

The Ability of Chlorophyll to Trap Carcinogen Aflatoxin B₁, a Theoretical Approach.

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Figure S8. a) RMSD, and b) radius of gyration of **chl a 2**, obtained by MD simulation.

Table S1. Natural atomic charges (e⁻) for **chl a 1**, **chl a 2**, AFB₁ and their complexes.

Table S2. Interaction energy, in kcal/mol, of complexes **chl a 1-AFB₁**, and **chl a 2-AFB₁** in the gas phase.

Table S3. Interaction energy, in kcal/mol, of complexes **chl a 1-2AFB₁**, and **chl a 2-2AFB₁** in water as solvent.

Table S4. Cartesian coordinates for the more stable optimized geometries.

AFB₁ = aflatoxin B₁

chl a 2 = folded chlorophyll a.

chl a 1- α -E-AFB₁ = **AFB₁** coordinated to unfolded chlorophyll a.

chl a 2- β -E-AFB₁ = **AFB₁** coordinated to folded chlorophyll a.

chl a 1-D-2AFB₁ = **2AFB₁** coordinate to unfolded chlorophyll a.

chl a 2-E-2AFB₁ = **2AFB₁** coordinate to folded chlorophyll a.

chl a 2-AFB_{1c} = an intermolecular hydrogen bond between **AFB₁** and chlorophyll a.

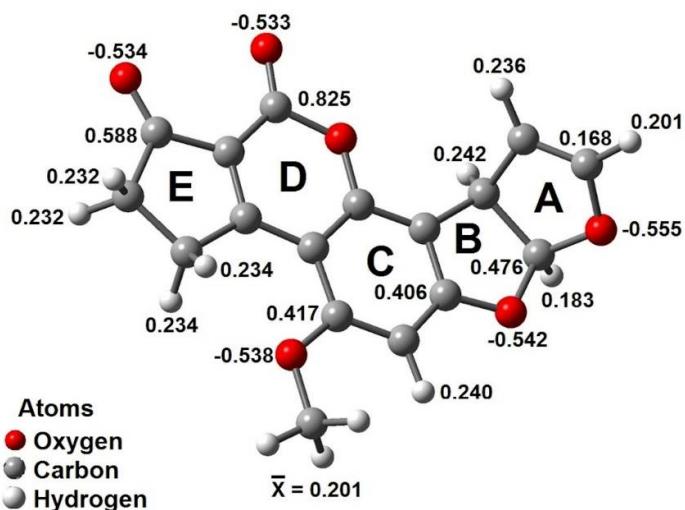


Figure S1. M06-2X/6-311G(d,p) optimized geometries of AFB₁. Atomic charges of selected atoms are in electron units.

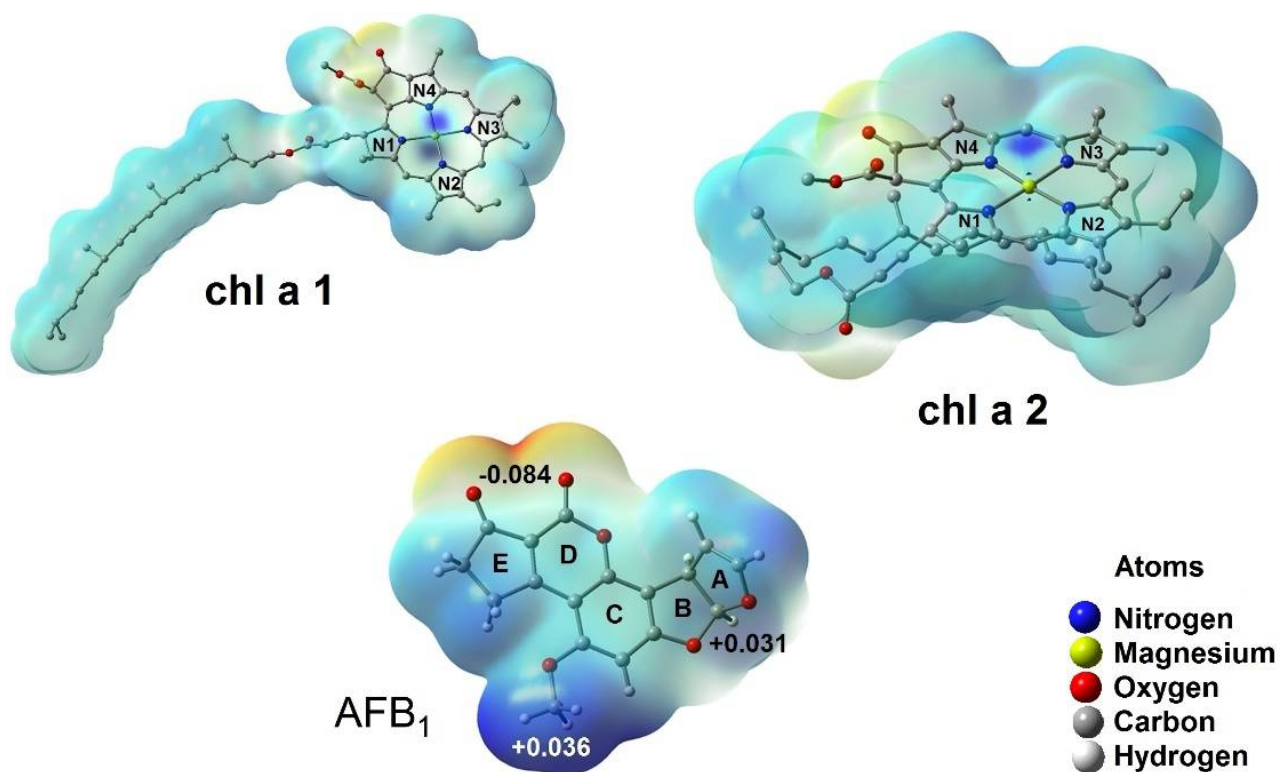


Figure S2. Reactive sites attained in the electrostatic potential map. The total electron density isosurface mapped with the molecular electrostatic potential of AFB₁, **chl a 1**, and **chl a 2**. The density = 0.0004 and isovalue = 0.02, at a level of calculation M06-2X/6-311G(d,p). The color scheme for the MEP is as follows: blue for electron-deficient, partially positive charge; light blue for the slightly electron-deficient region; yellow for a slightly electron-rich region and red for electron-rich, partial negative charge [1].

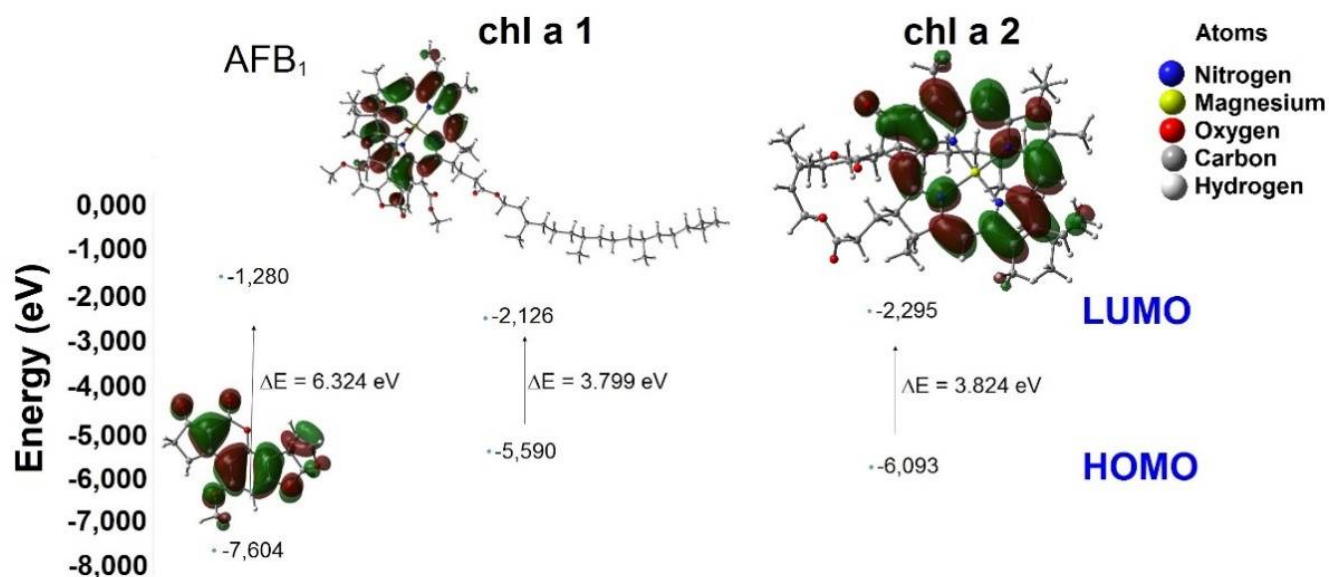


Figure S3. HOMO (eV), LUMO (eV) and the inter-frontier molecular orbital energy gaps of the target molecules.

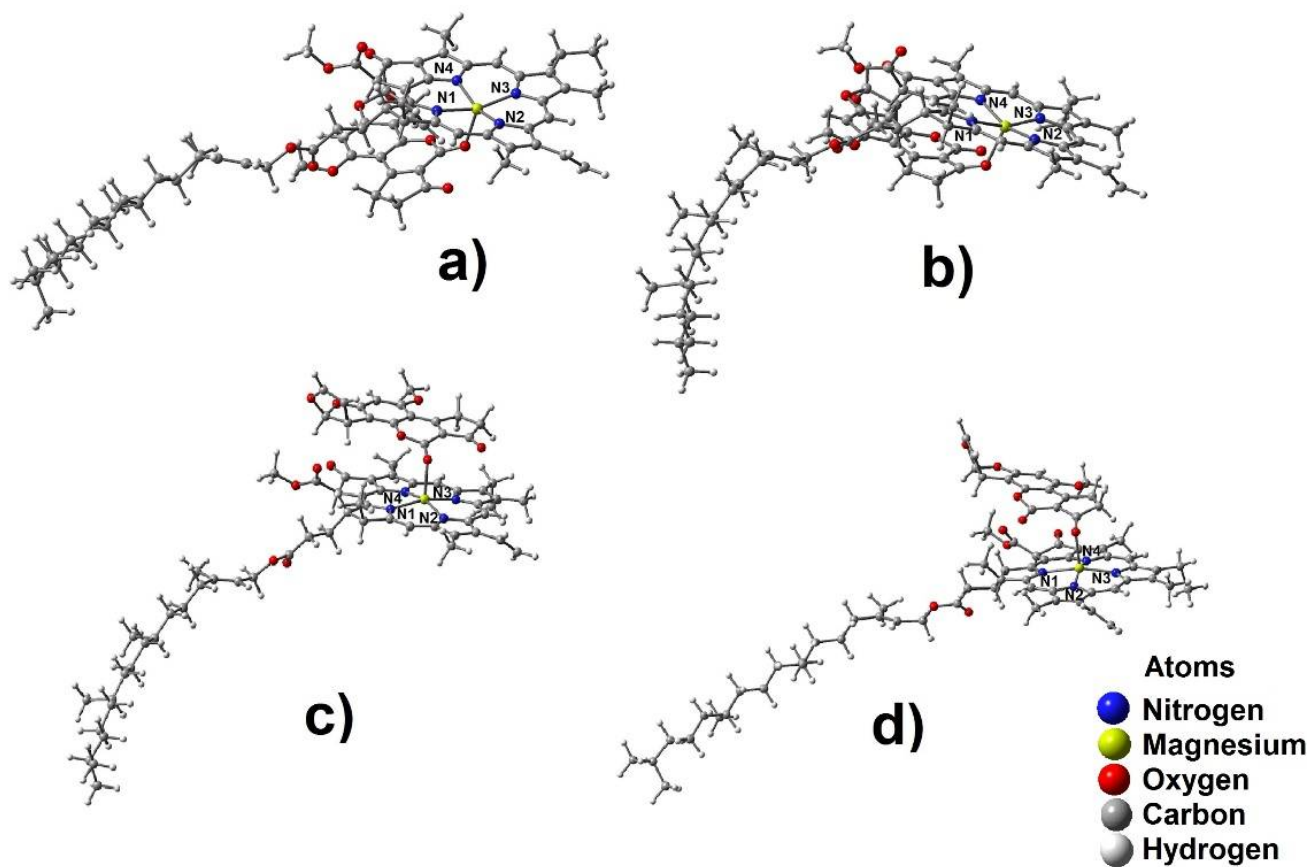


Figure S4. Optimized geometry: a) chl a 1- α -D-AFB₁, b) chl a 1- α -E-AFB₁, c) chl a 1- β -D-AFB₁, and d) chl a 1- β -E-AFB₁.

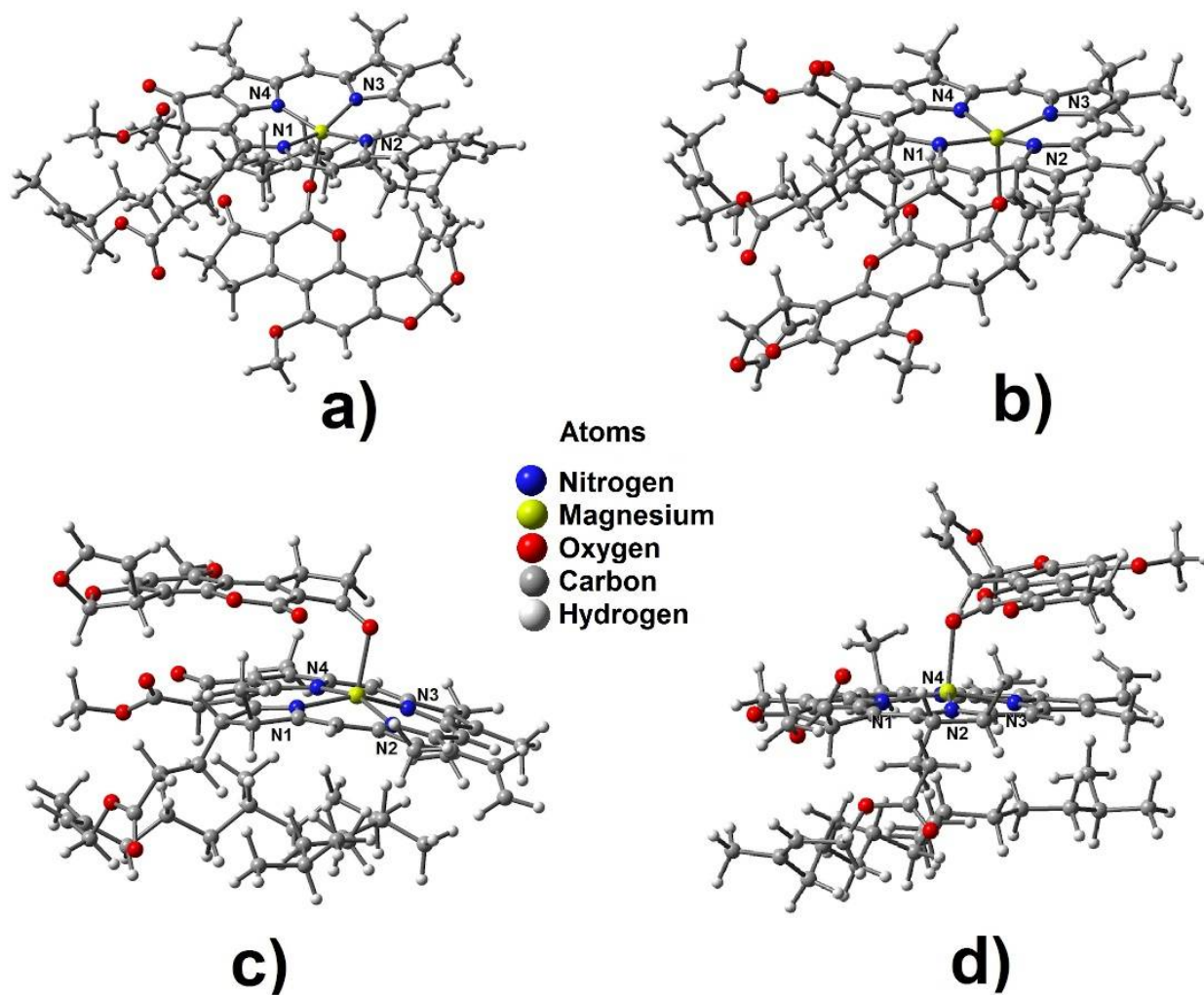


Figure S5. Optimized geometry: a) chl a 2- α -D-AFB₁, b) chl a 2- α -E-AFB₁, c) chl a 2- β -E-AFB₁, and d) chl a 2- β -D-AFB₁.

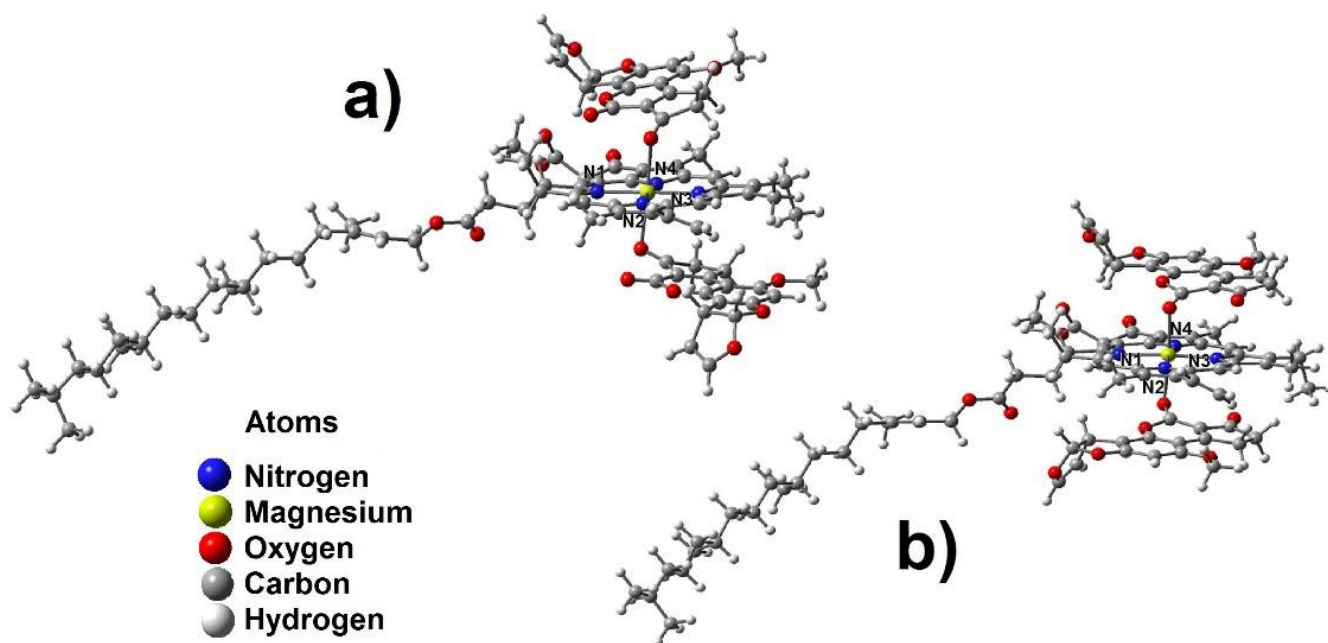


Figure S6. Optimized geometry of a) chl a 1-E-2AFB₁, and b) chl a 1-D-2AFB₁.

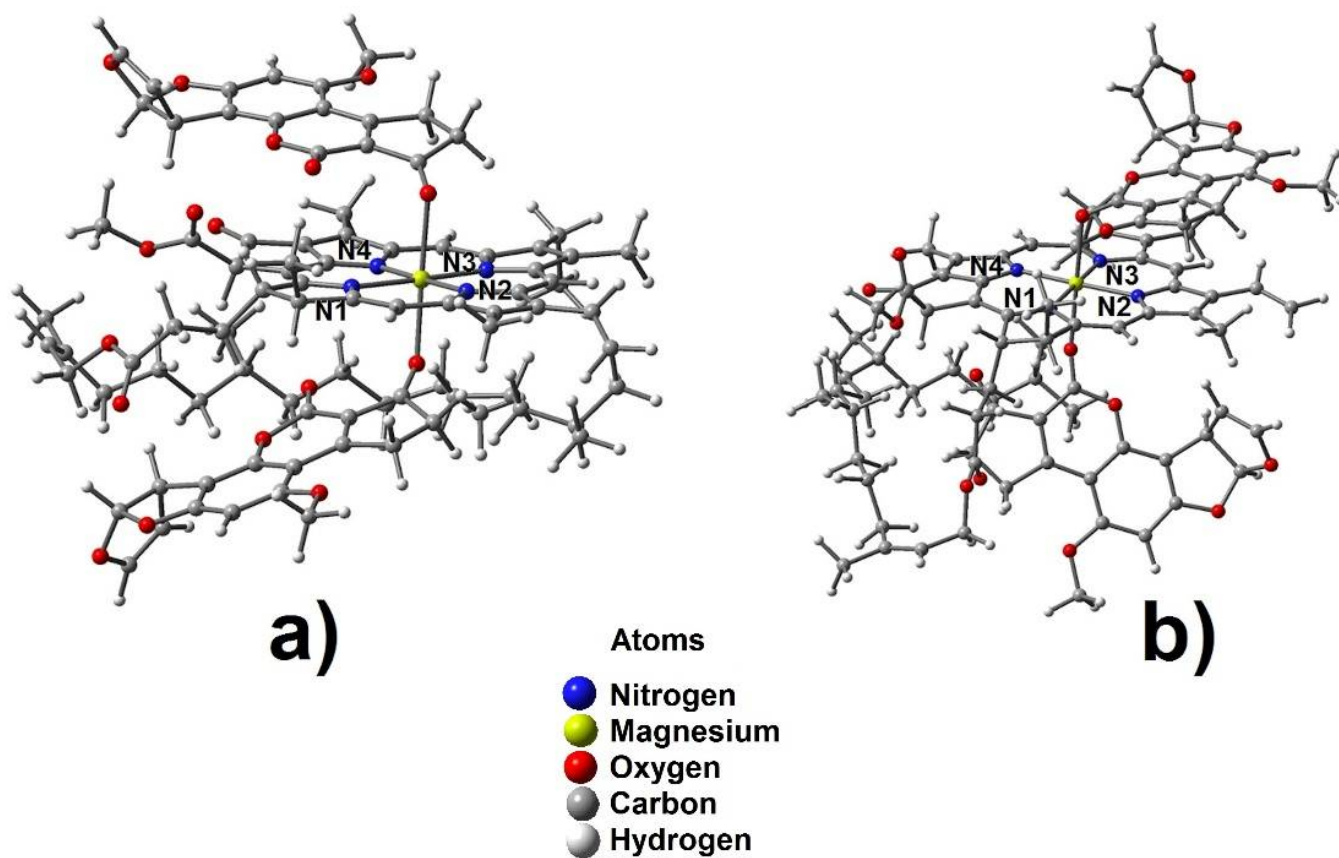


Figure S7. Optimized geometry of a) chl a 2-E-2AFB₁, and b) chl a 2-D-2AFB₁.

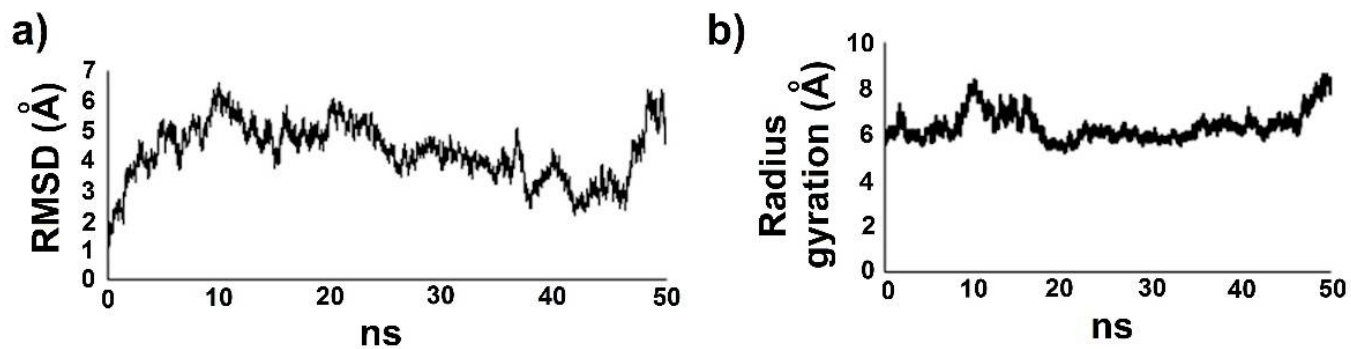


Figure S8. a) RMSD, and b) radius of gyration of chl a 2, obtained by MD simulation.

Table S1. Natural atomic charges (e⁻) for **chl a 1**, **chl a 2**, AFB₁ and their complexes.

	Mg ²⁺	N1	N2	N3	N4	O*	C*
chl a 2	1.762	-0.796	-0.745	-0.770	-0.726		
chl a 1	1.758	-0.792	-0.744	-0.769	-0.725		
chl a 1-α-D-AFB₁	1.759	-0.763	-0.708	-0.730	-0.693	-0.697	0.873
chl a 1-α-E-AFB₁	1.760	-0.698	-0.760	-0.706	-0.734	-0.695	0.646
chl a 1-β-E-AFB₁	1.758	-0.767	-0.716	-0.744	-0.691	-0.695	0.641
chl a 1-β-D-AFB₁	1.757	-0.763	-0.712	-0.745	-0.694	-0.699	0.868
chl a 2-α-D-AFB₁	1.760	-0.763	-0.723	-0.737	-0.690	-0.686	0.877
chl a 2-α-E-AFB₁	1.765	-0.746	-0.721	-0.742	-0.695	-0.687	0.648
chl a 2-β-E-AFB₁	1.766	-0.779	-0.718	-0.755	-0.704	-0.697	0.643
chl a 2-β-D-AFB₁	1.762	-0.768	-0.715	-0.747	-0.691	-0.691	0.870
chl a 1-D-2AFB₁	1.753	-0.763	-0.713	-0.727	-0.672	-0.664	0.869
						-0.612	0.853
chl a 1-E-2AFB₁	1.753	-0.751	-0.723	-0.737	-0.662	-0.650	0.638
						-0.622	0.624
chl a 2-D-2AFB₁	1.749	-0.737	-0.693	-0.711	-0.669	-0.631	0.866
						-0.640	0.865
chl a 2-E-2AFB₁	1.756	-0.737	-0.703	-0.726	-0.673	-0.643	0.638
						-0.644	0.642
				E		D	
				O	C	O	C
AFB ₁				-0.534	0.588	-0.533	0.825

E= specific oxygen atom of carbonyl group on ring E, D = specific oxygen atom of lactone group on ring D, O* and C*= atoms of AFB₁ when have coordinated to the Mg²⁺ ion.

Table S2. Interaction energy, in kcal/mol, of complexes **chl a 1-AFB₁**, and **chl a 2-AFB₁** in the gas phase.

Molecule	Interaction energy	Energy differences between chlorophylls	Energy difference considering carbonylic interaction
	kcal/mol		
chl a 1- α -D-AFB ₁	-32.1	4.3	3.6
chl a 1- α -E-AFB ₁	-36.4	0.0	0.0
chl a 1- β -E-AFB ₁	-35.5	0.9	0.9
chl a 1- β -D-AFB ₁	-35.7	0.7	0.0
chl a 2- α -D-AFB ₁	-33.3	5.9	0.0
chl a 2- α -E-AFB ₁	-36.5	2.7	2.7
chl a 2- β -E-AFB ₁	-39.2	0.0	0.0
chl a 2- β -D-AFB ₁	-32.8	6.4	0.4
chl a 1-AFB ₁ a	-17.0		
chl a 1-AFB ₁ b	-15.6		
chl a 2-AFB ₁ c	-29.6		
chl a 2-AFB ₁ d	-20.4		
chl a 1-E-2AFB ₁	-60.0	2.2	20.1
chl a 1-D-2AFB ₁	-62.3	0.0	11.2
chl a 2-E-2AFB ₁	-64.8	0.0	0.0
chl a 2-D-2AFB ₁	-58.0	6.7	0.0

a and b = three hydrogen bond interactions, c = one hydrogen bond interaction, d = two hydrogen bond interactions.

Table S3. Interaction energy, in kcal/mol, of complexes **chl a 1-2AFB₁**, and **chl a 2-2AFB₁** in water as solvent.

Chemical system	Interaction energy	Energy differences between chlorophylls	Energy difference considering carbonylic interaction
		kcal/mol	kcal/mol
chl a 1- α -D-AFB ₁	-22.6	3.9	3.8
chl a 1- α -E-AFB ₁	-26.5	0.0	0.0
chl a 1- β -E-AFB ₁	-23.6	2.9	2.9
chl a 1- β -D-AFB ₁	-26.4	0.1	0.0
chl a 2- α -D-AFB ₁	-23.4	4.5	3.3
chl a 2- α -E-AFB ₁	-27.6	0.3	0.3
chl a 2- β -E-AFB ₁	-27.9	0.0	0.0
chl a 2- β -D-AFB ₁	-26.6	1.3	0.0
chl a 1-AFB ₁ a	-10.3		
chl a 1-AFB ₁ b	-11.2		
chl a 2-AFB ₁ c	-23.0		
chl a 2-AFB ₁ d	-14.4		
chl a 1-E-2AFB ₁	-45.9	3.1	13.1
chl a 1-D-2AFB ₁	-49.0	0.0	8.7
chl a 2-E-2AFB ₁	-47.6	0.0	0.0
chl a 2-D-2AFB ₁	-46.3	1.3	0.0

a and b = three hydrogen bond interactions, c = one hydrogen bond interaction, d = two hydrogen bond interactions.

Table S4. Cartesian coordinate

AFB ₁				chl a 2				chl a 1- α -AFB ₁											
O1				O1				O1											
C	3.97611200	-1.04111000	0.09524100	Mg	-1.74528200	-0.15458700	-1.10972300	H	2.12533400	-5.23279100	-1.52614700	Mg	-5.04515200	2.62023600	0.24818000	H	5.46154600	-4.05418300	1.74152200
C	4.66777900	-2.02098700	0.08248700	N	-3.08474400	-1.65161200	-0.97568800	H	0.99765900	-4.45282500	-2.64258300	N	-4.55355000	4.59720000	0.30025000	H	7.01901200	-4.04542500	2.57864900
C	4.47256400	0.39949900	0.25272600	C	-4.45654900	-1.52729500	-1.10286700	C	3.15220800	-2.38470700	0.13230300	C	-5.38757900	5.62857900	-0.06678300	C	7.96219600	-1.64028700	1.87880100
C	5.17443000	0.61354400	-0.55520400	C	-5.05786300	-2.85027000	-0.88980800	H	2.68758500	-2.20283900	1.11967000	C	-4.61692000	6.87799900	-0.08526840	H	7.91616500	-0.54053500	1.83306700
H	5.02225400	0.47604600	1.19130600	C	-4.03165600	-3.72964200	-0.69769600	C	3.78170400	-1.49761400	-0.06882300	C	-3.35579000	5.56230200	0.32406000	H	8.21533300	-1.90779500	2.91304700
C	3.22487300	1.29070000	0.21338200	C	-2.80105500	-2.95694400	-0.77370300	C	4.06838300	-3.59282400	0.18075200	C	-3.33745700	5.18272000	0.54529200	C	9.02871600	-1.23734200	0.95534600
H	3.23447600	2.01391400	-0.60315100	C	-6.48461000	-3.17542400	-0.95313400	H	3.54429500	-4.48384000	0.52829100	C	-5.10438100	8.20610900	-0.46202000	H	8.81804300	-1.90032700	-0.08263800
H	3.07767600	1.86641000	1.12835500	H	-6.83396100	-3.94046000	-1.26300600	H	4.46668500	-3.77816000	-0.82221000	H	-4.75319100	9.03663200	0.14574500	H	15.31977000	-3.22687200	1.02786500
C	2.07363000	0.32835700	0.20258700	C	-7.35461500	-2.62735000	-0.80702400	C	5.22463600	-3.34074700	1.18017000	C	-5.90202200	8.43682900	-1.49768800	C	10.43397000	-0.54055200	1.30075500
C	2.50227600	-0.96072200	-0.03402800	H	-8.39701000	-2.92165600	-1.79818600	O	5.43780200	-3.91968300	2.14672600	H	-6.21539700	9.47648700	-1.72307600	H	10.29437300	-0.42102800	1.13927400
C	1.60571700	-2.09327600	-0.20051000	H	-7.04378000	-1.89267700	-2.53571000	O	5.98408000	-2.62859000	0.16637960	H	-6.24334900	7.76749400	-1.85068800	H	10.75427300	-1.85668500	2.28812200
O	1.86985100	-3.24957000	-0.25583500	C	-4.09854200	-5.20684400	-0.48434900	C	7.06058300	-1.90546900	1.53416300	C	-2.17689000	7.46533400	0.49010400	C	11.54747000	-1.76853300	0.28476900
O	0.25228000	-1.73577000	-0.30526000	H	-3.62196100	-5.48519300	0.46358600	H	7.72777000	-2.75590000	1.68870800	H	-1.35837600	7.16499900	-0.17008900	H	11.21615600	-1.37289900	-0.68554100
C	0.67733000	0.63841400	-0.07842100	C	-3.57636900	-5.74448900	-1.28252400	H	6.63086700	-3.87376000	2.50026400	H	-1.79966900	7.43468800	1.15615800	C	12.08947400	-1.00226500	0.68890200
C	-0.18872800	-0.48154500	-0.25480300	H	-5.10308900	-5.55612500	-0.46985300	C	7.74745900	-0.76469400	0.85860300	H	-2.43436200	5.49146000	0.25088300	H	12.53653900	0.04023300	0.91777000
C	-1.25721300	2.14418100	-0.13466800	C	-5.13612200	-0.32542000	-1.04273700	C	7.52467600	-0.52889000	1.10244400	H	-6.73885700	5.84729000	-0.29551000	H	13.17343000	-0.40253800	1.65454000
C	-1.54953100	-0.27026500	-0.38316400	H	-6.21658400	-0.38066700	-0.97896400	H	8.43728600	-1.03515700	0.06324400	H	-7.27878700	6.43012800	-0.52177400	C	13.94827400	-1.02513300	-0.32045000
H	-1.76504200	-0.32103600	-0.09099900	C	-3.25886600	1.23974400	-1.22710000	C	8.18194200	1.59351900	0.26784300	H	-7.051078500	3.18429000	0.08254400	H	13.57695700	-0.68048300	-1.42944000
C	-2.05129200	1.04959500	-0.31829700	C	-4.57763000	0.97493800	-1.08008800	H	7.42016900	1.16385500	-0.27877900	C	-7.52610700	6.34624300	-0.29262200	H	14.25793200	-0.25616900	-0.46355900
C	0.11250400	1.94021400	-0.01976300	C	-3.13265800	2.60776700	-1.29470300	H	8.87749800	1.16772300	-0.45670400	C	-8.12147000	2.25635000	0.08753500	C	15.11913400	-0.44235700	0.1183600
O	0.99362200	2.94701000	0.15263600	C	-4.43573800	3.22236200	-1.04746000	H	8.72696200	2.30231000	0.89767800	C	-9.32791000	2.98062300	-0.26769400	H	14.37248400	0.85045000	0.37260000
C	0.49297000	4.27225500	0.21944200	C	-5.34005900	2.18946700	-1.04577700	C	6.55098200	1.03760700	2.14060600	C	-8.95616700	4.28272300	-0.45685600	H	15.55083000	-0.55140000	1.03182900
H	-0.11819500	4.54245300	-0.70876600	C	-6.82388700	2.77812400	-0.84639900	H	6.51619300	0.37949100	3.01795300	C	-8.91305100	5.55494600	-0.85172000	C	15.73017000	0.03109400	-1.93172500
H	1.36150800	4.91008700	0.35814700	H	-7.13330100	1.77146000	0.05502200	H	6.89812100	2.01427200	2.50020300	H	-9.84215900	8.89447000	-1.79619700	H	15.77819100	-0.45756200	-1.83668400
H	-0.18832300	4.38899400	1.06649900	H	-7.35240400	1.79806900	-1.69477600	C	5.14150200	1.18437900	1.55409300	H	-9.76761000	7.24239500	-0.09627400	H	17.26860000	1.03043100	-0.40855500
O	-3.39296700	1.07027500	-0.44604000	H	-7.16066600	3.31642000	-0.90498200	H	4.82132300	0.19379500	1.02943500	H	-10.84510600	5.15760000	-0.97882900	H	16.74577900	1.95493800	-0.12663800
O	-4.80161600	-0.57873700	0.38475400	C	-4.68631900	4.69785000	-1.17318000	H	5.19638100	1.81565600	0.65391900	C	-10.67380700	2.35621000	-0.46629700	H	17.70910800	0.63301300	0.51652200
C	-3.03731200	-1.97295000	0.75221800	H	-5.72142000	4.90825600	-1.39810300	C	4.11246100	1.74089000	2.54376900	H	-11.45287300	3.11035700	-0.34665600	H	18.38751000	1.38126800	-1.38894800
H	-2.40863500	-2.71018300	1.27159900	C	-4.06282000	5.20952300	-1.85628300	C	3.89486000	0.19739400	3.29404100	H	-10.85474900	1.16038500	0.31335000	H	17.54534700	1.60676500	-2.35721800
C	-4.20450000	-1.52249000	1.18749400	C	-4.40502000	5.26504400	-0.28065700	H	4.54169900	2.51933500	3.06847000	C	-10.78951700	1.70391900	-1.85129700	H	19.02623300	0.50902200	-1.56177800
H	-4.77845600	-1.78905500	2.06201400	C	-4.56874600	6.34442000	-1.31060600	C	2.81289000	2.08326000	1.85765000	H	-11.76190400	1.22389300	-1.98238000	C	19.52523600	2.53482700	-0.88019400
C	-2.69127000	-1.23606600	-0.52301500	H	-3.37276400	5.06102900	0.57702700	H	2.54460700	1.42056200	1.18538000	H	-10.00545200	0.95632300	-1.193255900	H	18.64908500	3.45197500	-0.84536800
C	-2.5817800	-1.90080300	-1.38165300	H	-5.05683900	4.78397000	1.01870700	C	1.60836600	2.28773800	2.80266100	H	-10.69705000	2.50663200	-2.63313100	H	19.55007500	2.32331000	0.15347500
C	-3.89967800	-0.25926300	-0.64292000	H	-1.92188100	3.297299400	-1.47065600	H	1.68120600	1.53283300	3.59346500	C	-10.51674600	2.94346700	0.39707900	C	20.51367600	0.50052800	-1.70719900
H	-4.44188700	-0.25572600	-1.58587500	H	-1.98806200	4.27288900	-1.56932600	H	1.61676600	3.26399300	3.30459900	H	-8.98968800	0.34363700	0.33498000	C	21.12152400	1.89149300	-1.69903600
				C	-0.65904600	2.72024800	-1.53535300	C	0.28073400	2.09232400	2.06089800	C	-6.91757900	0.18396500	0.76427900	C	18.82057500	-2.26489900	0.12352400
				N	-0.45831500	1.35904700	-1.34977800	H	0.18315100	1.03062200	1.80461900	N	-5.67273200	0.78492200	0.86350500	H	10.93149000	-3.80343100	-0.20892800
				C	-1.51552100	-3.50395900	-0.65682600	H	0.29791500	2.63039300	1.10483300	C	-2.20408000	4.40350700	0.96219000	H	12.40645200	-3.69624900	1.07838700
				H	-1.46629700	-4.56890500	-0.46862100	C	-0.91948800	2.56987200	2.87572400	H	-1.30279400	4.98459800	1.12042200	H	10.60533500	-3.45322600	-0.61085100
				C	-0.29808900	-2.84552100	-0.27508000	H	-0.82029400	2.21746800	3.91204200	C	-2.10649700	3.04045500	1.27595600	H	6.87370700	-1.30649400	-1.30218200
				H	-0.1218200	-1.52865300	-0.98951500	H	-0.88137800	3.66467500	2.92425000	N	-1.30733900	1.21856600	1.06892000	H	16.14231900	-2.00804000	-1.70724700
				C	1.22357200	-1.23803800	-1.07905500	C	-2.29478900	2.13425200	2.35543400	C	-2.62299500	0.87492500	1.39400300	H	17.33029200	-1.76567700	-0.4184800
				C	0.84842200	1.16621200	-1.45065200	H	-2.36742800	2.38372100	1.28997200	C	-4.82453900	-0.18632100	1.15706200	H	17.65274900	-1.17912500	-2.05583100
				C	1.68421600	0.02973700	-1.3192100	C	-2.46898700	1.60872500	2.49759400	C	-3.42100800	-0.29246000	1.37261300	C	13.21387800	0.33046500	-1.07373300
				C	0.60002000	3.38035500	-1.77633700	H	-1.89977200	0.89289300	1.71083600	C	-8.16686800	-1.22331500	1.04382000	H	20.75808100	4.86367000	-1.0771200
				C	1.54421000	2.36830000	-1.72630300	H	-2.01042000	0.29155700	3.44437000	C	-5.46456700	1.44447200	1.21613100	H	22.25886100	1.01597000	-1.62247500
				C	0.82060300	1.83374900	-2.01229200	C	-3.91897400	0.10344000	2.47800000	C	-7.92176400	-2.22652300	0.96862600	H	21.58729600	3.69893800	-0.03695800
				H	0.25679000	5.18665000	-2.18849400	H	-4.42409300	0.47842800	1.58401500	H	-8.69767600	-2.00329800	1.70543700	C	20.18106200	3.14352700	-3.16189500
				H	1.87878600	5.0310000	-2.8171200	H	-4.44396900	5.52020400	3.34089700	H	-7.52390600	-3.22050600	1.17958300	H	19.51030940	4.10045300	-3.2061700
				H	0.49932300	5.42891000	-1.14904400	C	-4.00314100	-1.42263200	2.53650000	H	-8.38910000	-2.25277300	-0.02134800	H	19.69078800	2.31646600	-3.67400800
				C	2.														

chl a 2-β-E-AFB ₁					chl a 1-D-2AFB ₁										
O1					O1										
Mg	2.35789400	-0.91822900	-0.94294600	H	-6.50999900	147613200	4.57445000	Mg	-4.16600000	194300000	-0.51000000	H	15.97800000	-0.61800000	0.32700000
N	3.69091000	0.40553000	-1.70689000	H	-5.55632900	155005000	6.06630900	N	-3.87000000	3.91900000	-0.17900000	C	16.46800000	1.07700000	1.56200000
C	5.05653080	0.33218000	-1.6140900	C	-3.18853700	2.49872600	4.88779900	C	-4.80500000	4.90900000	-0.35500000	H	15.99900000	1.98800000	1.91600000
C	5.65223600	1.35767000	-2.48003600	H	-2.77247000	3.50933200	4.89376900	C	-4.27700000	6.16300000	0.20000000	H	16.68800000	0.46900000	2.45500000
C	4.62772300	1.99807400	-3.09885900	H	-3.17936700	2.15249700	5.92599900	C	-3.05500000	5.87300000	0.72500000	C	17.82600000	1.47000000	0.92300000
C	3.39650000	1.40673000	-2.48007000	C	-2.28208700	1.58467300	4.0538900	C	-2.87000000	4.45400000	0.46800000	H	17.60600000	0.30320000	0.00500000
C	7.08951000	1.57968000	-2.6345600	C	-2.30576900	1.94599000	3.01615900	C	-4.92800000	7.47600000	0.17700000	H	18.59800000	2.40000000	1.86800000
H	7.72363600	0.69720800	-2.59823000	H	-2.70470000	0.57585400	4.03573600	H	-4.83200000	8.07200000	1.08100000	H	17.95700000	3.25700000	2.11200000
C	7.65304400	2.77565400	-2.79589300	C	-0.83712100	1.55755800	4.55518300	C	-5.58600000	7.98800000	-0.86200000	H	18.78400000	1.87500000	2.81200000
H	8.72443000	2.88075000	-2.96562400	H	-0.38046100	2.53378500	4.35474000	H	-6.03700000	8.87000000	-0.80800000	C	19.92200000	2.92600000	1.33000000
H	7.06020300	3.68398000	-2.79238300	H	-0.83088300	1.43189700	5.64596900	H	-5.67000000	7.45000000	-1.79900000	H	19.75500000	3.31500000	0.30200000
C	4.66913200	3.04898900	-1.15994100	C	0.02350900	0.44901900	3.93092200	C	-2.08800000	6.78900000	1.40000000	H	20.64400000	2.10700000	1.27000000
H	4.52075000	4.05223000	-3.74985300	H	-0.15158000	0.44674800	2.84366600	H	-1.14600000	6.84000000	0.84500000	C	20.52800000	4.01400000	2.19900000
H	3.88829100	2.87689000	-4.90385800	C	1.52173500	0.76090000	4.1297200	H	-1.85600000	6.44000000	2.41000000	H	19.87400000	4.89700000	2.18800000
H	5.63606000	3.03220900	-4.66541500	H	1.72157500	1.76073100	3.87503400	H	-2.48700000	7.80100000	1.47000000	H	20.54400000	3.65700000	3.23600000
C	5.75675900	-0.54380900	-0.81755900	H	1.78085400	0.59887200	5.18288100	C	-6.04900000	4.73800000	-0.91600000	C	21.94500000	4.44500000	1.80800000
H	6.83650500	-0.44733500	-0.81277000	C	2.44748300	-0.18745400	3.27077900	H	-6.70500000	5.60000000	-0.94000000	H	22.59100000	3.56500000	1.86300000
N	3.80435200	-1.82239500	0.15070100	H	1.96302500	-0.33428400	2.28078600	N	-5.92500000	2.37900000	-1.56700000	C	13.52200000	-1.85500000	-0.22400000
C	5.21200100	-1.54602800	0.02767900	H	2.44277000	-1.19586300	3.78929500	C	-6.56800000	3.55400000	-1.50300000	H	12.76600000	-2.24500000	-0.90800000
C	3.79537000	-2.82755700	1.08868900	C	3.85764300	0.30746800	3.09138500	C	-6.75000000	1.50700000	-2.24400000	H	13.67400000	-2.59400000	0.57000000
C	5.11986300	-3.21003500	1.55780900	H	4.17625300	0.83624000	4.00096700	C	-7.87500000	2.18000000	-2.62000000	H	14.45200000	-1.76900000	-0.78700000
C	6.00394400	-2.39096400	0.90541900	H	4.52951000	-0.55308400	2.98178600	C	-7.86500000	3.46800000	-2.15000000	C	18.85600000	0.24200000	0.54700000
C	7.49155600	-2.3268600	1.04873300	C	4.08942200	1.21727100	1.87291000	C	-8.85800000	4.58300000	-2.21900000	H	18.10900000	-0.42900000	-0.18000000
H	7.80948000	-1.33623400	1.40100400	H	3.93056700	0.60678100	0.97743200	H	-8.42900000	5.47300000	-2.68800000	H	18.92600000	-0.32000000	1.44700000
H	7.9976400	-2.50663400	0.09406000	C	3.08823900	2.37975000	1.78535500	H	-9.18800000	4.86200000	-1.21400000	H	19.57900000	0.52300000	0.03600000
H	7.85254100	-3.06137000	1.76456600	H	2.08482900	1.96980400	1.67549000	H	-9.73700000	4.29600000	-2.79800000	C	22.47300000	5.48500000	2.79600000
C	5.40208100	-1.19723300	2.63982200	H	3.10544300	2.95087700	2.72349000	C	-9.09000000	1.59000000	-3.42200000	H	21.85700000	6.38900000	2.76200000
H	6.40489200	-1.41616000	2.50512900	C	3.35717600	3.34654800	0.62707600	H	-10.02600000	2.10900000	-3.19400000	H	23.50100000	5.77300000	2.55800000
C	4.71034000	-3.60458000	1.78191000	H	3.64518700	2.78155000	-0.26587200	H	-9.24200000	0.54500000	-3.12900000	H	22.45600000	1.01600000	3.82000000
C	5.28753600	-3.56376600	4.03322000	H	4.19371500	4.00483000	0.87212900	C	-8.87000000	1.86300000	-4.93000000	C	22.00200000	5.98100000	0.38100000
H	5.46665300	-4.29828300	4.82028600	C	2.10716600	4.18494400	0.32600600	H	-9.63400000	1.21900000	-5.50300000	H	21.31000000	4.89400000	2.69000000
H	4.29349800	-3.13394800	4.17833800	H	1.41558300	3.56968500	-0.26584800	H	-7.89200000	1.13700000	-5.17700000	H	21.73200000	4.23400000	-0.35800000
H	6.00357500	-2.75506200	4.14385000	H	1.58947200	4.44520000	1.26648600	H	-8.69900000	2.70200000	-5.24000000	H	23.00600000	5.34800000	0.14000000
C	2.60509100	-3.34755900	1.57179300	C	2.35913000	5.49933500	-0.41584500	C	-6.46000000	0.17200000	-2.49500000	C	-8.10700000	1.27000000	0.63200000
H	2.68288400	-1.13636500	2.34482700	H	2.97258500	6.13521900	0.23551800	H	-7.20500000	-0.40300000	-3.03800000	D	-7.96100000	2.43900000	0.88300000
C	1.31009100	-2.88409300	1.78191000	C	-0.36120400	-0.91860200	4.49985800	C	-5.29500000	-0.50700000	-2.11010000	C	-9.39300000	0.62300000	0.11800000
N	1.08962400	-1.84249400	0.38536100	H	-1.3879100	-1.19138200	4.28258200	N	-4.25400000	0.37200000	-1.44900000	H	-9.74200000	1.17000000	-0.75800000
C	2.10887500	1.18805500	-2.82962400	H	-0.23588500	-0.90970700	5.58797100	C	-1.65200000	1.32000000	0.83400000	H	-10.15300000	0.70200000	0.89700000
C	2.03427200	2.7007800	-3.22553600	C	0.27403200	-1.70664300	4.08269400	H	-0.89200000	4.35400000	1.34200000	C	-9.04600000	-0.84400000	-0.16300000
C	0.92416700	1.51038800	-2.19449500	C	5.54186000	1.69449400	1.85157000	C	-1.33600000	2.43400000	0.59600000	H	-9.02600000	-1.07400000	-1.23300000
N	0.75229300	0.4872500	-1.34874900	H	6.22205000	0.84352200	1.95627800	N	-2.11000000	1.52400000	-0.00900000	H	-9.73400000	-1.55800000	0.29100000
C	-0.52120700	0.53208400	-0.80398800	H	5.73510000	2.38767900	2.67877000	C	-1.43300000	0.32900000	-0.13600000	C	-7.65000000	-0.02200000	0.38200000
C	-0.18369700	-1.49714800	0.54965260	H	5.78920800	2.18963700	0.91455400	C	-3.30600000	-0.78100000	-1.30000000	C	-7.12800000	0.17200000	0.79500000
C	-0.97578500	-0.40723100	0.07518700	C	1.03601700	6.21634200	-0.68494900	C	-1.98500000	-0.76200000	-0.74100000	C	-5.75200000	0.29200000	1.18300000
C	0.10101500	-2.32759000	1.95187400	H	0.47523000	5.62350500	-1.36835100	C	-4.97700000	-1.88600000	-2.33900000	D	-5.12000000	1.33000000	1.29000000
C	-0.84574600	-2.33268000	1.47254900	H	1.19575500	7.19328100	-1.14878670	C	-3.69700000	-2.04100000	-1.81600000	D	-5.09600000	-0.88000000	1.41500000
C	-0.09083700	-4.29837800	2.98740000	H	0.46083200	6.36279000	0.232710500	C	-5.83700000	-2.90400000	-3.01800000	C	-6.87300000	-2.21400000	0.45200000
H	-0.05493100	-5.29937800	2.54353200	C	3.12438300	5.2765100	-1.71783800	H	-6.87200000	-2.84000000	-2.67000000	C	-5.59400000	-2.08500000	1.03400000
H	-1.06094600	-4.76107000	3.46986100	H	2.53930100	4.63006400	-2.38480800	H	-5.45900000	-3.90600000	-2.81400000	C	-6.43700000	-4.61040000	0.12700000
H	0.68606800	-1.25768000	3.75505800	H	4.08232500	4.77566200	-1.53794900	H	-5.84000000	-2.75300000	-4.10100000	C	-4.77400000	-3.17300000	1.20400000
C	-2.20486800	-1.95139900	1.68009400	H	3.31589200	6.21596700	-2.23438600	C	-2.64100000	-3.01000000	-1.54200000	H	-6.69800000	-5.58600000	-0.23700000
D	-3.14078100	-2.34878700	2.23970200	C	1.1818100	-3.05783900	-2.36538600	D	-2.64100000	-4.20600000	-1.59900000	C	-5.20200000	-4.40600000	0.74500000
C	-2.20892700	-0.42618600	0.98567900	D	2.19538800	-2.37920500	-2.47073000	C	-1.39900000	-2.12300000	-1.08600000	C	-7.26500000	-3.50000000	-0.10100000
H	-1.97383200	0.210084200	1.85763000	C	11124300	-4.43844700	-1.74553500	H	-0.78700000	-2.05800000	-1.99500000	D	-8.47000000	-3.55000000	-0.62000000
C	-3.55502000	0.08240600	0.55340500	H	1.84742700	-4.52250600	-0.94583300	C	-0.55300000	-2.83100000	-0.06000000	C	-8.89200000	-4.79200000	-1.15800000
D	-4.36904800	0.1608400	1.60273600	H	1.35764400	-5.16673300	-2.52234300	D	0.15900000	-3.80500000	-0.62700000	H	-8.19300000	-5.13700000	-1.92600000
C	-5.68331800	0.64877700	1.32286100	C	-0.34630500	-4.56440400	-1.27778900	C	0.93200000	-4.60400000	0.27000000	H	-9.86500000	-4.80600000	-1.16060000
H	-5.62273100	1.66660600	0.93433700	H	-0.42940900	-4.46798300	-0.19407280	H	1.40900000	-5.35900000	-0.34800000	H	-9.98500000	-5.54700000	-0.37400000
H	-6.20498900	0.63745600	2.27590500	H	-0.82801900	-5.50501500	-1.54890600	H	0.28300000	-5.06800000	1.01600000	D	-4.29400000	0.91800000	0.91800000
H	-6.17701800	-0.10713200	0.60326000	C	-1.05452600	-3.48787200	-1.93646500	H	1.67700000						

chl a 2-E-2AFB₁

01

Mg -2.50845900 -0.47355000 -0.88537000
N -3.40087500 -1.91142000 -1.99757600
C -4.72343600 -2.24999900 -1.93675700
C -4.95352800 -3.40836000 -2.80987500
C -3.75269500 -3.70992600 -3.38656000
C -2.78793800 -2.74276500 -2.85264600
C -6.24847400 -4.08300000 -2.99602000
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C -6.42454500 -5.37578600 -3.13004000
H -7.41028700 -5.79870900 -3.27405600
H -5.58026700 -6.06808300 -3.07365300
C -3.43437900 -4.76282900 -4.39759900
H -2.96309200 -5.63786200 -3.93723200
H -2.74557700 -4.37661900 -5.15216200
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C -5.68337500 -1.61855500 -1.17455000
H -6.69257700 -2.00369200 -1.26446500
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C -5.50843600 -0.53403000 -0.26757700
C -4.55133300 0.02949000 0.93873000
C -5.97939000 1.16545000 1.20343000
C -6.57465000 0.14715000 0.45164600
C -8.04378800 -0.14844000 0.38205500
H -8.28036700 -0.86078900 -0.40888600
H -8.61083000 0.76364000 0.17933400
H -4.40308000 -0.56702400 1.32288500
C -6.62928200 2.04549600 2.17610400
H -7.6859500 2.51335900 1.9155500
H -8.1895300 3.04530300 2.10224300
C -6.50540800 1.53489000 3.67933000
H -6.99480000 2.20868200 4.32378400
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C -3.58654500 1.92550700 1.40053300
H -8.93437400 2.70569700 2.07321000
C -2.22925300 1.95008500 1.05736600
C -1.85387600 0.88897700 2.40075900
C -1.4734800 -2.72038400 -3.16515300
H -1.07908900 -3.45986200 -3.88285700
C -0.43513400 -1.84233400 -2.89555500
N -0.59898000 -0.87824000 -1.79137700
C -0.58983500 -0.23526300 -1.54353800
C -0.39317400 1.32915400 0.09330300
C -0.6947500 0.810408700 -0.65723200
C -1.24866900 2.94402200 1.41843900
C -0.08165400 2.52148400 0.8071000
C -1.46644600 4.1763700 2.24204800
H -2.2803900 4.7723200 1.84009800
H -0.56368200 4.78408300 2.27407600
H -1.07296800 3.91658500 3.27782400
C 1.32334700 2.83750700 0.59212600
D 1.97442100 3.7872200 0.95798300
C 1.90772700 1.61548000 -0.22105100
H 2.46055300 1.01616800 0.54316300
C 2.93887900 2.04203200 -1.23598700
C 0.04655200 2.4698800 -0.60680900
C 5.10736300 2.8848200 -1.46228400
H 5.52227300 2.30376300 -1.98845700
H 5.85523100 3.3307200 -0.80412800
H 4.7478100 3.63084400 -2.18374500
D 2.83949600 2.0160600 -2.43285300
C 1.01018920 1.19692700 -3.19849700
H 1.42396900 -2.8947900 -2.92346700
C 1.69854800 -0.78806700 -2.41276300
H 1.99520700 0.01652700 -3.09368700
C 1.09235300 -1.5772500 -4.74095500
H 0.57742700 -2.52469400 -5.25400000
H 2.13750800 -1.7863700 -5.03476800
H 0.63723000 -0.7737000 -4.99139000
C 2.92263500 -1.23481600 -1.60829300
H 7.1273700 -2.19504200 -1.02877700
C 1.83262700 -0.52193800 -0.80882300
H 4.83395500 -1.36982500 -2.47348000
H 4.14136700 -2.22101200 -3.15319600
H 4.31452900 -0.44783400 -3.04699800
C 5.38749700 -1.58733400 -1.58887600
C 5.95577500 -2.6160500 -1.38009600
D 5.7348800 -0.4700400 -0.99202700
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chl a 2-AFB₁c

01

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Reference

1.- Politzer, P.; Murray, J.S. The fundamental nature and role of the electrostatic potential in atoms and molecules. *Theor Chim Acta* **2002**, 108,134-142.