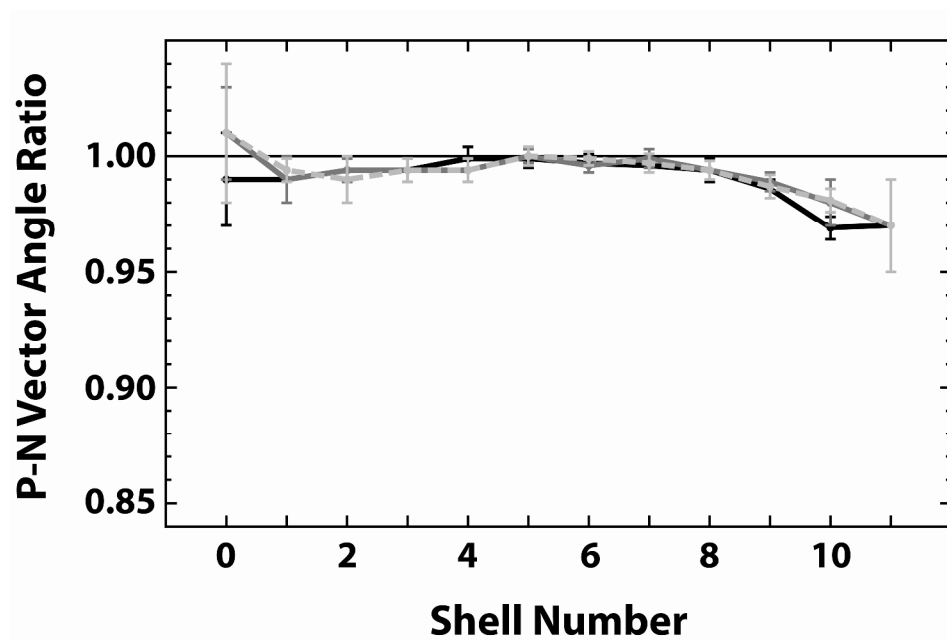


Figure S5. DPPC headgroups show no preferential orientation in the leaflet opposite the probe. P-N vector angle ratio by shell for opposite leaflet lipids in bilayers containing DiI-C18:0 (black), DiI-C18:2 (dark gray) and DiI-C12:0 (light gray dashed) compared to the pure DPPC simulations. Error bars indicate 95% confidence intervals.