

Chemistry–A European Journal

Supporting Information

Quest for the Most Aromatic Pathway in Charged Expanded Porphyrins

Irene Casademont-Reig, Tatiana Woller, Victor García, Julia Contreras-García,
William Tiznado, Miquel Torrent-Sucarrat,* Eduard Matito,* and Mercedes Alonso*

1 Conformational and topological switches induced by (de)protonation

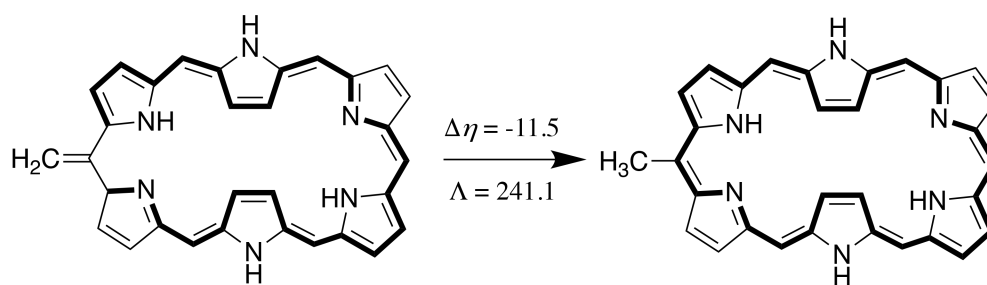


Figure S1: Reaction used to evaluate several global aromaticity descriptors in hexaphyrins. $\Delta\eta$ is given in $\text{kcal}\cdot\text{mol}^{-1}$ and Λ in $\text{ppm}\cdot\text{cgs}$.

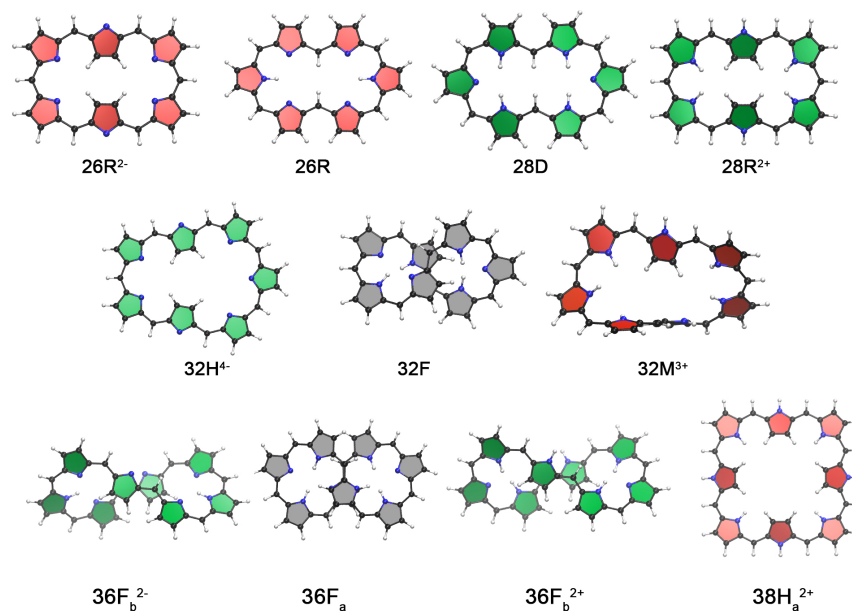


Figure S2: Lowest-energy conformations for neutral and charged hexa-, hepta-, and octaphyrins. Aromatic and antiaromatic macrocycles are colored as red and green, respectively. Nonaromatic macrocycles are colored in grey.

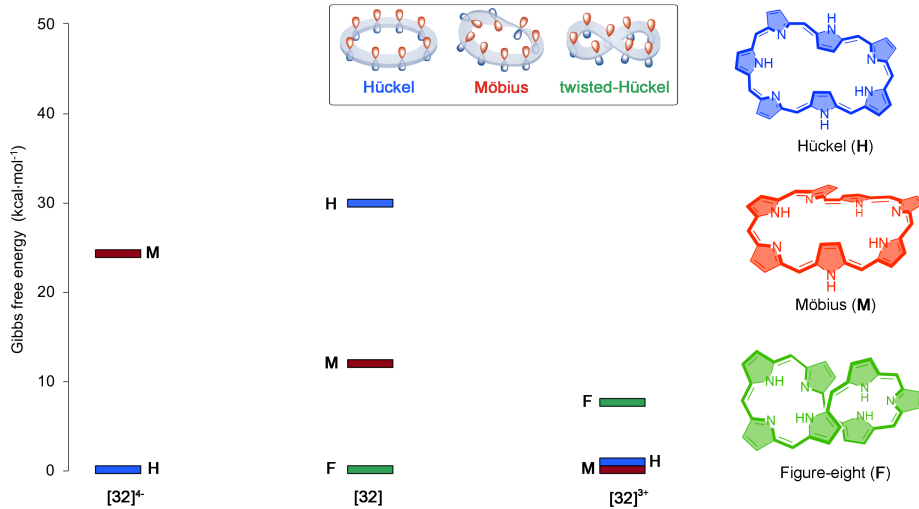


Figure S3: Evolution of the relative Gibbs free energy of selected heptaphyrins with oxidation state upon protonation or deprotonation of the macrocycles.

Table S1: Relative Gibbs free energies (kcal·mol⁻¹), hydrogen bond index (N_H) and ring strain (Φ_p in $^\circ$) of hexaphyrins and heptaphyrins.^a

[26]	ΔG	N_H	Φ_p	[28]	ΔG	N_H	Φ_p	[32]	ΔG	N_H	Φ_p
F	17.9	3	33.4	F	8.8	3	25.3	F	0.0	3.5	19.0
M	21.5	2	32.6	M	5.9	2.5	32.5	M	11.9	3	34.6
R	7.6	2	12.6	R	7.6	2	11.0	H	29.8	1	24.8
D	0.0	3	3.1	D	0.0	3	11.2	F ⁴⁻	-	-	-
T	27.8	0	26.4	T	28.7	0	24.1	M ⁴⁻	24.2	0	28.8
F ²⁻	50.3	0	37.1	F ²⁺	-	-	-	H ⁴⁻	0.0	0	0.0
M ²⁻	-	0	-	M ²⁺	3.8	0	34.4	F ³⁺	7.5	0	31.2
R ²⁻	0.0	0	6.4	R ²⁺	0.0	0	17.6	M ³⁺	0.0	0	34.7
D ²⁻	27.7	0	0.0	D ²⁺	8.1	0	16.3	H ³⁺	0.8	0	24.8
T ²⁻	7.1	0	24.4	T ²⁺	9.5	0	32.8				

^a Gibbs free energies at the M06-2X/6-311+G(d,p)//CAM-B3LYP/6-311G(d,p) level of theory.

Table S2: Relative Gibbs free energies (kcal·mol⁻¹), hydrogen bond index (N_H) and ring strain (Φ_p in $^\circ$) of octaphyrins.^a

[36]	ΔG	N_H	Φ_p	[36]	ΔG	N_H	Φ_p	[38]	ΔG	N_H	Φ_p	[36]	ΔG	N_H	Φ_p
F _a	0.0	4	15.4	F _a ²⁻	12.6	3	15.1	F _a ²⁺	2.3	0	17.7	F _a ²⁺	1.5	3	16.0
F _b	14.5	3	17.7	F _b ²⁻	0.0	3	20.1	F _b ²⁺	9.7	0	24.7	F _b ²⁺	0.0	3	19.8
M _a	27.9	2.5	24.5	M _a ²⁻	1.2	2.5	23.8	M _a ²⁺	10.9	0	26.4	M _a ²⁺	4.6	2	24.8
M _b	49.7	0	30.1	M _b ²⁻	21.4	0	31.2	M _b ²⁺	11.0	0	35.8	M _b ²⁺	27.9	0	31.0
H _a	33.4	0	6.0	H _a ²⁻	12.2	0	3.8	H _a ²⁺	0.0	0	16.7	H _a ²⁺	13.7	0	11.1
H _b	49.8	0	26.7	H _b ²⁻	17.4	0	22.1	H _b ²⁺	5.8	0	27.8	H _b ²⁺	31.6	0	27.6

^a Gibbs free energies at the M06-2X/6-311+G(d,p)//CAM-B3LYP/6-311G(d,p) level of theory.

Table S3: Relative Gibbs free energies (kcal·mol⁻¹) and RMSD values of expanded porphyrins optimized with CAM-B3LYP and different basis sets.

system	ΔG	ΔG	RMSD	system	ΔG	ΔG	RMSD
	D3 dispersion	No dispersion			6-311+G(d,p)	6-311G(d,p)	
26R ²⁻	0	0	0.017	28R	9.9	7.6	0.008
26D ²⁻	26.9	27.7	0.030	28D	0	0	0.057
26F ²⁻	51.1	50.4	0.052	28T	30.2	28.7	0.011
26M ²⁻	5.5	7.1	0.026	28M	7.4	5.9	0.015
26M ²⁻	<i>a</i>	<i>a</i>	<i>a</i>	28F	10.3	8.8	0.017
28R ²⁺	0	0	0.107	32F ⁴⁻	<i>a</i>	<i>a</i>	<i>a</i>
28M ²⁺	3.5	3.8	0.143	32H ⁴⁻	0	0	0.017
28T ²⁺	9.8	9.5	0.022	32M ⁴⁻	22.8	24.2	0.018
28D ²⁺	7.7	8.1	0.043				
28F ²⁺	<i>a</i>	<i>a</i>	<i>a</i>				

^a Those topologies could not be optimized as an energy minimum since the conformation changes upon the optimization.

Table S4: Pearson coefficients between several descriptors of aromaticity for unsubstituted expanded porphyrins (n=20).

	$\Delta\eta$	Λ	NICS(0)	NICS _{zz} (1)
$\Delta\eta$	1			
Λ	-0.672 ^a	1		
NICS(0)	-0.641 ^a	0.915 ^a	1	
NICS _{zz} (1)	-0.670 ^a	0.962 ^a	0.923 ^a	1

^a The correlation is significant at the 0.01 level (two-tailed).

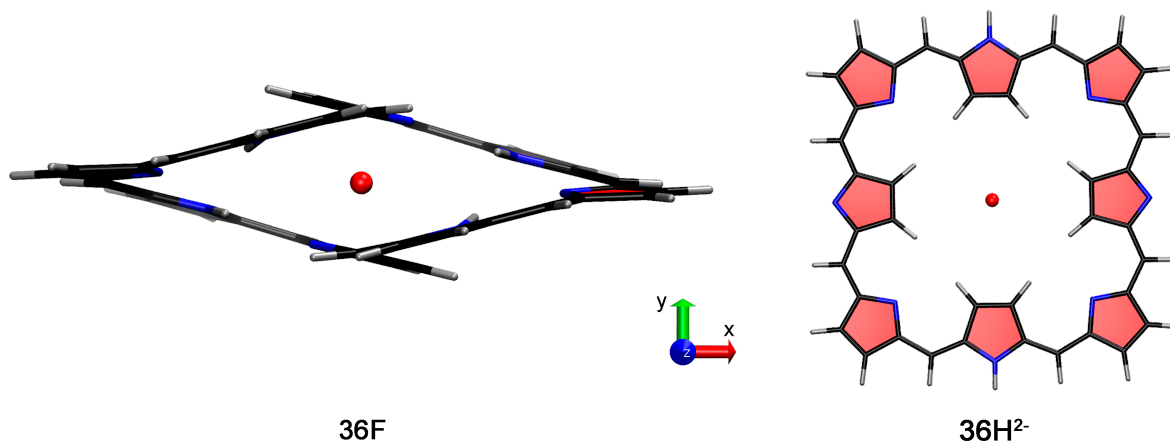


Figure S4: Orientation of the twisted-Hückel and Hückel octaphyrin structures employed in the calculation of the NICS-based indices. The magnetic field is oriented along the z axis.

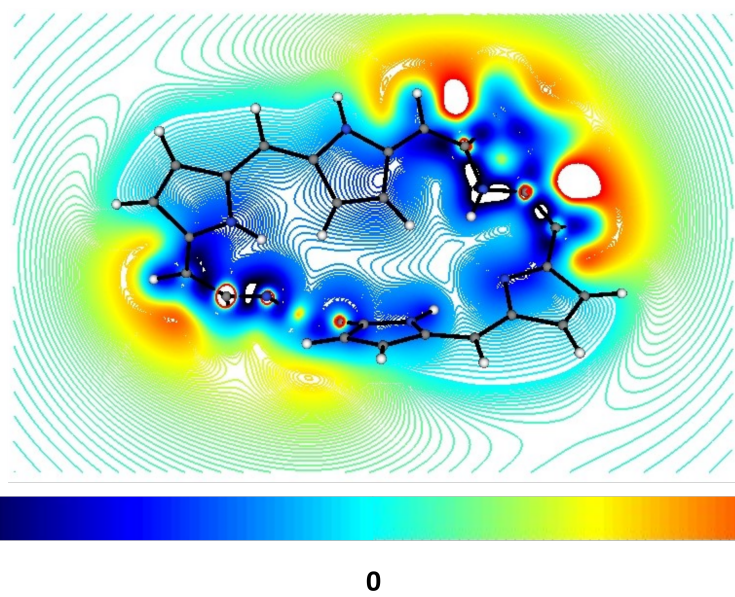


Figure S5: 2D-NICS_{zz}(1) plots (in ppm) of the magnetic shielding of **28M** in the xy plane 1Å situated above the molecular plane.

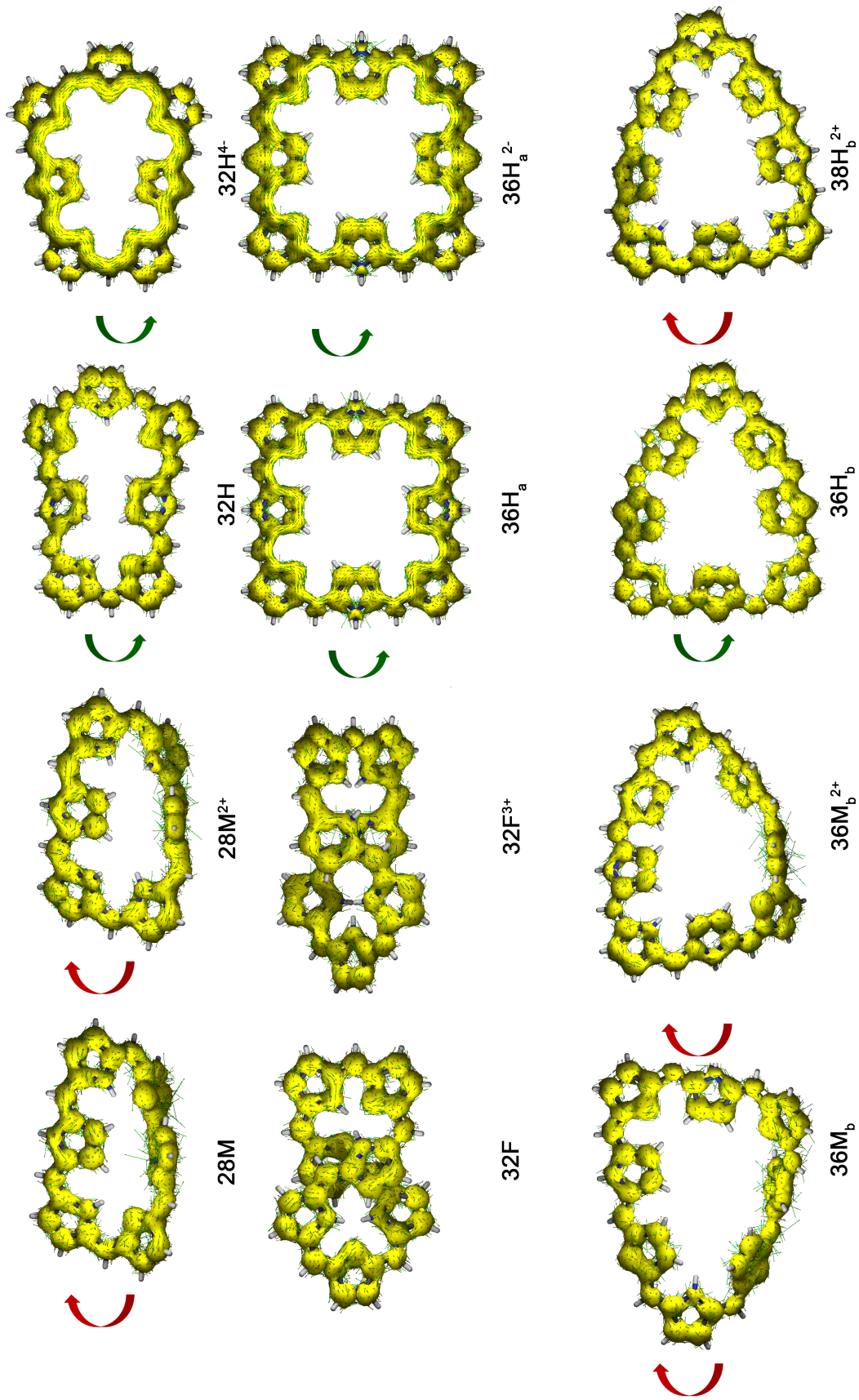


Figure S6: AICD plots of selected neutral, protonated, and deprotonated expanded porphyrins (28M, 28M²⁺, 32H, 32H⁴⁻, 32F, 32F³⁺, 36H_a, 36H_a²⁻, 36M_b, 36M_b²⁺, 36H_b, and 38H_b²⁺) (isovalue of 0.06 au).

2 Magnetically induced currents in neutral and charged expanded porphyrins

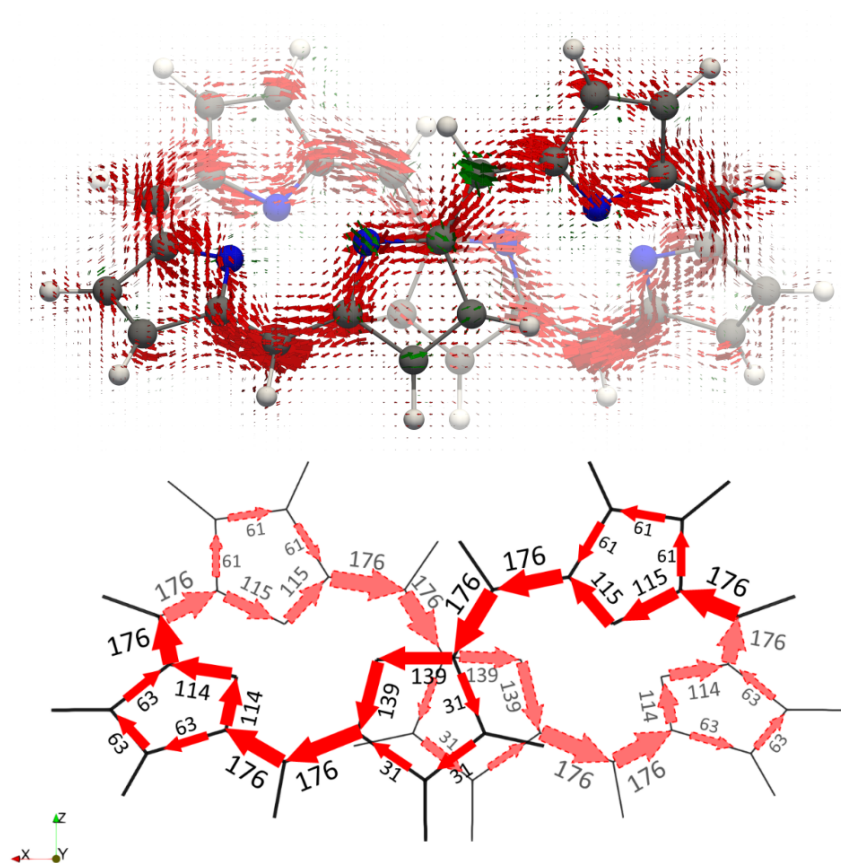


Figure S7: Visualization of the current density plots using a vector representation in $26\mathbf{F}^{2-}$. The percentage ratios are with respect to the net bond current strength in pyrrole ($11.8 \text{ nA}\cdot\text{T}^{-1}$). The external magnetic field is parallel to the symmetry z axis. The percentage ratios are with respect to the net bond current strength in pyrrole ($11.8 \text{ nA}\cdot\text{T}^{-1}$).

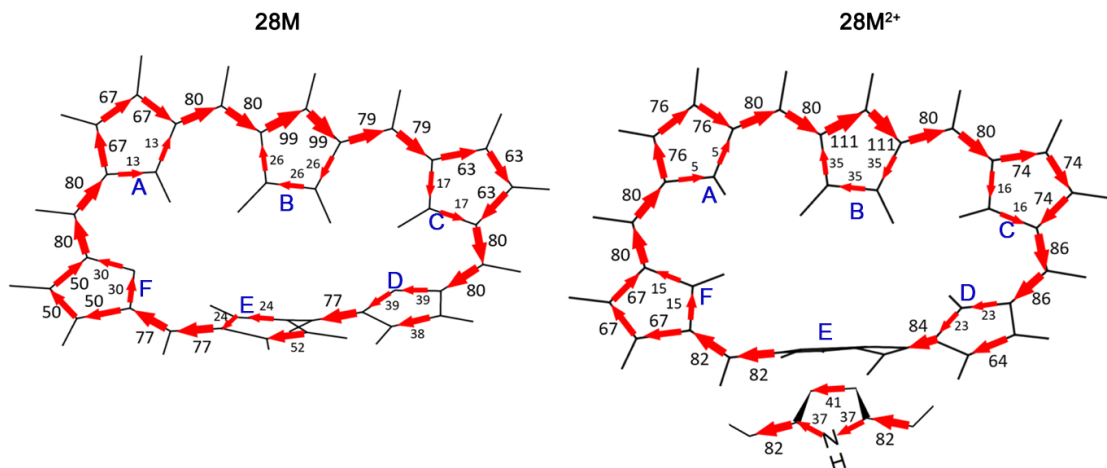


Figure S8: Percentage ratios of the current strength in **28M** and **28M²⁺** with respect to the net bond current strength in pyrrole (11.8 nA·T⁻¹).

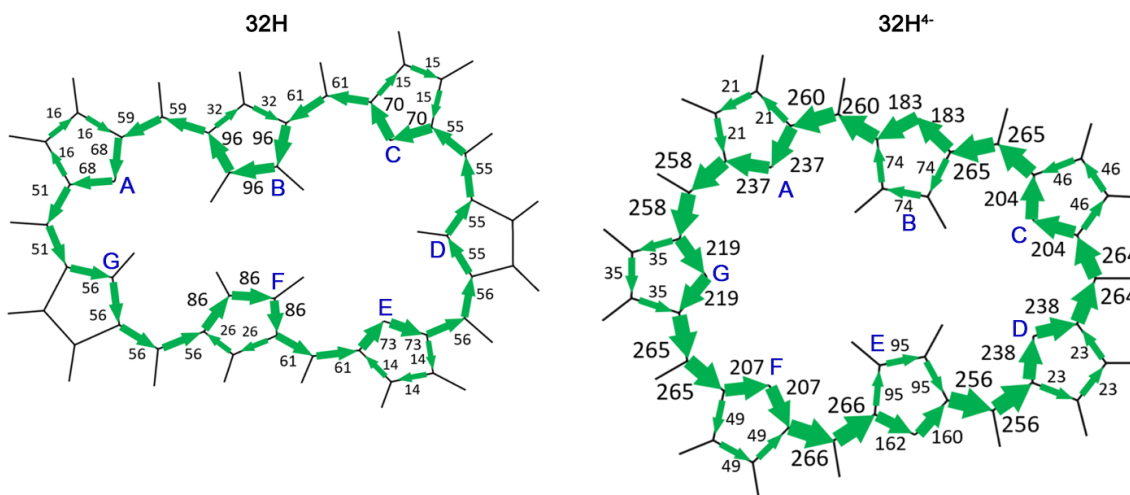


Figure S9: Percentage ratios of the current strength in **32H** and **32H⁴⁻** with respect to the net bond current strength in pyrrole (11.8 nA·T⁻¹).

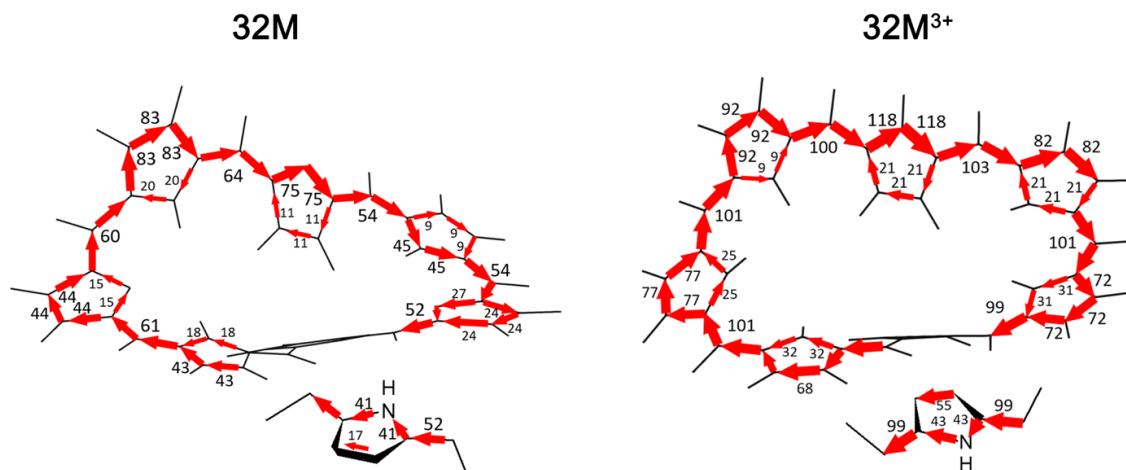


Figure S10: Percentage ratios of the current strength in **32M** and **32M³⁺** with respect to the net bond current strength in pyrrole ($11.8 \text{ nA}\cdot\text{T}^{-1}$).

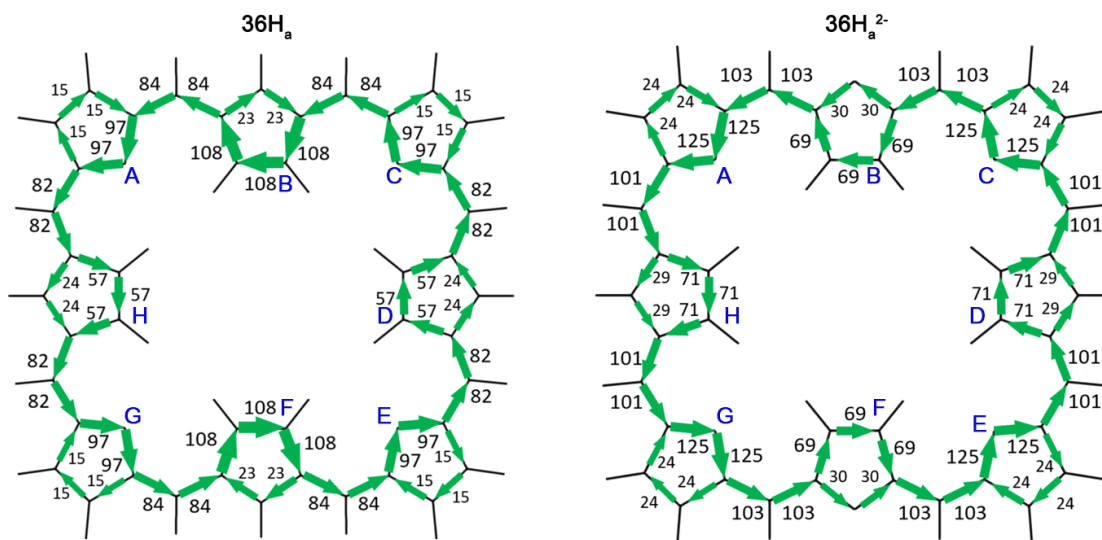


Figure S11: Percentage ratios of the current strength in **36H_a** and **36H_a²⁻** with respect to the net bond current strength in pyrrole ($11.8 \text{ nA}\cdot\text{T}^{-1}$).

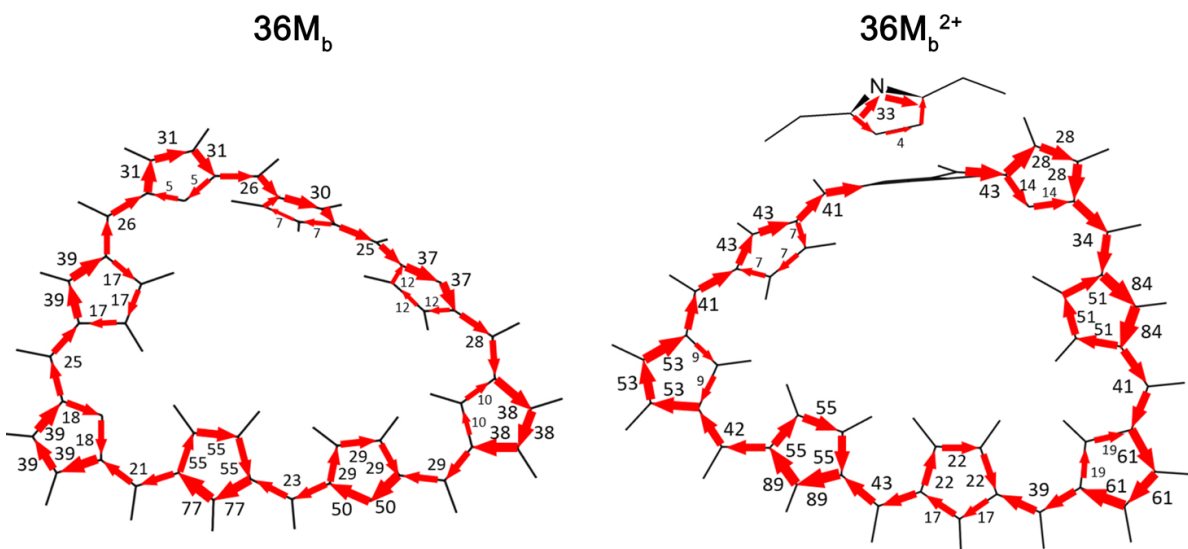


Figure S12: Percentage ratios of the current strength in $36M_b$ and $36M_b^{2+}$ with respect to the net bond current strength in pyrrole ($11.8 \text{ nA}\cdot\text{T}^{-1}$).

3 Local aromaticity indices in neutral and charged expanded porphyrins

Table S5: Evolution of the aromaticity indices for the annulene pathway of expanded porphyrins upon the protonation of the macrocycle.^a

system	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV_{min}
28R	28	0.021	0.300	0.620	0.067	1.31	0.42
28R²⁺	28	0.020	0.264	0.682	0.051	1.07	0.16
28M	28	0.018	0.269	0.677	0.059	1.45	0.60
28M²⁺	28	0.018	0.253	0.718	0.048	1.14	0.01
32F	32	0.020	0.292	0.632	0.065	1.28	0.39
32F³⁺	32	0.020	0.251	0.695	0.048	1.15	0.15
32M	32	0.020	0.292	0.664	0.065	1.40	0.53
32M³⁺	32	0.017	0.234	0.749	0.045	1.17	0.00
36M_b	36	0.025	0.330	0.582	0.076	1.16	0.35
36M_b²⁺	38	0.024	0.313	0.593	0.069	1.48	0.15
36H_b	36	0.025	0.323	0.579	0.076	1.15	0.21
38H_b²⁺	38	0.019	0.250	0.702	0.024	1.48	0.04

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

Table S6: Evolution of the aromaticity indices for the inner pathway of expanded porphyrins upon the protonation of the macrocycle.^a

system	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV_{min}
28R	24	0.020	0.251	0.719	0.054	0.07	0.03
28R²⁺	24	0.020	0.219	0.735	0.043	-0.07	0.05
28M	24	0.016	0.215	0.784	0.043	0.13	0.01
28M²⁺	24	0.019	0.217	0.750	0.041	-0.16	0.01
32F	28	0.017	0.238	0.759	0.053	0.38	0.02
32F³⁺	28	0.020	0.227	0.732	0.044	0.11	0.10
32M	28	0.018	0.238	0.765	0.051	0.33	0.01
32M³⁺	28	0.018	0.211	0.783	0.038	0.10	0.00
36M_b	32	0.024	0.287	0.662	0.064	0.43	0.04
36M_b²⁺	32	0.022	0.259	0.701	0.057	0.16	0.03
36H_b	32	0.024	0.286	0.662	0.064	0.43	0.00
38H_b²⁺	32	0.018	0.208	0.783	0.038	-0.34	0.02

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

Table S7: Evolution of the aromaticity indices for the outer pathway of expanded porphyrins upon the protonation of the macrocycle.^a

system	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV_{min}
28R	30	0.026	0.324	0.487	0.069	1.45	0.08
28R²⁺	30	0.022	0.295	0.604	0.063	1.59	0.20
28M	30	0.024	0.311	0.515	0.067	1.48	0.04
28M²⁺	30	0.020	0.283	0.642	0.059	1.67	0.22
32F	35	0.027	0.330	0.443	0.070	1.39	0.01
32F³⁺	35	0.021	0.156	0.613	0.060	1.54	0.12
32M	35	0.026	0.324	0.475	0.069	1.41	0.06
32M³⁺	35	0.019	0.267	0.684	0.055	1.63	0.20
36M_b	40	0.032	0.362	0.315	0.079	1.14	0.03
36M_b²⁺	40	0.027	0.323	0.464	0.057	0.16	0.03
36H_b	40	0.032	0.363	0.300	0.080	1.12	0.02
38H_b²⁺	40	0.019	0.003	0.665	0.000	1.87	0.27

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

Table S8: Evolution of the aromaticity indices for the annulene pathway of expanded porphyrins upon deprotonation of the macrocycles.^a

system	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV_{min}
26R	26	0.019	0.278	0.716	0.065	1.56	0.60
26R²⁻	26	0.017	0.218	0.659	0.057	1.58	0.38
26F	26	0.023	0.317	0.619	0.074	1.27	0.35
26F²⁻	26	0.015	0.124	0.720	0.044	1.66	0.66
32M	32	0.020	0.292	0.664	0.065	1.40	0.53
32M⁴⁻	32	0.023	0.261	0.450	0.066	1.30	0.02
32H	32	0.022	0.309	0.611	0.070	1.21	0.42
32H⁴⁻	32	0.024	0.280	0.423	0.071	1.11	0.01
36H_a	36	0.022	0.304	0.636	0.068	1.21	0.40
36H_a²⁻	34	0.020	0.265	0.659	0.067	1.36	0.54

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

Table S9: Evolution of the aromaticity indices for the inner pathway of expanded porphyrins upon deprotonation.^a

system	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV _{min}
26R	24	0.017	0.250	0.785	0.057	1.05	0.12
26R ²⁻	24	0.009	0.142	0.888	0.045	1.73	1.23
26F	24	0.020	0.281	0.712	0.067	0.86	0.43
26F ²⁻	24	0.006	0.018	0.971	0.029	1.83	1.54
32M	28	0.018	0.238	0.765	0.051	0.33	0.01
32M ⁴⁻	28	0.010	0.137	0.840	0.044	0.86	0.21
32H	28	0.020	0.258	0.730	0.056	0.40	0.01
32H ⁴⁻	28	0.012	0.175	0.793	0.052	0.82	0.09
36H_a	32	0.020	0.260	0.738	0.056	0.56	0.03
36H_a ²⁻	32	0.017	0.219	0.773	0.049	0.93	0.11

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

Table S10: Evolution of the aromaticity indices for the outer pathway of expanded porphyrins upon deprotonation.^a

system	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV _{min}
26R	30	0.031	0.351	0.346	0.078	1.02	0.01
26R ²⁻	30	0.031	0.334	0.286	0.077	1.13	0.01
26F	30	0.034	0.369	0.256	0.080	1.00	0.00
26F ²⁻	30	0.029	0.296	0.345	0.069	1.25	0.18
32M	35	0.026	0.324	0.475	0.069	1.41	0.06
32M ⁴⁻	35	0.028	0.311	0.305	0.070	1.51	0.02
32H	35	0.029	0.348	0.389	0.077	1.25	0.01
32H ⁴⁻	35	0.029	0.322	0.280	0.072	1.47	0.01
36H_a	40	0.030	0.351	0.360	0.076	1.20	0.04
36H_a ²⁻	40	0.027	0.314	0.456	0.068	1.70	0.22

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

Table S11: Aromaticity indices of the three most delocalized circuits of anionic expanded porphyrins.^a

system	Circuit ^b	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV _{min}
26R ²⁻	iiiiii	24	0.009	0.142	0.888	0.045	1.73	1.23
26R ²⁻	iiiiio	25	0.012	0.167	0.812	0.047	1.78	0.59
26R ²⁻	oiiiiio	26	0.015	0.19	0.741	0.048	1.82	0.59
26F ²⁻	iiiiii	24	0.006	0.018	0.971	0.029	1.83	1.53
26F ²⁻	iiiiio	25	0.010	0.074	0.841	0.037	1.74	0.66
26F ²⁻	iiiioio	26	0.014	0.124	0.732	0.044	1.66	0.63
32H ⁴⁻	iooioio	32	0.021	0.26	0.503	0.060	1.16	0.19
32H ⁴⁻	ioiioio	31	0.021	0.256	0.516	0.064	0.96	0.19
32H ⁴⁻	ioioiio	30	0.021	0.254	0.531	0.068	0.73	0.19
32M ⁴⁻	oiooooo	32	0.020	0.234	0.535	0.056	1.54	0.35
32M ⁴⁻	oiooioo	32	0.019	0.23	0.543	0.055	1.53	0.35
32M ⁴⁻	oiooioi	31	0.019	0.225	0.565	0.059	1.34	0.35
36H_a ²⁻	iiiioiio	34	0.020	0.265	0.659	0.067	1.36	0.54
36H_a ²⁻	iiiioioo	35	0.021	0.284	0.628	0.068	1.29	0.54
36H_a ²⁻	ioioioio	36	0.023	0.302	0.599	0.069	1.23	0.54

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

^b i denotes the inner route (C-N-C) and o the outer route (C-C) in a pyrrole ring. The labelling of the conjugation pathway starts from the top right five-membered ring.

Table S12: Aromaticity indices of the three most delocalized circuits of cationic expanded porphyrins.^a

system	Circuit ^b	πe^-	FLU	BOA	HOMA	BLA	AV1245	AV _{min}
28R ²⁺	oooooo	30	0.022	0.295	0.604	0.063	1.59	0.20
28R ²⁺	iooooo	29	0.021	0.263	0.642	0.174	1.34	0.16
28M ²⁺	oooooo	30	0.020	0.283	0.642	0.059	1.67	0.22
28M ²⁺	ooioooo	29	0.019	0.266	0.685	0.052	1.46	0.20
32F ³⁺	ioooooo	32	0.017	0.232	0.761	0.039	1.38	0.20
32F ³⁺	iooioooo	33	0.018	0.25	0.713	0.045	1.38	0.21
32F ³⁺	ioooooo	34	0.020	0.273	0.655	0.054	1.41	0.17
32M ³⁺	ooiooio	32	0.016	0.222	0.787	0.036	1.25	0.28
32M ³⁺	ooiooio	33	0.017	0.235	0.760	0.042	1.43	0.32
32M ³⁺	ooioooo	34	0.018	0.252	0.724	0.049	1.52	0.24
36M_b ²⁺	oooooio	38	0.024	0.313	0.593	0.069	1.48	0.15
36M_b ²⁺	ooooiio	37	0.025	0.313	0.577	0.072	1.28	0.15
36M_b ²⁺	ooioiio	36	0.021	0.285	0.674	0.060	1.12	0.12
38H_b ²⁺	oooooooo	40	0.019	0.266	0.665	0.055	1.87	0.27
38H_b ²⁺	oooioooo	39	0.018	0.249	0.716	0.048	1.73	0.21
38H_b ²⁺	ooooooooio	39	0.018	0.249	0.708	0.048	1.71	0.19

^a All indices were calculated at the CAM-B3LYP/6-311+G(d,p) level of theory.

^b i denotes the inner route (C-N-C) and o the outer route (C-C). The labelling of the conjugation pathway starts from the top right five-membered ring, except for 32M and 36M in which the first labelled five-membered ring corresponds to the top left one.

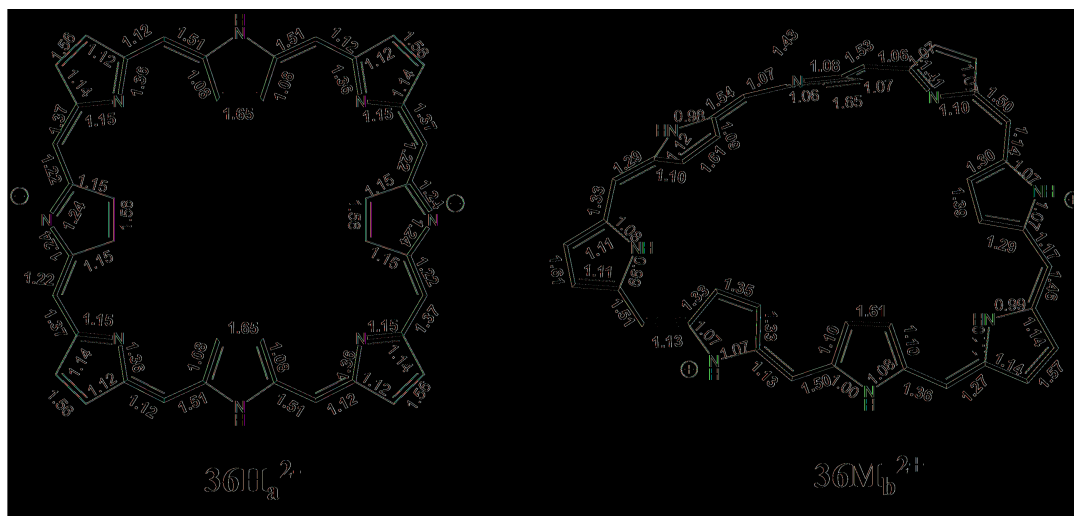


Figure S13: Delocalization indices (DIs) for $36M_a^{2-}$ and $36M_b^{2+}$ macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold bonds.

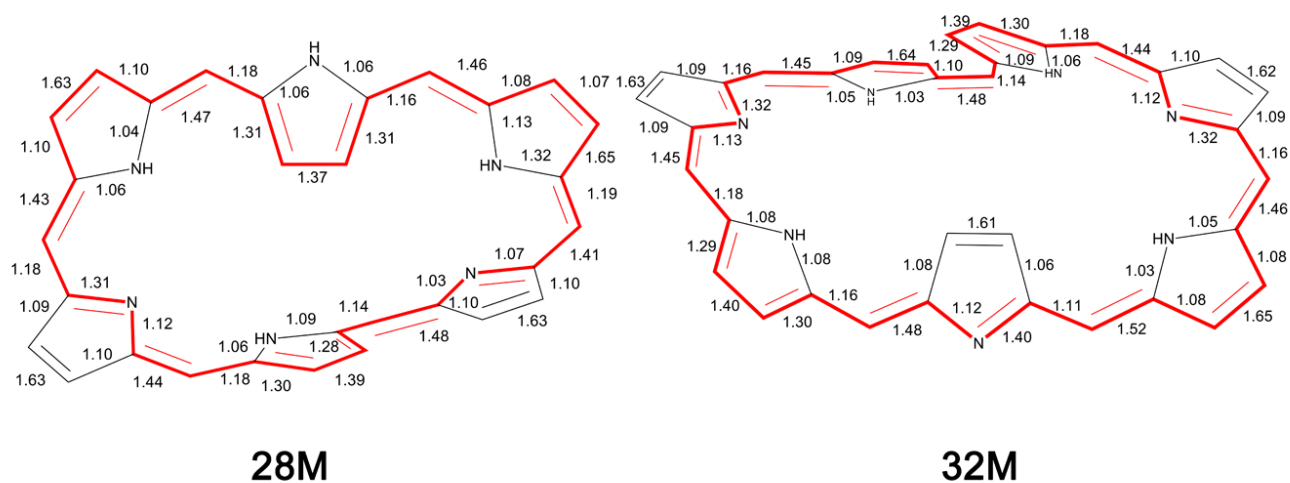


Figure S14: Delocalization indices (DIs) for $28M$ and $32M$ macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold red bonds.

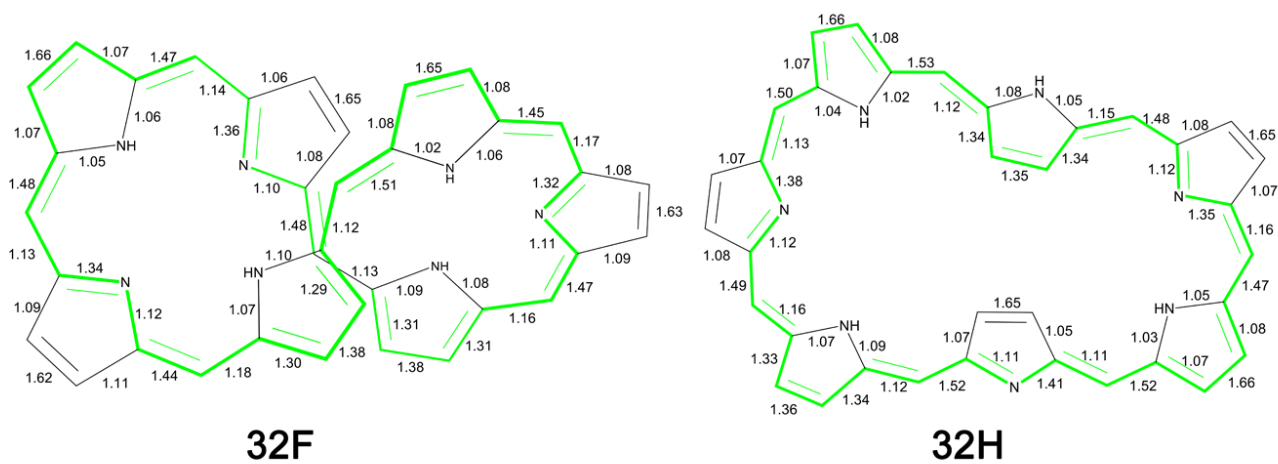


Figure S15: Delocalization indices (DIs) for **32F** and **32H** macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold bonds.

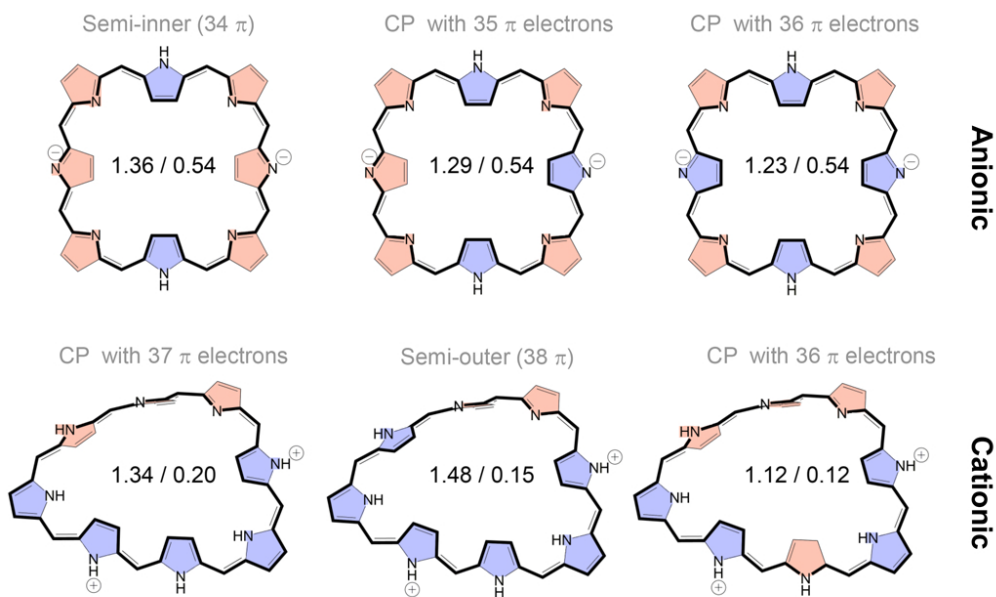


Figure S16: Illustration of the most delocalized pathways in $\mathbf{36H}_a^{2-}$ and $\mathbf{36M}_b^{2+}$.

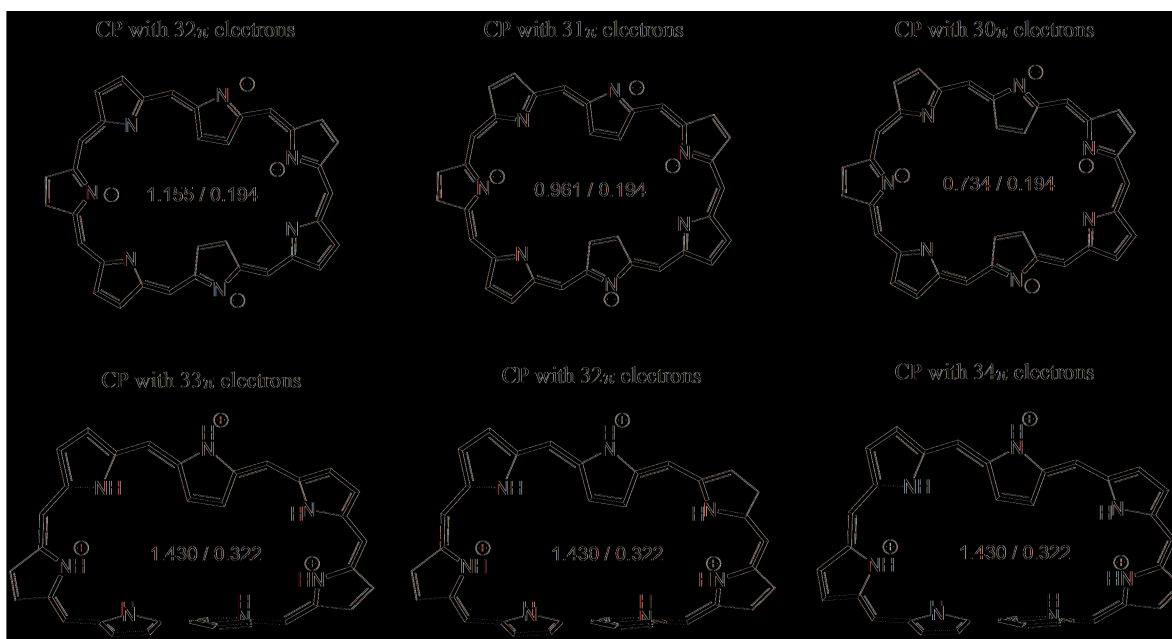


Figure S17: Illustration of the most delocalized pathways in 32H^{4-} and 32M^{3+} .

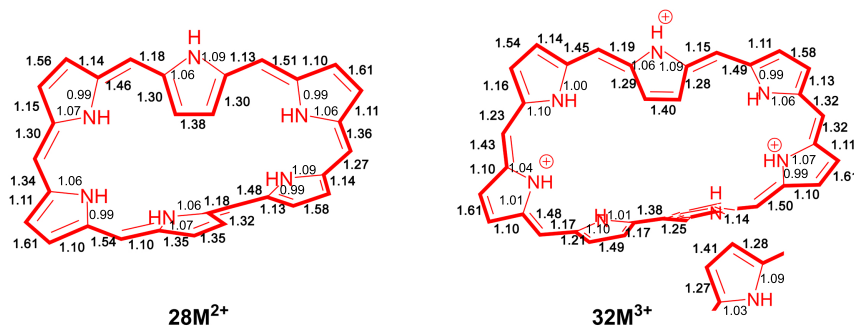


Figure S18: Delocalization indices (DIs) for 28M^{2+} and 32M^{3+} macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold bonds.

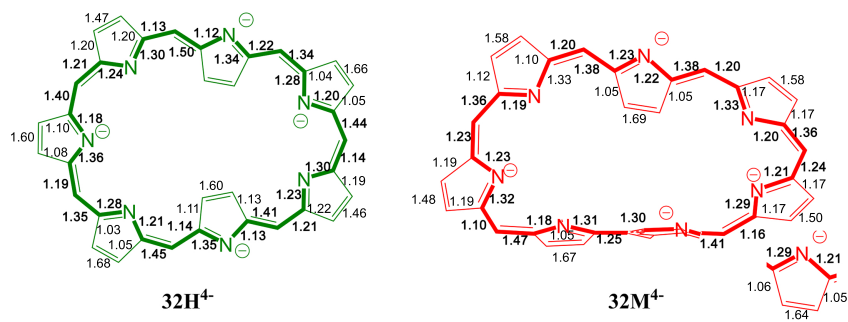


Figure S19: Delocalization indices (DIs) for 32H^{4-} and 32M^{4-} macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold bonds.

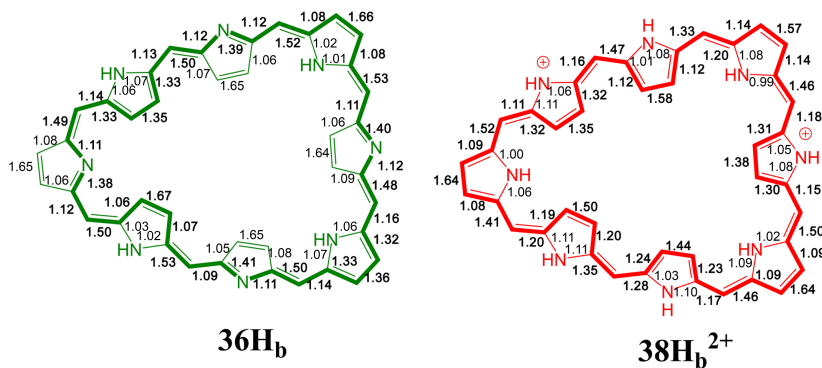


Figure S20: Delocalization indices (DIs) for **36H_b** and **38H_b²⁺** macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold bonds.

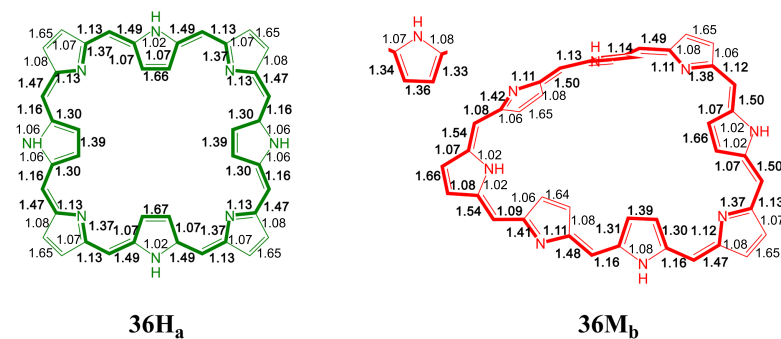


Figure S21: Delocalization indices (DIs) for **36H_a** and **36M_b** macrocycles computed at the CAM-B3LYP/6-311G(d,p) level of theory. The main conjugation pathway is depicted with bold bonds.