

Application of Engineered Myoglobins for Biosynthesis of Clofazimine by Integration with Chemical Synthesis

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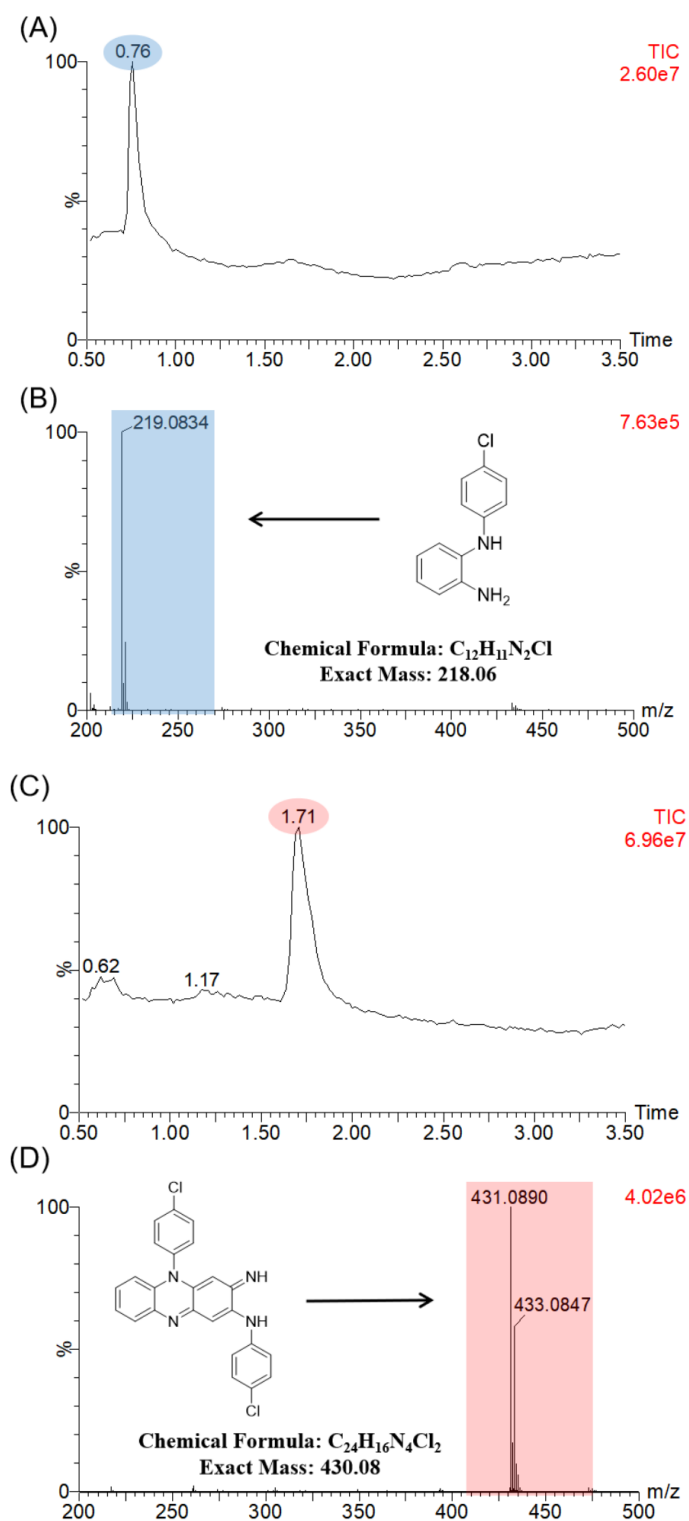


Figure S1. UPLC-ESI-MS traces of (A), (B) substrate (0.25 mM, calculated, 218.06 Da; observed, 219.08 Da, $[M+H]^+$), and (C), (D) the oxidative coupling product catalyzed by F43Y/T67R Mb in the presence of 2 mM H_2O_2 at 25 °C (calculated, 430.08 Da; observed, 431.09 Da, $[M+H]^+$).

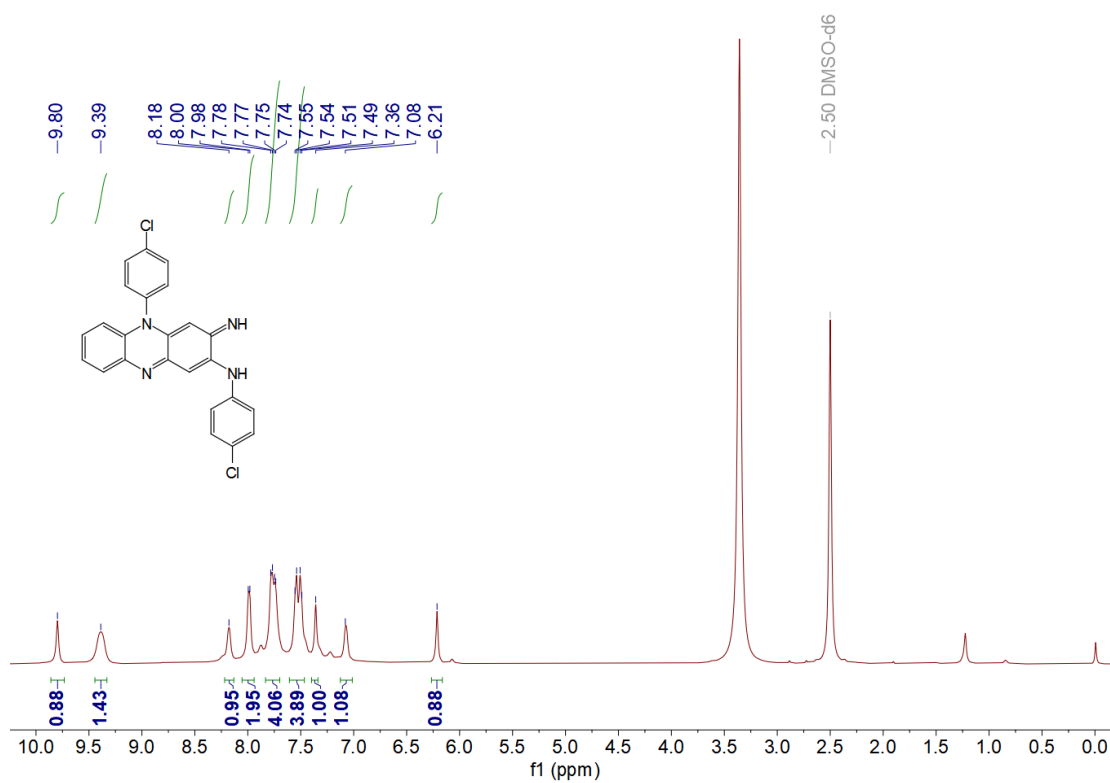


Figure S2. $^1\text{H-NMR}$ (500 Hz) spectrum of N-5-CCPIPA in DMSO- d_6 at room temperature. $\delta(\text{ppm}) = 9.80$ (s, 1H), 9.39 (s, 1H), 8.18 (s, 1H), 7.99 (d, $J = 7.7$ Hz, 2H), 7.83 – 7.70 (m, 4H), 7.61 – 7.47 (m, 4H), 7.36 (s, 1H), 7.07 (d, $J = 7.3$ Hz, 1H), 6.21 (s, 1H).

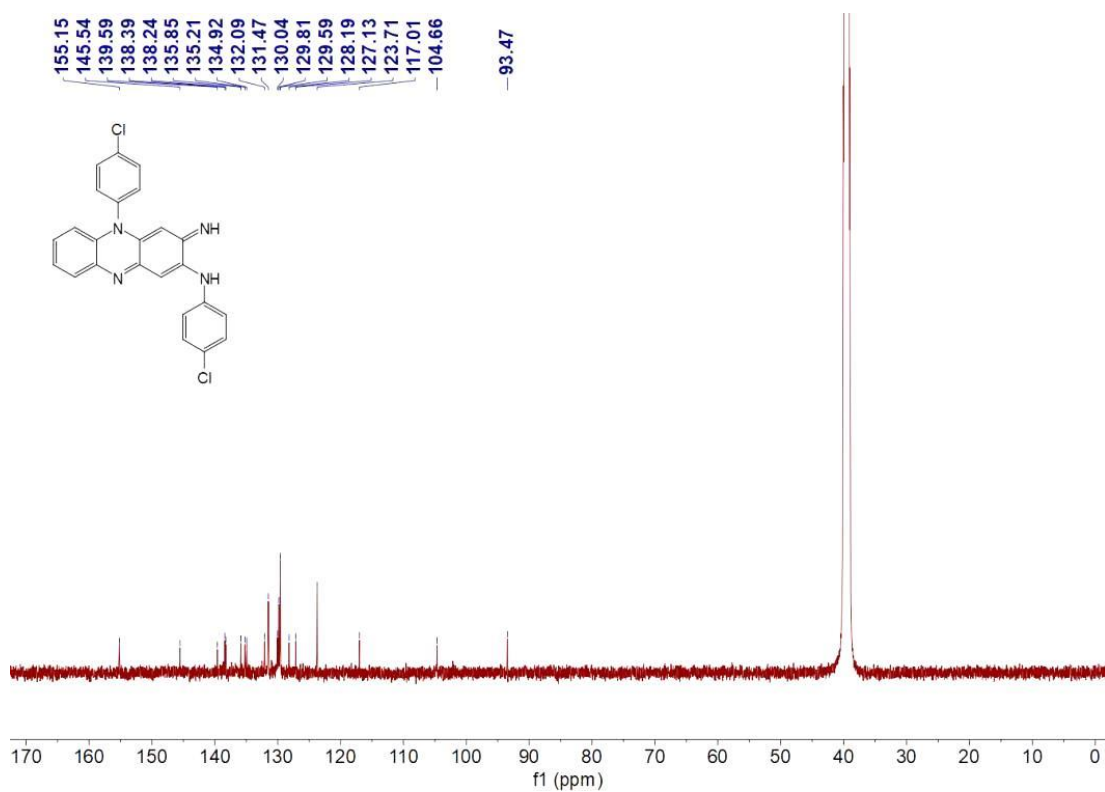


Figure S3. ^{13}C -NMR (126 Hz) spectrum of N-5-CCPIPA in DMSO- d_6 at room temperature. $\delta(\text{ppm}) = 155.16, 145.54, 139.59, 138.39, 138.24, 135.85, 135.21, 134.92, 132.09, 131.47, 130.04, 129.81, 129.59, 128.19, 127.13, 123.72, 117.01, 104.66, 93.47$.

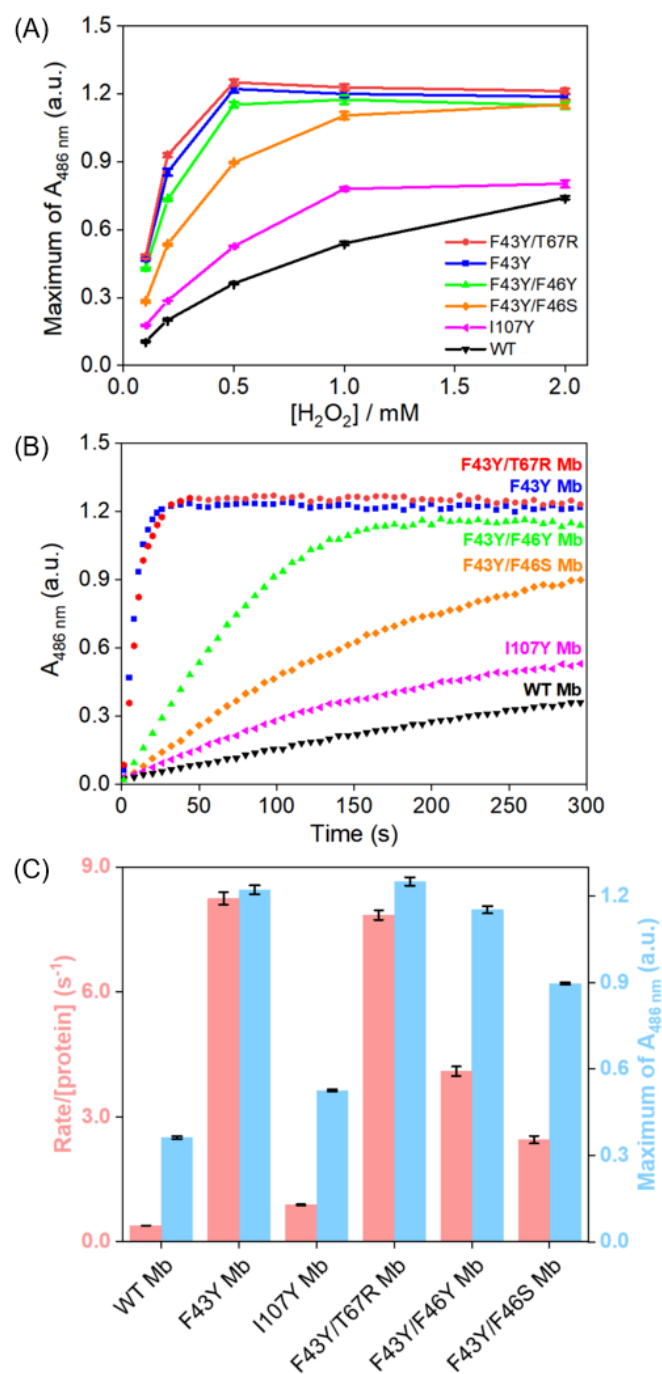


Figure S4. (A) Plots of the maximum absorbance of the oxidative coupling product of N-4-CPBDA catalyzed by WT Mb and its mutants at different concentrations of H₂O₂ within 5 min (pH 6.0, 25 °C). (B) Representative time-dependent absorbance changes at 486 nm, and (C) Comparison of the initial rates and the maximum absorbance of the reactions catalyzed by WT Mb and its mutants with 0.5 mM H₂O₂. All assays were carried out in triplicate.

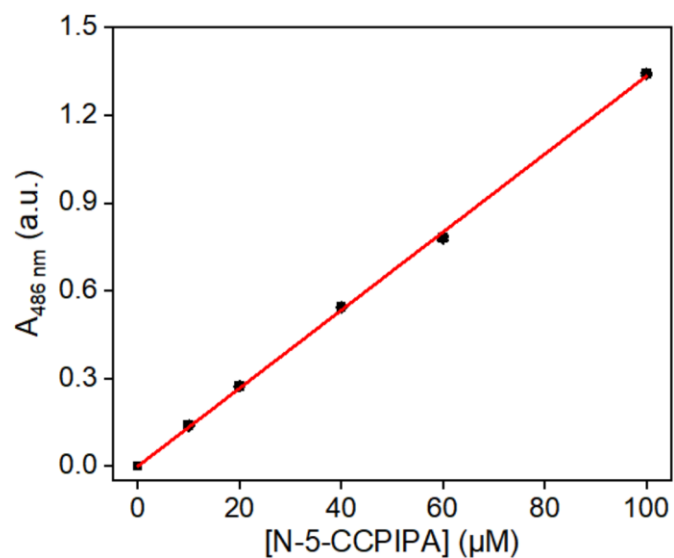


Figure S5. Standard curve of N-5-CCPIPA monitored at 486 nm using Agilent 8453 diode array spectrometer. The extinction coefficient was measured to be $\epsilon_{486 \text{ nm}} = 13 \text{ mM}^{-1} \text{ cm}^{-1}$ (100 mM potassium phosphate buffer containing 5% methanol (v/v), pH 6.0). All assays were carried out in triplicate.

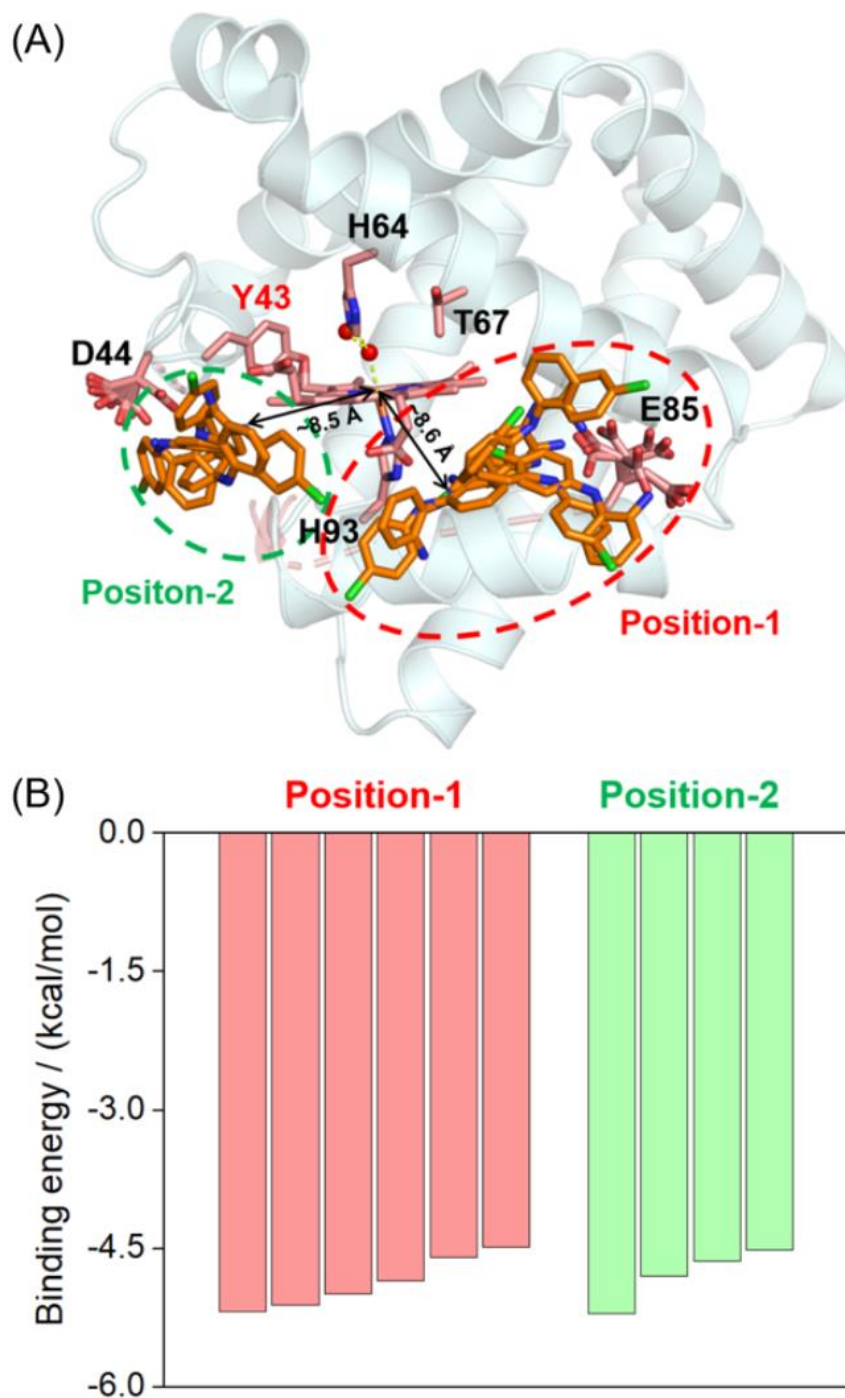


Figure S6. (A) Docking structures of N-4-CPBDA binding to F43Y Mb with the lowest 10 binding energies. Dotted arrows indicate the closest distance between the heme iron and the N-atom of the secondary amine of N-4-CPBDA in both positions. The N-4-CPBDA binding positions are indicated by circle lines. (B) Comparison of the binding energies between those bound to the position-1 and position-2.

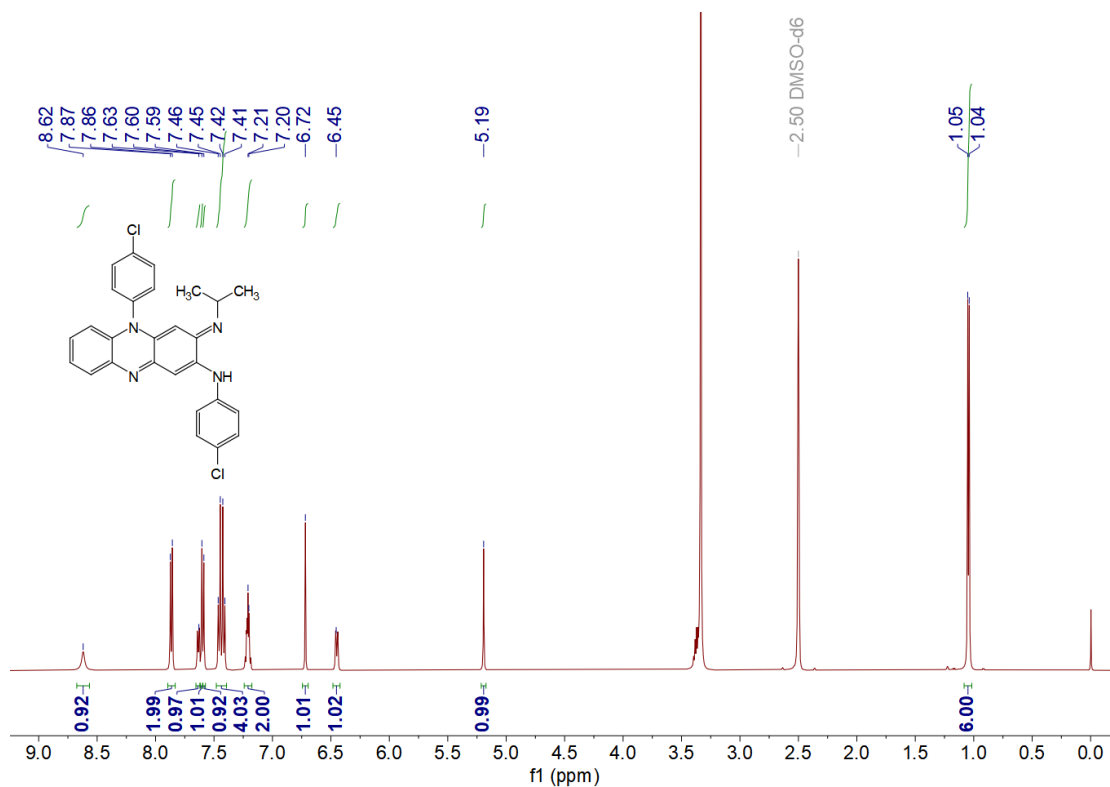


Figure S7. $^1\text{H-NMR}$ (500 Hz) spectrum of CFZ in DMSO- d_6 at room temperature. $\delta(\text{ppm}) = 8.62$ (s, 1H), 7.86 (d, $J = 8.6$ Hz, 2H), 7.63 (s, 1H), 7.60 (s, 1H), 7.59 (s, 1H), 7.48 – 7.39 (m, 4H), 7.21 (d, $J = 3.6$ Hz, 2H), 6.72 (s, 1H), 6.45 (s, 1H), 5.19 (s, 1H), 1.04 (d, $J = 6.2$ Hz, 6H).

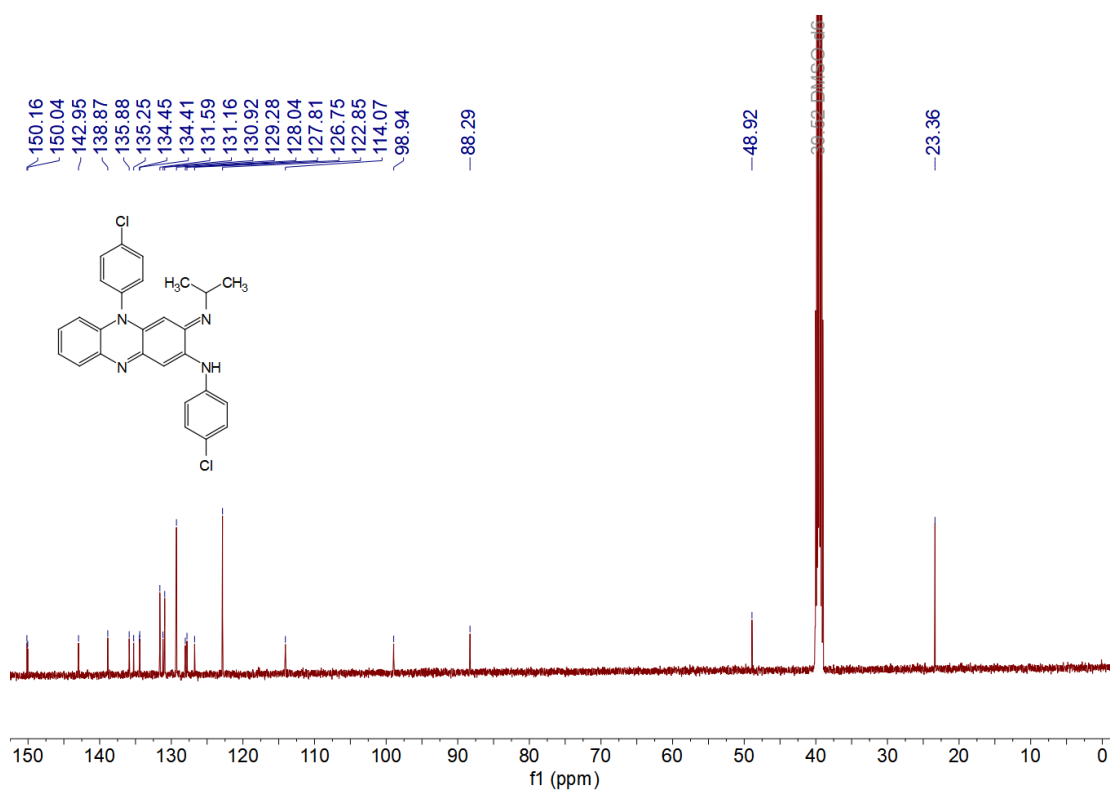


Figure S8. ¹³C-NMR (126 MHz) spectrum of CFZ in DMSO-d₆ at room temperature. $\delta(\text{ppm}) = 150.16, 150.04, 142.95, 138.87, 135.88, 135.25, 134.45, 134.41, 131.59, 131.16, 130.92, 129.28, 128.04, 127.81, 126.75, 122.85, 114.07, 98.94, 88.29, 48.92, 23.36.$

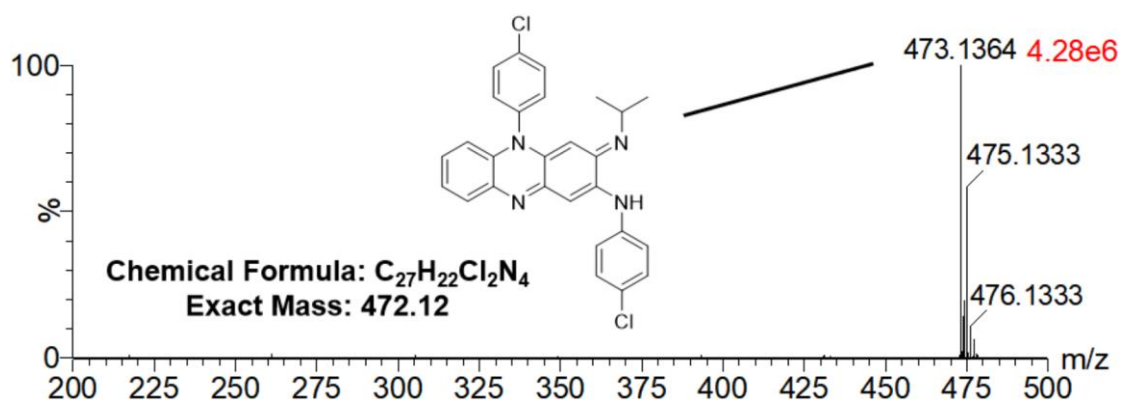


Figure S9. ESI-MS spectrum of the synthesized CFZ (calculated, 472.12 Da; observed, 473.14 Da, $[M+H]^+$).

Table S1. Docking results of N-4-CPBDA binding to F43Y/T67R Mb.

Model	E_{binding}^a	$E_{\text{inter-mol}}^b$	E_{vdw}^c	E_{elec}^d	E_{total}^e	$E_{\text{torsional}}^f$
1	-5.49	-6.39	-2.11	-0.21	-11.24	0.89
2	-4.99	-5.88	-2.57	-0.14	-13.68	0.89
3	-4.88	-5.78	-3.68	-0.18	-13.12	0.89
4	-4.61	-5.5	-2.4	-0.03	-13.62	0.89
5	-4.58	-5.47	-1.82	-0.05	-14.62	0.89
6	-4.18	-5.07	-0.72	-0.02	-13.93	0.89
7	-4.15	-5.04	-1.63	-0.04	-13.85	0.89
8	-4.09	-4.98	-1.69	-0.01	-13.01	0.89
9	-4.08	-4.97	-0.39	-0.01	-13.4	0.89
10	-4.06	-4.96	-2.64	-0.06	-15.38	0.89

^a Binding energy. ^b Intermolecular energy. ^c van der Waals energies. ^d Electrostatic interaction. ^e Total energy of the complex. ^f Torsional free energy. Unit: kcal/mol.

Table S2. Docking results of N-4-CPBDA binding to F43Y Mb.

Model	E_{binding}^a	E_{inter-mol}^b	E_{vdw}^c	E_{elec}^d	E_{total}^e	E_{torsional}^f
1	-5.21	-6.1	-3.46	-0.1	-11.73	0.89
2	-5.19	-6.09	-2.36	-0.3	-12.73	0.89
3	-5.12	-6.01	-2.99	-0.25	-12.04	0.89
4	-4.99	-5.88	-2.3	-0.16	-11.65	0.89
5	-4.85	-5.74	-1.9	-0.01	-13.49	0.89
6	-4.8	-5.7	-2.12	-0.06	-12.74	0.89
7	-4.64	-5.54	-3.43	-0.13	-12.5	0.89
8	-4.6	-5.5	-1.98	-0.08	-12.43	0.89
9	-4.52	-5.41	-1.86	-0.02	-11.85	0.89
10	-4.49	-5.39	-1.81	-0.01	-12.56	0.89

^a Binding energy. ^b Intermolecular energy. ^c van der Waals energies. ^d Electrostatic interaction. ^e Total energy of the complex. ^f Torsional free energy. Unit: kcal/mol.