

Supporting Information

Quantum Mechanics/Molecular Mechanics Study on the Photoreactions of Dark- and Light-Adapted States of a Blue-Light YtvA LOV Photoreceptor

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Contents

- 1. Computational Methods
 - 1.1 System Setup
 - 1.2 QM/MM Method
 - 1.2.1 Vertical Excitation Energies
 - 1.2.2 Geometric Parameters of Key Minima
- 2. Inter-Conversion Paths
- 3. Minimum-Energy Paths from AU, AD, and BU
- 4. Tables
- 5. Comparison with Previous Theoretical Work
- 6. References
- 7. Cartesian Coordinates of All Optimized Structures

1. <u>Computational Methods</u>

1.1 System Setup

The initial geometry was constructed from an X-ray structure of YtvA LOV domains (PDB ID: 2PR5, chain A). [1] In this structure, there are two main conformations with different orientations of the Cys62 residue, which are labeled as conf-A and conf-B in Fig. S1. Both are used in our work in order to study possible conformation-dependent spectroscopic properties and excited-state reactions.

The constructed system consisted of 102 amino acids, 252 crystal waters, and a flavin mononucleotide (FMN) cofactor. It was first solvated in an equilibrated water box with a radius of 35 Å (with its origin at the center of mass of the FMN cofactor). The protonation states of ionizable groups were determined using the PDB2PQR web server. [2] The system was then neutralized by adding nine sodium ions on the protein surface near negatively charged residues. A spherical quartic boundary potential was applied to the outer water molecules to prevent them from diffusing away. [3]

To maintain the proper protein structure in the initial MM minimizations, all non-hydrogen atoms of the protein and cofactor were restrained with harmonic potentials and only their hydrogen atoms and the water molecules were allowed to move. The solvated system was first relaxed by performing energy minimizations of 500 steps. Then, successive MD simulations of 15 ps were carried out to heat the system gradually to 300 K, while the force constants of the harmonic restraints to the protein were gradually reduced from 90 to 20 kcal \cdot mol⁻¹ \cdot Å⁻². Finally, all restraints were removed and a production NVT MD of 2 ns was run with a 1 fs time step (T=300K). [3]

2

In MM minimizations and MD simulations, the protein residues and the FMN cofactor were described with the CHARMM22 all-atom force field [4], and the water molecules were represented by the TIP3P [5] model.



Figure S1. Overview (left) and view (right) into the FMN binding pocket of the YtvA LOV domain (PDB code 2PR5). Shown are the two conformations of the cysteine residue, conf-A and conf-B.



Figure S2. Illustration of binding pocket (left) and selected QM region (right) employed in the QM/MM geometry optimizations.

1.2 QM/MM Method

The QM region in our QM/MM calculations consisted of lumiflavin and the thiol group of the Cys62 residue. The remaining atoms made up the MM region (see Fig. S2). The QM/MM partitioning involved cuts through two covalent bonds between the QM and the MM subsystems. To explore possible conformation-dependent effects, photochemical reactions were

initiated from both conformations (conf-A and conf-B) considering two different orientations of the H atom of the thiol group. Hence, a total of four conformers were studied (AU, AD, BU, and BD).

The QM region was described using the complete-active-space selfconsistent field (CASSCF) method. Two-state averaged and state-specific CASSCF calculations were carried out for singlet state (S₁) and for the singlet state (S_0) and triplet state (T_1) , respectively. All CASSCF computations made use of an active space of 10 electrons distributed over 8 molecular orbitals (see Fig. S3). Since the CASSCF approach does not treat dynamic correlation, the complete-active-space second-order perturbation (CASPT2) approach [6, 7] was applied to re-evaluate the energies of all optimized structures and paths. The MS-CASPT2 computations employed a larger active space of 12 electrons in 9 molecular orbitals (see Fig. S3). In addition, we used the Cholesky decomposition technique with unbiased auxiliary basis sets for accurate two-electron integral approximations [8] and the imaginary shift technique (0.2 au) to avoid intruder-state issues. [9] The ionization potential electron affinity shift was set to zero. [10, 11] According to our experience, the CASPT2 method without IPEA correction performs better for organic systems, as also shown by González and coworkers. [11]

The MM region was described by the all-atom CHARMM22 force field (protein residues and FMN cofactor) and the TIP3P model (water molecules). [5, 12, 13] The electrostatic embedding scheme [14] was adopted in QM/MM calculations. No cutoffs were imposed for nonbonding QM/MM interactions. Hydrogen link atoms were used at the QM/MM boundary in combination with the charge shift model. [15, 16]

4

In all QM/MM geometry optimizations, the QM atoms and all MM atoms within 12 Å from the center of mass of the FMN core chromophore were allowed to move. The other atoms were frozen after the 2 ns MD simulations (see above). To obtain detailed mechanistic information, minimum-energy reaction pathways in ground and excited states were computed using constrained geometry optimizations along selected reaction coordinates.

In addition, we employed the QM(UB3LYP)/MM method [17, 18] to study inter-conversion paths of the four conformers in the T₁ state (AU, AD, BD, and BU). Specifically, we optimized T₁ minimum-energy paths (MEPs) that connect either AU and AD or BU and BD, while the conversion between AU and BD was studied with the nudged elastic band (NEB) method. [19-21]

The QM(MS-CASPT2)/MM method was used to calculate vertical excitation energies and oscillator strengths at optimized QM/MM geometries.

The 6-31G* basis set [22, 23] was chosen for all QM(CASSCF)/MM and QM(UB3LYP)/MM calculations. The larger cc-pVTZ basis set [24] was employed for all QM(MS-CASPT2)/MM single-point calculations.

The following codes were used: QM/MM calculations, ChemShell3.5 package; [25] QM(CASSCF), GAUSSIAN09 package; [26] QM(MS-CASPT2), MOLCAS8.0 [27, 28]; MM DL_POLY module [29] as implemented in the ChemShell3.5. [25]

5



Figure S3. Active spaces used to obtain the CASSCF (top) and MS-CASPT2 (bottom) results (see text). Shown are the natural orbitals from CASSCF calculations.

1.2.1 Vertical Excitation Energies

Table S1. Vertical Excitation Energies (E_⊥, kcal/mol), Oscillator Strengths (f), and Singly Occupied Orbitals Involved in the $S_0 \rightarrow S_1$ ($^1\pi\pi^*$) Electronic Transition; Computed with the 5-Root State-Averaged QM(MS-CASPT2)/MM Method at the QM(CASSCF)/MM Optimized S₀ Minima.

	E	f	singly occupied orbitals
AU	63.7 (448.8 nm)	0.2879	
AD	63.5 (450.2 nm)	0.2839	
BU	63.7(448.8 nm)	0.3112	
BD	63.5 (450.2 nm)	0.3037	
INT	13.2 (2162.5 nm)	_	
Ρ	70.4 (406 nm)	0.2555	

1.2.2 Geometric Parameters of Key Minima

Table S2. Selected Key Geometric Parameters of Stationary Structures Optimized by the QM(CASSCF)/MM Method (Distances in Å and Angles in °).

		Distance		А	ngle	Dihedral Angle
		N5-H	C4a-S	N5-H-S	H-N5-C4a	S-H-N5-C4a
	AU-S0	4.569	4.601	60.5	104.8	-96.5
	AU-S1	4.706	4.606	52.3	105.5	-99.3
	AU-T1	4.520	4.546	57.7	104.3	-98.1
	AD-S0	4.910	4.561	40.1	105.7	104.4

AD-S1	4.900	4.589	41.0	105.8	102.5
AD-T1	4.897	4.607	41.1	106.1	102.3
BU-S0	3.799	3.647	71.3	61.4	175.3
BU-S1	3.779	3.677	72.9	61.5	174.6
BU-T1	3.886	3.569	65.9	59.4	177.3
BD-S0	3.291	3.675	91.6	105.4	8.4
BD-S1	3.260	3.703	93.3	105.7	8.6
BD-T1	3.264	3.669	92.6	104.3	8.8
INT-S0	1.030	3.372	115.4	116.9	72.3
INT-S1	0.997	3.727	83.3	118.3	82.2
INT-T1	1.000	3.720	85.8	118.5	75.4
P-S0	0.997	1.928	67.8	114.1	33.9
P-S1	1.003	1.939	69.6	115.9	37.5
P-T1	0.998	1.906	70.7	114.4	33.3

2. Inter-Conversion Paths



Figure S4. QM(UB3LYP)/MM computed T_1 inter-conversion path with respect to the rotation along the H-C-S-H dihedral angle, which connects the AD and AU conformers.



Figure S5. QM(UB3LYP)/MM computed T_1 inter-conversion path with respect to the rotation along the H-C-S-H dihedral angle, which connects the BD and BU conformers.



Figure S6. QM(UB3LYP)/MM computed T₁ NEB path connecting AU and BD.

3. Minimum-Energy Paths from AU, AD, and BU



Figure S7. QM(MS-CASPT2//CASSCF)/MM computed minimum-energy reaction paths from AU along the N5-H distance in the S_0 , S_1 , and T_1 states.



Figure S8. QM(MS-CASPT2//CASSCF)/MM computed minimum-energy reaction paths of AD along the N5-H distance in the S_0 , S_1 , and T_1 states.



Figure S9. QM(MS-CASPT2//CASSCF)/MM computed minimum-energy reaction paths of BU along the N5-H distance in the S_0 , S_1 , and T_1 states.

4. Tables

Tables S3-S17 document the energies obtained in the reaction path calculations. The total QM/MM energy can be decomposed into a QM contribution that contains the electrostatic QM-MM interaction energy (denoted as CASSCF A.E. or MS-CASPT2 A.E.) and an MM contribution that contains the van-der-Waals QM-MM interaction energy (denoted as MM). These quantities are given as absolute energies (in Hartree). Also listed are the relative energies at the MS-CASPT2 level (ΔE , in kcal/mol) with respect to the lowest ground-state conformer AU-S0.

Table S3. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from AU-S1 (Reaction Coordinate: N5-H Distance). The Total QM/MM Energy Profiles are Plotted in Figure S7 (Red Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₁-min (4.706)				
Root1 (S ₀)	-3091.10440	-1308.65920		3.6
Root2 (S ₁)	-3091.00480	-1308.58177	-105.503029	52.2
Root3		-1308.55160		71.2
S ₁ (4.70)				
Root1 (S ₀)	-3091.10440	-1308.65916		3.7
Root2 (S ₁)	-3091.00480	-1308.58175	-105.50302	52.2
Root3		-1308.55157		71.2
S ₁ (4.60)			·	·
Root1 (S ₀)	-3091.10500	-1308.65914		3.7
Root2 (S ₁)	-3091.00400	-1308.58149	-105.50391	52.4
Root3		-1308.55070		71.7
S ₁ (4.40)			·	·
Root1 (S ₀)	-3091.10400	-1308.65820		4.3
Root2 (S ₁)	-3091.00220	-1308.58035	-105.50462	53.1
Root3		-1308.54992		72.2
S ₁ (4.20)				
Root1 (S ₀)	-3091.10060	-1308.65594		5.7
Root2 (S ₁)	-3090.99940	-1308.57871	-105.50489	54.2
Root3		-1308.54875		73.0
S ₁ (3.80)				
Root1 (S ₀)	-3091.09740	-1308.65263		7.8
Root2 (S ₁)	-3090.99480	-1308.57743	-105.50364	55.0
Root3		-1308.55019		72.1
S ₁ (3.40)				
Root1 (S ₀)	-3091.09810	-1308.64859		10.3
Root2 (S ₁)	-3090.99430	-1308.57080	-105.50070	59.1
Root3		-1308.53988		78.5
S ₁ (3.00)				
Root1 (S ₀)	-3091.09770	-1308.64609		11.9
Root2 (S ₁)	-3090.99390	-1308.56891	-105.49868	60.3
Root3		-1308.53740		80.1
S ₁ (2.60)				
Root1 (S ₀)	-3091.09730	-1308.64540		12.3
Root2 (S ₁)	-3090.99610	-1308.56927	-105.49678	60.1
Root3		-1308.53655		80.6

S ₁ (2.20)				
Root1 (S ₀)	-3091.09040	-1308.64363		13.4
Root2 (S ₁)	-3090.99360	-1308.57015	-105.49695	59.5
Root3		-1308.53808		79.7
S ₁ (1.80)				
Root1 (S ₀)	-3091.07500	-1308.63660		17.8
Root2 (S ₁)	-3090.98200	-1308.56545	-105.49611	62.5
Root3		-1308.53405		82.2
S ₁ (1.60)		·	· · · · · · · · · · · · · · · · · · ·	
Root1 (S ₀)	-3091.06880	-1308.63580		18.3
Root2 (S ₁)	-3090.97490	-1308.56056	-105.50117	65.5
Root3		-1308.53692		80.4
S ₁ (1.50)		·		
Root1 (S ₀)	-3091.05730	-1308.62959		22.2
Root2 (S ₁)	-3090.96710	-1308.55684	-105.50131	67.9
Root3		-1308.53719		80.2
S ₁ (1.40)		·	· · · · · · · · · · · · · · · · · · ·	
Root1 (S ₀)	-3091.04500	-1308.62191		27.0
Root2 (S ₁)	-3090.95850	-1308.55135	-105.50140	71.3
Root3		-1308.54639		74.4
S ₁ (1.30)				
Root1 (S ₀)	-3091.03120	-1308.61578		30.9
Root2 (S ₁)	-3090.95110	-1308.56463	-105.50117	63.0
Root3		-1308.54800		73.4
S ₁ (1.20)				
Root1 (S ₀)	-3091.01710	-1308.60902		35.1
Root2 (S ₁)	-3090.94710	-1308.57653	-105.50027	55.5
Root3		-1308.53833		79.5
S ₁ (1.10)				
Root1 (S ₀)	-3091.00720	-1308.60194		39.6
Root2 (S ₁)	-3090.94770	-1308.59124	-105.49857	46.3
Root3		-1308.54742		73.8
S ₁ (1.00)				
Root1 (S ₀)	-3091.06940	-1308.62198		27.0
Root2 (S ₁)	-3091.06540	-1308.61778	-105.50200	29.6
Root3		-1308.55220	-	70.8

Table S4. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from AU-T1 (Reaction Coordinate: N5-H Distance).The Corresponding Energy Profiles are Plotted in Figure S7 (Blue Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
T₁-min (4.520)				
Root1 (T ₁)	-3091.04080	-1308.58753		48.6
Root2		-1308.56233	-105.50746	64.4
Root3		-1308.54523		75.2
T ₁ (4.50)		·	·	·
Root1 (T ₁)	-3091.04090	-1308.58722		48.8
Root2		-1308.56206	-105.50746	64.6
Root3		-1308.54506		75.3
T ₁ (4.40)		·	·	·
Root1 (T ₁)	-3091.04000	-1308.58735		48.7
Root2		-1308.56180	-105.50796	64.8
Root3		-1308.54458		75.6
T ₁ (4.20)				
Root1 (T ₁)	-3091.03740	-1308.58589		49.6
Root2		-1308.56006	-105.50830	65.9
Root3		-1308.54288		76.6
T ₁ (3.80)				
Root1 (T ₁)	-3091.03360	-1308.58403		50.8
Root2		-1308.57084	-105.50637	59.1
Root3		-1308.52407		88.4
T ₁ (3.40)				
Root1 (T ₁)	-3091.03210	-1308.57722		55.1
Root2		-1308.55166	-105.50395	71.1
Root3		-1308.53503		81.6
T ₁ (3.00)				
Root1 (T ₁)	-3091.03050	-1308.57546		56.2
Root2		-1308.54952	-105.50281	72.5
Root3		-1308.53239		83.2
T ₁ (2.60)				
Root1 (T ₁)	-3091.03120	-1308.57402		57.1
Root2		-1308.54806	-105.50084	73.4
Root3		-1308.52975		84.9
T ₁ (2.20)				
Root1 (T ₁)	-3091.02660	-1308.57428		56.9
Root2		-1308.54933	-105.50043	72.6
Root3		-1308.52990		84.8

T ₁ (1.80)				
Root1 (T ₁)	-3091.02000	-1308.57222		58.2
Root2		-1308.54798	-105.49881	73.4
Root3		-1308.52735		86.4
T ₁ (1.60)				
Root1 (T ₁)	-3091.00730	-1308.56784		61.0
Root2		-1308.54434	-105.49885	75.7
Root3		-1308.53202		83.5
T₁ (1.50)				
Root1 (T ₁)	-3090.99850	-1308.56201		64.6
Root2		-1308.53920	-105.49880	78.9
Root3		-1308.53188		83.5
T ₁ (1.40)				
Root1 (T ₁)	-3090.99330	-1308.56969		59.8
Root2		-1308.54328	-105.50522	76.4
Root3		-1308.53386		82.3
T₁ (1.30)				
Root1 (T ₁)	-3090.98700	-1308.57112		58.9
Root2		-1308.55354	-105.50315	69.9
Root3		-1308.53778		79.8
T₁ (1.20)				
Root1 (T ₁)	-3090.98190	-1308.58039		53.1
Root2		-1308.56137	-105.50260	65.0
Root3		-1308.54954		72.5
T ₁ (1.10)				
Root1 (T ₁)	-3091.06580	-1308.62072		27.8
Root2		-1308.61423	-105.50109	31.9
Root3		-1308.52486		87.9
T ₁ (1.00)				
Root1 (T ₁)	-3091.07250	-1308.62498		25.1
Root2		-1308.61834	-105.50141	29.3
Root3		-1308.53120		84.0

Table S5. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from AU-S0 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S7 (Green Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₀-min (4.569)				
Root1 (S ₀)	-3091.13640	-1308.66500		0.0
Root2 (S ₁)		-1308.56356	-105.50711	63.7
Root3		-1308.53980		78.6
S ₀ (4.50)				
Root1 (S ₀)	-3091.13570	-1308.66443		0.4
Root2 (S ₁)		-1308.56291	-105.50739	64.1
Root3		-1308.53952		78.7
S ₀ (4.40)				
Root1 (S ₀)	-3091.13480	-1308.66374		0.8
Root2 (S ₁)		-1308.56222	-105.50754	64.5
Root3		-1308.53898		79.1
S ₀ (4.20)				
Root1 (S ₀)	-3091.13270	-1308.66211		1.8
Root2 (S ₁)		-1308.56079	-105.50760	65.4
Root3		-1308.53783		79.8
S ₀ (3.80)				
Root1 (S ₀)	-3091.12900	-1308.65797		4.4
Root2 (S ₁)		-1308.55658	-105.50643	68.0
Root3		-1308.53360		82.5
S ₀ (3.40)				
Root1 (S ₀)	-3091.12740	-1308.65407		6.9
Root2 (S ₁)		-1308.55247	-105.50426	70.6
Root3		-1308.52875		85.5
S ₀ (3.00)				
Root1 (S ₀)	-3091.12550	-1308.65089		8.9
Root2 (S ₁)		-1308.54982	-105.50362	72.3
Root3		-1308.52562		87.5
S ₀ (2.60)				
Root1 (S ₀)	-3091.12620	-1308.64934		9.8
Root2 (S ₁)		-1308.54865	-105.50179	73.0
Root3		-1308.52349		88.8
S ₀ (2.20)				
Root1 (S ₀)	-3091.12200	-1308.64842		10.4
Root2 (S ₁)		-1308.54949	-105.50098	72.5
Root3		-1308.52503		87.8

S ₀ (1.80)				
Root1 (S ₀)	-3091.10950	-1308.64182		14.5
Root2 (S ₁)		-1308.53210	-105.50012	83.4
Root3		-1308.52375		88.6
S ₀ (1.60)		·	·	
Root1 (S ₀)	-3091.09900	-1308.63812		16.9
Root2 (S ₁)		-1308.53005	-105.49849	84.7
Root3		-1308.52558		87.5
S ₀ (1.50)				
Root1 (S ₀)	-3091.08920	-1308.63237		20.5
Root2 (S ₁)		-1308.53290	-105.49883	82.9
Root3		-1308.52688		86.7
S ₀ (1.40)				
Root1 (S ₀)	-3091.07710	-1308.62796		23.2
Root2 (S ₁)		-1308.54729	-105.49879	73.9
Root3		-1308.53636		80.7
S ₀ (1.30)				
Root1 (S ₀)	-3091.06370	-1308.62194		27.0
Root2 (S ₁)		-1308.55415	-105.49875	69.6
Root3		-1308.53254		83.1
S ₀ (1.20)				
Root1 (S ₀)	-3091.05520	-1308.61768		29.7
Root2 (S ₁)		-1308.56700	-105.50334	61.5
Root3		-1308.53889		79.1
S ₀ (1.10)				
Root1 (S ₀)	-3091.05150	-1308.61623		30.6
Root2 (S ₁)		-1308.59343	-105.49812	44.9
Root3		-1308.56599		62.1
S ₀ (1.00)				
Root1 (S ₀)	-3091.04240	-1308.61612		30.7
Root2 (S ₁)		-1308.60072	-105.50033	40.3
Root3		-1308.55969	[66.1

Table S6. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from AD-S1 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S8 (Red Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₁-min (4.900)				
Root1 (S ₀)	-3091.10810	-1308.65130		8.6
Root2 (S ₁)	-3091.00480	-1308.57346	-105.50419	57.4
Root3		-1308.54162		77.4
S ₁ (4.80)				
Root1 (S ₀)	-3091.10760	-1308.65157		8.4
Root2 (S ₁)	-3091.00460	-1308.57398	-105.50405	57.1
Root3		-1308.54228		77.0
S ₁ (4.60)				
Root1 (S ₀)	-3091.10660	-1308.65317		7.4
Root2 (S ₁)	-3091.00500	-1308.57599	-105.50246	55.9
Root3		-1308.54442		75.7
S ₁ (4.40)				
Root1 (S ₀)	-3091.10530	-1308.65336		7.3
Root2 (S ₁)	-3091.00430	-1308.57664	-105.50244	55.5
Root3		-1308.54564		74.9
S ₁ (4.20)				
Root1 (S ₀)	-3091.10490	-1308.65393		7.0
Root2 (S ₁)	-3091.00430	-1308.57724	-105.50225	55.1
Root3		-1308.54651		74.4
S ₁ (3.80)				
Root1 (S ₀)	-3091.10240	-1308.65346		7.2
Root2 (S ₁)	-3091.00260	-1308.57722	-105.50276	55.1
Root3		-1308.54635		74.5
S ₁ (3.40)				
Root1 (S ₀)	-3091.10220	-1308.65279		7.7
Root2 (S ₁)	-3091.00260	-1308.57676	-105.50320	55.4
Root3		-1308.54509		75.2
S ₁ (3.00)				
Root1 (S ₀)	-3091.10380	-1308.65212		8.1
Root2 (S ₁)	-3091.00370	-1308.57213	-105.50339	58.3
Root3		-1308.54373		76.1
S ₁ (2.60)				
Root1 (S ₀)	-3091.10290	-1308.65085		8.9
Root2 (S ₁)	-3091.00260	-1308.57136	-105.50369	58.8
Root3		-1308.54272		76.7

S ₁ (2.20)				
Root1 (S ₀)	-3091.09660	-1308.64866		10.3
Root2 (S ₁)	-3090.99790	-1308.57019	-105.50319	59.5
Root3		-1308.54217		77.1
S ₁ (1.80)				
Root1 (S ₀)	-3091.08240	-1308.64402		13.2
Root2 (S ₁)	-3090.98840	-1308.56818	-105.50203	60.8
Root3		-1308.54067		78.0
S ₁ (1.60)				
Root1 (S ₀)	-3091.07220	-1308.63673		17.7
Root2 (S ₁)	-3090.98070	-1308.56740	-105.50067	61.2
Root3		-1308.53459		81.8
S ₁ (1.50)				
Root1 (S ₀)	-3091.06160	-1308.63125		21.2
Root2 (S ₁)	-3090.97330	-1308.55436	-105.50075	69.4
Root3		-1308.54138		77.6
S ₁ (1.40)				
Root1 (S ₀)	-3091.04760	-1308.62273		26.5
Root2 (S ₁)	-3090.96260	-1308.55244	-105.50323	70.6
Root3		-1308.54795		73.5
S ₁ (1.30)				
Root1 (S ₀)	-3091.03410	-1308.61714		30.0
Root2 (S ₁)	-3090.95530	-1308.56564	-105.50321	62.4
Root3		-1308.55125		71.4
S ₁ (1.20)				
Root1 (S ₀)	-3091.02460	-1308.61207		33.2
Root2 (S ₁)	-3090.95100	-1308.57542	-105.50448	56.2
Root3		-1308.53836		79.5
S ₁ (1.10)				
Root1 (S ₀)	-3091.01630	-1308.61256		32.9
Root2 (S ₁)	-3090.95320	-1308.59377	-105.50082	44.7
Root3		-1308.55410		69.6
S ₁ (1.00)				
Root1 (S ₀)	-3091.07030	-1308.62333		26.2
Root2 (S ₁)	-3091.06580	-1308.61877	-105.50725	29.0
Root3		-1308.55534		68.8

Table S7. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from AD-T1 (Reaction Coordinate: N5-H Distance).The Corresponding Energy Profiles are Plotted in Figure S8 (Blue Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
T₁-min (4.897)				
Root1 (T ₁)	-3091.03690	-1308.58297		51.5
Root2		-1308.56861	-105.50232	60.5
Root3		-1308.54294		76.6
T ₁ (4.80)				
Root1 (T ₁)	-3091.03650	-1308.58371		51.0
Root2		-1308.56914	-105.50194	60.2
Root3		-1308.54313		76.5
T ₁ (4.60)				·
Root1 (T ₁)	-3091.03650	-1308.58523		50.1
Root2		-1308.57090	-105.50164	59.1
Root3		-1308.54421		75.8
T ₁ (4.40)				·
Root1 (T ₁)	-3091.03680	-1308.58684		49.1
Root2		-1308.57230	-105.50071	58.2
Root3		-1308.54554		75.0
T ₁ (4.20)				
Root1 (T ₁)	-3091.03480	-1308.58714		48.9
Root2		-1308.57255	-105.50067	58.0
Root3		-1308.54545		75.0
T ₁ (3.80)				
Root1 (T ₁)	-3091.03520	-1308.58798		48.3
Root2		-1308.57301	-105.50070	57.7
Root3		-1308.54538		75.1
T ₁ (3.40)				
Root1 (T ₁)	-3091.03520	-1308.58688		49.0
Root2		-1308.57104	-105.50108	59.0
Root3		-1308.54383		76.0
T ₁ (3.00)				
Root1 (T ₁)	-3091.03680	-1308.58499		50.2
Root2		-1308.56885	-105.50080	60.3
Root3		-1308.54212		77.1
T ₁ (2.60)				
Root1 (T ₁)	-3091.03540	-1308.58502		50.2
Root2		-1308.56886	-105.50109	60.3
Root3		-1308.54122		77.7

T ₁ (2.20)				
Root1 (T ₁)	-3091.03110	-1308.58612		49.5
Root2		-1308.57029	-105.50077	59.4
Root3		-1308.54211		77.1
T ₁ (1.80)				
Root1 (T ₁)	-3091.01570	-1308.57979		53.5
Root2		-1308.56204	-105.50091	64.6
Root3		-1308.53094		84.1
T ₁ (1.60)				
Root1 (T ₁)	-3091.01180	-1308.57903		53.9
Root2		-1308.55829	-105.49840	67.0
Root3		-1308.53135		83.9
T ₁ (1.50)				
Root1 (T ₁)	-3091.00400	-1308.57232		58.2
Root2		-1308.55677	-105.49860	67.9
Root3		-1308.54306		76.5
T ₁ (1.40)				
Root1 (T ₁)	-3090.99610	-1308.56989		59.7
Root2		-1308.56131	-105.49855	65.1
Root3		-1308.54418		75.8
T ₁ (1.30)				
Root1 (T ₁)	-3090.99010	-1308.57537		56.2
Root2		-1308.56774	-105.49880	61.0
Root3		-1308.54805		73.4
T ₁ (1.20)				
Root1 (T ₁)	-3090.98730	-1308.58098		52.7
Root2		-1308.56826	-105.49856	60.7
Root3		-1308.56132		65.1
T ₁ (1.10)				
Root1 (T ₁)	-3090.98760	-1308.59838		41.8
Root2		-1308.58395	-105.49799	50.9
Root3		-1308.55184	ŀ	71.0
T ₁ (1.00)				
Root1 (T ₁)	-3090.98680	-1308.62518		25.0
Root2		-1308.61738	-105.49699	29.9
Root3		-1308.53195	Ē	83.5

Table S8. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from AD-S0 (Reaction Coordinate: N5-H Distance).The Corresponding Energy Profiles are Plotted in Figure S8 (Green Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₀-min (4.910)				
Root1 (S ₀)	-3091.13820	-1308.65770		4.6
Root2 (S ₁)		-1308.55653	-105.50662	68.1
Root3		-1308.53173		83.6
S ₀ (4.80)				
Root1 (S ₀)	-3091.13750	-1308.65754		4.7
Root2 (S ₁)		-1308.55679	-105.50675	67.9
Root3		-1308.53223		83.3
S ₀ (4.60)				
Root1 (S ₀)	-3091.13760	-1308.65843		4.1
Root2 (S ₁)		-1308.55796	-105.50637	67.2
Root3		-1308.53396		82.2
S ₀ (4.40)		·	•	
Root1 (S ₀)	-3091.13470	-1308.65798	-105.50625	4.4
Root2 (S ₁)		-1308.55787		67.2
Root3		-1308.53424		82.1
S ₀ (4.20)		·	•	
Root1 (S ₀)	-3091.13560	-1308.65965		3.4
Root2 (S ₁)		-1308.55897	-105.50653	66.5
Root3		-1308.53480		81.7
S ₀ (3.80)		·	•	
Root1 (S ₀)	-3091.13460	-1308.65915		3.7
Root2 (S ₁)		-1308.55893	-105.50664	66.6
Root3		-1308.53497		81.6
S ₀ (3.40)		·	·	·
Root1 (S ₀)	-3091.13330	-1308.65722		4.9
Root2 (S ₁)		-1308.55732	-105.50701	67.6
Root3		-1308.53282		83.0
S ₀ (3.00)				
Root1 (S ₀)	-3091.13660	-1308.65451		6.6
Root2 (S ₁)		-1308.55433	-105.50664	69.5
Root3		-1308.52804		86.0
S ₀ (2.60)				
Root1 (S ₀)	-3091.13300	-1308.65368		7.1
Root2 (S ₁)		-1308.55444	-105.50764	69.4
Root3		-1308.52818		85.9

S ₀ (2.20)				
Root1 (S ₀)	-3091.12890	-1308.65327		7.4
Root2 (S ₁)		-1308.55502	-105.50667	69.0
Root3		-1308.52891		85.4
S ₀ (1.80)				
Root1 (S ₀)	-3091.11680	-1308.64744		11.0
Root2 (S ₁)		-1308.53767	-105.50525	79.9
Root3		-1308.52591		87.3
S ₀ (1.60)				
Root1 (S ₀)	-3091.10300	-1308.63999		15.7
Root2 (S ₁)		-1308.53306	-105.50552	82.8
Root3		-1308.52659		86.9
S ₀ (1.50)		·		
Root1 (S ₀)	-3091.09300	-1308.63517		18.7
Root2 (S ₁)		-1308.54033	-105.50558	78.2
Root3		-1308.53942		78.8
S ₀ (1.40)		·		
Root1 (S ₀)	-3091.08070	-1308.62903	-105.50554	22.6
Root2 (S ₁)		-1308.54767		73.6
Root3		-1308.53862		79.3
S ₀ (1.30)		·		
Root1 (S ₀)	-3091.06770	-1308.62480		25.2
Root2 (S ₁)		-1308.55530	-105.50483	68.8
Root3		-1308.52791		86.0
S ₀ (1.20)		·		
Root1 (S ₀)	-3091.05880	-1308.61889		28.9
Root2 (S ₁)		-1308.56866	-105.50813	60.5
Root3		-1308.54042		78.2
S ₀ (1.10)		·		
Root1 (S ₀)	-3091.05200	-1308.61497		31.4
Root2 (S ₁)		-1308.59183	-105.50463	45.9
Root3		-1308.56488		62.8
S ₀ (1.00)		·		
Root1 (S ₀)	-3091.05150	-1308.62264		26.6
Root2 (S ₁)		-1308.59981	-105.50319	40.9
Root3		-1308.56417		63.3

Table S9. Absolute Energies (A.E., Hartree), Relative Energies (Δ E, kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from BU-S1 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S9 (Red Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₁-min (3.779)				
Root1 (S ₀)	-3091.10120	-1308.64901		10.0
Root2 (S ₁)	-3090.99420	-1308.57422	-105.50836	57.0
Root3		-1308.54378		76.1
S ₁ (3.60)				
Root1 (S ₀)	-3091.10090	-1308.65203		8.1
Root2 (S ₁)	-3090.99400	-1308.56964	-105.50829	59.8
Root3		-1308.53937		78.8
S ₁ (3.40)				
Root1 (S ₀)	-3091.09930	-1308.65167		8.4
Root2 (S ₁)	-3090.99360	-1308.56973	-105.50754	59.8
Root3		-1308.53924		78.9
S ₁ (3.00)				
Root1 (S ₀)	-3091.09650	-1308.65059		9.0
Root2 (S ₁)	-3090.99210	-1308.56904	-105.50570	60.2
Root3		-1308.53896		79.1
S ₁ (2.60)				
Root1 (S ₀)	-3091.09140	-1308.64705		11.3
Root2 (S ₁)	-3090.98920	-1308.56689	-105.50528	61.6
Root3		-1308.53652		80.6
S ₁ (2.20)				
Root1 (S ₀)	-3091.08770	-1308.64560		12.2
Root2 (S ₁)	-3090.98630	-1308.56617	-105.50387	62.0
Root3		-1308.53559		81.2
S ₁ (1.80)		·		
Root1 (S ₀)	-3091.07640	-1308.64061		15.3
Root2 (S ₁)	-3090.97790	-1308.56333	-105.50368	63.8
Root3		-1308.53503		81.6
S ₁ (1.60)				
Root1 (S ₀)	-3091.06120	-1308.63307		20.0
Root2 (S ₁)	-3090.96620	-1308.55840	-105.50413	66.9
Root3		-1308.53287		82.9
S ₁ (1.50)				
Root1 (S ₀)	-3091.05000	-1308.62510		25.0
Root2 (S ₁)	-3090.95900	-1308.54754	-105.50412	73.7
Root3		-1308.53721		80.2

S ₁ (1.40)				
Root1 (S ₀)	-3091.04400	-1308.62523	-105.50173	25.0
Root2 (S ₁)	-3090.95450	-1308.55087		71.6
Root3		-1308.54781		73.5
S ₁ (1.30)				
Root1 (S ₀)	-3091.03070	-1308.61867		29.1
Root2 (S ₁)	-3090.94760	-1308.56559	-105.50194	62.4
Root3		-1308.55081		71.7
S ₁ (1.20)				
Root1 (S ₀)	-3091.01720	-1308.61269		32.8
Root2 (S ₁)	-3090.94370	-1308.57621	-105.50131	55.7
Root3		-1308.53925		78.9
S₁ (1.10)				
Root1 (S ₀)	-3091.01080	-1308.61485		31.5
Root2 (S ₁)	-3090.94790	-1308.59514	-105.49579	43.8
Root3		-1308.55641		68.1

Table S10. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from BU-T1 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S9 (Blue Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
T₁-min (3.886)				
Root1 (T ₁)	-3091.04450	-1308.58701		48.9
Root2		-1308.56492	-105.51578	62.8
Root3		-1308.54274		76.7
T ₁ (3.60)		·		
Root1 (T ₁)	-3091.04420	-1308.58603		49.6
Root2		-1308.56425	-105.51542	63.2
Root3		-1308.54097		77.8
T ₁ (3.40)		·		
Root1 (T ₁)	-3091.04230	-1308.58607		49.5
Root2		-1308.56568	-105.51544	62.3
Root3		-1308.54273		76.7
T ₁ (3.00)		·		
Root1 (T ₁)	-3091.04020	-1308.58066		52.9
Root2		-1308.55477	-105.51435	69.2
Root3		-1308.54076		78.0
T ₁ (2.60)				

Root1 (T ₁)	-3091.03870	-1308.58484		50.3
Root2		-1308.56402	-105.51224	63.4
Root3		-1308.52003		91.0
T ₁ (2.20)				
Root1 (T ₁)	-3091.03690	-1308.57857		54.2
Root2		-1308.55491	-105.51086	69.1
Root3		-1308.54275		76.7
T ₁ (1.80)				
Root1 (T ₁)	-3091.02780	-1308.57537		56.2
Root2		-1308.55165	-105.51124	71.1
Root3		-1308.49741		105.2
T ₁ (1.60)				
Root1 (T ₁)	-3091.01640	-1308.57141		58.7
Root2		-1308.54618	-105.51165	74.6
Root3		-1308.51292		95.4
T ₁ (1.50)				
Root1 (T ₁)	-3091.00900	-1308.56926	-105.51175	60.1
Root2		-1308.53707		80.3
Root3		-1308.53389		82.3
T₁ (1.40)				
Root1 (T ₁)	-3091.00050	-1308.56850		60.6
Root2		-1308.54393	-105.51169	76.0
Root3		-1308.53194		83.5
T ₁ (1.30)				
Root1 (T ₁)	-3090.99180	-1308.57149		58.7
Root2		-1308.55506	-105.51164	69.0
Root3		-1308.53823		79.6
T ₁ (1.20)				
Root1 (T ₁)	-3090.98730	-1308.58243		51.8
Root2		-1308.56305	-105.51038	64.0
Root3		-1308.54998		72.2
T ₁ (1.10)				
Root1 (T ₁)	-3091.06640	-1308.62293		26.4
Root2		-1308.61589	-105.50740	30.8
Root3		-1308.52889		85.4

Table S11. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from BU-S0 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S9 (Green Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₀-min (3.799)				
Root1 (S ₀)	-3091.13080	-1308.65591		5.7
Root2 (S ₁)		-1308.55442	-105.51084	69.4
Root3		-1308.52690		86.7
S ₀ (3.60)				•
Root1 (S ₀)	-3091.12970	-1308.65604		5.6
Root2 (S ₁)		-1308.55481	-105.51096	69.1
Root3		-1308.52823		85.8
S ₀ (3.40)				
Root1 (S ₀)	-3091.12850	-1308.65563		5.9
Root2 (S ₁)		-1308.55485	-105.51088	69.1
Root3		-1308.52832		85.8
S ₀ (3.00)				
Root1 (S ₀)	-3091.12530	-1308.65419		6.8
Root2 (S ₁)		-1308.55388	-105.51026	69.7
Root3		-1308.52722		86.5
S ₀ (2.60)				
Root1 (S ₀)	-3091.12250	-1308.65163		8.4
Root2 (S ₁)		-1308.55198	-105.50858	70.9
Root3		-1308.52501		87.9
S ₀ (2.20)				
Root1 (S ₀)	-3091.11850	-1308.64934		9.8
Root2 (S ₁)		-1308.55035	-105.50727	71.9
Root3		-1308.52258		89.4
S ₀ (1.80)				
Root1 (S ₀)	-3091.10790	-1308.64591		12.0
Root2 (S ₁)		-1308.54931	-105.50696	72.6
Root3		-1308.52072		90.5
S ₀ (1.60)				
Root1 (S ₀)	-3091.09480	-1308.63868		16.5
Root2 (S ₁)		-1308.53015	-105.50699	84.6
Root3		-1308.52118		90.3
S ₀ (1.50)				
Root1 (S ₀)	-3091.08530	-1308.63330		19.9
Root2 (S ₁)		-1308.53207	-105.50702	83.4
Root3		-1308.52525		87.7

S ₀ (1.40)				
Root1 (S ₀)	-3091.07270	-1308.62768	-105.50711	23.4
Root2 (S ₁)		-1308.54736		73.8
Root3		-1308.52818		85.9
S ₀ (1.30)				
Root1 (S ₀)	-3091.06580	-1308.62778		23.4
Root2 (S ₁)		-1308.55527	-105.50445	68.9
Root3		-1308.52514		87.8
S ₀ (1.20)				
Root1 (S ₀)	-3091.05110	-1308.61975		28.4
Root2 (S ₁)		-1308.56851	-105.50463	60.6
Root3		-1308.54016		78.3
S ₀ (1.10)				
Root1 (S ₀)	-3091.04790	-1308.61694		30.2
Root2 (S ₁)		-1308.59297	-105.50026	45.2
Root3		-1308.56664		61.7

Table S12. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from BD-S1 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S10 (Red Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₁-min (3.260)				
Root1 (S ₀)	-3091.10440	-1308.64520		12.4
Root2 (S ₁)	-3090.99350	-1308.56290	-105.51260	64.1
Root3		-1308.53104		84.1
S ₁ (3.10)		·		
Root1 (S ₀)	-3091.10300	-1308.64576	-105.51199	12.1
Root2 (S ₁)	-3090.99370	-1308.56398		63.4
Root3		-1308.53203		83.4
S ₁ (2.90)		·		
Root1 (S ₀)	-3091.10130	-1308.64527		12.4
Root2 (S ₁)	-3090.99340	-1308.56429	-105.51176	63.2
Root3		-1308.53214		83.4
S ₁ (2.70)		•		
Root1 (S ₀)	-3091.10040	-1308.64596	105 51114	12.0
Root2 (S ₁)	-3090.99300	-1308.56540	-105.51144	62.5

Root3		-1308.53308		82.8
S ₁ (2.50)		·		
Root1 (S ₀)	-3091.10000	-1308.64679		11.4
Root2 (S ₁)	-3090.99410	-1308.56652	-105.51040	61.8
Root3		-1308.53464		81.8
S ₁ (2.30)				
Root1 (S ₀)	-3091.09910	-1308.64722		11.2
Root2 (S ₁)	-3090.99510	-1308.56768	-105.50977	61.1
Root3		-1308.53637		80.7
S ₁ (2.10)				
Root1 (S ₀)	-3091.09370	-1308.64518		12.4
Root2 (S ₁)	-3090.99170	-1308.56686	-105.51026	61.6
Root3		-1308.53562	100101020	81.2
S ₁ (1.90)		·		
Root1 (S ₀)	-3091.08620	-1308.64167		14.6
Root2 (S ₁)	-3090.98660	-1308.56458	-105.51076	63.0
Root3		-1308.53525		81.4
S ₁ (1.70)				
Root1 (S ₀)	-3091.07450	-1308.63614		18.1
Root2 (S ₁)	-3090.97800	-1308.56087	-105.51107	65.3
Root3		-1308.53330		82.7
S ₁ (1.60)				
Root1 (S ₀)	-3091.06510	-1308.63069		21.5
Root2 (S ₁)	-3090.97210	-1308.55768	-105.51107	67.3
Root3		-1308.53175		83.6
S ₁ (1.50)				
Root1 (S ₀)	-3091.05920	-1308.62819		23.1
Root2 (S ₁)	-3090.96700	-1308.55572	-105.51076	68.6
Root3		-1308.53566		81.2
S ₁ (1.40)				
Root1 (S ₀)	-3091.04590	-1308.61924		28.7
Root2 (S ₁)	-3090.95890	-1308.54933	-105.51081	72.6
Root3		-1308.54426		75.8
S ₁ (1.30)				
Root1 (S ₀)	-3091.03180	-1308.61323		32.5
Root2 (S ₁)	-3090.95130	-1308.55927	-105.51076	66.4
Root3		-1308.54120		77.7
S ₁ (1.20)				
Root1 (S ₀)	-3091.01840	-1308.60670		36.6
Root2 (S ₁)	-3090.94840	-1308.57421	-105.50993	57.0
Root3		-1308.53610		80.9
S ₁ (1.10)				
Root1 (S ₀)	-3091.00880	-1308.60117		40.1
Root2 (S ₁)	-3090.94960	-1308.58936	-105.50858	47.5
Root3		-1308.54741		73.8

Table S13. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from BD-T1 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S10 (Blue Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
T₁-min (3.264)				
Root1 (T ₁)	-3091.04220	-1308.57619		55.7
Root2		-1308.55114	-105.51359	71.5
Root3		-1308.54145		77.5
T ₁ (3.10)		·	•	
Root1 (T ₁)	-3091.04190	-1308.57650		55.5
Root2		-1308.55182	-105.51372	71.0
Root3		-1308.54056		78.1
T ₁ (2.90)		·	•	
Root1 (T ₁)	-3091.04110	-1308.57664		55.5
Root2		-1308.55197	-105.51376	70.9
Root3		-1308.54008		78.4
T ₁ (2.70)		·	•	
Root1 (T ₁)	-3091.04160	-1308.57714		55.1
Root2		-1308.55219	-105.51340	70.8
Root3		-1308.54033		78.2
T ₁ (2.50)				
Root1 (T ₁)	-3091.04060	-1308.57702		55.2
Root2		-1308.54925	-105.51320	72.6
Root3		-1308.53111		84.0
T ₁ (2.30)		·	•	
Root1 (T ₁)	-3091.03930	-1308.57744		55.0
Root2		-1308.54977	-105.51303	72.3
Root3		-1308.53077		84.2
T ₁ (2.10)		·	•	
Root1 (T ₁)	-3091.03600	-1308.57663		55.5
Root2		-1308.54905	-105.51331	72.8
Root3		-1308.52903		85.3
T ₁ (1.90)				
Root1 (T ₁)	-3091.03090	-1308.57369		57.3
Root2		-1308.55041	105.51371	71.9
Root3		-1308.49560		106.3
T ₁ (1.70)				
Root1 (T ₁)	-3091.02290	-1308.57149		58.7
Root2		-1308.54739	-105.51387	73.8
Root3		-1308.50031		103.4

T ₁ (1.60)				
Root1 (T ₁)	-3091.01660	-1308.56966		59.8
Root2		-1308.54510	-105.51394	75.2
Root3		-1308.52329		88.9
T ₁ (1.50)		·		
Root1 (T ₁)	-3091.00840	-1308.56738		61.3
Root2		-1308.53960	-105.51405	78.7
Root3		-1308.53197		83.5
T ₁ (1.40)				
Root1 (T ₁)	-3091.00080	-1308.56786		61.0
Root2		-1308.54329	-105.51387	76.4
Root3		-1308.53008		84.7
T ₁ (1.30)				
Root1 (T ₁)	-3090.99160	-1308.57188		58.4
Root2		-1308.55267	-105.51384	70.5
Root3		-1308.53737		80.1
T ₁ (1.20)				
Root1 (T ₁)	-3090.98690	-1308.58128		52.5
Root2		-1308.56103	-105.51244	65.2
Root3		-1308.54776		73.6
T ₁ (1.10)				
Root1 (T ₁)	-3091.06850	-1308.62097		27.6
Root2		-1308.61441	-105.50906	31.8
Root3		-1308.52694		86.6

Table S14. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for N5-H Bond Formation from BD-S0 (Reaction Coordinate: N5-H Distance). The Corresponding Energy Profiles are Plotted in Figure S10 (Green Line).

(N5-H)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₀-min (3.291)				
Root1 (S ₀)	-3091.13630	-1308.65013		9.3
Root2 (S ₁)		-1308.54893	-105.51498	72.8
Root3		-1308.52006		91.0
S ₀ (3.10)				
Root1 (S ₀)	-3091.13570	-1308.65021		9.3
Root2 (S ₁)		-1308.54932	-105.51497	72.6
Root3		-1308.52085		90.5
S ₀ (2.90)				

Root1 (S ₀)	-3091.13590	-1308.65058		9.1
Root2 (S ₁)		-1308.54988	-105.51462	72.2
Root3		-1308.52104		90.3
S ₀ (2.70)				
Root1 (S ₀)	-3091.13420	-1308.65042		9.2
Root2 (S ₁)		-1308.54999	-105.51463	72.2
Root3		-1308.52111		90.3
S ₀ (2.50)				
Root1 (S ₀)	-3091.13400	-1308.65074		9.0
Root2 (S ₁)		-1308.55073	-105.51422	71.7
Root3		-1308.52141		90.1
S ₀ (2.30)				
Root1 (S ₀)	-3091.13070	-1308.64971		9.6
Root2 (S ₁)		-1308.55049	-105.51443	71.9
Root3		-1308.52127		90.2
S ₀ (2.10)				
Root1 (S ₀)	-3091.12840	-1308.64875		10.2
Root2 (S ₁)		-1308.53632	-105.51422	80.8
Root3		-1308.52004		91.0
S ₀ (1.90)				
Root1 (S ₀)	-3091.12140	-1308.64606		11.9
Root2 (S ₁)		-1308.53485	-105.51458	81.7
Root3		-1308.52001		91.0
S ₀ (1.70)				
Root1 (S ₀)	-3091.11170	-1308.64173		14.6
Root2 (S ₁)		-1308.53256	-105.51453	83.1
Root3		-1308.52081		90.5
S ₀ (1.60)				
Root1 (S ₀)	-3091.10320	-1308.63697		17.6
Root2 (S ₁)		-1308.52985	-105.51466	84.8
Root3		-1308.52275		89.3
S₀ (1.50)				
Root1 (S ₀)	-3091.09340	-1308.63163		20.9
Root2 (S ₁)		-1308.53501	-105.51476	81.6
Root3		-1308.52518		87.7
S ₀ (1.40)				
Root1 (S ₀)	-3091.08100	-1308.62617		24.4
Root2 (S ₁)		-1308.54803	-105.51480	73.4
Root3		-1308.52561		87.5
S ₀ (1.30)				
Root1 (S ₀)	-3091.07370	-1308.62697		23.9
Root2 (S ₁)		-1308.55708	-105.51064	67.7
Root3		-1308.52466		88.1
S ₀ (1.20)				
Root1 (S ₀)	-3091.06130	-1308.62124		27.5
Root2 (S ₁)		-1308.57157	-105.51066	58.6
Root3		-1308.54206		77.1

S ₀ (1.10)				
Root1 (S ₀)	-3091.05310	-1308.61464		31.6
Root2 (S ₁)		-1308.59067	-105.50960	46.6
Root3		-1308.55920		66.4

Table S15. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM energies (Hartree) of Optimized Structures for C4a-S Bond Formation from INT-S1 (Reaction Coordinate: C4a-S Distance). The Corresponding Energy Profiles are Plotted in Figure 3 (Red Line).

(C4a-S)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₁-min (3.727)				
Root1 (S ₀)	-3091.06960	-1308.62204		27.0
Root2 (S ₁)	-3091.06560	-1308.61784	-105.50212	29.6
Root3		-1308.55210		70.8
S ₁ (3.70)			•	
Root1 (S ₀)	-3091.06980	-1308.62239		26.7
Root2 (S ₁)	-3091.06590	-1308.61814	-105.50233	29.4
Root3		-1308.55296		70.3
S ₁ (3.50)			•	
Root1 (S ₀)	-3091.06810	-1308.62297		26.4
Root2 (S ₁)	-3091.06500	-1308.61839	-105.50358	29.3
Root3		-1308.55594		68.4
S ₁ (3.30)		·	•	·
Root1 (S ₀)	-3091.06570	-1308.62388		25.8
Root2 (S ₁)	-3091.06290	-1308.61842	-105.50401	29.2
Root3		-1308.55912		66.4
S ₁ (3.10)				
Root1 (S ₀)	-3091.06200	-1308.62536		24.9
Root2 (S ₁)	-3091.05890	-1308.61744	-105.50420	29.9
Root3		-1308.56080		65.4
S ₁ (2.90)				
Root1 (S ₀)	-3091.06010	-1308.62854		22.9
Root2 (S ₁)	-3091.05340	-1308.61673	-105.50352	30.3
Root3		-1308.56177		64.8
S ₁ (2.70)				
Root1 (S ₀)	-3091.05930	-1308.62827		23.1
Root2 (S ₁)	-3091.04240	-1308.60975	-105.50336	34.7
Root3		-1308.55685		67.9
S ₁ (2.50)				
Root1 (S ₀)	-3091.05980	-1308.62571	105 50279	24.7
Root2 (S ₁)	-3091.02200	-1308.59464	-103.30370	44.2

Root3		-1308.53768		79.9
S ₁ (2.30)				
Root1 (S ₀)	-3091.06630	-1308.63122		21.2
Root2 (S ₁)	-3090.99640	-1308.57892	-105.50313	54.0
Root3		-1308.53869		79.3
S ₁ (2.10)				
Root1 (S ₀)	-3091.05250	-1308.62154		27.3
Root2 (S ₁)	-3090.96300	-1308.57245	-105.50342	58.1
Root3		-1308.52678		86.7
S ₁ (1.90)				
Root1 (S ₀)	-3091.03290	-1308.61128		33.7
Root2 (S ₁)	-3090.93200	-1308.56566	-105.50374	62.3
Root3		-1308.51041		97.0

Table S16. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for C4a-S Bond Formation from INT-T1 (Reaction Coordinate: C4a-S Distance). The Corresponding Energy Profiles are Plotted in Figure 3 (Blue Line).

(C4a-S)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
	QM+QM-MM (ele)	QM+QM-MM (ele)	MM + QM- MM (vdw)	
T₁-min (3.720)				
Root1 (T ₁)	-3091.07250	-1308.62499		25.1
Root2		-1308.61834	-105.50141	29.3
Root3		-1308.53120		84.0
T₁ (3.70)				
Root1 (T ₁)	-3091.07240	1308.62499		25.1
Root2		-1308.61838	-105.50143	29.3
Root3		-1308.53126		83.9
T ₁ (3.50)				
Root1 (T ₁)	-3091.07030	-1308.62527		24.9
Root2		-1308.61792	-105.50260	29.5
Root3		-1308.53170		83.6
T ₁ (3.30)				
Root1 (T ₁)	-3091.06770	-1308.62482		25.2
Root2		-1308.61782	-105.50308	29.6
Root3		-1308.53246		83.2
T ₁ (3.10)				
Root1 (T ₁)	-3091.06410	-1308.62493	-105.50363	25.2

Root2		-1308.61766		29.7
Root3		-1308.53534		81.4
T ₁ (2.90)				
Root1 (T ₁)	-3091.05850	-1308.62062		27.9
Root2		-1308.61201	-105.50344	33.3
Root3		-1308.53109		84.0
T ₁ (2.70)		·		
Root1 (T ₁)	-3091.05170	-1308.61862		29.1
Root2		-1308.60563	-105.50328	37.3
Root3		-1308.53094		84.1
T ₁ (2.50)		·		
Root1 (T ₁)	-3091.03880	-1308.61386		32.1
Root2		-1308.59239	-105.50327	45.6
Root3		-1308.52552		87.5
T ₁ (2.30)				
Root1 (T ₁)	-3091.01810	-1308.60428		38.1
Root2		-1308.57093	-105.50312	59.0
Root3		-1308.53936		78.8
T ₁ (2.10)				
Root1 (T ₁)	-3090.99190	-1308.59626		43.1
Root2		-1308.55462	-105.50176	69.3
Root3		-1308.53567		81.2
T ₁ (1.90)		·		
Root1 (T ₁)	-3090.99630	-1308.58099		52.7
Root2		-1308.53455	-105.49963	81.9
Root3		-1308.52322		89.0

Table S17. Absolute Energies (A.E., Hartree), Relative Energies (ΔE , kcal/mol), and MM Energies (Hartree) of Optimized Structures for C4a-S Bond Formation from INT-S0 (Reaction Coordinate: C4a-S Distance). The Corresponding Energy Profiles are Plotted in Figure 3 (Green Line).

(C4a-S)	CASSCF A.E.	MS-CASPT2 A.E.	ММ	MS-CASPT2 ∆E
S₀-min (3.372)				
Root1 (S ₀)	-3091.04460	-1308.61977		28.4
Root2 (S ₁)		-1308.59870	-105.50016	41.6
Root3		-1308.56020		65.8
S ₀ (3.10)				
Root1 (S ₀)	-3091.04090	-1308.61996		28.3
Root2 (S ₁)		-1308.59628	-105.50173	43.1
Root3		-1308.55773		67.3

S ₀ (2.90)				
Root1 (S ₀)	-3091.04760	-1308.62686		23.9
Root2 (S ₁)		-1308.60095	-105.49963	40.2
Root3		-1308.55873		66.7
S ₀ (2.70)				
Root1 (S ₀)	-3091.07400	-1308.62980		22.1
Root2 (S ₁)		-1308.59564	-105.50264	43.5
Root3		-1308.54615		74.6
S ₀ (2.50)				
Root1 (S ₀)	-3091.08080	-1308.63161		21.0
Root2 (S ₁)		-1308.57746	-105.50288	54.9
Root3		-1308.53143		83.8
S ₀ (2.30)				
Root1 (S ₀)	-3091.09170	-1308.63893		16.4
Root2 (S ₁)		-1308.55375	-105.50294	69.8
Root3		-1308.53809		79.6
S ₀ (2.10)				
Root1 (S ₀)	-3091.10640	-1308.65158		8.4
Root2 (S ₁)		-1308.54295	-105.50411	76.6
Root3		-1308.52785		86.1
S ₀ (1.90)				
Root1 (S ₀)	-3091.10990	-1308.65142		8.5
Root2 (S ₁)		-1308.53769	-105.50470	79.9
Root3		-1308.51004		97.2

5. Comparison with Previous Theoretical Work

On the computational side, early TD-DFT and MCQDPT2 calculations addressed model systems of a photo-LOV1 domain from *Chlamydomonas reinhardtii*. [30, 31] The formation of the C4a-S covalent adduct in this photo-LOV1 domain was explored in the T₁ state at the QM/MM level using the restricted open-shell Hartree-Fock method. [32] This process was later reinvestigated in the S₀ and T₁ states using high-level CASSCF and MCQDPT2 calculations on a model system consisting of lumiflavin and thiomethanol, without considering the protein environment. [33] The photophysics and photochemistry of structurally modified deaza flavin derivatives in the YtvA LOV domain were investigated both experimentally and at the QM/MM level. [34] Recent QM/MM studies also addressed the photophysical properties of a LOV-based fluorescent protein (iLOV-Q489K). [35, 36] In the realm of the

YtvA photoreceptors, the spectroscopic properties of the YtvA LOV domain from *Bacillus subtilis* were explored at the TD-DFT and DFT/MRCI levels. [37]

At the QM(CASSCF)/MM level, we optimized four S₀ conformers of the dark-adapted state, which are referred to as AU, AD, BU, and BD in Fig. 2. In AU and AD, the cysteine residue is oriented toward the dimethyl substituted part of the flavin core, while it points toward the two carbonyl groups of the flavin in BU and BD. At the QM(MS-CASPT2)/MM level, AU is predicted to be most stable (Table S18), in agreement with previous work. [37] We also optimized the corresponding S₁ and T₁ conformers at the QM(CASSCF)/MM level. In the S₁ state, AU is still the most stable conformer, about 5 kcal/mol lower than the others at the QM(MS-CASPT2)/MM level, while BU is the lowest minimum in the T₁ state, 4.9 kcal/mol lower than AU (see Table S18). One should note that only the BD conformer was considered in previous theoretical calculations of photo-LOV1 domains, [30–33] although it is obviously not the most stable in our YtvA photoreceptor.

Table S18. QM(MS-CASPT2(12,9)/cc-pVTZ//CASSCF(10,8)/6-31G*)/MM Calculated Relative Energies (kcal/mol) of Reactants, Intermediates, Products, and Transition States in the S_0 , T_1 , and S_1 Electronic States.

	AU	AD	BU	BD	INT	Р
S ₀	0.0	4.9	3.4	4.4	32.7	10.0
$S_0(TS)$	36.3	33.0	34.9	36.3	-	-
T ₁	48.4	54.5	43.5	51.7	28.7	57.3
T₁(TS)	63.2	65.1	57.7	56.9	-	-
S_1	54.8	59.3	59.6	60.6	32.7	62.9
S ₁ (TS)	76.0	73.1	75.0	70.3	-	-

Only one intermediate and one adduct product could be located in the S_0 state at the QM(CASSCF)/MM level. In photo-LOV1 domains, the S_0 adduct product was previously estimated to be 8.7 kcal/mol higher in energy than the BD conformer at the QM(RHF)/MM level [32] which is close to our QM(MS-CASPT2)/MM value of 5.6 kcal/mol. In MCQDPT2//CASSCF (2,2) model calculations, this S_0 adduct was computed to be 12.8 kcal/mol lower than BD; [33] this qualitative discrepancy might be caused by the neglect of polarization effects from the surroundings. The energy of the T₁ intermediate was previously found to be 17.6 kcal/mol relative to BD at the QM(ROHF)/MM

level [32] which is smaller than our QM(MS-CASPT2)/MM value of 24.3 kcal/mol (see Table S18).

The computed $S_0 \rightarrow S_1$ vertical excitation energies at the four S_0 reactant minima are nearly the same, 63.5 kcal/mol for AD and BD, and 63.7 kcal/mol for AU and BU. They are very close to the experimentally measured absorption band maximum of 447 nm [63.7 kcal/mol] [38] and previous QM/MM results. [37] Model calculations on the isolated chromophore gave higher $S_0 \rightarrow S_1$ vertical excitation energies of 75.9 and 77.6 kcal/mol [33] at the MCQDPT2//CASSCF(2,2) and MCQDPT2//CASSCF(4,4) levels, respectively.

The computed $S_0 \rightarrow S_1$ vertical excitation energy of 70.4 kcal/mol [406 nm] at the S_0 product minimum is slightly lower than the experimentally measured absorption band maximum of 390 nm [73.3 kcal/mol]. [38] Model calculations on the isolated chromophore again gave higher values of 86.3 and 77.7 kcal/mol at the MCQDPT2//CASSCF(2,2) and MCQDPT2//CASSCF(4,4) levels, respectively. [33]

The photochemical reaction from the dark-adapted starts with an excitedstate hydrogen transfer that produces a diradical intermediate. We computed the corresponding minimum-energy pathways for all four conformers in the S₁ and T₁ states. The energies of the computed transition states range between 70.3-76.0 kcal/mol in the S₁ case and between 56.9-65.1 kcal/mol in the T₁ case at the QM(MS-CASPT2)/MM level (see Table S18). We note that previous theoretical calculations for photo-LOV1 domains at the QM(ROHF)/MM and MCQDPT2//CASSCF(2,2) levels predicted the T₁ energy of the BD transition state to be 81.0 and 61.4 kcal/mol, respectively, relative to the S₀ BD minimum. [32, 33]

6. References

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7. Cartesian Coordinates of All Optimized Structures (QM part) In xyz format (unit: angström)

CASSCF method

AU-S0

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