

Supplementary information of

**Computational study on donor-acceptor optical
markers for Alzheimer's disease: a game of charge
transfer and electron delocalization.**

by

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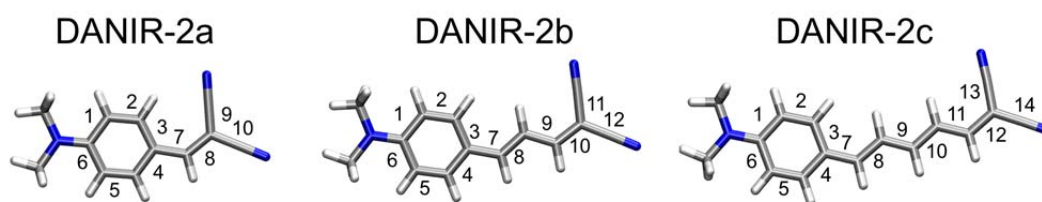
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Table S1. Bond lengths (Å) of DANIR-2a, -2b and -2c at the S₀ and S₁ optimized geometries.



Bond lengths (Å) S ₀	2a	2b	2c	Bond lengths (Å) S ₀	2a	2b	2c
	1	1.42	1.42		1.41	8	1.36
2	1.38	1.38	1.38	9	1.43	1.43	1.43
3	1.41	1.41	1.40	10	1.43	1.36	1.36
4	1.41	1.41	1.40	11	-	1.43	1.42
5	1.38	1.38	1.38	12	-	1.43	1.36
6	1.41	1.41	1.41	13	-	-	1.43
7	1.46	1.45	1.45	14	-	-	1.43
Bond lengths (Å) S ₁	2a	2b	2c	Bond lengths (Å) S ₁	2a	2b	2c
1	1.42	1.42	1.42	8	1.42	1.39	1.39
2	1.37	1.37	1.37	9	1.41	1.40	1.40
3	1.43	1.43	1.42	10	1.42	1.40	1.39
4	1.43	1.42	1.42	11	-	1.42	1.40
5	1.37	1.37	1.37	12	-	1.42	1.40
6	1.43	1.43	1.42	13	-	-	1.42
7	1.42	1.42	1.42	14	-	-	1.42

Table S2. PBE-D2 interaction energies and Grimme dispersion contribution (kJ/mol) of NIAD-4, DANIR-2a and DANIR-2c for selected marker-amyloid fibril complex models.

	I PELE	II PELE	I PBE-D	II PBE-D	I D	II D
	(OPLS)	(OPLS)				
NIAD-4	-337	-253	-339	-242	-255	-175
DANIR-2a	-179	-121	-195	-132	-195	-111
DANIR-2c	-251	-222	-265	-283	-232	-220

■ 20 ps / 80 snapshots x axis: wavelength (nm)
■ 10 ps / 40 snapshots y axis: cumulative oscillator strength

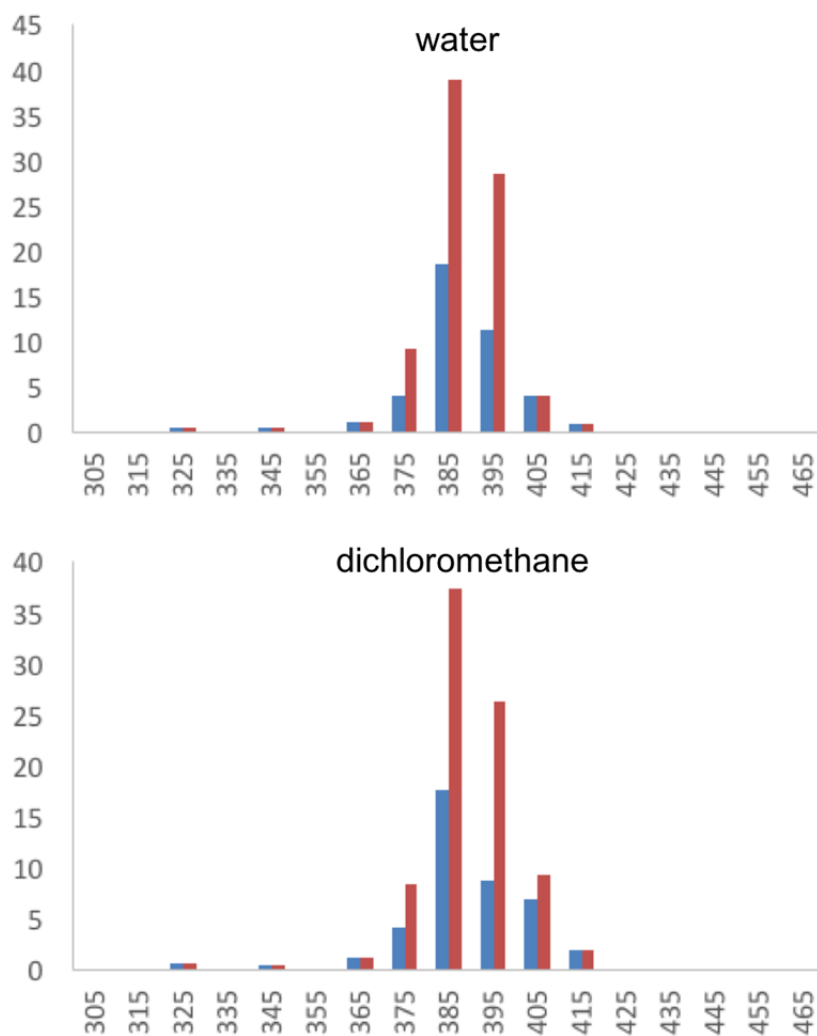
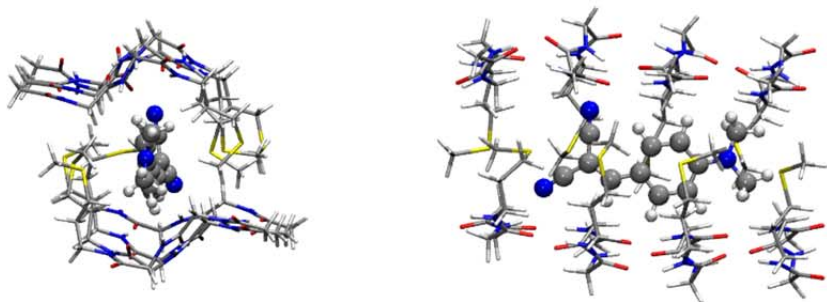
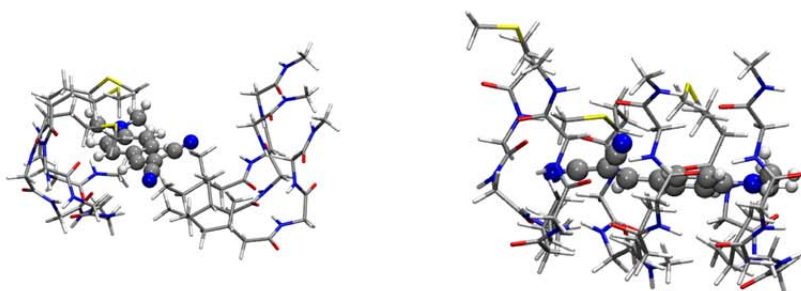


Figure S1. Computed TD-DFT(CAM-B3LYP) spectra from *ab initio* molecular dynamics simulations (PBE-D) of DANIR-2a in water and dichloromethane. Simulation time: 20 ps (80 evenly distributed snapshots) – red bars; 10 ps (40 evenly distributed snapshots) – blue bars.

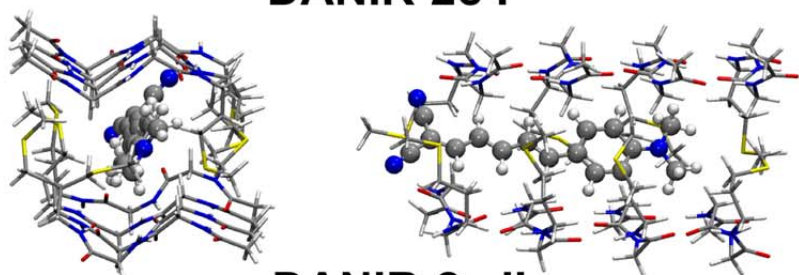
DANIR-2a I



DANIR-2a II



DANIR-2c I



DANIR-2c II

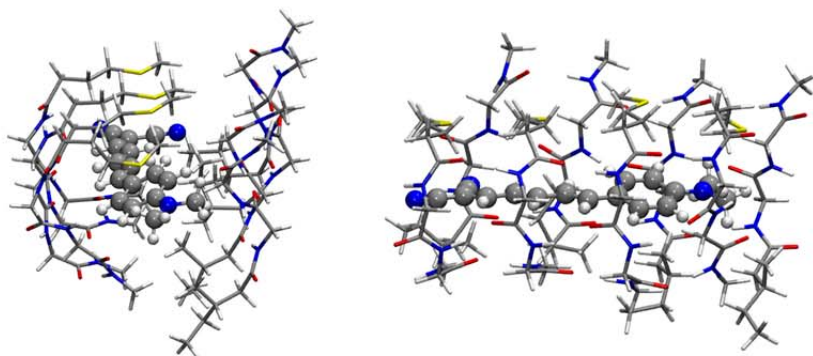


Figure S2. Selected models of fibril-marker interactions.

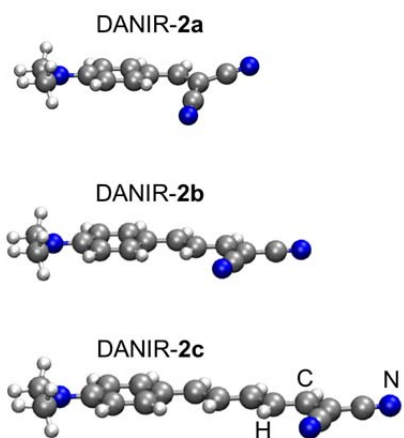
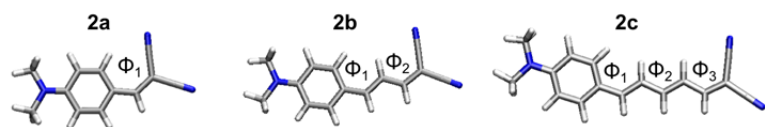


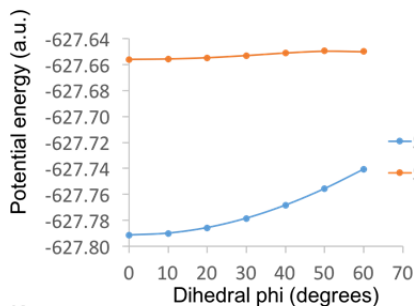
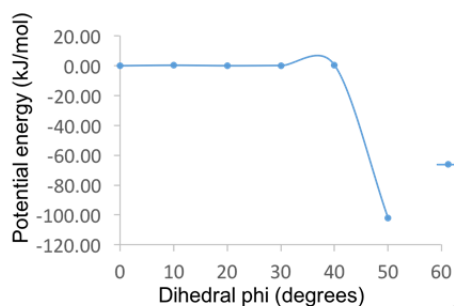
Figure S3. Optimized S_1 geometries of DANIR-2a, -2b and -2c.



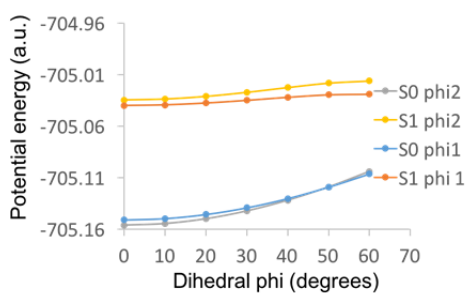
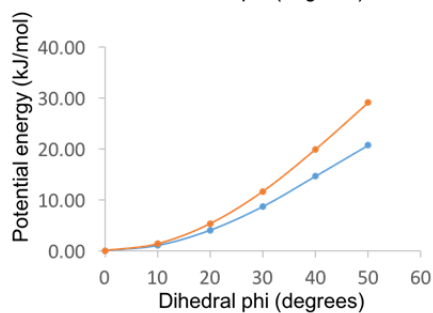
a) Relaxed scans, E(S1)

b) Rigid scans E(S0), E(S1)

DANIR-2a



DANIR-2b



DANIR-2c

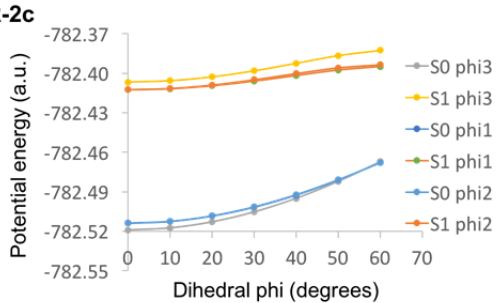
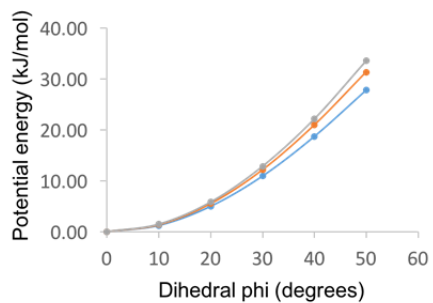


Figure S4 a) Relaxed and b) rigid potential energy scans around the double bonds of DANIR-2a, 2b and 2c.

