

Supporting Information (32 pages)

for

**A Theoretical Study on the Nature of On- and Off-States of Reversibly
Photoswitching Fluorescent Protein Dronpa: Absorption, Emission,
Protonation and Raman**

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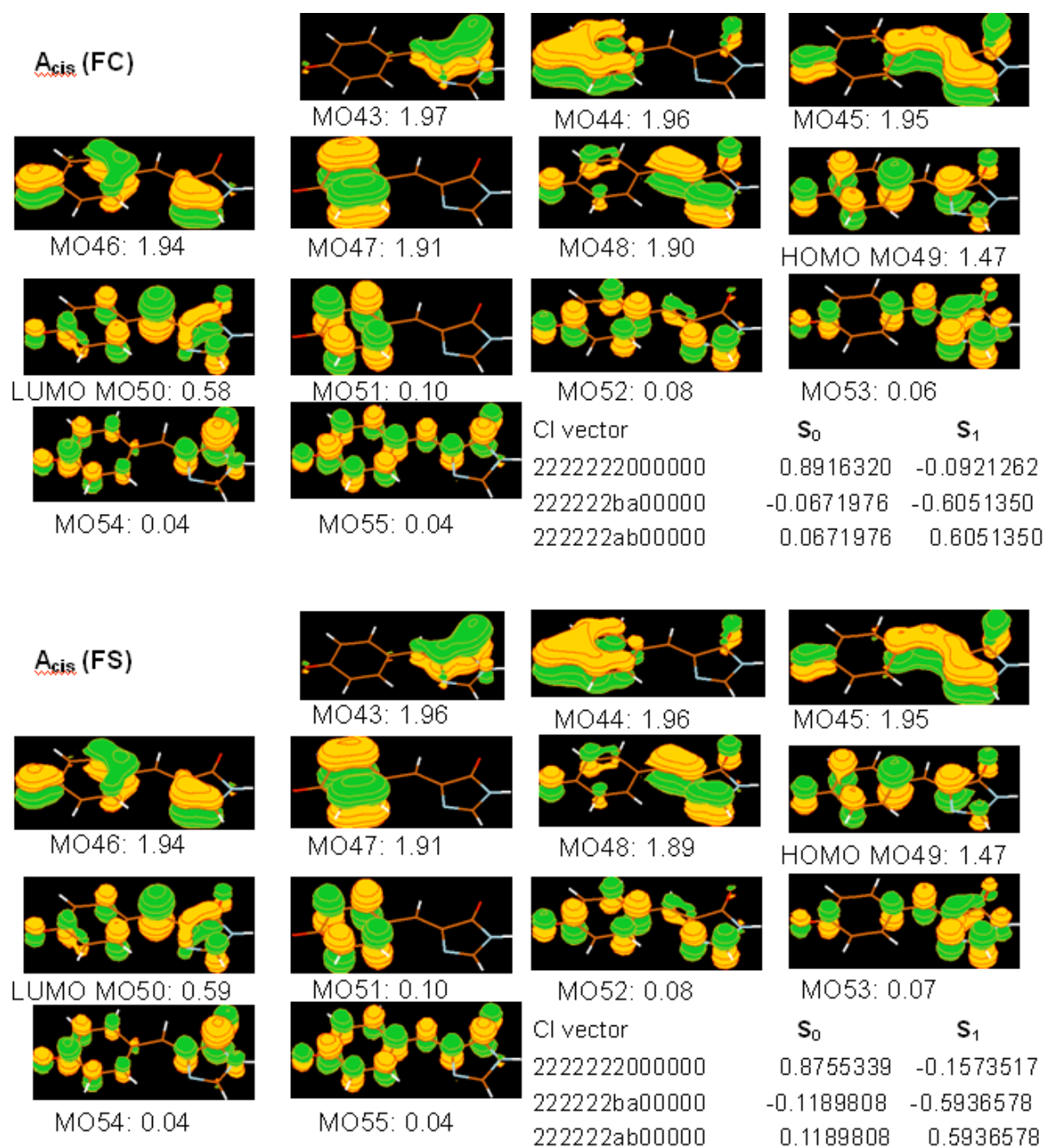


Figure S1. The active space of the ground state (FC) and excited state (FS) **A_{cis}** for the SA2-CASSCF(14e,13o)/6-31G(d) calculation. The occupation numbers and major electronic configurations are also included. In this Supporting Information, **FC** (Frank-Condon) and **FS** (fluorescence state) were designated as the nearly planar ground-state and excited-state minimum, respectively.

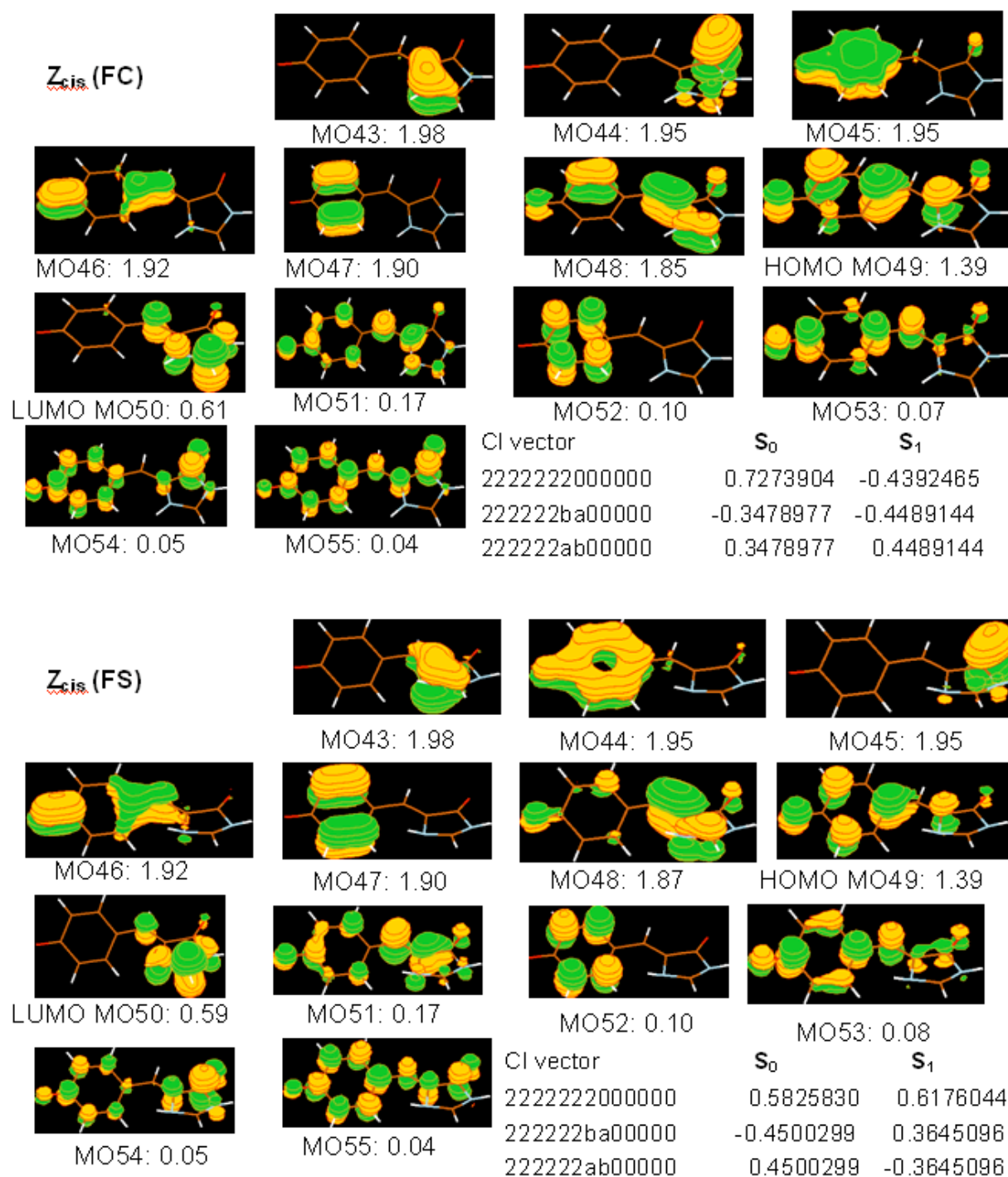


Figure S2. The active space of the ground state (FC) and excited state (FS) **Z_{cis}** for the SA2-CASSCF(14e,13o)/6-31G(d) calculation. The occupation numbers and major electronic configurations are also included.

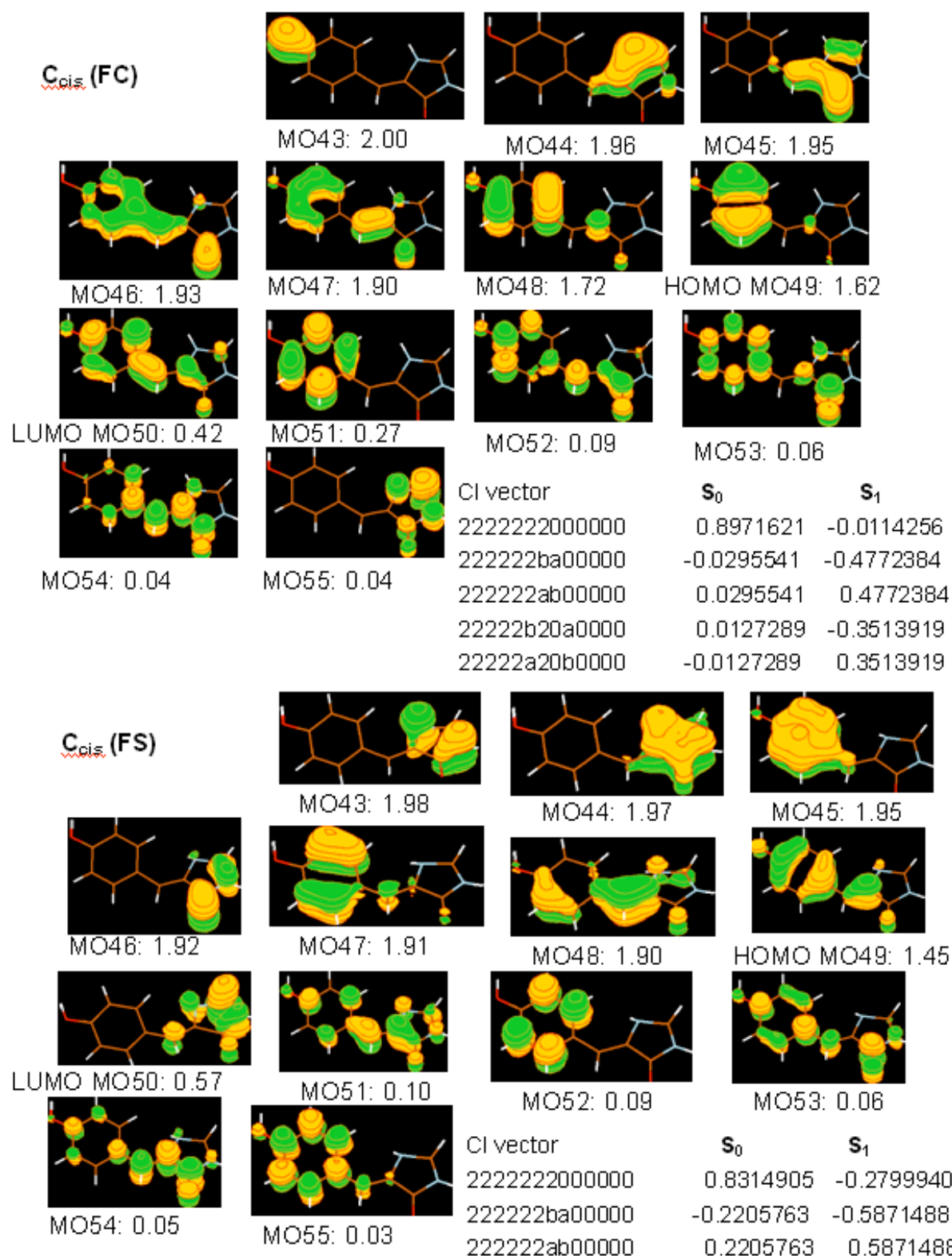


Figure S3. The active space of the ground state (FC) and excited state (FS) C_{cis} for the SA2-CASSCF(14e,13o)/6-31G(d) calculation. The occupation numbers and major electronic configurations are also included.

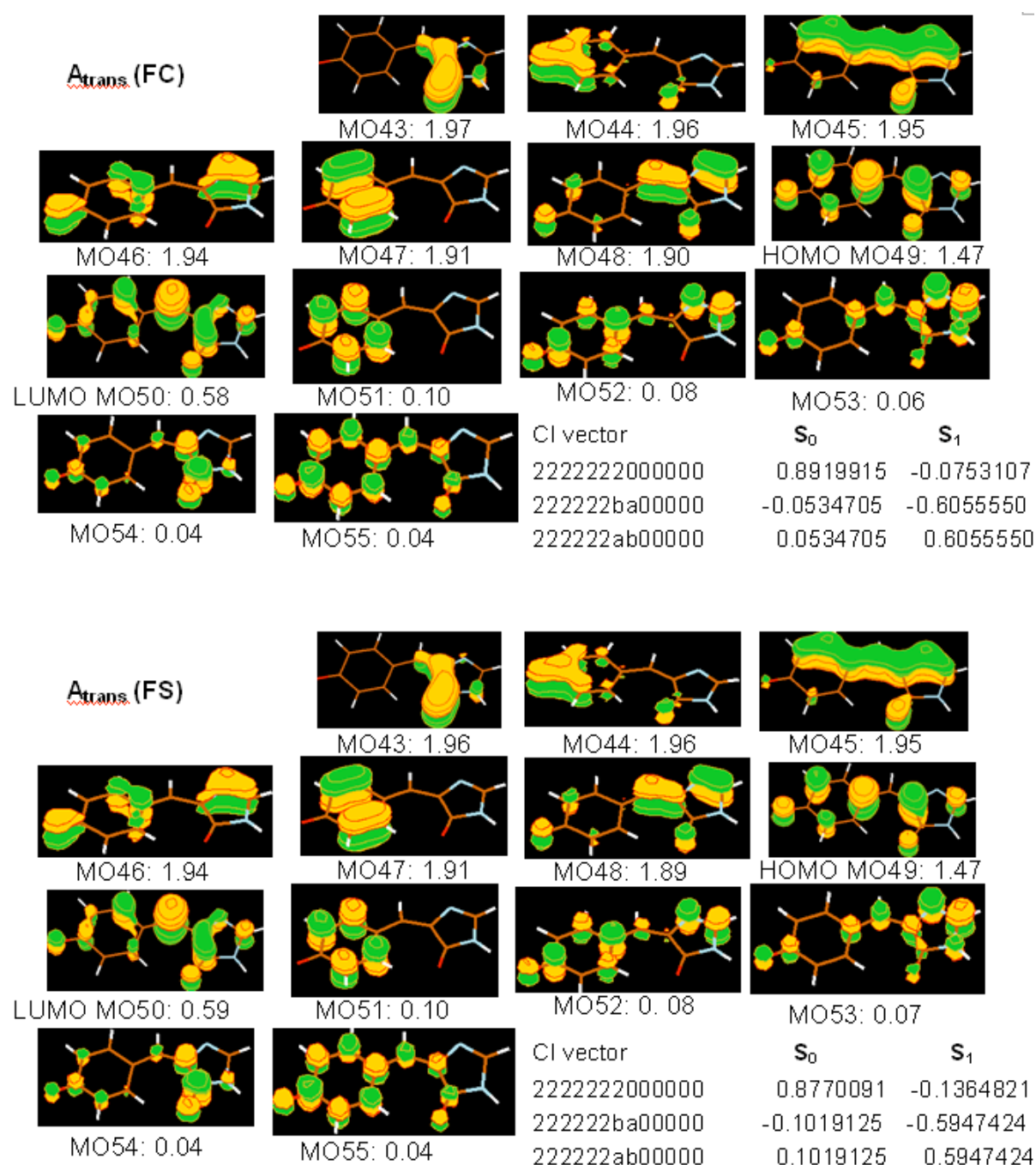


Figure S4. The active space of the ground state (FC) and excited state (FS) **A_{trans}** for the SA2-CASSCF(14e,13o)/6-31G(d) calculation. The occupation numbers and major electronic configurations are also included.

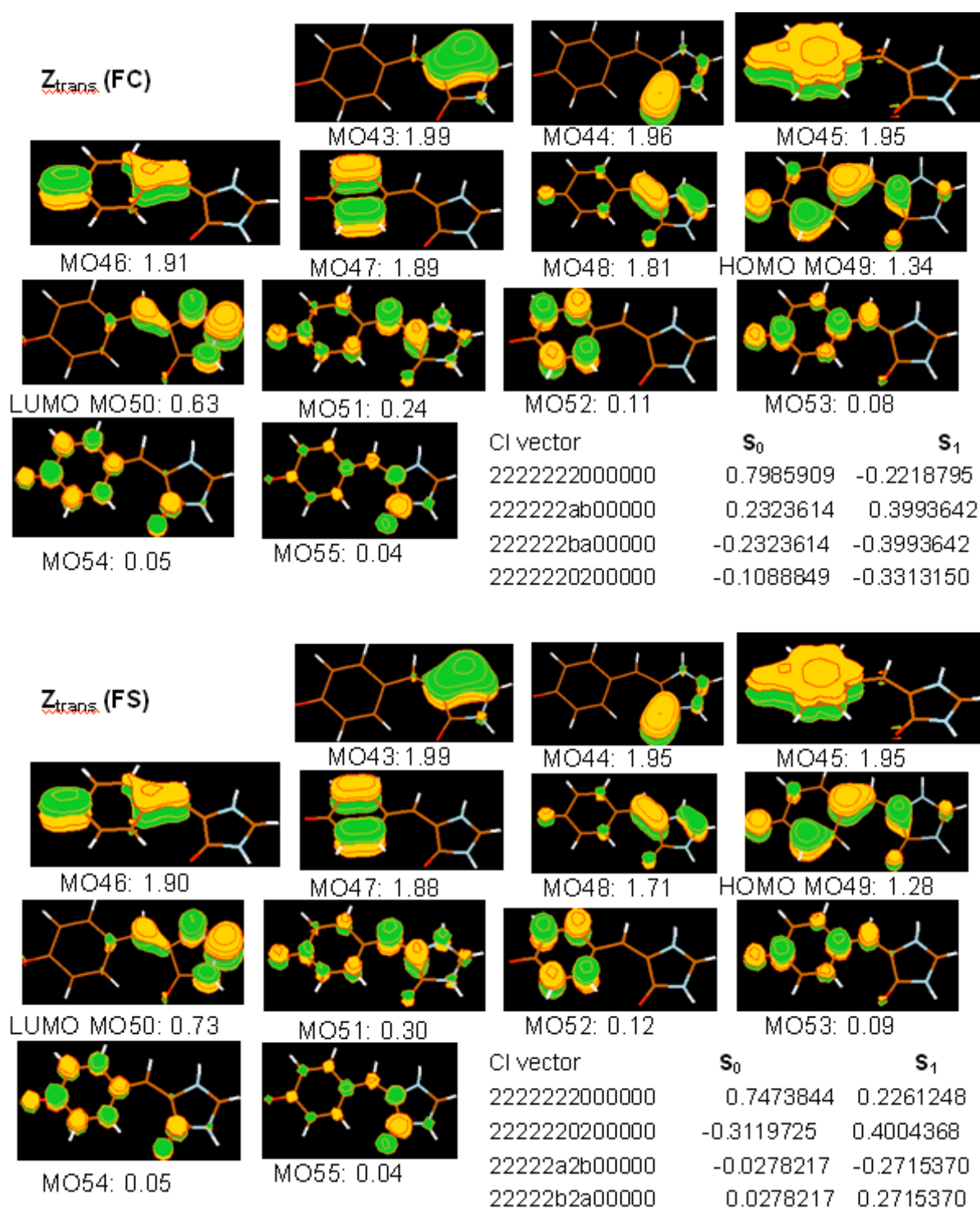


Figure S5. The active space of the ground state (FC) and excited state (FS) Z_{trans} for the SA2-CASSCF(14e,13o)/6-31G(d) calculation. The occupation numbers and major electronic configurations are also included.

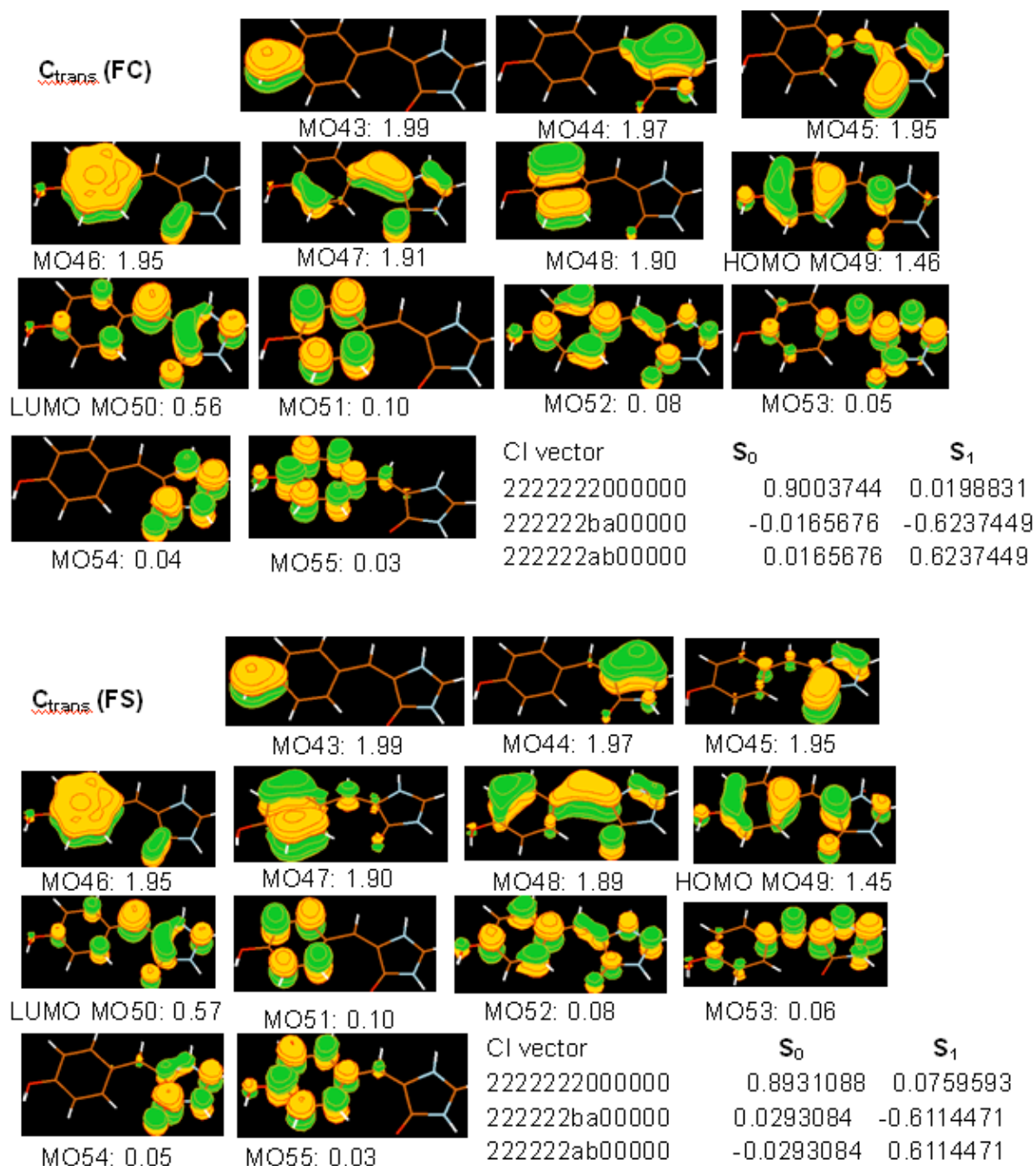


Figure S6. The active space of the ground state (FC) and excited state (FS) C_{trans} for the SA2-CASSCF(14e,13o)/6-31G(d) calculation. The occupation numbers and major electronic configurations are also included.

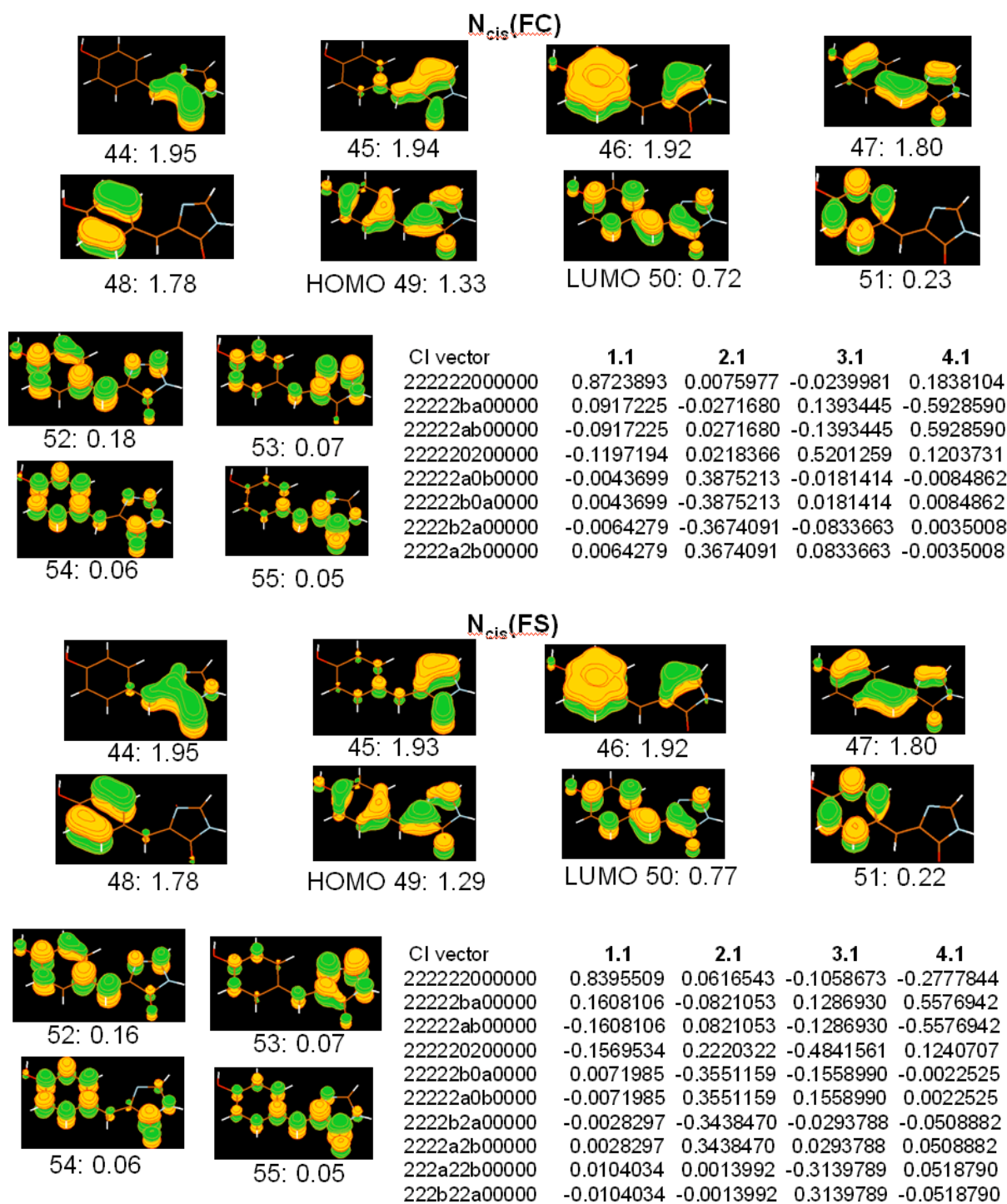


Figure S7. The active space of the ground state (FC) and excited state (FS) N_{cis} for the SA4-CASSCF(12e,12o)/6-31G(d) calculation. The ground and excited-state chromophore are optimized by SA2-CASSCF(12e,12o)/6-31G(d) and SA4-CASSCF(12e,12o)/6-31G(d) levels, respectively. The occupation numbers and major electronic configurations are also included.

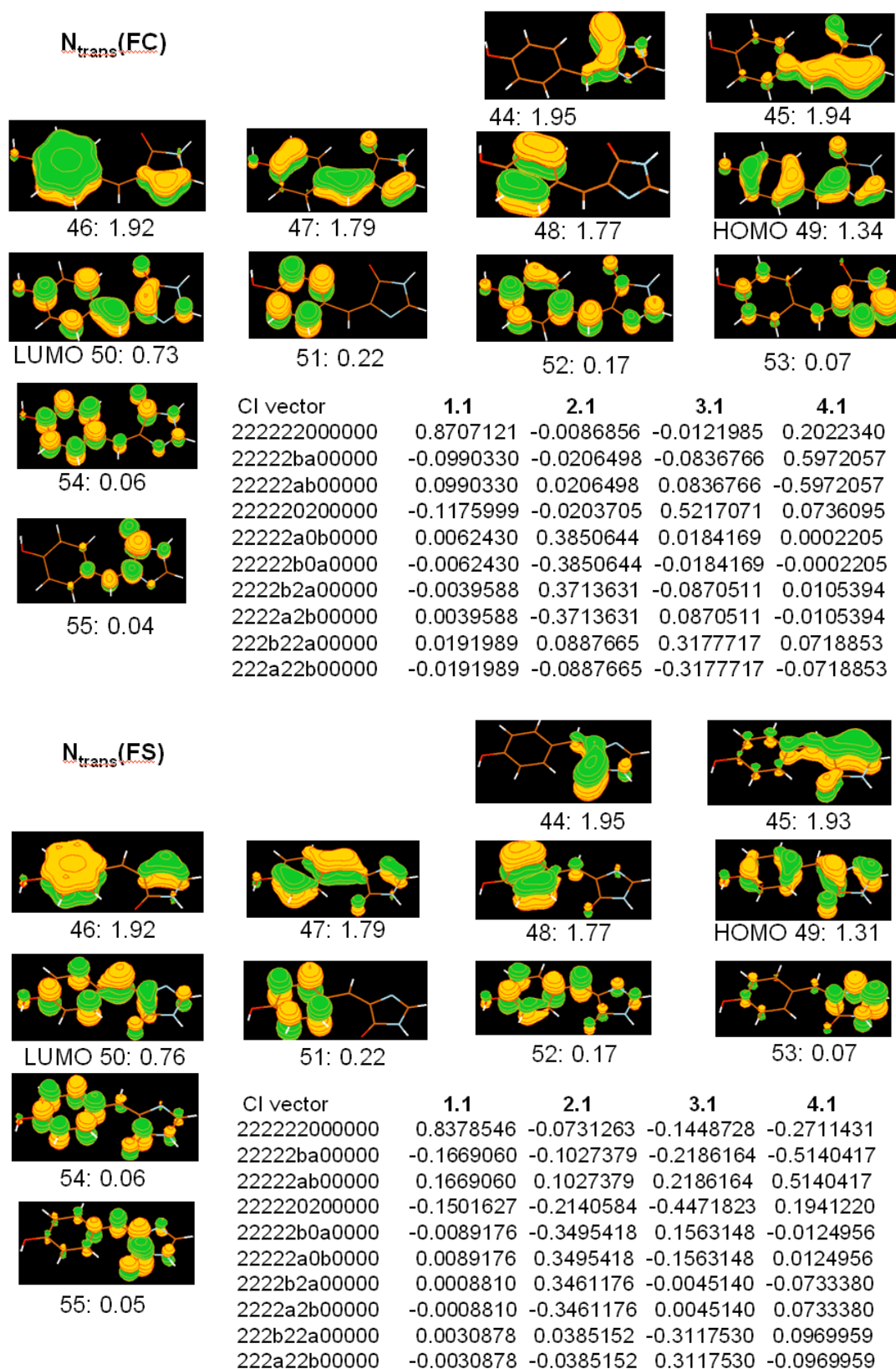


Figure S8. The active space of the ground state (FC) and excited state (FS) N_{trans} for the SA4-CASSCF(12e,12o)/6-31G(d) calculation. The ground and excited-state chromophore are optimized by SA2-CASSCF(12e,12o)/6-31G(d) and SA4-CASSCF(12e,12o)/6-31G(d) levels, respectively. The occupation numbers and major electronic configurations are also included.

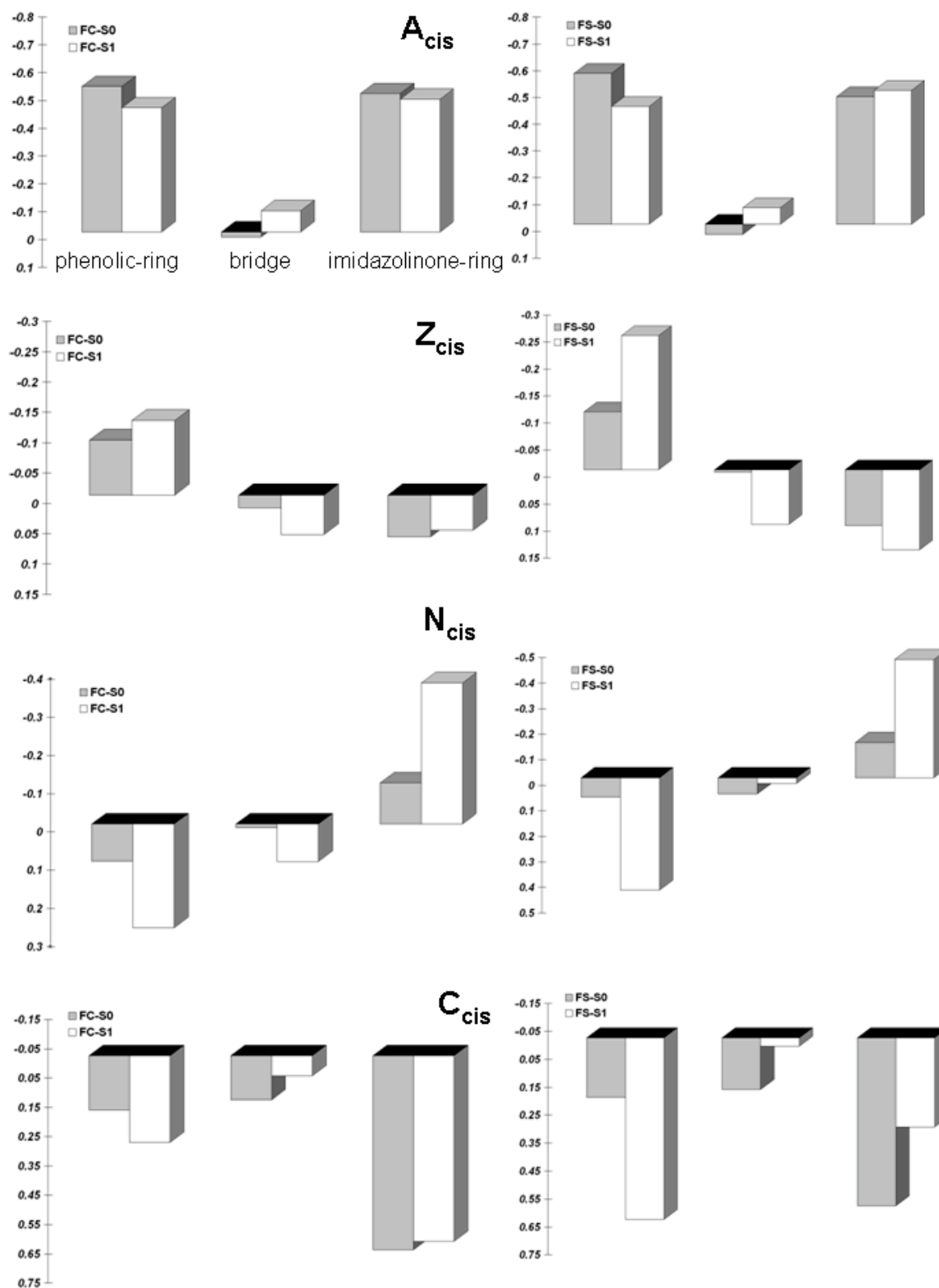


Figure S9. The S₀ and S₁ Mulliken charge analysis at FS and FC of the *cis* chromophores in four protonation states calculated by the SA2-CASSCF(14e,13o)/6-31G(d) method, except for N_{cis} by SA4-CASSCF(12e,12o)/6-31G(d).

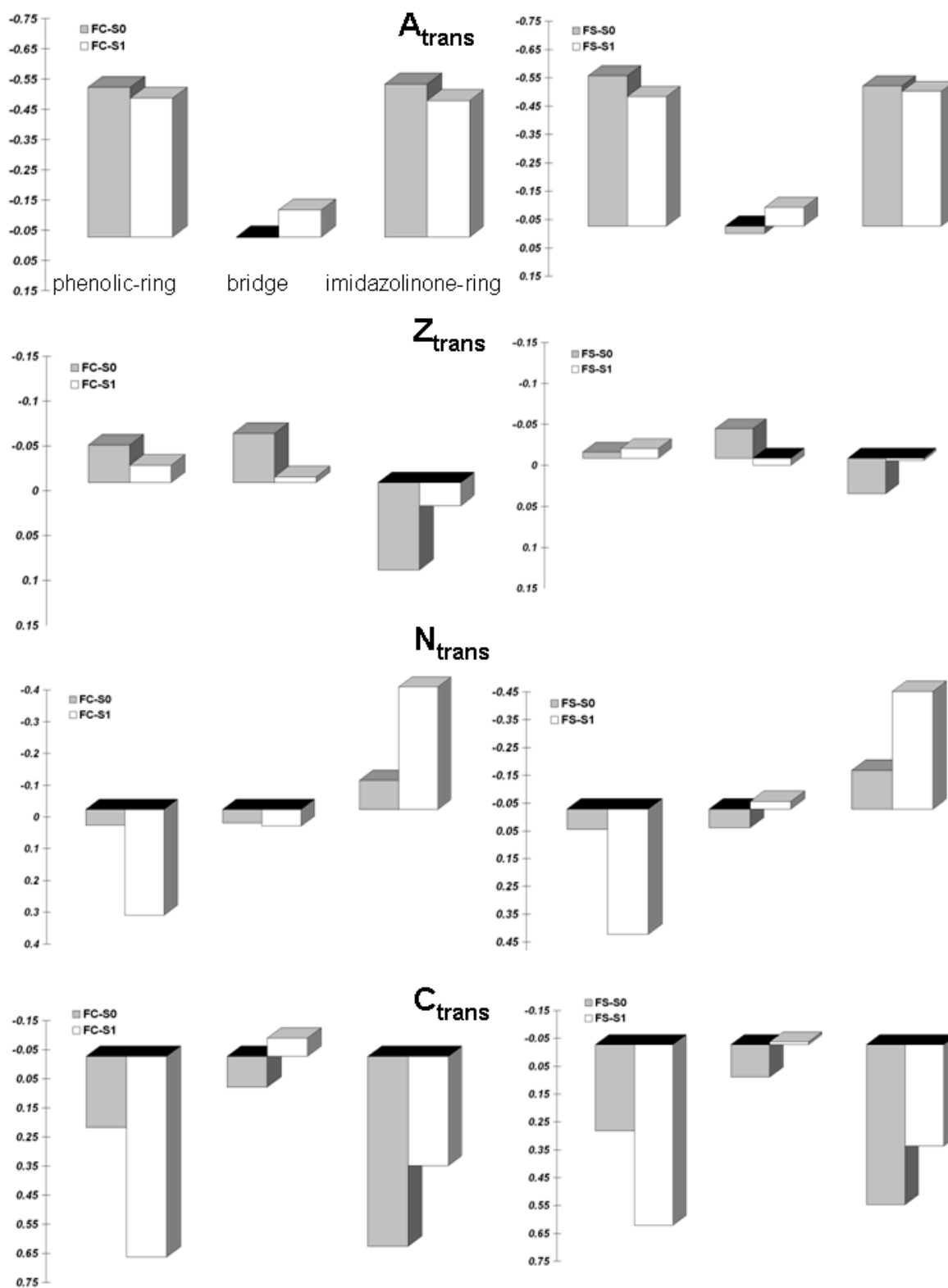


Figure S10. The S_0 and S_1 Mulliken charge analysis at **FS** and **FC** of the *trans* chromophores in four protonation states calculated by the SA2-CASSCF(14e,13o)/6-31G(d) method, except for N_{trans} by SA4-CASSCF(12e,12o)/6-31G(d).

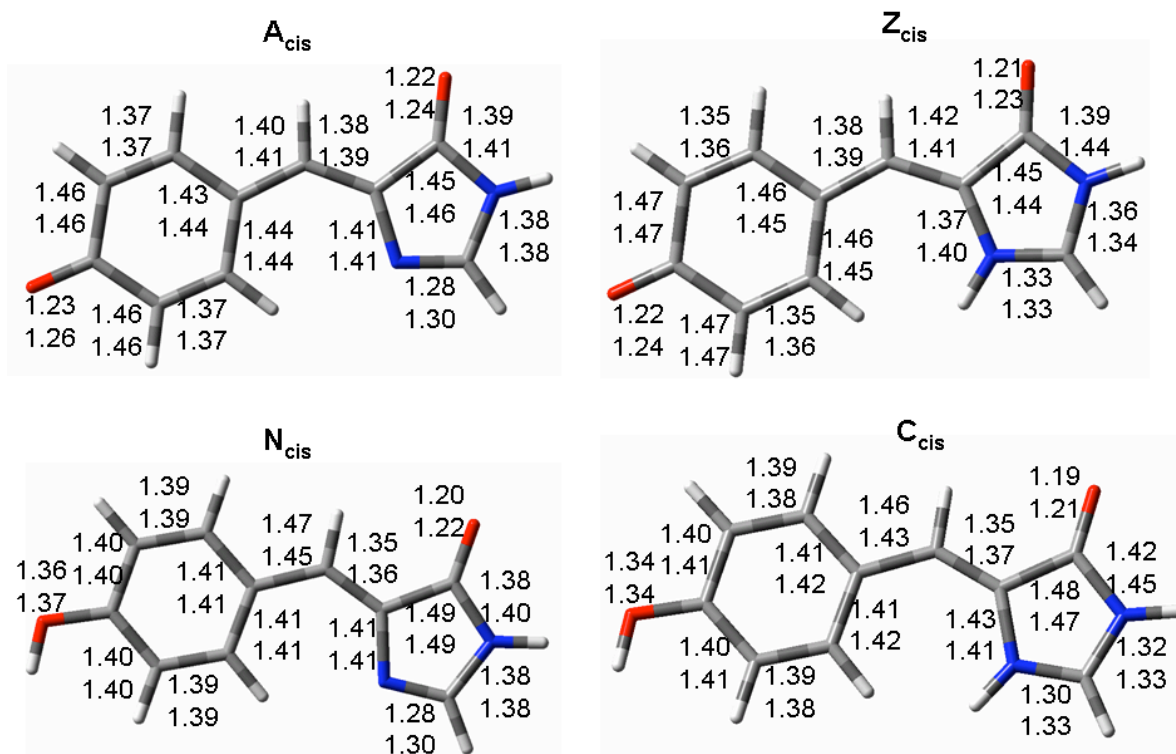


Figure S11. The bond lengths of the *cis* conformers at FC optimized by the SA2-CASSCF(14e,13o)/6-31G(d) (upper) and B3LYP/6-31+G(d,p) methods (lower).

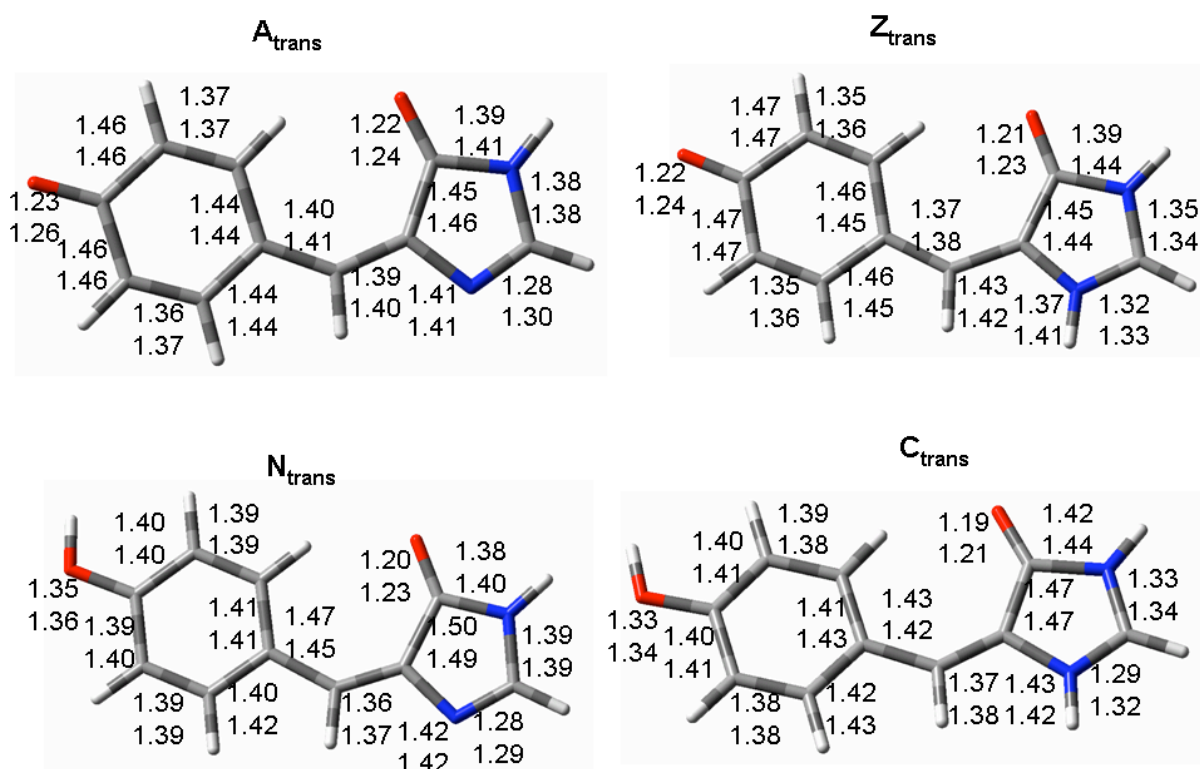


Figure S12. The bond lengths of the *trans* conformers at FC optimized by the SA2-CASSCF(14e,13o)/6-31G(d) (upper) and B3LYP/6-31+G(d,p) methods (lower).

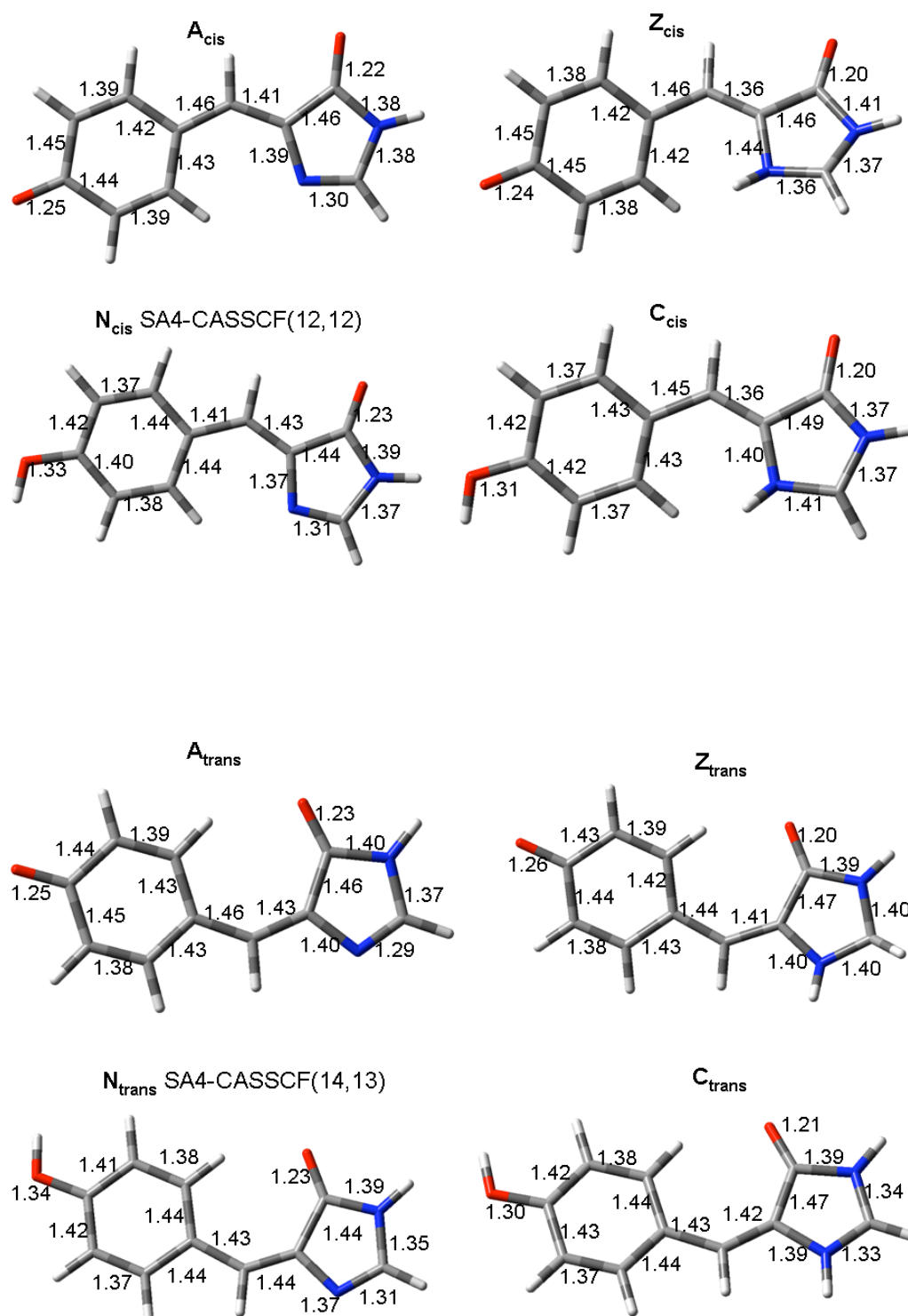


Figure S13. The bond lengths of the *cis* and *trans* conformers at FS optimized by the SA2-CASSCF(14e,13o)/6-31G(d) method. The N_{cis} and N_{trans} were optimized by the SA4-CASSCF(12e,12o)/6-31G(d) and SA4-CASSCF(14e,13o)/6-31G(d) methods, respectively.

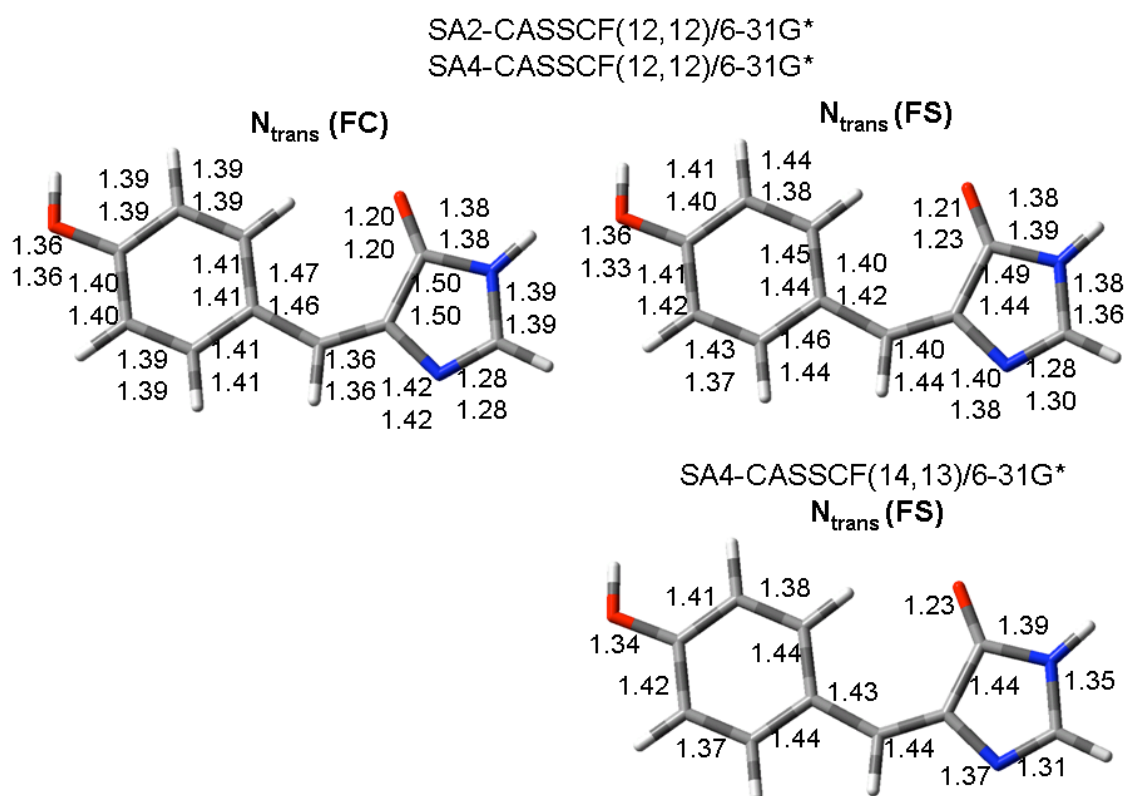


Figure S14. The bond lengths of N_{trans} at FC and FS optimized by the SA2-CASSCF(12e,12o)/6-31G(d), SA4-CASSCF(12e,12o)/6-31G(d) and SA4-CASSCF(14e, 13o)/6-31G(d) methods, respectively.

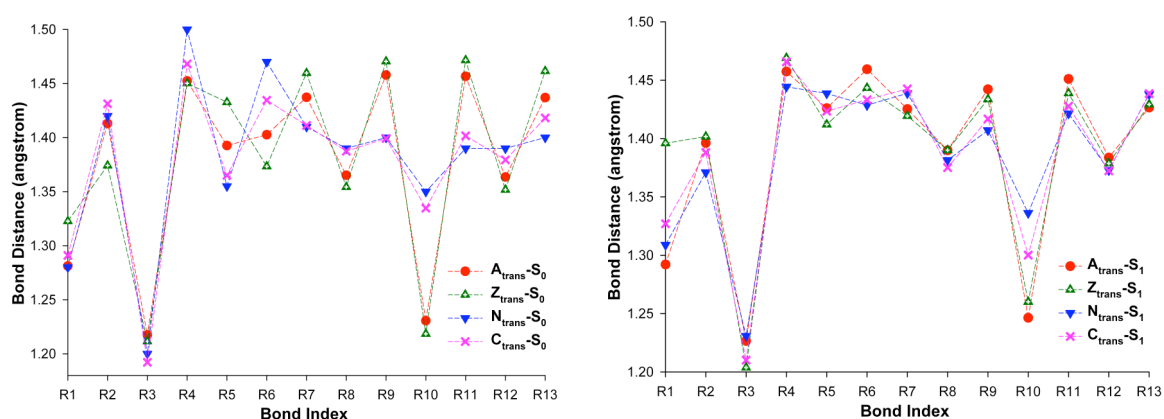


Figure S15. The key bond lengths of the ground-state (left) and excited-state (right) *trans* chromophore at the different protonation states in the gas phase calculated by the SA2-CASSCF(14e,13o)/6-31G(d) method, except for the excited-state N_{trans} by SA4-CASSCF(14e,13o)/6-31G(d) method.

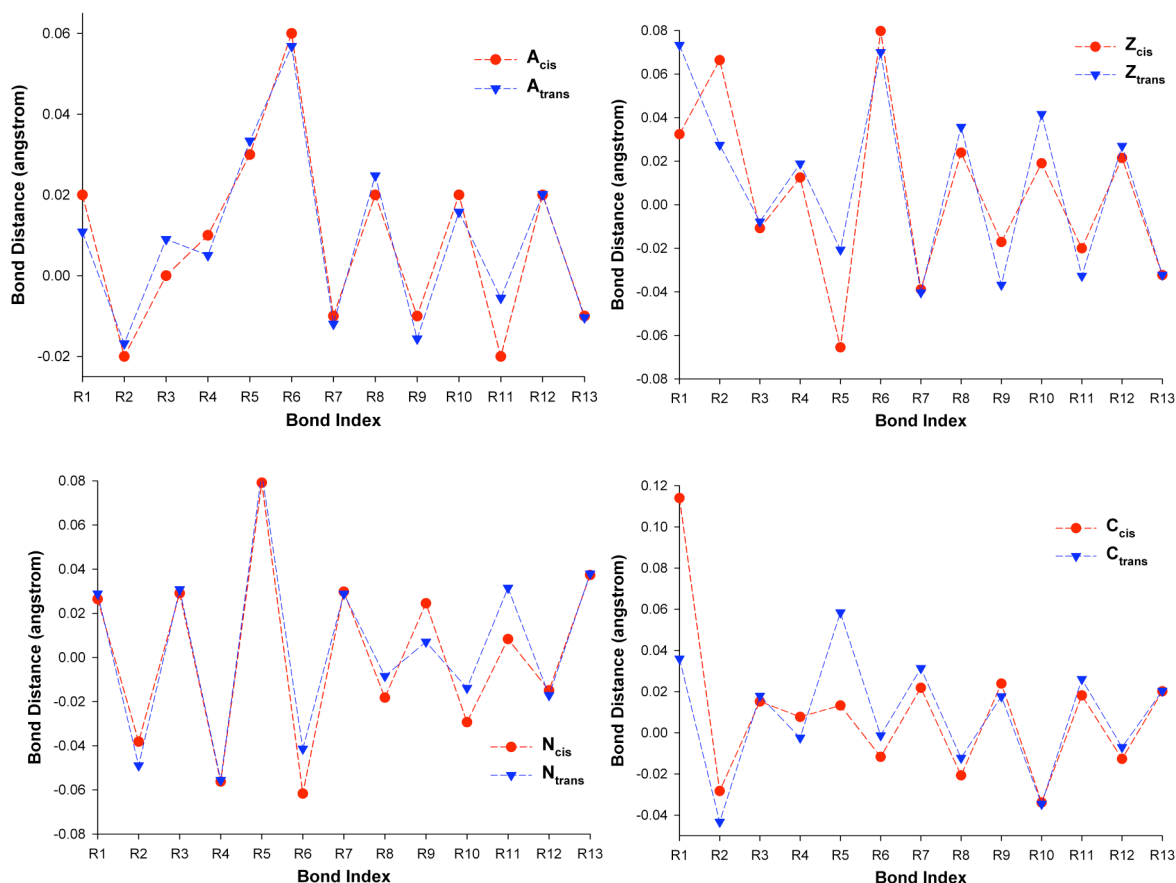


Figure S16. The bond length differences between the excited state and ground state chromophores in the gas phase calculated by the SA2-CASSCF(14e,13o)/6-31G(d) method, except for the excited-state N_{cis} and N_{trans} by SA4-CASSCF(12e,12o)/6-31G(d) and SA4-CASSCF(14e,13o)/6-31G(d) methods, respectively.

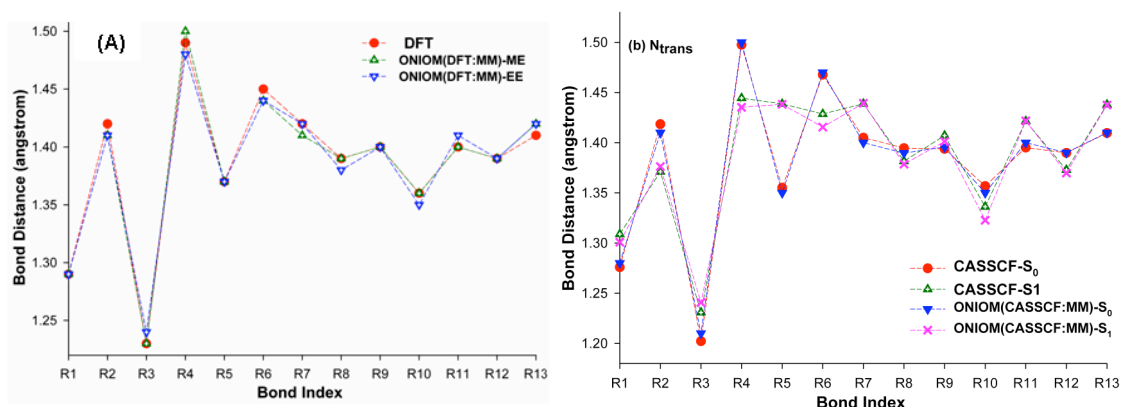


Figure S17. Calculated key bond lengths of N_{trans} : (A) the ground state chromophore in the gas phase by the B3LYP/6-31+G(d,p) method and in the protein by the ONIOM(B3LYP/6-31+G(d,p):AMBER)-ME and -EE methods. (B) the ground state (S_0) and excited state (S_1) chromophore in the gas phase by the state-average (SA) CASSCF(12e,12o)/6-31G(d) method and in the protein by the ONIOM(SA-CASSCF(12e,12o)/6-31G(d):AMBER)-ME method. Two-state-average (SA2) for the ground state and four-state-average (SA4) for the excited state.

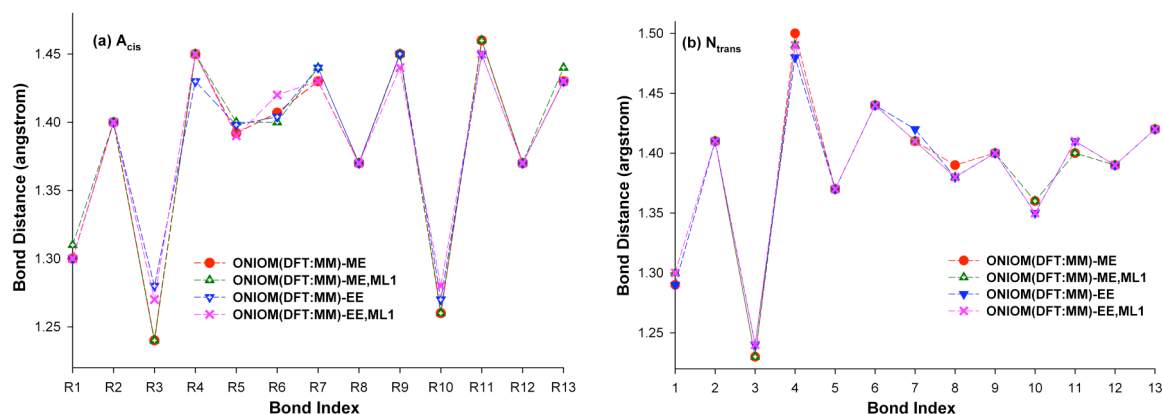


Figure S18. The key bond lengths of the ground state chromophores in the proteins calculated by the ONIOM(B3LYP/6-31+G(d,p):AMBER)-EE and -ME methods for both small QM model and larger QM model.

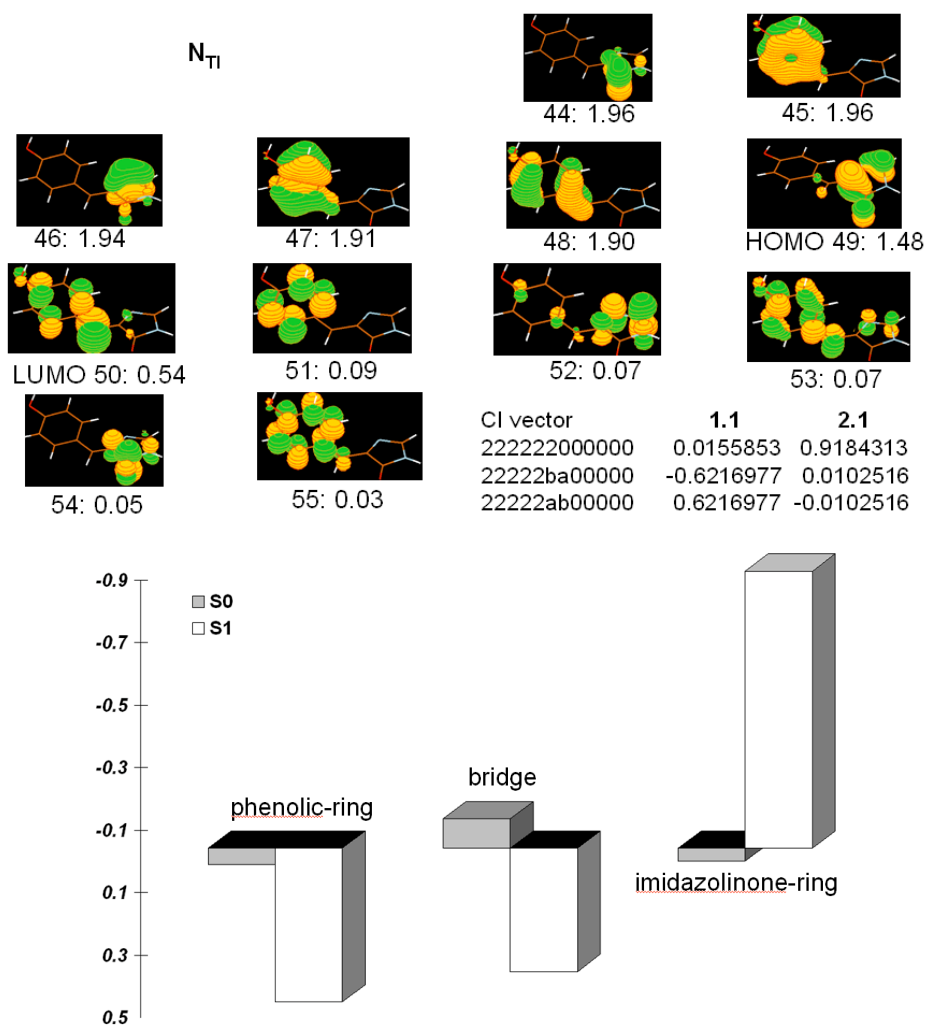


Figure S19. The active space of N_{Ti} in the SA2-CASSCF(12e,12o)/6-31G(d) calculation (top). The occupation numbers and major electronic configurations are also included. The Mulliken charge analysis at N_{Ti} by the SA2-CASSCF(12e,12o)/6-31G(d) method (bottom).

Raman Spectra: As shown in Figures S11-S12, the chromophore skeleton is partly altered by the protonation state of the chromophore. Such structural changes should be reflected by IR and Raman spectrum. For the simple illustration, the calculated Raman spectra for **A_{cis}**, **Z_{cis}** and **N_{trans}** in the gas-phase are showed in Figure S20. The experimental Raman spectra of the anionic and neutral GFP analogues in solution^{1,2} show different Raman markers for the different protonated states, which provide a clear signature in the double bond stretching region between 1500 and 1700 cm⁻¹. The Raman spectra calculated in the gas-phase show different dominant bands for **A_{cis}** (1592.8 cm⁻¹), **N_{trans}** (1593.3 cm⁻¹) and **Z_{cis}** (1616.9 cm⁻¹). **Z_{cis}** has a higher wave number for the marker band of the C=O, C=C and C=N stretching mode than **A_{cis}**. Therefore, the Raman spectra can also help assign the protonation state of Dronpa and other fluorescence proteins, such as asFP595.³

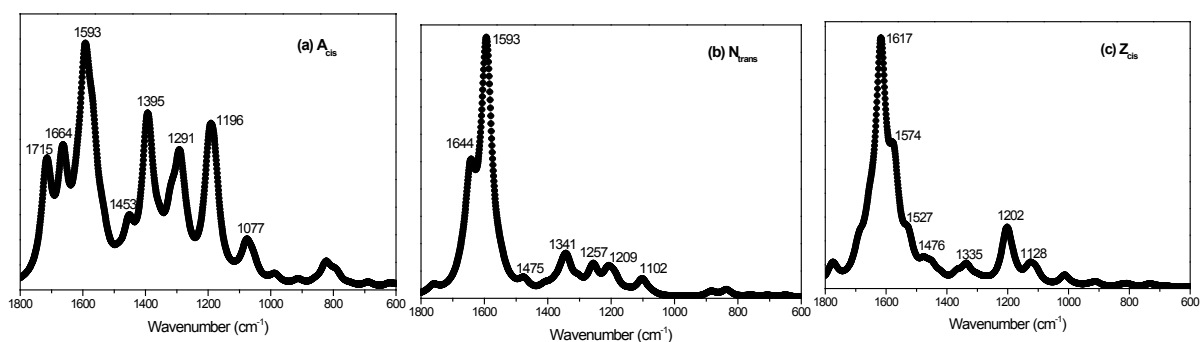


Figure S20. Raman spectra for (a) **A_{cis}**, (b) **N_{trans}** and (c) **Z_{cis}** in the gas phase calculated by the B3LYP/6-31+G(d,p) method.

¹ (b) Altoe, P.; Bernardi, F.; Garavelli, M.; Orlandi, G.; Negri, F. *J. Am. Chem. Soc.* **2005**, *127*, 3952.

² Bell, A. F.; He, X.; Wachter, R. M.; Tonge, P. J. *Biochemistry* **2000**, *39*, 4423.

³ During our manuscript preparation, the preresonant spectroscopy and model calculations on the GFP and its variants have also recently been employed to study photochromic fluorescent proteins. The Raman spectra shows *cis-trans* isomerization and a change of protonation state of the chromophore in the studied fluorescent proteins: Luin, S.; Voliani, V.; Lanza, G.; Bizzarri, R.; Nifosi, R.; Amat, P.; Tozzini, V.; Serresi, M.; Beltram, F. *J. Am. Chem. Soc.* **2009**, *131*, 96.

Table S1. Absorption and emission energies (in eV) of the chromophore evaluated by the CASPT2/6-31G(d)//CASSCF/6-31G(d) method with the smaller active space [(12e,11o) or (12e,12o)]. Two- and three-state average approaches (SA2 and SA3, respectively) were used.

Method	<i>Cis</i>		<i>Trans</i>	
	CASPT2// SA2-CASSCF	CASPT2// SA3-CASSCF	CASPT2// SA2-CASSCF	CASPT2// SA3-CASSCF
Absorption				
Anionic	2.72	2.72	2.73	2.72
Zwitterionic	3.12	3.05	3.50	3.44
Neutral	4.28	4.24	4.16	4.24
Cationic	3.14	- ^a	2.91	2.95
Emission				
Anionic	2.48	2.46	2.50	2.48
Zwitterionic	-	-	2.17	2.14
Neutral	3.77	3.74	3.75	3.73
Cationic	2.11	2.14	2.59	2.66

^aThe CASPT2 calculation failed to converge.

Table S2. Absorption and emission energies of the different forms of the chromophore calculated by the SAC-CI(Level2)/D95(d)//SA2-CASSCF/6-31G(d) method with the small active space [(12e,11o) or (12e,12o)].

Method	<i>Cis</i>	<i>Trans</i>
	SAC-CI// SA2-CASSCF	SAC-CI// SA2-CASSCF
Absorption		
Anionic	2.55	2.56
Zwitterionic	2.49	2.78
Neutral	3.67	3.63
Cationic	3.00	2.75
Emission		
Anionic	2.31	2.29
Zwitterionic	-	2.25
Neutral	3.27	3.18
Cationic	1.27	2.31

Table S3(a). Absorption and emission energies of the chromophore in the gas phase calculated by SA4-CASSCF, SS-CASPT2 and MS-CASPT2 with ANO-S basis sets and IPEA zero-order Hamiltonian.

Method	SA4-CASSCF	SS-CASPT2	MS-CASPT2
N_{cis} (absorption)^a			
S ₁	4.56 ^e /4.56 ^f	4.53 ^e /4.52 ^f	4.07 ^e /3.90 ^f
S ₂	5.42 ^e /5.28 ^f	5.08 ^e /4.94 ^f	4.60 ^e /4.55 ^f
S ₃	5.78 ^e /5.56 ^f	4.18 ^e /3.90 ^f	5.33 ^e /4.99 ^f
N_{cis} (emission)^{b,e}			
S ₁	4.18	4.15	3.32
S ₂	4.30	4.25	4.31
S ₃	4.87	3.42	4.46
N_{trans} (absorption)^a			
S ₁	4.55 ^e /4.56 ^f /4.54 ^{b,e}	4.53 ^e /4.52 ^f /4.51 ^{b,e}	4.10 ^e /3.77 ^f /4.09 ^{b,e}
S ₂	5.43 ^e /5.26 ^f /5.39 ^{b,e}	5.16 ^e /4.82 ^f /5.16 ^{b,e}	4.59 ^e /4.54 ^f /4.58 ^{b,e}
S ₃	5.69 ^e /5.51 ^f /5.66 ^{b,e}	4.06 ^e /3.99 ^f /4.02 ^{b,e}	5.29 ^e /5.09 ^f /5.25 ^{b,e}
N_{trans} (emission)			
S ₁	4.20 ^{b,e} /4.07 ^{d,f}	4.15 ^{b,e} /3.72 ^{d,f}	3.15 ^{b,e} /2.97 ^{d,f}
S ₂	4.30 ^{b,e} /4.23 ^{d,f}	4.13 ^{b,e} /4.22 ^{d,f}	4.35 ^{b,e} /4.28 ^{d,f}
S ₃	4.76 ^{b,e} /4.69 ^{d,f}	3.50 ^{b,e} /3.57 ^{d,f}	4.60 ^{b,e} /4.45 ^{d,f}
Z_{cis} (absorption)^{c,f}			
S ₁	3.33	3.17	2.88
S ₂	3.86	3.29	4.16
S ₃	4.48	4.49	4.71

^aGeometry optimized by SA2-CASSCF(12e,12o)/6-31G(d). ^bGeometry optimized by SA4-CASSCF(12e,12o)/6-31G(d).

^cGeometry optimized by SA2-CASSCF(14e,13o)/6-31G(d). ^dGeometry optimized by SA4-CASSCF(14e,13o)/6-31G(d).

^e(12e,12o) was used in these calculations. ^f(14e,13o) was used in these calculations.

Table S3(b). Absorption and emission energies of the chromophore in the gas phase

calculated by SS-CASPT2 and MS-CASPT2 with ANO-S basis sets and the original zero-order Hamiltonian.

Method	SS-CASPT2	MS-CASPT2
N_{cis} (absorption)^a		
S ₁	4.14 ^e /4.13 ^f	3.62 ^e /3.49 ^f
S ₂	4.65 ^e /4.47 ^f	4.22 ^e /4.18 ^f
S ₃	3.77 ^e /3.49 ^f	4.97 ^e /4.52 ^f
N_{cis} (emission)^{b,e}		
S ₁	3.72	2.98
S ₂	3.85	3.94
S ₃	3.05	4.11
N_{trans} (absorption)^a		
S ₁	4.13 ^e /4.13 ^f /4.12 ^{b,e}	3.69 ^e /3.31 ^f /3.70 ^{b,e}
S ₂	4.72 ^e /4.33 ^f /4.72 ^{b,e}	4.21 ^e /4.17 ^f /4.20 ^{b,e}
S ₃	3.66 ^e /3.57 ^f /3.62 ^{b,e}	4.89 ^e /4.68 ^f /4.84 ^{b,e}
N_{trans} (emission)		
S ₁	3.72 ^{b,e} /3.34 ^{d,f}	2.78 ^{b,e} /2.54 ^{d,f}
S ₂	3.73 ^{b,e} /3.81 ^{d,f}	3.99 ^{b,e} /3.91 ^{d,f}
S ₃	3.11 ^{b,e} /3.16 ^{d,f}	4.31 ^{b,e} /4.17 ^{d,f}

^aGeometry optimized by SA2-CASSCF(12e,12o)/6-31G(d). ^bGeometry optimized by SA4-CASSCF(12e,12o)/6-31G(d).

^cGeometry optimized by SA2-CASSCF(14e,13o)/6-31G(d). ^dGeometry optimized by SA4-CASSCF(14e,13o)/6-31G(d).

^e(12e,12o) was used in these calculations. ^f(14e,13o) was used in these calculations.

Table S4. The calculated absorption energies (in eV) and oscillator strengths (in the parentheses) of the chromophore in the proteins by ONIOM(TD-B3LYP/6-31++G(d,p):AMBER)-EE at ONIOM(B3LYP/6-31+G(d,p):AMBER)-EE optimized structures.

Conformation in proteins	Cis (On-state)		Trans (Off-state)	
QM model	MS	ML1	MS	ML1
Anionic, A	3.11(0.84)	3.04(0.98)	3.05(0.68)	2.98(0.81)
Zwitterionic, Z	3.09(0.79)	2.96(0.95)	3.20(0.85)	3.03(1.03)
Neutral, N	3.39(0.71)	3.30(0.37) ^a	3.33(0.53)	3.17(0.27) ^b
Cationic, C	3.18(0.73)	3.26(0.54)	3.17(0.69)	3.11(0.81)

^aThe vertical absorption energy and oscillator strength for another excitation (HOMO-2 → LUMO) is 3.55 eV and 0.33, respectively. ^bThe vertical absorption energy and oscillator strength for another excitation (HOMO-1 → LUMO) is 3.40 eV and 0.36, respectively.

Table S5(a). Absorption and emission energies of the chromophore in the proteins calculated

by ONIOM-EE, SA4-CASSCF, SS-CASPT2 and MS-CASPT2 with ANO-S basis sets and the IPEA zero-order Hamiltonian.

Method	SA4-CASSCF	SS-CASPT2	MS-CASPT2
N_{trans} (absorption)^a			
S₁	4.61 ^c /4.61 ^d	4.54 ^c /4.53 ^d	3.68 ^c /3.62 ^d
S₂	5.26 ^c /5.17 ^d	4.03 ^c /3.89 ^d	4.60 ^c /4.57 ^d
S₃	5.71 ^c /5.49 ^d	5.06 ^c /4.79 ^d	5.53 ^c /5.14 ^d
N_{trans} (emission)^b			
S₁	3.92 ^c /3.81 ^d	3.00 ^c /2.94 ^d	3.22 ^c /3.12 ^d
S₂	4.22 ^c /4.21 ^d	4.14 ^c /4.12 ^d	4.31 ^c /4.28 ^d
S₃	4.44 ^c /4.35 ^d	4.32 ^c /4.14 ^d	4.52 ^c /4.37 ^d

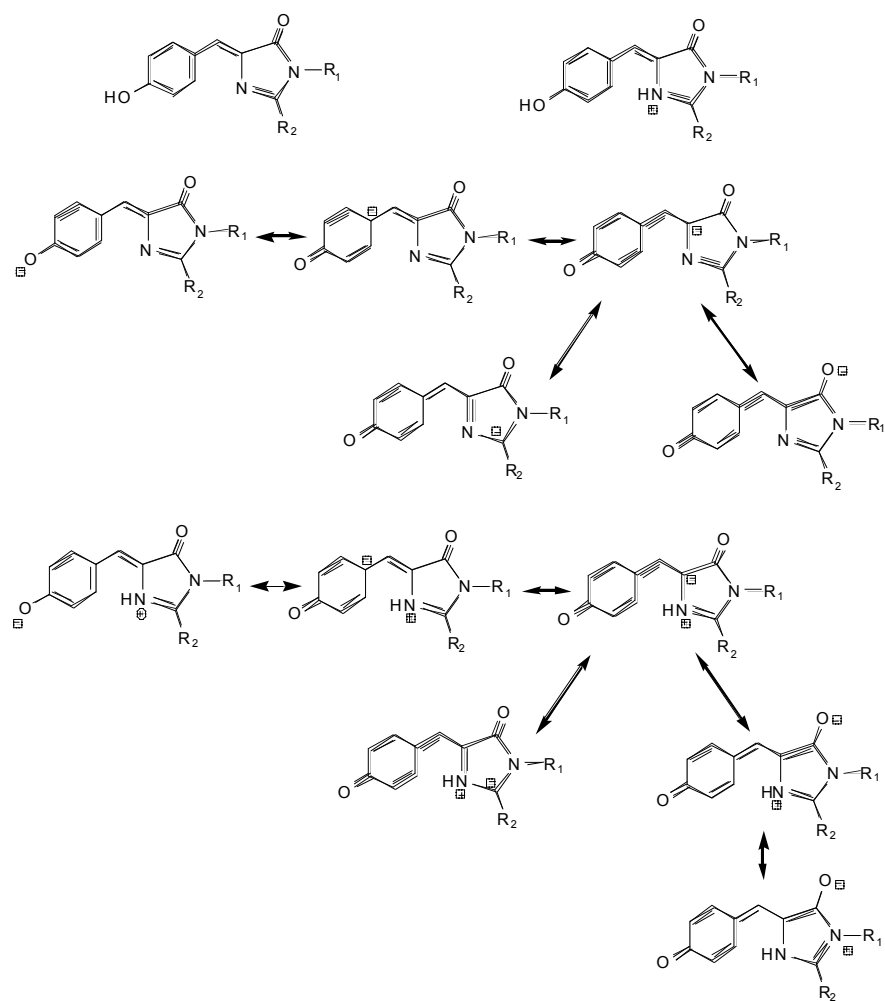
^aGeometry optimized by SA2-CASSCF(12e,12o)/6-31G(d). ^bGeometry optimized by SA4-CASSCF(12e,12o)/6-31G(d). ^c(12e,12o) was used in these calculations. ^d(14e,13o) was used in these calculations.

Table S5(b). Absorption and emission energies of the chromophore in the proteins calculated by ONIOM-EE, SS-CASPT2 and MS-CASPT2 with ANO-S basis sets and the original zero-order Hamiltonian.

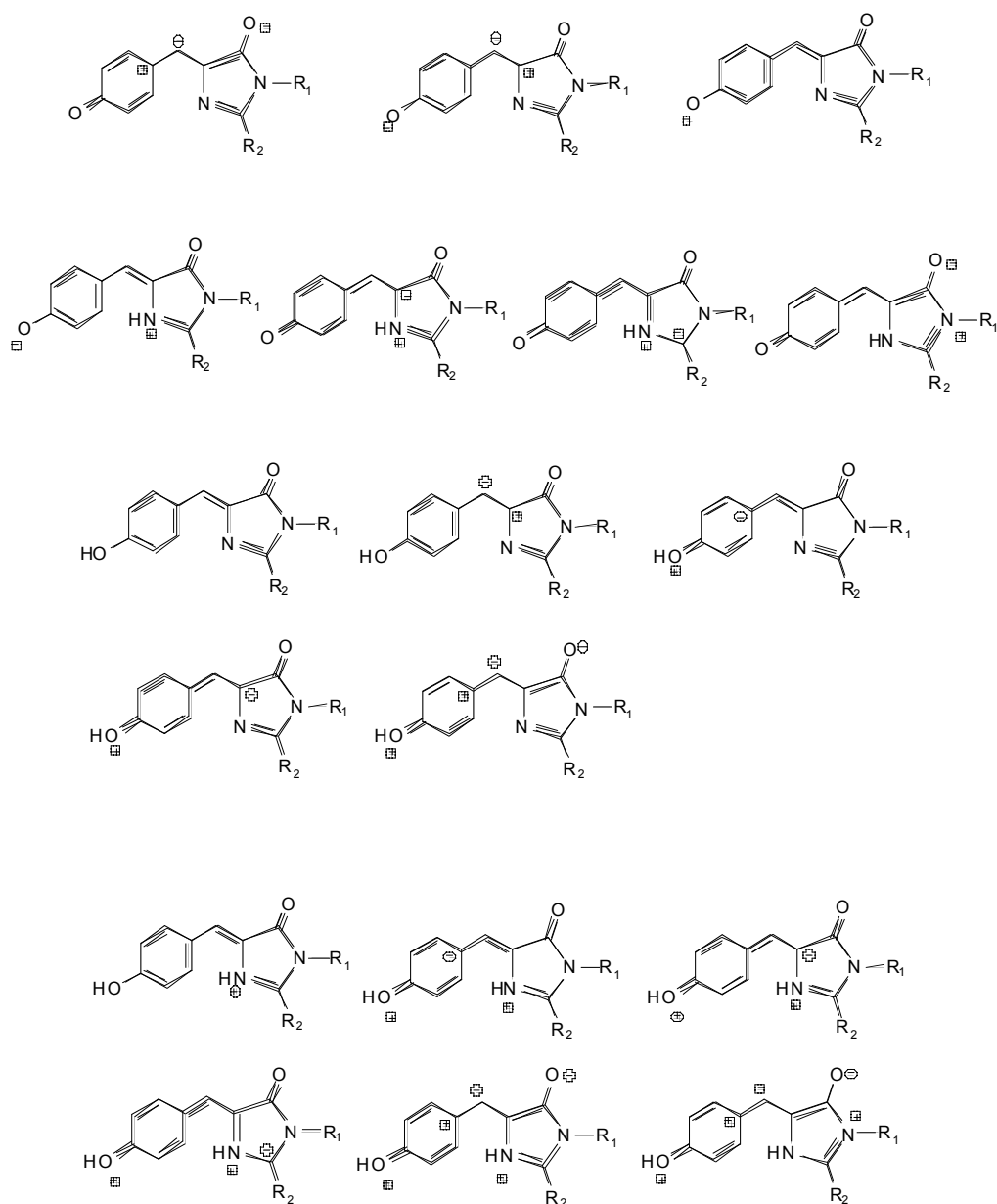
Method	SS-CASPT2	MS-CASPT2
N_{trans} (absorption)^a		
S₁	4.07 ^c /4.05 ^d	3.24 ^c /3.16 ^d
S₂	3.50 ^c /3.36 ^d	4.11 ^c /4.11 ^d
S₃	4.29 ^c /4.28 ^d	4.60 ^c /4.55 ^d
N_{trans} (emission)^b		
S₁	2.69 ^c /2.58 ^d	2.84 ^c /2.92 ^d
S₂	3.66 ^c /3.72 ^d	3.76 ^c /3.95 ^d
S₃	3.71 ^c /3.86 ^d	3.83 ^c /4.10 ^d

^aGeometry optimized by SA2-CASSCF(12e,12o)/6-31G(d). ^bGeometry optimized by SA4-CASSCF(12e,12o)/6-31G(d). ^c(12e,12o) was used in these calculations. ^d(14e,13o) was used in these calculations.

Scheme S1. Resonance structures for the ground-state chromophores



Scheme S2. Resonance structures for the excited state chromophores



Cartesian coordinates (in Å) and absolute energies (a.u.) of the CASSCF(14e,13o)-optimized structures

A_{cis} (FC):

CASSCF/6-31G*:

FC(S0): -641.543786001379 hartree

FC(S1): -641.421781551197 hartree

CASPT2/6-31G*:

FC(S0): -643.300066895141 hartree

FC(S1): -643.200683429753 hartree

C	-1.4360465408	-0.4452400853	-0.1744435878
N	-0.3359541545	0.1802773366	0.0356563030
N	-1.2539358789	-1.6796300574	-0.7735617078
C	0.1099727274	-1.8734843912	-0.9712534098
O	0.5950433033	-2.8668134345	-1.4788257727
C	0.6915088192	-0.6535695678	-0.4393923714
C	2.0537530210	-0.4041498897	-0.4277686841
C	2.8254049495	0.6781199183	0.0238087855
C	2.3012106814	1.8699757447	0.6278755216
C	4.2513446542	0.6248561593	-0.1122556756
C	3.1083967970	2.8883635679	1.0464555635
C	5.0790073316	1.6296778752	0.2983745025
C	4.5598644638	2.8431528370	0.9124209403
O	5.2880100042	3.7614580213	1.2897235549
H	1.2379473277	1.9525250653	0.7494824082
H	4.6843922074	-0.2554213398	-0.5619840126
H	2.6913359242	3.7732726580	1.4970294088
H	6.1468180669	1.5559446501	0.1804832037
H	-1.9516290339	-2.3367392211	-1.0276809596
H	-2.4125298733	-0.0793007823	0.0750983780
H	2.6167920428	-1.2184554246	-0.8568326587

A_{cis} (FS):

CASSCF/6-31G*:

FS(S0): -641.537469826444 hartree

FS(S1): -641.427393494410n hartree

CASPT2/6-31G*:

FS(S0): -643.298038044879 hartree

FS(S1): -643.207459526269 hartree

C	-1.4133479548	-0.3985835707	-0.1539123392
N	-0.2660097020	0.1795460362	0.0299705307
N	-1.2831350687	-1.6377377589	-0.7512232194
C	0.0613288956	-1.8846975335	-0.9728354804
O	0.5195193458	-2.8959596332	-1.4870201339

C	0.7036510453	-0.6837259193	-0.4546255685
C	2.1048027026	-0.4948568371	-0.4747584615
C	2.8683832738	0.6528724714	0.0085166266
C	2.2921272853	1.8162742594	0.6030703549
C	4.2864708601	0.6280715979	-0.1132959525
C	3.0825231817	2.8718517404	1.0405755392
C	5.0822199363	1.6759035901	0.3200986871
C	4.5205263940	2.8688170954	0.9275900369
O	5.2407470977	3.8090618279	1.3158809690
H	1.2282767354	1.8667074897	0.7095188429
H	4.7516912996	-0.2364533343	-0.5579786937
H	2.6274693103	3.7398123312	1.4861043545
H	6.1527893015	1.6327037651	0.2165279568
H	-2.0085577980	-2.2695870013	-0.9916528808
H	-2.3649387281	0.0112615856	0.1143418992
H	2.6641694266	-1.3064625622	-0.9024833379

Z_{cis} (FC):

CASSCF/6-31G*:

FC(S0): -642.055344910342 hartree

FC(S1): -641.929258892667 hartree

CASPT2/6-31G*:

FC(S0): -643.802381741722 hartree

FC(S1): -643.689397172129 hartree

C	-1.5485747904	-0.5039053769	-0.1886540709
N	-0.3896187718	0.1069188330	0.0106108775
N	-1.2624235327	-1.6911164324	-0.7777628354
C	0.1090341631	-1.8553501645	-0.9651273098
O	0.6304389529	-2.8224751331	-1.4678736301
C	0.6763820176	-0.6352143286	-0.4284089593
C	2.0710533696	-0.3599052609	-0.4077699624
C	2.8277128356	0.7007728855	0.0340569473
C	2.3206374242	1.9202931093	0.6473430505
C	4.2771278893	0.6223610313	-0.1140380163
C	3.1359065219	2.9205751157	1.0570763322
C	5.1129234512	1.6078514403	0.2873973034
C	4.5980497041	2.8357291767	0.9062711464
O	5.3292366311	3.7372867181	1.2763794949
H	1.2643429381	2.0514552539	0.7876674729
H	4.6816447861	-0.2666441474	-0.5645278763
H	2.7470988302	3.8151347328	1.5076612695
H	6.1776611643	1.5293083568	0.1690381972
H	-1.9287344894	-2.3762485628	-1.0519230719
H	-2.5102565950	-0.1279543048	0.0687914976

H	2.6150394276	-1.1815984181	-0.8387844196
H	-0.3259583676	1.0024883160	0.4344787726

Z_{cis} (FS):

CASSCF/6-31G*:

FS(S0): -642.026557991865 hartree

FS(S1): -641.959875944131 hartree

CASPT2/6-31G*:

FS(S0): -643.783756910114 hartree

FS(S1): -643.727666413347 hartree

C	-1.3088012873	-0.5838975617	0.2652186989
C	-0.0185978585	-1.7104399036	-1.2019104460
C	0.7522531175	-0.6237700308	-0.6010771917
C	2.0805710853	-0.3775716378	-0.7240988953
C	2.8398808142	0.7220211756	-0.1464491986
C	2.2839718496	2.0043145712	0.1249611565
C	4.2164124526	0.5247163604	0.1414977887
C	3.0381599936	3.0152103852	0.6738849346
C	4.9930767984	1.5151132590	0.6999771153
C	4.4381882396	2.8203252943	1.0059800074
N	-0.1668056073	0.1461965206	0.1918287743
N	-1.3345982502	-1.5305530417	-0.7255172689
O	0.3195028190	-2.5901934053	-1.9402502435
O	5.1192374296	3.7185678523	1.5154920268
H	1.2653829870	2.2085358849	-0.1512631843
H	4.6613366172	-0.4305155531	-0.0792270652
H	2.6150592656	3.9844799991	0.8631537473
H	6.0328511365	1.3537708844	0.9158794333
H	-1.9842424274	-2.2853929369	-0.7272152475
H	-2.1767152213	-0.2712147975	0.8058878675
H	2.6290608408	-1.1191113300	-1.2788612335
H	0.1815481755	0.5790707444	1.0262116007

N_{cis} (FC):

SA2-CASSCF/6-31G*:

FC(S0): -642.110276142151 hartree

FC(S1): -641.94022971076 hartree

CASPT2/6-31G*:

FC(S0): -643.843450351084 hartree

FC(S1): -643.686889492736 hartree

C	-1.3981046793	-0.4293370736	-0.1699081544
C	0.0840348189	-1.9021512486	-0.9824141448
C	0.7203710527	-0.6632918806	-0.4461749614

C	2.0539605416	-0.4570891724	-0.4529053387
C	2.8400533438	0.6853005109	0.0259784222
C	2.2840247910	1.8351758775	0.6127050710
C	4.2386669118	0.6264223616	-0.1105873783
C	3.1009450988	2.8790663483	1.0426171881
C	5.0579279598	1.6661895646	0.3172099360
C	4.4865125230	2.7989476553	0.8970530781
N	-0.2970846581	0.1881942960	0.0361595030
N	-1.2618258871	-1.6702862017	-0.7677072412
O	0.5773673843	-2.8755416967	-1.4812701313
O	5.3209213647	3.7879713114	1.2996758280
H	1.2228016704	1.9128664521	0.7319176395
H	4.6899071105	-0.2422154649	-0.5562537866
H	2.6564265288	3.7523324062	1.4899242061
H	6.1250518191	1.6140587860	0.2096179502
H	-1.9904281987	-2.3007970828	-1.0084959549
H	-2.3645681254	-0.0450804701	0.0874296022
H	2.6238615372	-1.2643680033	-0.8792655022
H	4.8302116324	4.5051861748	1.6768246894

N_{cis} (FS):

SA4-CASSCF(12e,12o)/6-31G*:

FC(S0): -642.24963812 hartree

FC(S1): -642.09612443 hartree

FC(S2): -642.091444947 hartree

FC(S3): -642.07072206 hartree

SS-CASPT2/6-31G*:

FS(S0): -644.1802100797 hartree

FS(S1): -644.0275844091 hartree

FC(S2): -644.0238563567 hartree

FC(S3): -644.0545383565 hartree

MS-CASPT2/6-31G*:

FS(S0): -644.18261424 hartree

FS(S1): -644.0606074 hartree

FC(S2): -644.02433089 hartree

FC(S3): -644.01863667 hartree

C	-1.3838143164	-0.3544061954	-0.1359326732
C	0.0831991149	-1.8362981776	-0.9510266157
C	0.7198753907	-0.6543935888	-0.4419252870
C	2.1386050811	-0.4942784572	-0.4768884954
C	2.88358079300	6.044214963	-0.0155464721
C	2.26744353421	7.6525011550	5.812507982
C	4.31553533790	6.051029780	-0.1268278304

C 3.03110734902.82527055441.0228617689
 C 5.06885067801.66424043980.3151751975
 C 4.42833521772.78950628150.8970674429
 N -0.22418717920.21749061290.0439636413
 N -1.2604538664-1.5840441460-0.7269572865
 O 0.5492420668-2.8500174296-1.4666295201
 O 5.21308506893.77777690531.3031211337
 H 1.20198175881.78283024440.6725956988
 H 4.7998255025-0.2457293292-0.5668015923
 H 2.55927942143.68478660971.4659675909
 H 6.13891886631.67347616810.2361487090
 H -1.9923774518-2.2106061269-0.9658956494
 H -2.32817093420.06580156840.1364077696
 H 2.6755969219-1.3179821269-0.9081809559
 H 4.71557618494.49335505341.6801831476

C_{cis} (FC):

CASSCF/6-31G*:

FC(S0): -642.466861661966 hartree

FC(S1): -642.299746417262 hartree

CASPT2/6-31G*:

FC(S0): -644.202640041297 hartree

FC(S1): -644.054443751618 hartree

C -1.5059455479 -0.5372577638 -0.2106230682
 C 0.1098117643 -1.8982674717 -0.9836417474
 C 0.7181028322 -0.6595280129 -0.4444830656
 C 2.0376438177 -0.3919365180 -0.4218047684
 C 2.8320644242 0.7359639761 0.0499560803
 C 2.3174640543 1.9084886118 0.6430108426
 C 4.2267572122 0.6369755561 -0.1038103405
 C 3.1553083355 2.9311620320 1.0617301546
 C 5.0747897978 1.6580938466 0.3134921082
 C 4.5403734140 2.8105868328 0.8987530847
 N -0.4000340041 0.1062367953 0.0068922614
 N -1.2858237372 -1.7097868997 -0.7840370034
 O 0.6075328585 -2.8555617419 -1.4767754818
 O 5.4038785357 3.7609266406 1.2813669520
 H 1.2635985923 2.0518565851 0.7907459407
 H 4.6513344814 -0.2425353692 -0.5518168289
 H 2.7375939525 3.8146227633 1.5106872704
 H 6.1380315873 1.5790716076 0.1946302151
 H -1.9867574470 -2.3729912290 -1.0430496571
 H -2.4760570123 -0.1613514308 0.0445459801
 H 2.6090548307 -1.1978526294 -0.8485016123

H 4.9671567275 4.5087653725 1.6678705317
 H -0.3639610998 1.0072521366 0.4320611720

C_{cis} (FS):

CASSCF/6-31G*:

FS(S0): -642.423536519463 hartree

FS(S1): -642.335353742397 hartree

CASPT2/6-31G*:

FS(S0): -644.18496941396 hartree

FS(S1): -644.102679581711 hartree

C -1.4756898608 -0.5986270103 0.1171901050
 C 0.0474792226 -1.7485214852 -1.1180099762
 C 0.7314299717 -0.6053311370 -0.4520833460
 C 2.0707964572 -0.3962555141 -0.5595359535
 C 2.8483573805 0.7015017795 -0.0283125580
 C 2.2773207783 1.9473532803 0.3844765931
 C 4.2680773999 0.5588772911 0.0396311534
 C 3.0687672249 2.9758176648 0.8362081857
 C 5.0663170392 1.5737291684 0.4998119818
 C 4.4740300525 2.8008010258 0.9087772627
 N -0.2658663476 0.1236332853 0.2069468574
 N -1.2728046859 -1.6258630195 -0.7707749919
 O 0.5452014280 -2.5915483998 -1.8145122149
 O 5.2861675634 3.7276322629 1.3412272051
 H 1.2218000627 2.1044564689 0.2955256669
 H 4.7134298507 -0.3653520009 -0.2756112354
 H 2.6334453040 3.9136646457 1.1294590008
 H 6.1329102174 1.4780213100 0.5617332802
 H -1.9774172747 -2.2624566310 -1.0745057716
 H -2.4227277173 -0.0995490997 0.1853607163
 H 2.6057138166 -1.1604399704 -1.0913977883
 H 4.8511637323 4.5353628238 1.5958880652
 H -0.0559876154 0.4960272615 1.1097067622

A_{trans} (FC):

CASSCF/6-31G*:

FC(S0): -641.539158191781 hartree

FC(S1): -641.417074979664 hartree

CASPT2/6-31G*:

FC(S0): -643.2965982 hartree

FC(S1): -643.196832 hartree

C 0.4623103467 1.0316605525 -0.5041088026
 C 2.1387282974 -0.3106827122 0.1845695602
 C 2.4404554509 1.0970396978 0.3764755893

C	3.5425724017	1.7734529505	0.8934707717
C	4.7870732801	1.4135426462	1.4310748796
C	5.6643928117	2.4661172229	1.8638338383
C	5.2900434164	0.0777262475	1.5994204677
C	6.8969019286	2.2405522198	2.4017190486
C	6.5206238112	-0.1710518505	2.1354997481
C	7.4164617015	0.8910241900	2.5769612600
N	1.3422597408	1.8604379503	-0.0795030503
N	0.8667712965	-0.2841847212	-0.3784182034
O	2.7404703163	-1.3451088058	0.4086889541
O	8.5257836558	0.6664424658	3.0601186771
H	5.3243549088	3.4844379343	1.7547730885
H	4.6729280482	-0.7428730079	1.2916901508
H	7.5272090062	3.0555318723	2.7150348065
H	6.8745471452	-1.1815439670	2.2510357136
H	3.3758540309	2.8390169824	0.8633172773
H	-0.4921852415	1.2933937893	-0.9168795149
H	0.3670363070	-1.1013306569	-0.6351709600

A_{trans} (FS):

CASSCF/6-31G*:

FS(S0): -641.532843026794 hartree

FS(S1): -641.422575710067 hartree

CASPT2/6-31G*:

FS(S0): -643.2948968 hartree

FS(S1): -643.20371 hartree

C	0.4436467638	0.9817444151	-0.5140849917
C	2.1502066018	-0.3101982096	0.1893823669
C	2.4197454450	1.1109889823	0.3676790733
C	3.5160464561	1.8621876820	0.8848349276
C	4.8033401643	1.4562618278	1.4398389333
C	5.6885160159	2.4868001903	1.8753168654
C	5.2615254636	0.1147740028	1.5882032287
C	6.9324252997	2.2138674003	2.4165161546
C	6.5095543962	-0.1678465949	2.1308452482
C	7.4208404352	0.8569113194	2.5775826624
N	1.3173975507	1.8345849414	-0.0913747026
N	0.8747154044	-0.3121174287	-0.3760454816
O	2.7686600994	-1.3438651229	0.4206500572
O	8.5417326121	0.6138221801	3.0654886063
H	5.3736917022	3.5131724766	1.7776814332
H	4.6286297404	-0.6888198621	1.2740610680
H	7.5789599018	3.0131021047	2.7364949307
H	6.8324334812	-1.1897471792	2.2322584645

H	3.3407735575	2.9221584275	0.8506676962
H	-0.5130225303	1.2292780917	-0.9281094377
H	0.3947740991	-1.1434586446	-0.6242838028

Z_{trans} (FC):

CASSCF/6-31G*:

FC(S0): -642.055432085499 hartree

FC(S1): -641.920335059568 hartree

CASPT2/6-31G*:

FC(S0): -643.8029741 hartree

FC(S1): -643.6777492 hartree

C	0.4063389693	0.9840526205	-0.5311035357
C	2.1454203655	-0.3061167862	0.1874312599
C	2.4601864959	1.0956713013	0.3849102849
C	3.5735747125	1.8292496172	0.9092620538
C	4.7870021827	1.4551521844	1.4325976761
C	5.6906409296	2.5142368895	1.8772602397
C	5.2634768155	0.0841907028	1.5878968930
C	6.9076541606	2.2586325019	2.4071683826
C	6.4805801974	-0.1850867379	2.1172881612
C	7.3920973336	0.8781076655	2.5657199853
N	1.3601454292	1.7788540175	-0.0750913648
N	0.8734527974	-0.2782268961	-0.3758532019
O	2.7515069878	-1.3303380664	0.4141304617
O	8.4869519804	0.6347155176	3.0418697551
H	5.3614105372	3.5338853667	1.7729003959
H	4.6263019947	-0.7161478607	1.2721059131
H	7.5623658680	3.0469592583	2.7302011624
H	6.8272640620	-1.1958930901	2.2297355514
H	3.4086244665	2.8958625432	0.8798371012
H	-0.5318013751	1.2816564264	-0.9351820542
H	0.3904990091	-1.1111275766	-0.6245165111
H	1.2863165603	2.7717471412	-0.0669107595

Z_{trans} (FS):

CASSCF/6-31G*:

FS(S0): -642.036664189999 hartree

FS(S1): -641.952478498332 hartree

CASPT2/6-31G*:

FS(S0): -643.7830184 hartree

FS(S1): -643.7058272 hartree

C	0.3575783154	0.9727246295	-0.5819474740
C	2.2206352383	-0.2619942024	-0.0258197843
C	2.4722198542	1.1474554800	0.3034900690

C	3.5860302874	1.8647037887	0.7922597595
C	4.8299297518	1.4567787278	1.4004186038
C	5.7583239755	2.4916858554	1.7318899039
C	5.1876118012	0.1181432668	1.7083227543
C	6.9699600128	2.2185049053	2.3304387458
C	6.4047395280	-0.1724011371	2.3135788649
C	7.3382259486	0.8637576384	2.6459301017
N	1.3256129746	1.8506176555	-0.0909526008
N	0.8962012859	-0.3101767340	-0.4355262352
O	2.9527643712	-1.2168272803	0.0104058989
O	8.4408026017	0.6001349578	3.1960599427
H	5.5019041061	3.5116913798	1.5065444243
H	4.5221277012	-0.6804814024	1.4583428518
H	7.6626840038	3.0011122584	2.5771312181
H	6.6760587929	-1.1852032885	2.5452215668
H	3.4615108529	2.9311433965	0.7132941846
H	-0.6767590895	1.1474741413	-0.3397488311
H	0.5845282380	-1.0833002076	-0.9788568021
H	1.0373199281	2.6544929116	0.4211806872

N_{trans} (FC):

CASSCF/6-31G*:

FC(S0): -642.104913607878 hartree

FC(S1): -641.935023589486 hartree

CASPT2/6-31G*:

FC(S0): -643.84759696283 hartree

FC(S1): -643.692838715794 hartree

C	0.4727218124	1.0173326628	-0.5000650223
C	2.1438001477	-0.3195048796	0.1864907694
C	2.4445868940	1.1347072708	0.3795221916
C	3.5085119672	1.8087378759	0.8794043126
C	4.8058303703	1.4111236067	1.4391312940
C	5.6561834650	2.4535610294	1.8597655315
C	5.2672919094	0.0927301948	1.5897358966
C	6.9091392450	2.2026090594	2.4059274518
C	6.5239594284	-0.1631419426	2.1373401681
C	7.3471856113	0.8851176920	2.5461931898
N	1.3256814810	1.8715774655	-0.0868307667
N	0.8835100090	-0.2995596404	-0.3713533266
O	2.7792091243	-1.3112857283	0.4273482298
O	8.5746301074	0.6845738624	3.0831056044
H	5.3309754072	3.4734370794	1.7573633041
H	4.6569666474	-0.7300385904	1.2847273890
H	7.5484390311	3.0050242862	2.7229452237

H	6.8567361010	-1.1820719923	2.2431348902
H	8.7698712793	-0.2412169206	3.1316960878
H	3.3559579978	2.8734712294	0.8553278269
H	-0.4846046804	1.2603897444	-0.9148417185
H	0.3787764372	-1.1150982239	-0.6296224561

N_{trans} (FS):

SA4-CASSCF(14,13)/6-31G*:

FS(S0): -642.24789891 hartree

FS(S1): -642.09815708 hartree

FS(S2): -642.09241505 hartree

FS(S3): -642.07563622 hartree

SS-CASPT2(14,13)/6-31G*:

FS(S0): -644.1780749128 hartree

FS(S1): -644.0412683258 hartree

FS(S2): -644.0229212441 hartree

FS(S3): -644.0469612608 hartree

MS-CASPT2(14,13)/6-31G*:

FS(S0): -644.17982261 hartree

FS(S1): -644.07078919 hartree

FS(S2): -644.02240459 hartree

FS(S3): -644.01620936 hartree

C	0.47128316050.9389170970-0.5040600724
C	2.1981501295-0.29216777440.2115940558
C	2.43548772421.12320061960.3752394461
C	3.52782890561.90377436680.8921548894
C	4.78461047471.49323968991.4332860900
C	5.69469914222.51330464751.8792841984
C	5.22312586120.12993069291.5719689560
C	6.92316168962.21077842902.4122922140
C	6.4601597667-0.17027723372.1088799604
C	7.32631634700.85325497962.5355143988
N	1.34439053531.8171975813-0.0804019364
N	0.9243798658-0.3258416404-0.3550623695
O	2.8359564970-1.31695039260.4524016354
O	8.53238549070.62071281793.0614591858
H	5.39772265803.54118138591.7896632107
H	4.5765512685-0.65950310341.2524317596
H	7.59763679542.97736480652.7432588569
H	6.7647312828-1.19839619042.2018627255
H	8.7174025446-0.30885307843.1054106905
H	3.34689349662.96044514370.8552675007
H	-0.48621742521.1715628631-0.9194156562

H 0.4587035828-1.1704005670-0.5965836687

C_{trans} (FC):

CASSCF/6-31G*:

FC(S0): -642.461052586423 hartree

FC(S1): -642.321496000667 hartree

CASPT2/6-31G*:

FC(S0): -644.2103264 hartree

FC(S1): -644.1069738 hartree

C	0.4367788193	0.9850666508	-0.5176747003
C	2.1513792072	-0.3034233022	0.1904175211
C	2.4847375632	1.1114585800	0.3963448958
C	3.5458262136	1.8098351955	0.8960654313
C	4.8135784733	1.4203557580	1.4429685066
C	5.6694258199	2.4690060257	1.8661782092
C	5.2700340668	0.0932186851	1.5910755598
C	6.9120728179	2.2131692698	2.4075300780
C	6.5197517806	-0.1647880617	2.1354708470
C	7.3457394253	0.8874530225	2.5455993352
N	1.3320992244	1.8092401635	-0.0866207101
N	0.8502405468	-0.2669870897	-0.3854492292
O	2.7402774105	-1.3163752357	0.4101821115
O	8.5558501084	0.6960418127	3.0751618886
H	5.3480054019	3.4899576514	1.7654288327
H	4.6622171129	-0.7315202720	1.2871906899
H	7.5575727705	3.0083318259	2.7269119319
H	6.8524617504	-1.1820156454	2.2413718088
H	8.7791867585	-0.2240978834	3.1364952342
H	3.3999955629	2.8775386491	0.8751357035
H	-0.5092565146	1.2770934124	-0.9255421586
H	0.3378026957	-1.0832154575	-0.6463098694
H	1.2250005846	2.8035672849	-0.0934311676

C_{trans} (FS):

CASSCF/6-31G*:

FS(S0): -642.451309143477 hartree

FS(S1): -642.332084723135 hartree

CASPT2/6-31G*:

FS(S0): -644.2085234 hartree

FS(S1): -644.1152357 hartree

C	0.4073733259	0.9350986900	-0.5329254807
C	2.1697879094	-0.2832383311	0.1993888238
C	2.4571818412	1.1419352795	0.3855502631
C	3.5469353006	1.8992449561	0.9004447296

C	4.8071352224	1.4847299396	1.4429319674
C	5.7073939600	2.5160423119	1.8848891548
C	5.2434047231	0.1165798945	1.5805478533
C	6.9366878085	2.2203486330	2.4185312451
C	6.4743659822	-0.1840029931	2.1147266847
C	7.3376714582	0.8555656668	2.5406121808
N	1.3241101465	1.7866937074	-0.0909624863
N	0.8997637538	-0.3015556435	-0.3651551965
O	2.8143711729	-1.2775462577	0.4446556063
O	8.5121386471	0.6356080936	3.0530179308
H	5.4111315634	3.5438722681	1.7954870215
H	4.6026360357	-0.6778151292	1.2635015177
H	7.6145357720	2.9823723372	2.7507048383
H	6.7881291282	-1.2077623744	2.2120655796
H	8.7371419254	-0.2890473120	3.1150081647
H	3.3929158772	2.9630450103	0.8757274538
H	-0.5415100642	1.1926319736	-0.9435889001
H	0.4347981174	-1.1496866711	-0.6064366728
H	1.2026779931	2.7757969902	-0.1042215289

N_{rr}:

CASSCF(12e,12o)/6-31G*:

S0: -642.008729038378 hartree

S1: -641.951904128644 hartree

C	1.2659810139	0.2630304999	-1.1373258093
C	1.3728632892	0.7278423734	1.0733713922
C	2.5108852675	1.1497358043	0.3498734208
C	3.5369355866	2.0249470050	0.8672138733
C	4.7571755953	1.6156052494	1.4022000317
C	5.6981227359	2.5714898600	1.8885831775
C	5.0770766302	0.2334630299	1.4873922007
C	6.8953257087	2.1689816031	2.4241089085
C	6.2784256695	-0.1728374760	2.0260391004
C	7.1858139068	0.7906051312	2.4943637759
N	2.3929154895	0.8806063661	-1.0176459078
N	0.6189613696	0.1149356486	0.0693250104
O	1.0506883072	0.8080247870	2.2540316602
O	8.3560609958	0.4596757401	3.0257380138
H	5.4575165903	3.6166359490	1.8308741044
H	4.3551497302	-0.4705642336	1.1264248634
H	7.6200942266	2.8662309859	2.7971033160
H	6.5234724224	-1.2175501549	2.0957815843
H	8.4847842611	-0.4807468168	3.0493418999
H	3.3904096679	3.1021264843	0.8748806253

H 0.8670127015 -0.1181981774 -2.0547475555
H -0.2829140555 -0.2700010437 0.2228623698

CASSCF(14e,13o)/6-31G*:

S0: -642.01343339166 hartree

S1: -641.955459240170 hartree

C 1.2718585332 0.2564521206 -1.1326777705
C 1.3753656775 0.7331657473 1.0741386109
C 2.5083325113 1.1626920440 0.3472195348
C 3.5370539890 2.0351420805 0.8657757173
C 4.7552863374 1.6214807446 1.4009596742
C 5.6985835969 2.5738617704 1.8899645430
C 5.0715611464 0.2381558419 1.4836255981
C 6.8943029553 2.1671430483 2.4254908243
C 6.2715334200 -0.1723333377 2.0219000999
C 7.1812677991 0.7878472149 2.4926734608
N 2.3969433923 0.8831130616 -1.0154486989
N 0.6246315876 0.1162005970 0.0709029882
O 1.0530745188 0.8094823822 2.2564374002
O 8.3504591117 0.4527503927 3.0233936075
H 5.4606862110 3.6197327345 1.8343654104
H 4.3485761109 -0.4637558358 1.1207497857
H 7.6206060405 2.8616558078 2.8005845096
H 6.5139830072 -1.2177854384 2.0894469927
H 8.4766993584 -0.4880588724 3.0453996409
H 3.3923687149 3.1121189389 0.8733426031
H 0.8763135216 -0.1336986173 -2.0477691346
H -0.2667295410 -0.2913234256 0.2293136022
H 5.4575165903 3.6166359490 1.8308741044
H 4.3551497302 -0.4705642336 1.1264248634
H 7.6200942266 2.8662309859 2.7971033160
H 6.5234724224 -1.2175501549 2.0957815843
H 8.4847842611 -0.4807468168 3.0493418999
H 3.3904096679 3.1021264843 0.8748806253
H 0.8670127015 -0.1181981774 -2.0547475555
H -0.2829140555 -0.2700010437 0.2228623698