

Supplementary Information

Synthesis, spectroscopic characterization and DNA/HSA binding study of (phenyl/naphthyl)ethenyl-substituted 1,3,4-oxadiazolyl-1,2,4- oxadiazoles

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^1H and ^{13}C NMR spectra of compounds 7aa-be.

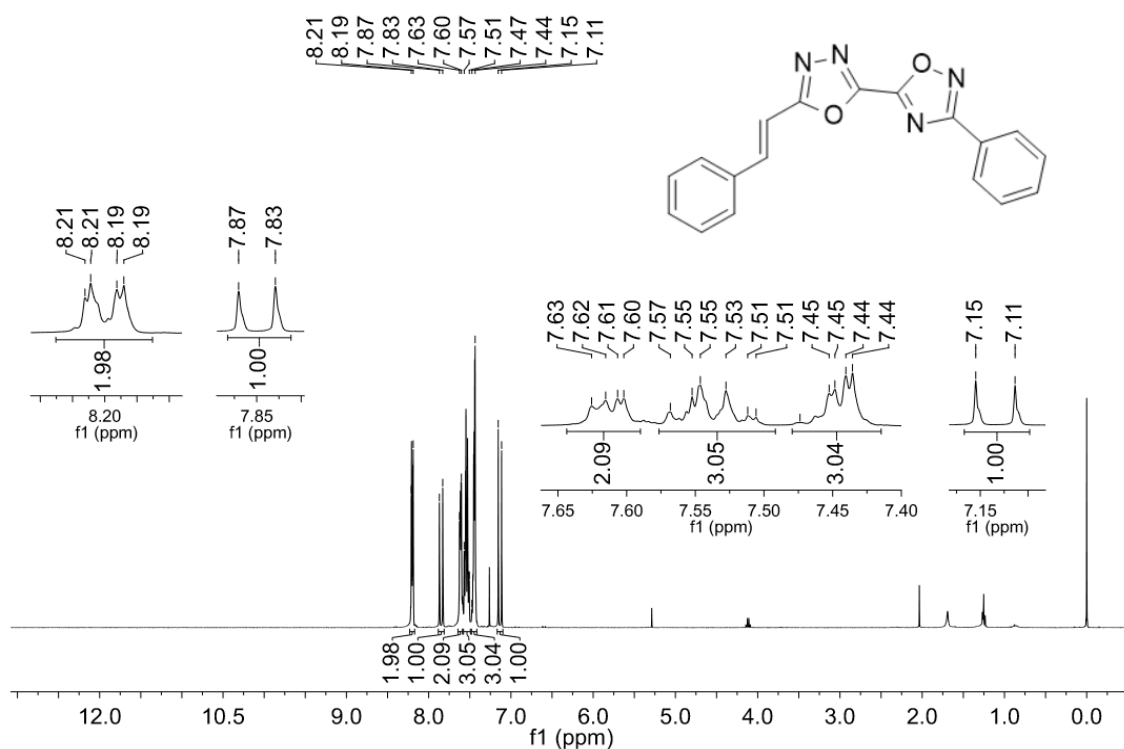


Figure S1. ^1H NMR spectrum (CDCl_3 at 400 MHz) of compound 7aa.

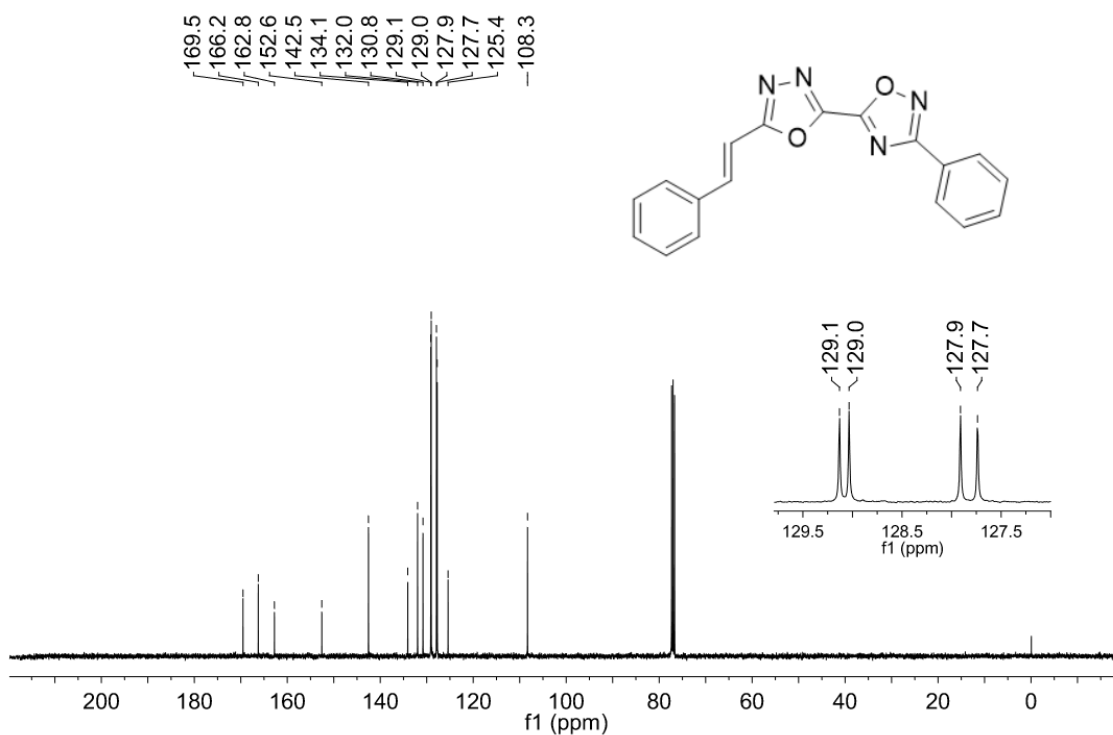


Figure S2. ^{13}C NMR spectrum (CDCl_3 at 100 MHz) of compound 7aa.

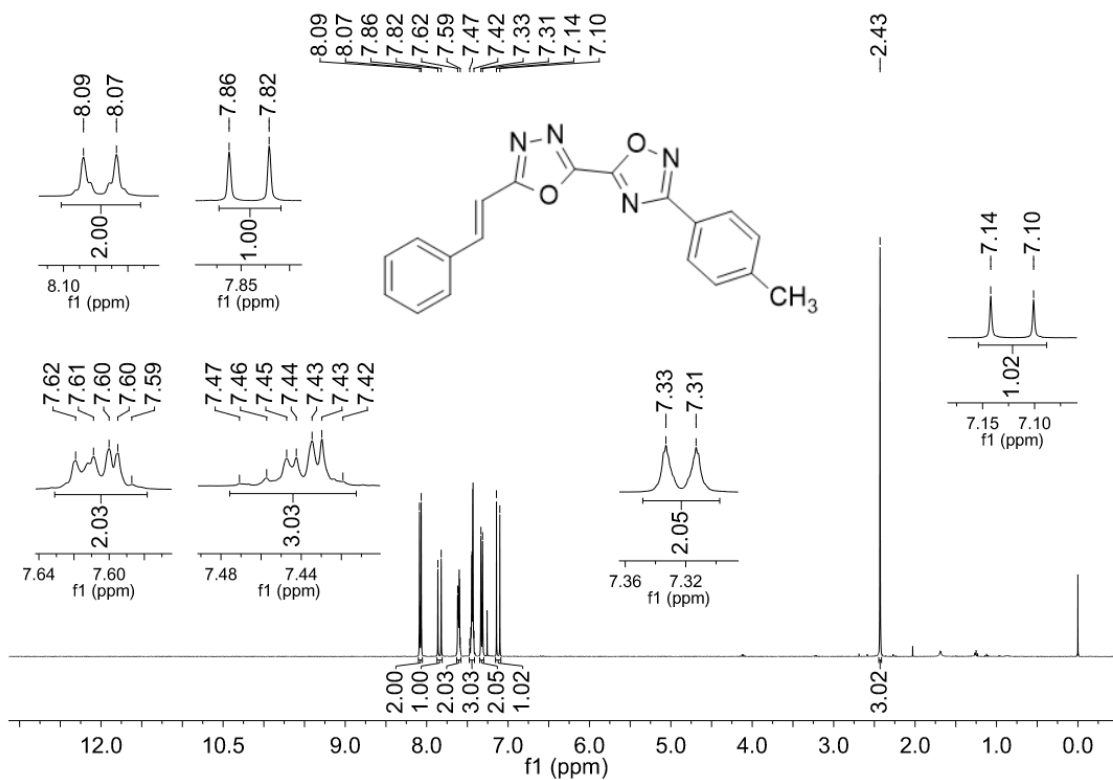


Figure S3. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7ab**.

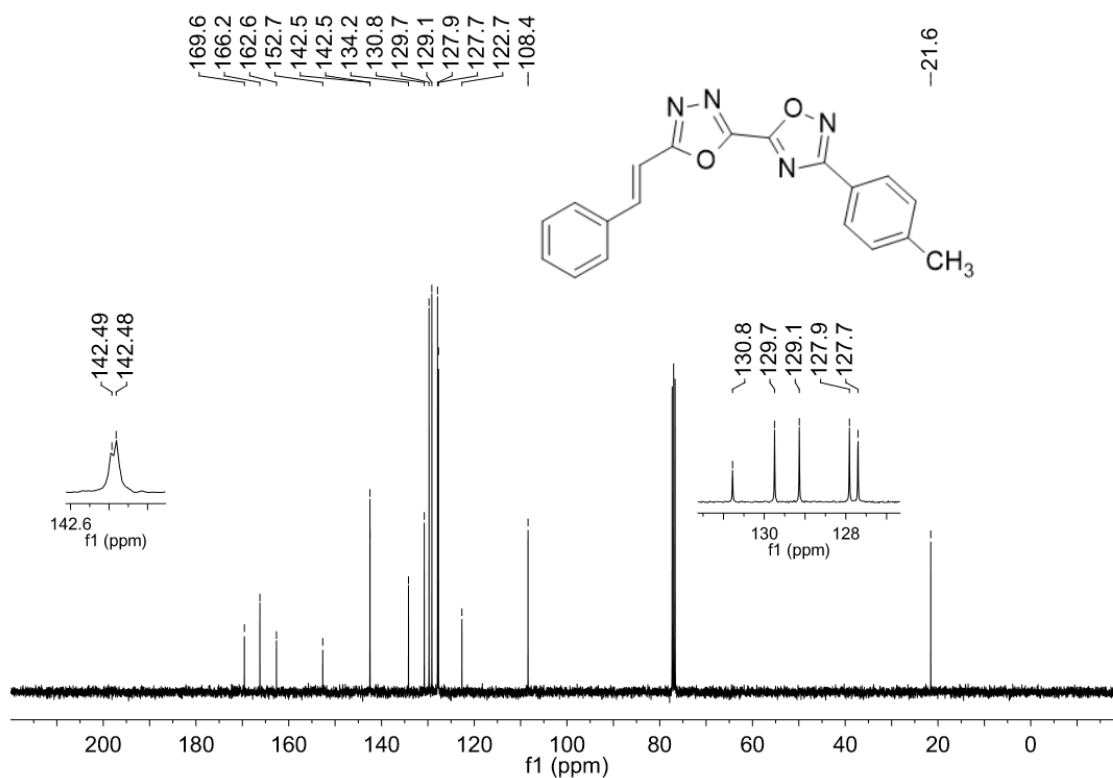


Figure S4. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7ab**.

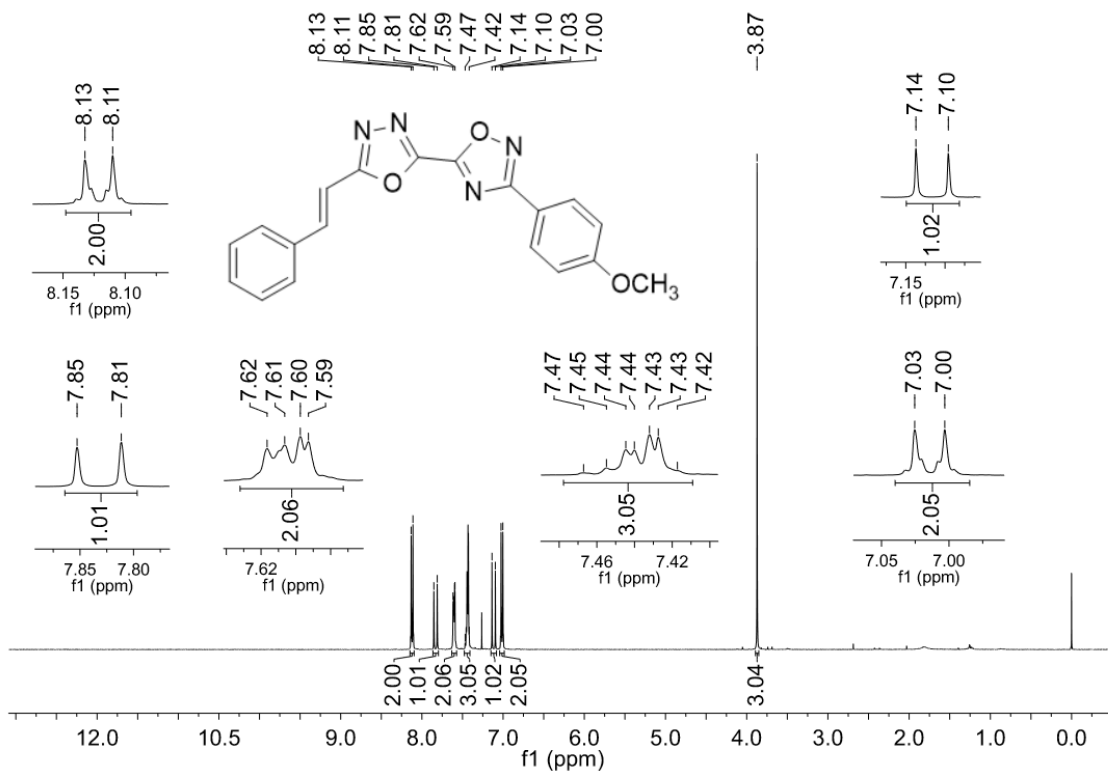


Figure S5. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7ac** (δ, TMS).

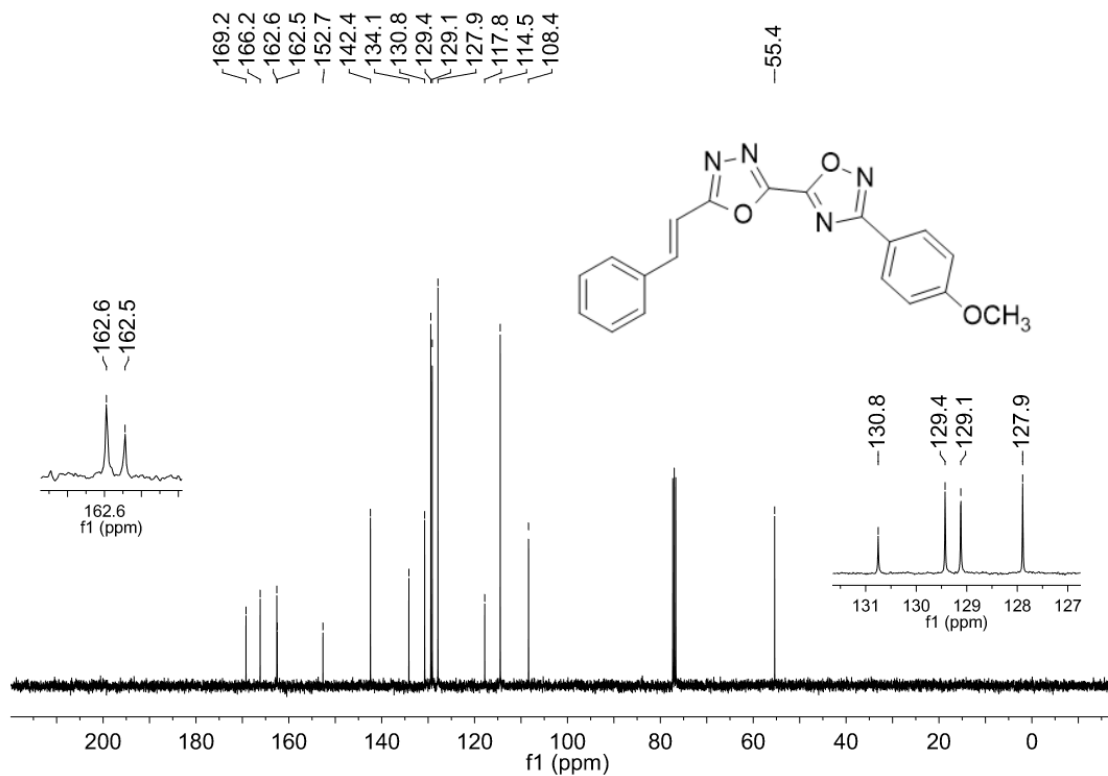


Figure S6. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7ac**.

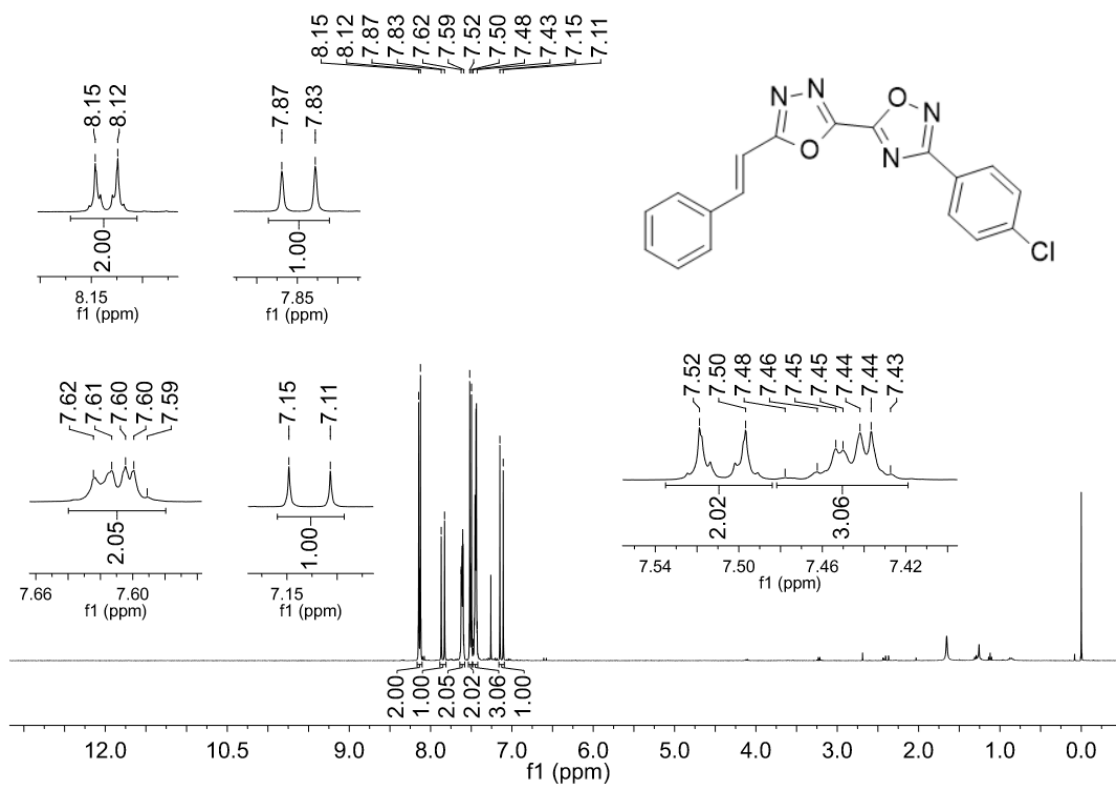


Figure S7. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7ad**.

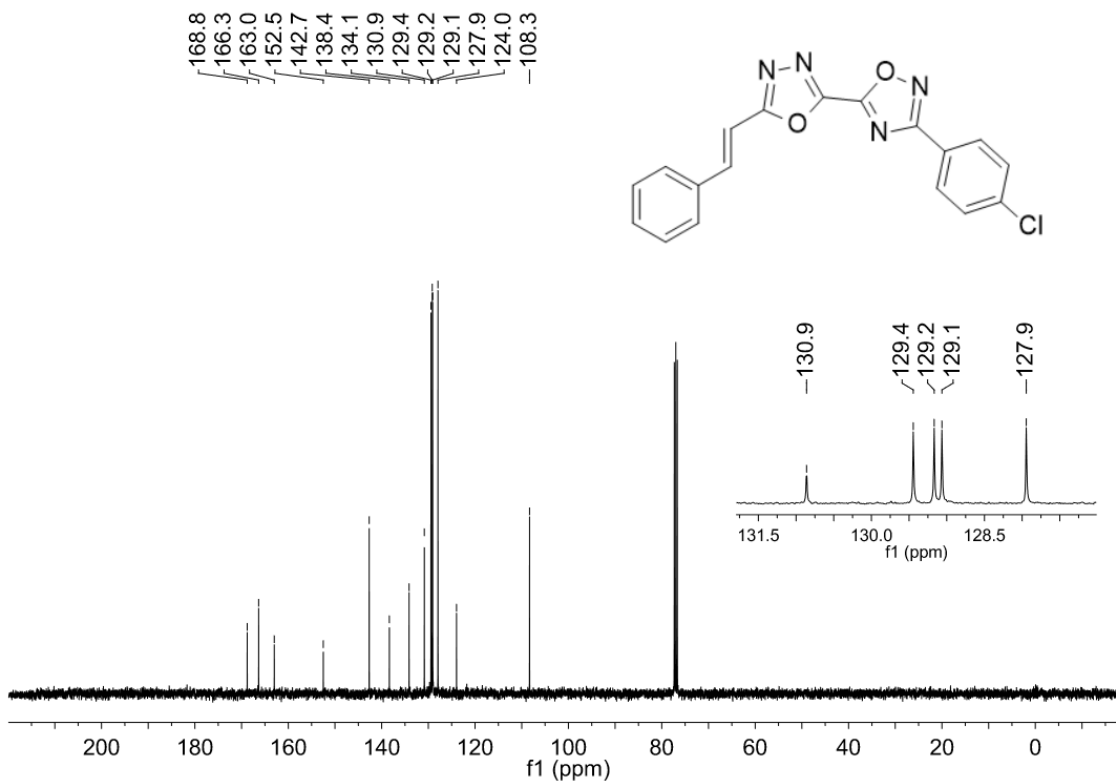


Figure S8. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7ad**.

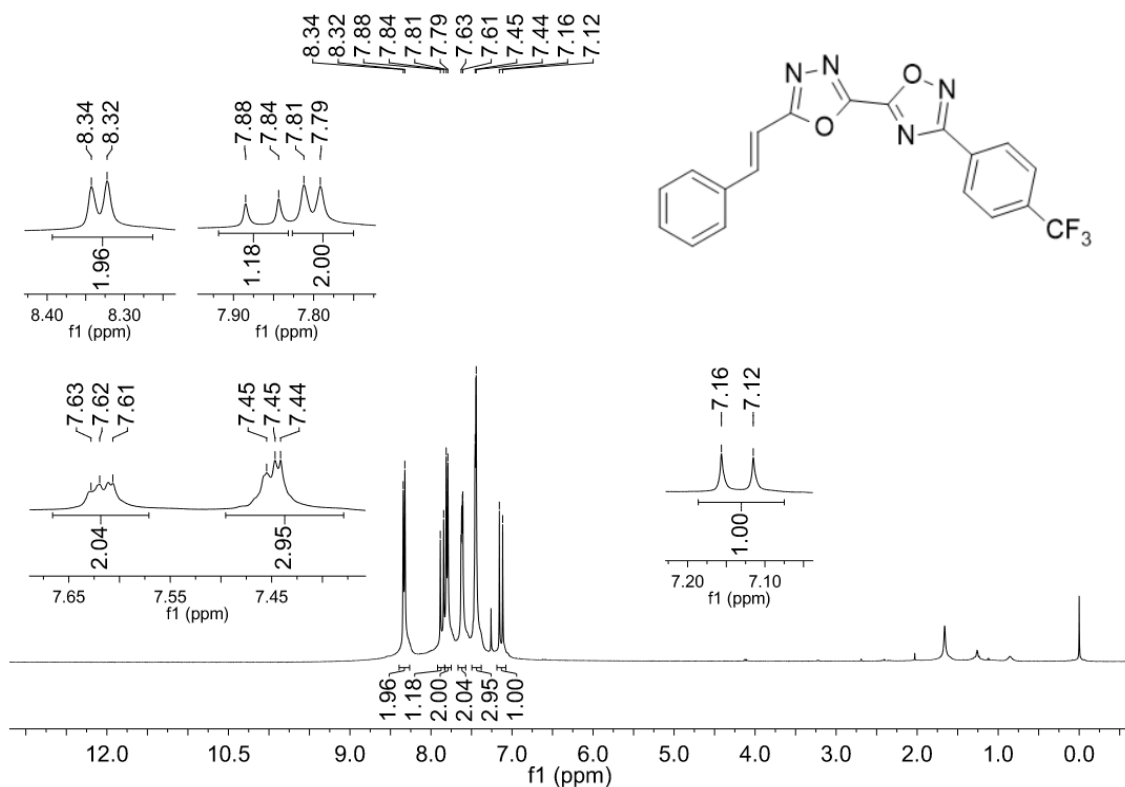


Figure S9. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7ae**.

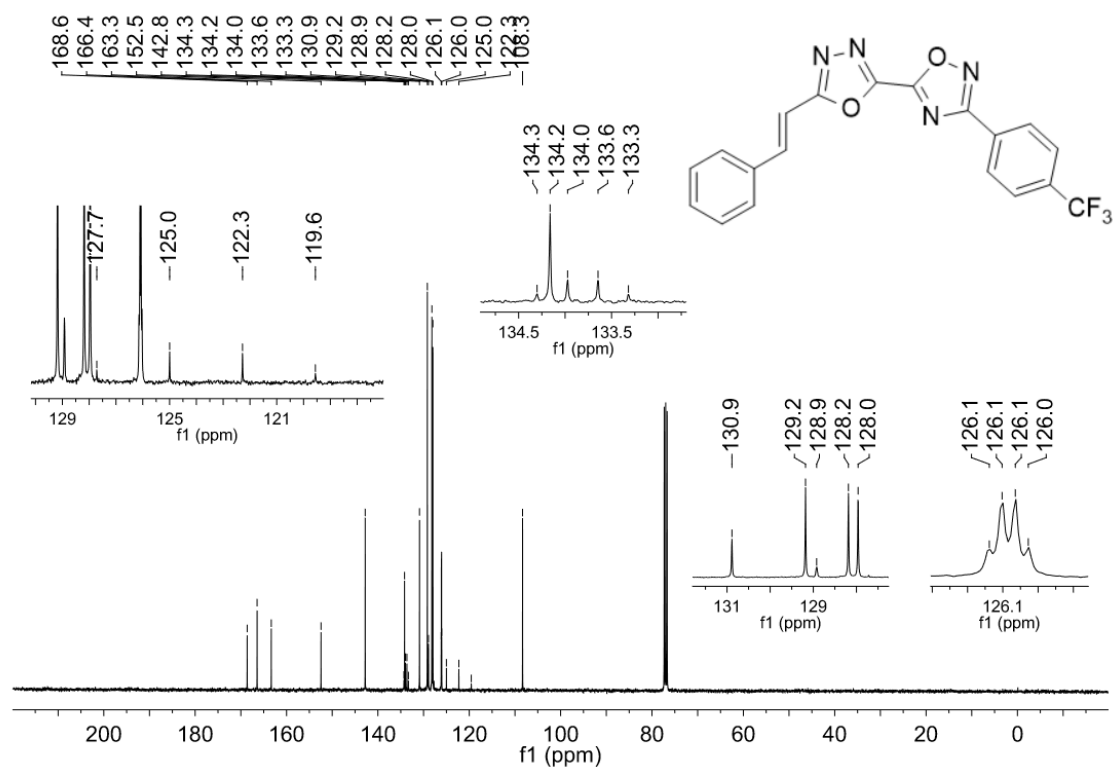


Figure S10. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7ae**.

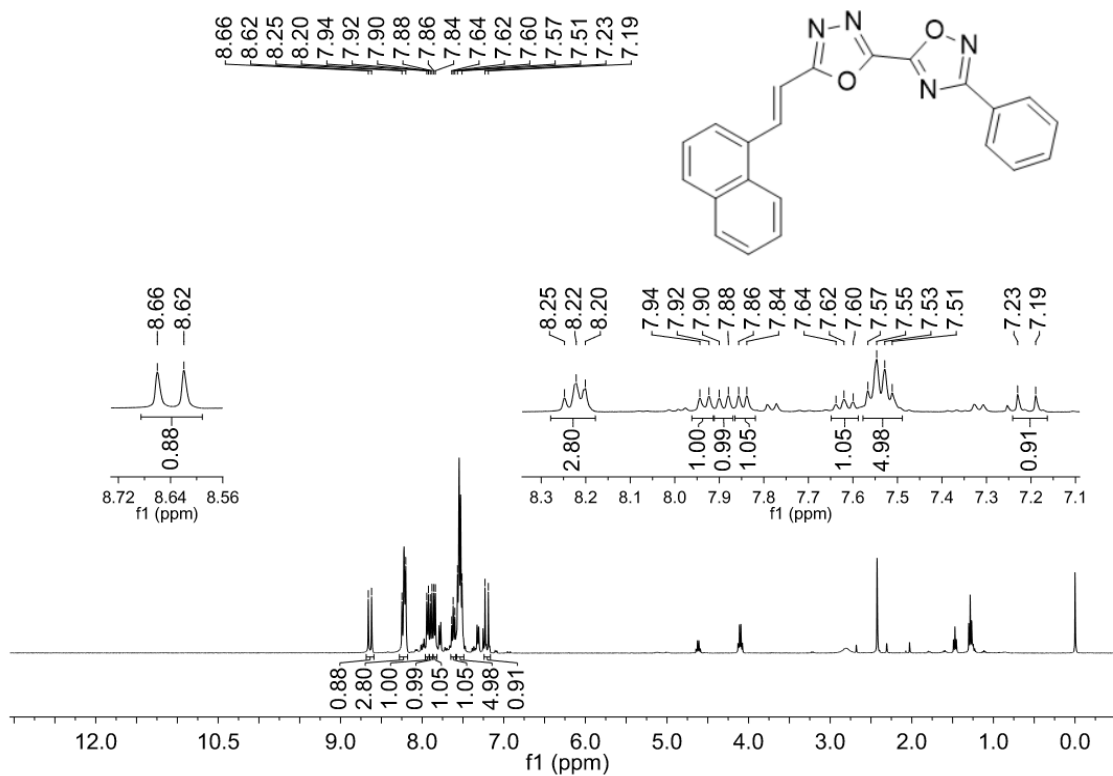


Figure S11. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7ba**.

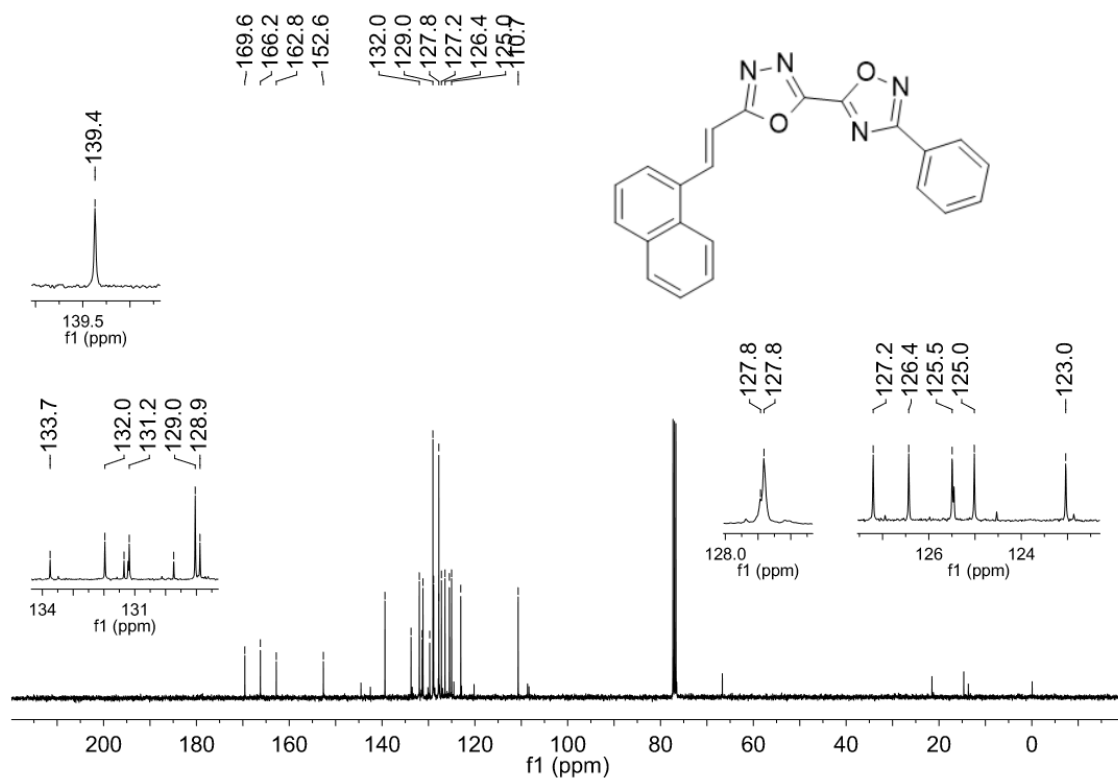


Figure S12. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7ba**.

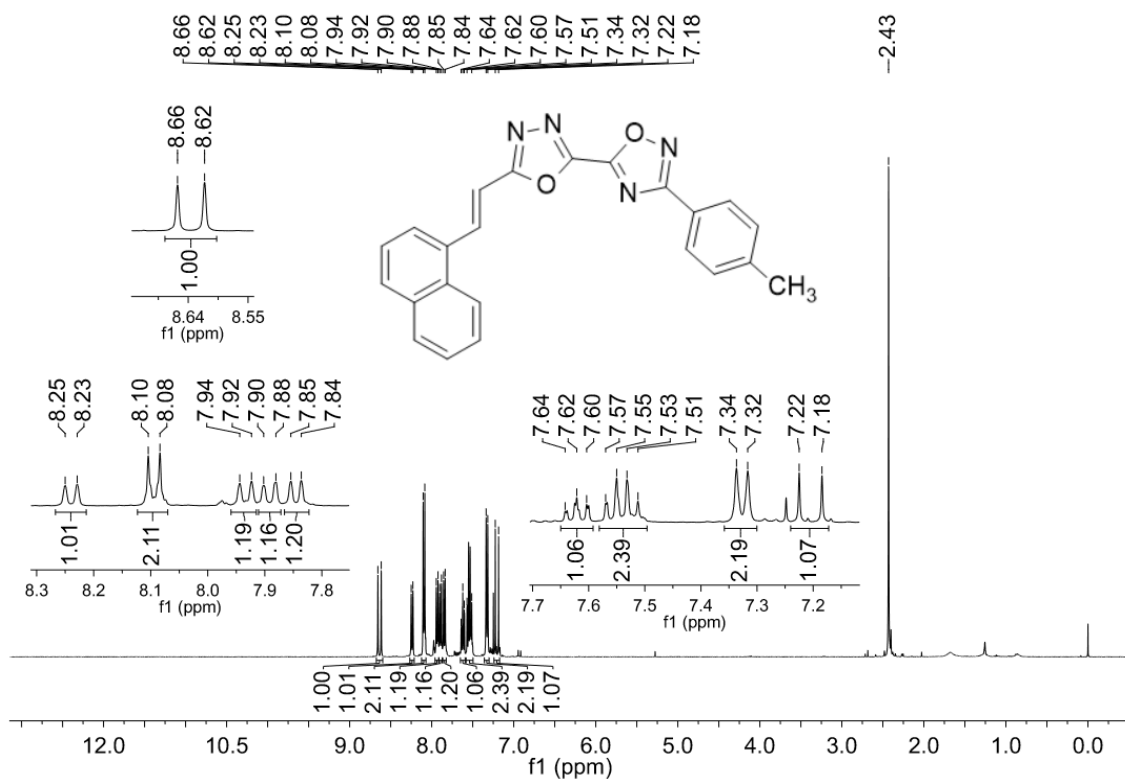


Figure S13. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7bb**.

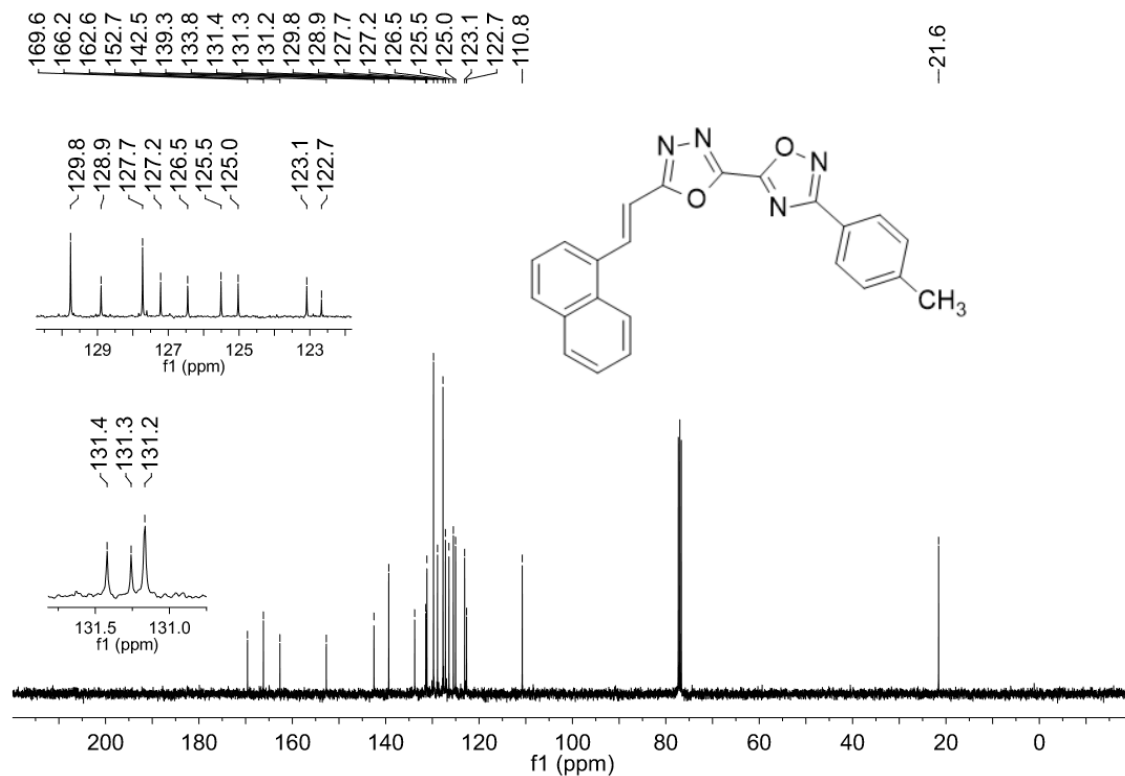


Figure S14. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7bb**.

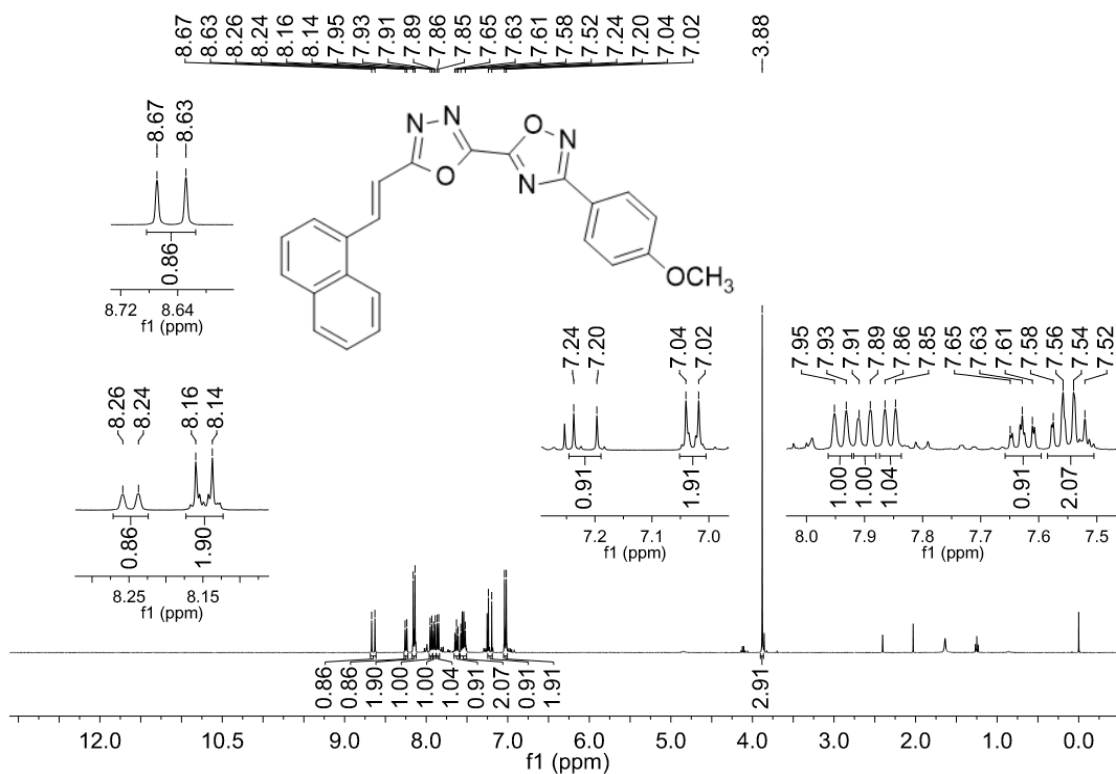


Figure S15. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7bc**.

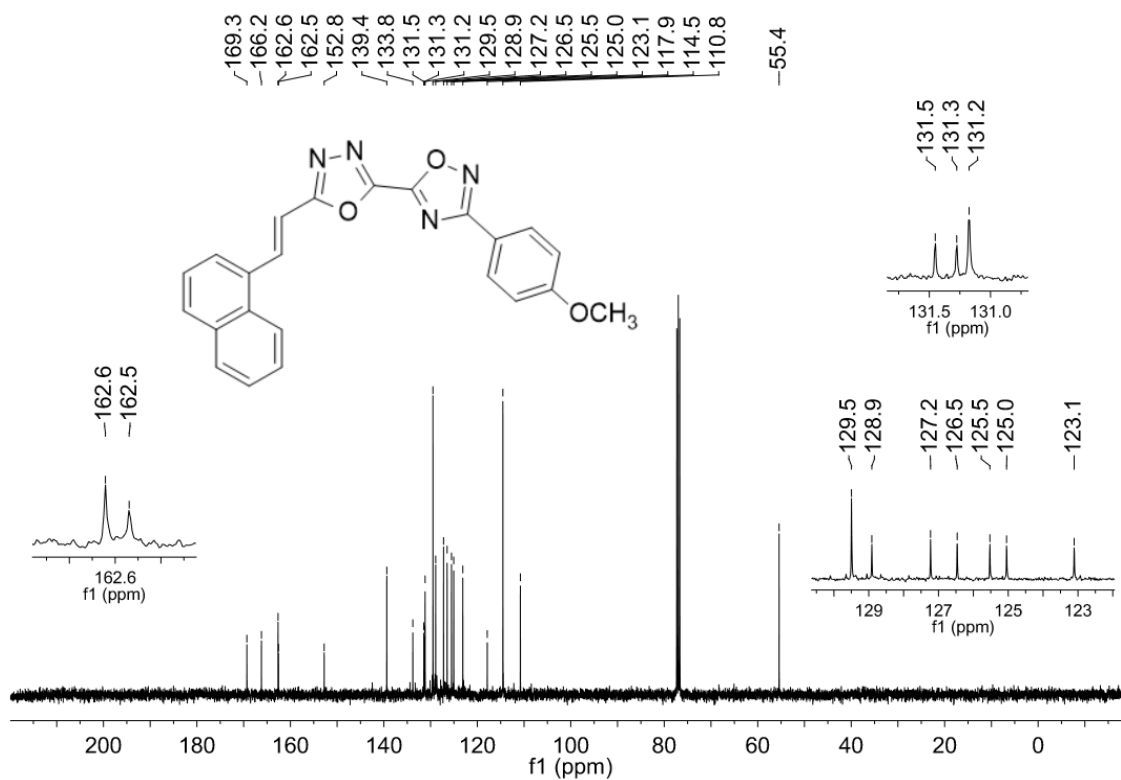


Figure S16. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7bc**.

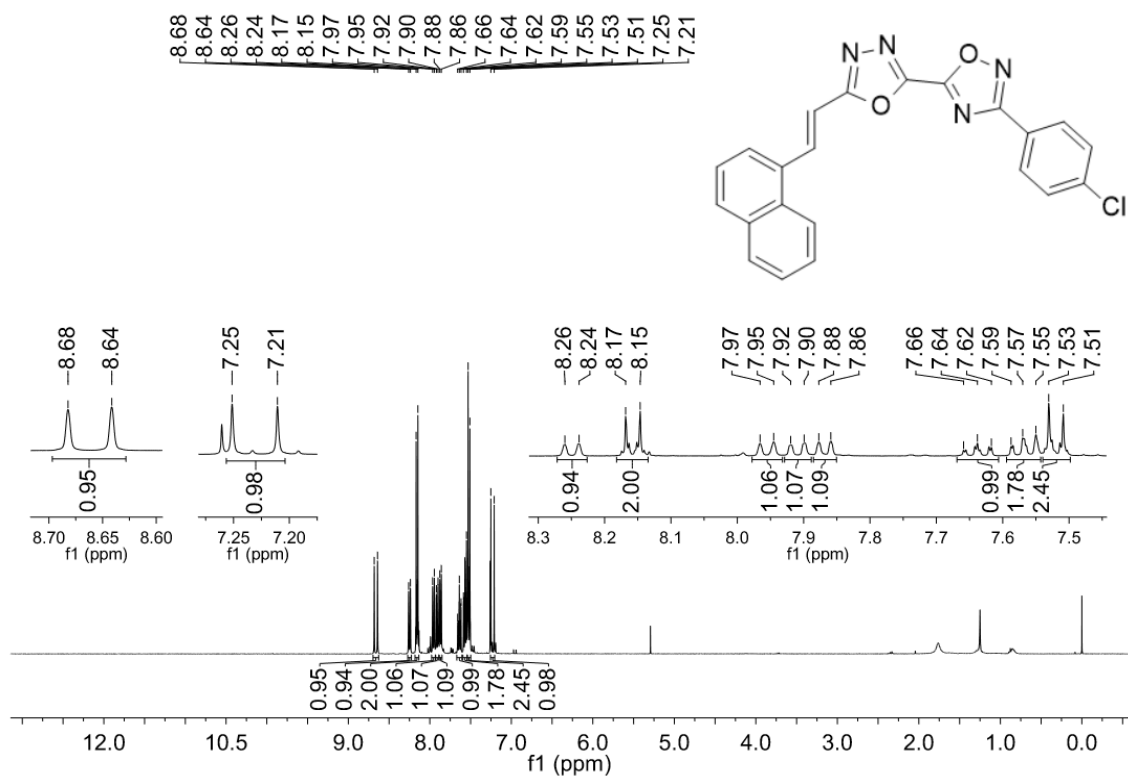


Figure S17. ¹H NMR spectrum (CDCl₃ at 400 MHz) of compound **7bd**.

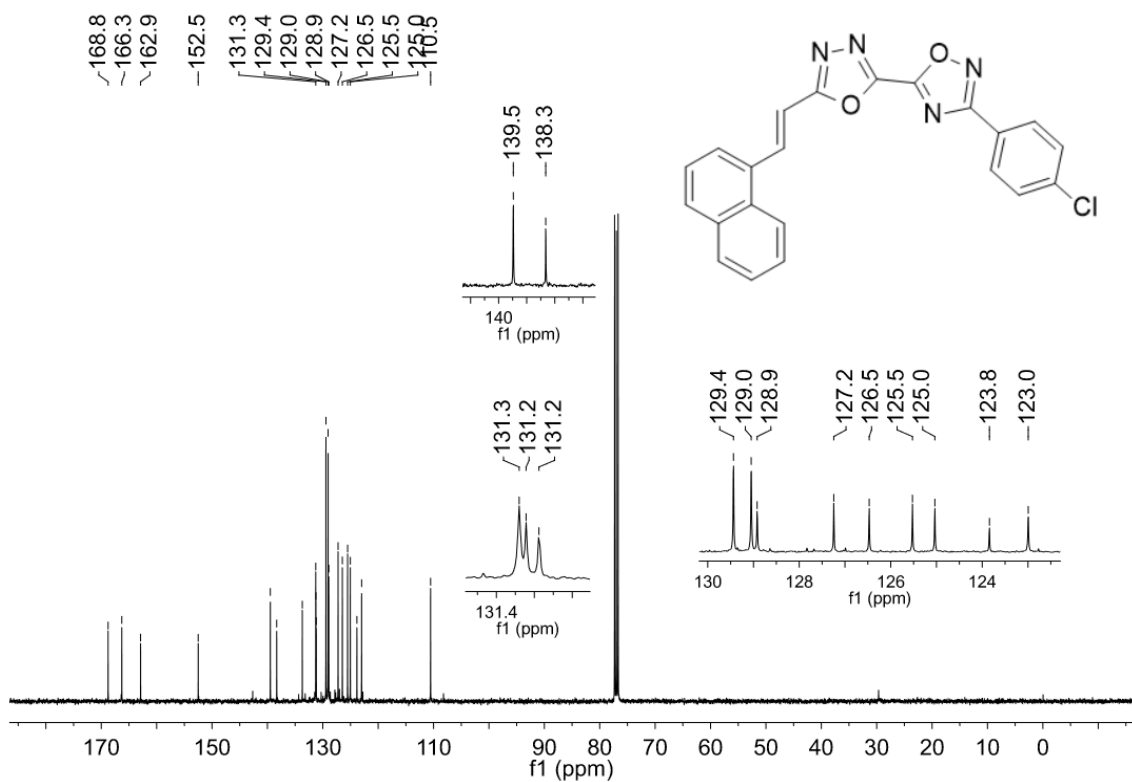


Figure S18. ¹³C NMR spectrum (CDCl₃ at 100 MHz) of compound **7bd**.

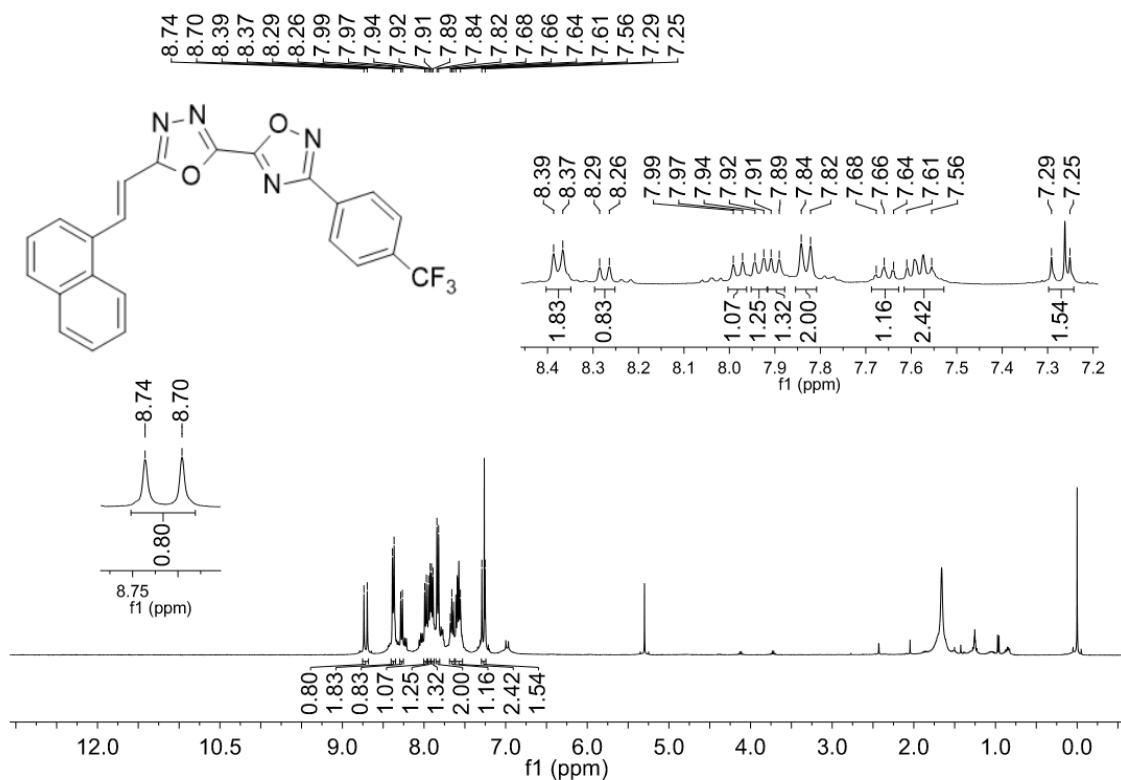


Figure S19. ^1H NMR spectrum (CDCl₃ at 400 MHz) of compound **7be**.

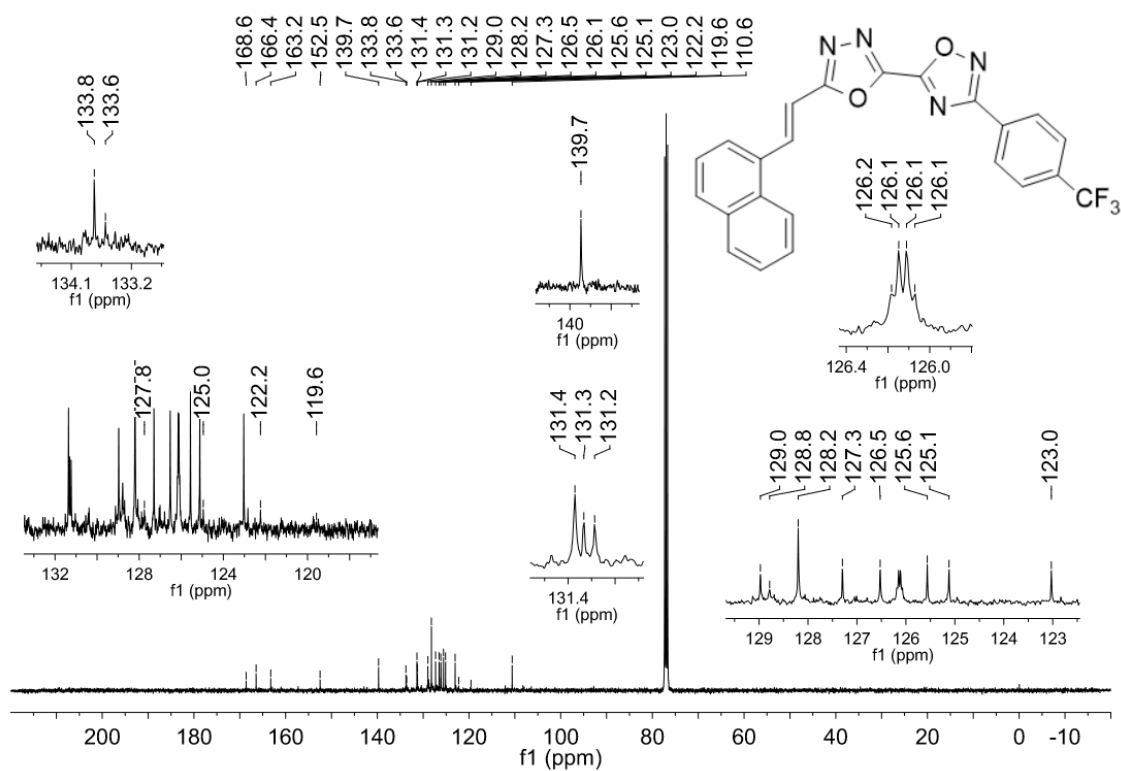


Figure S20. ^{13}C NMR spectrum (CDCl₃ at 100 MHz) of compound **7be**.

HRMS Spectra of compounds 7aa-be

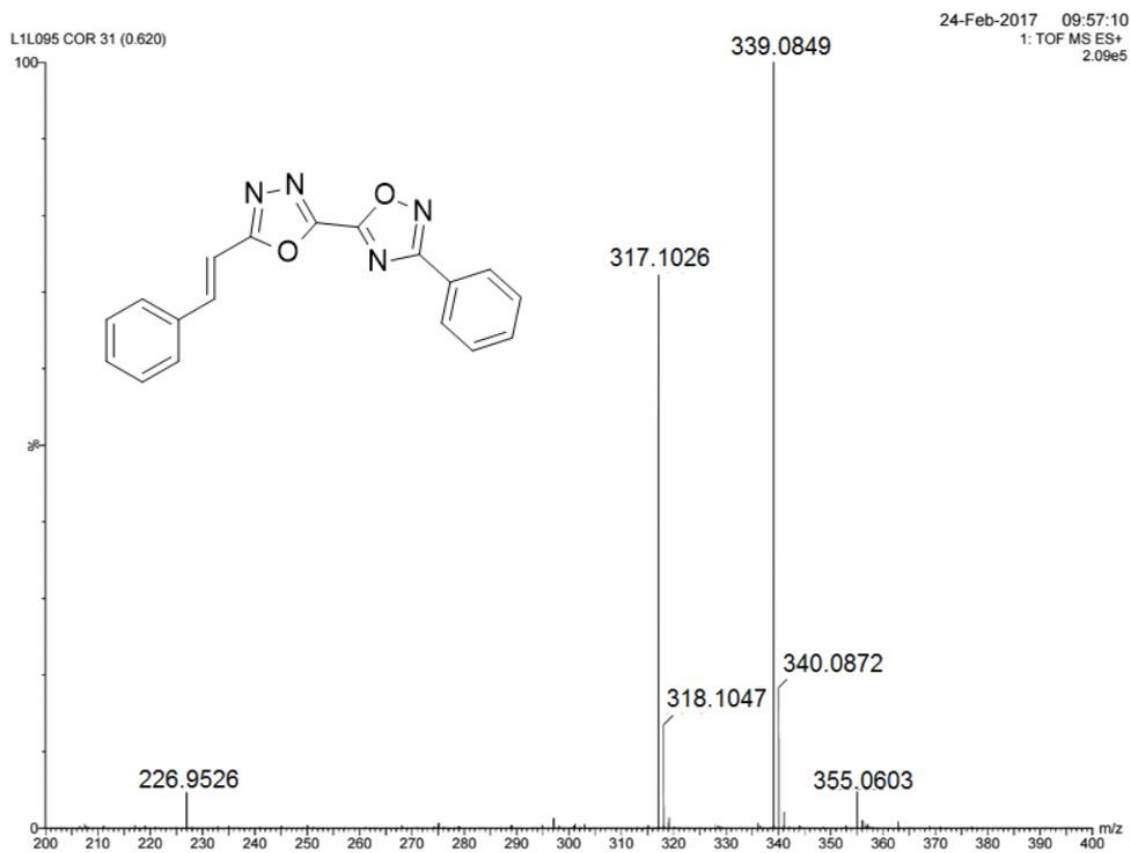


Figure S21. HRMS Spectrum of compound **7aa** (ESI+).

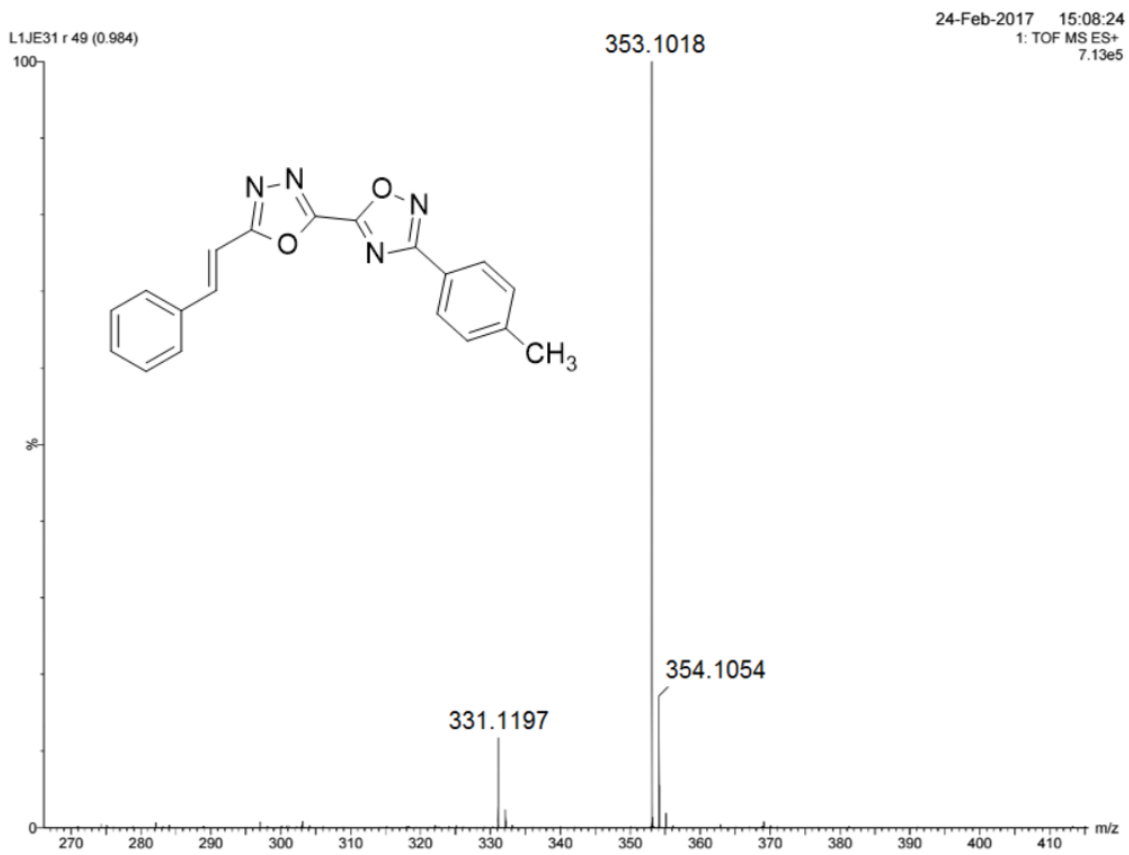


Figure S22. HRMS Spectrum of compound **7ab** (ESI+).

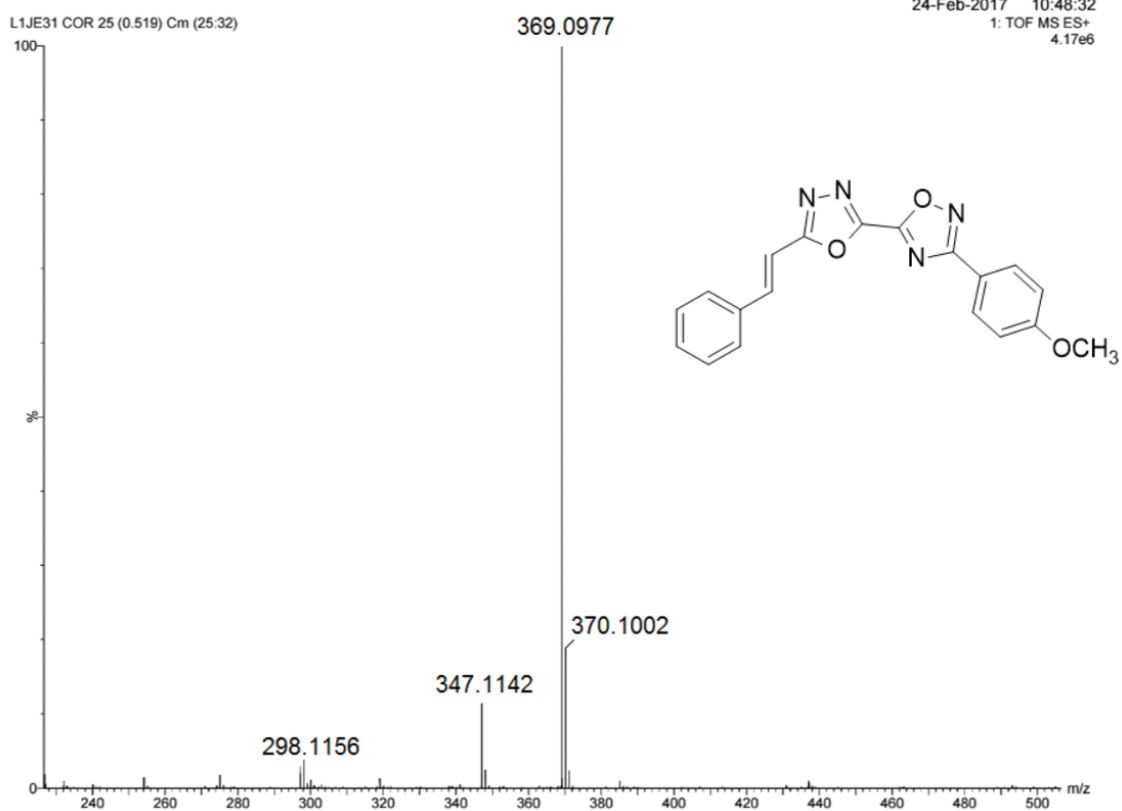


Figure S23. HRMS Spectrum of compound **7ac** (ESI+).

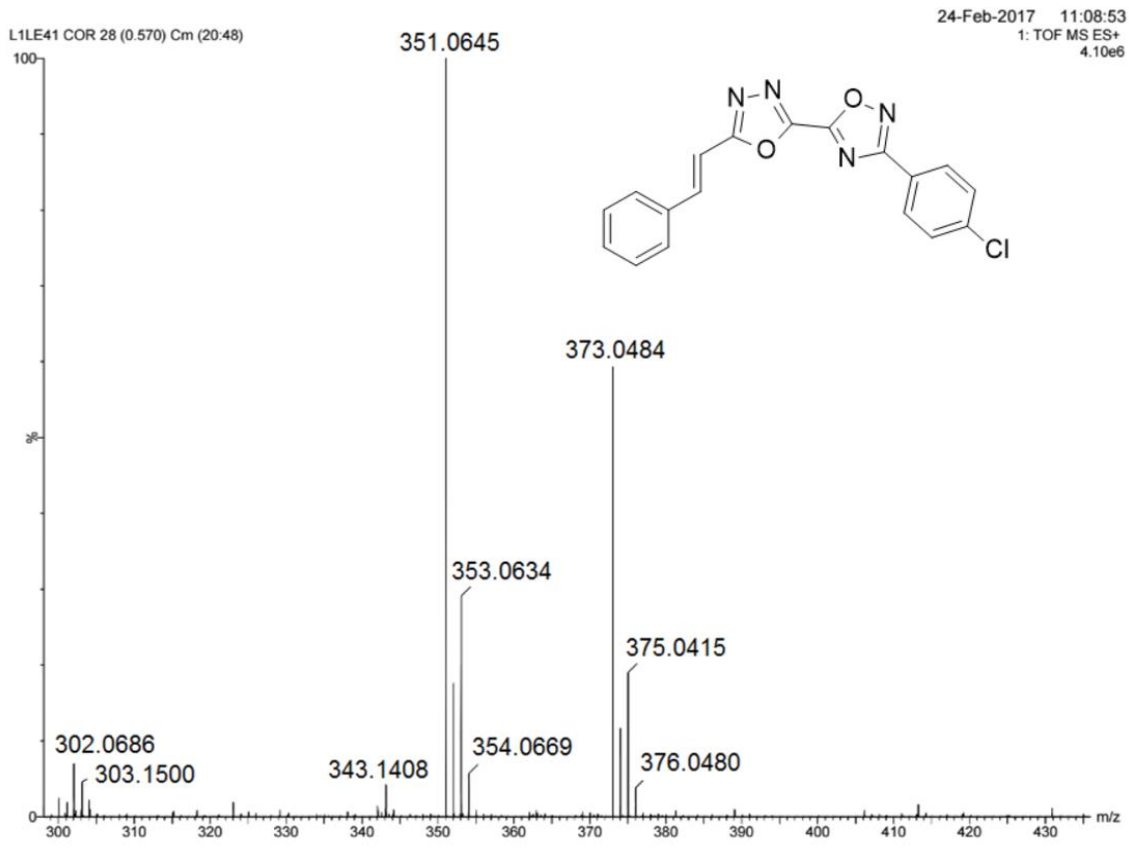


Figure S24. HRMS Spectrum of compound **7ad** (ESI+).

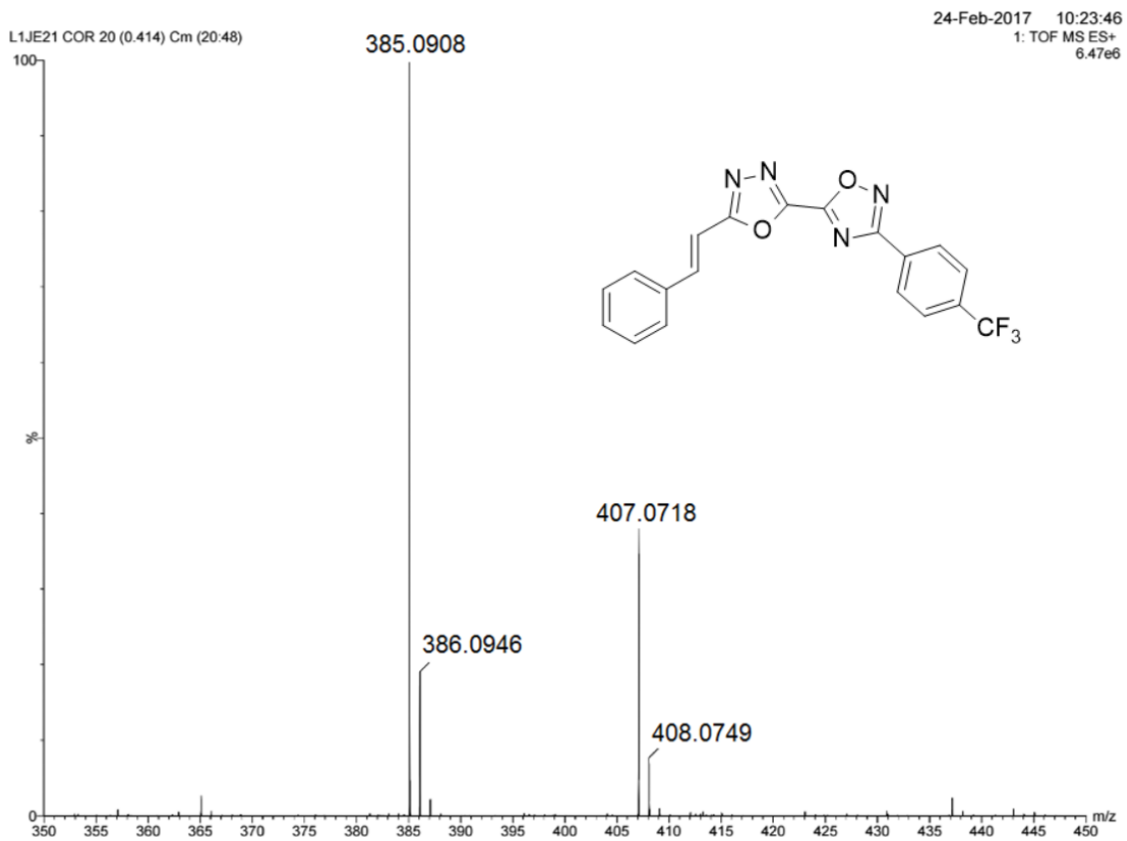


Figure S25. HRMS Spectrum of compound **7ae** (ESI+).

Waters
L1JE14 117 (1.995)

24-Feb-2017 11:39:39
TOF MS ES+
4.53e5

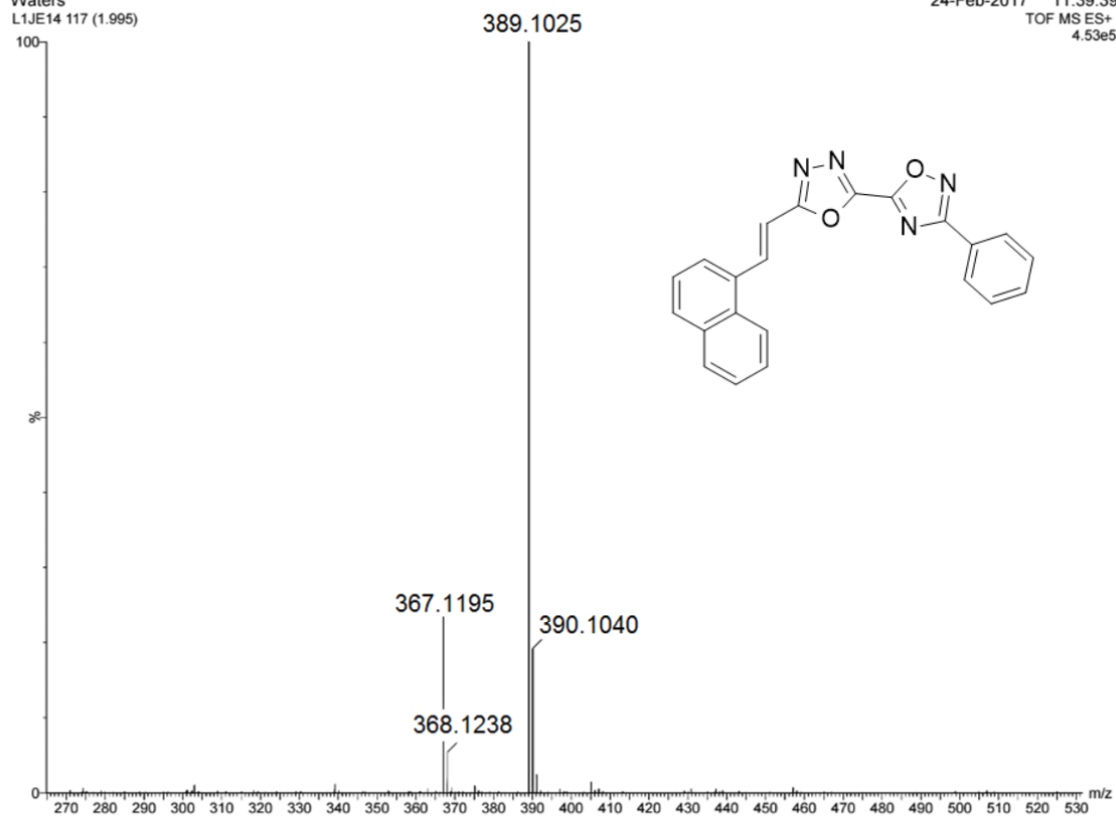


Figure S26. HRMS Spectrum of compound **7ba** (ESI+).

Waters
L1JE34 2402 5 (0.101) Cm (1.94)

24-Feb-2017 12:20:55
TOF MS ES+
5.20e7

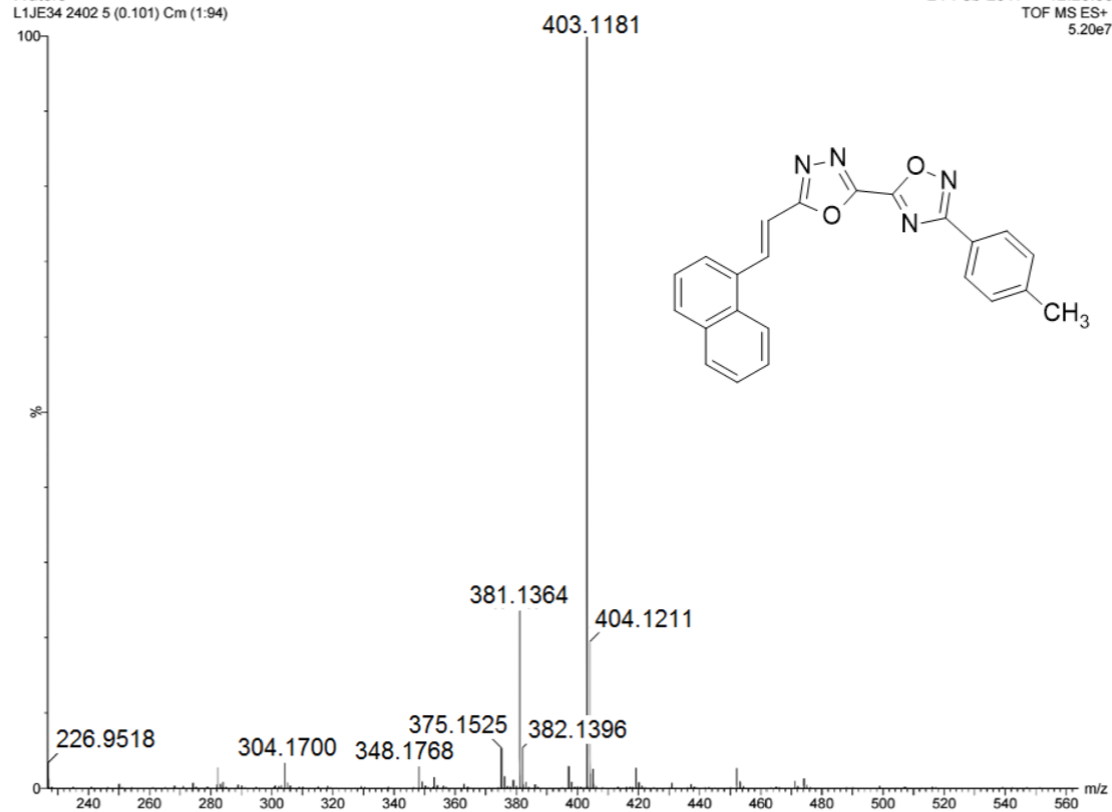


Figure S27. HRMS Spectrum of compound **7bb** (ESI+).

Waters
L1AE44 67 (1.150) Cm (2-114)

24-Feb-2017 12:47:40
TOF MS ES+
4.31e7

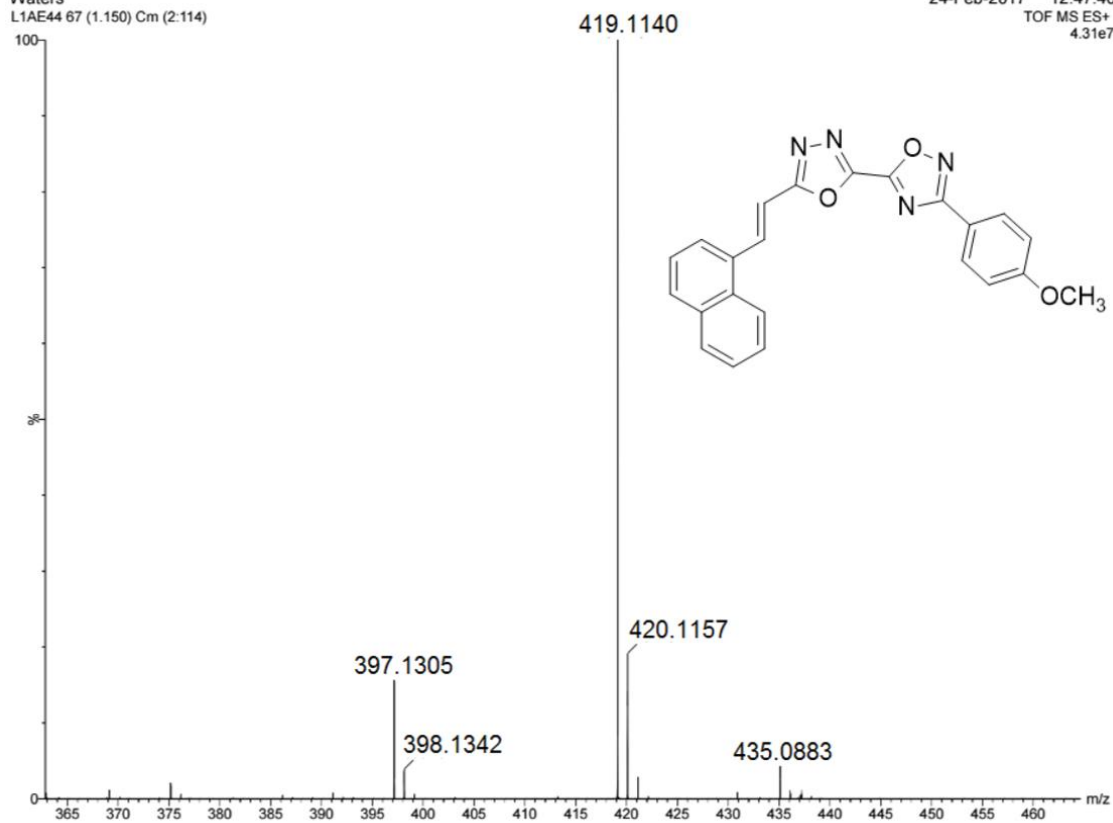


Figure S28. HRMS Spectrum of compound **7bc** (ESI+).

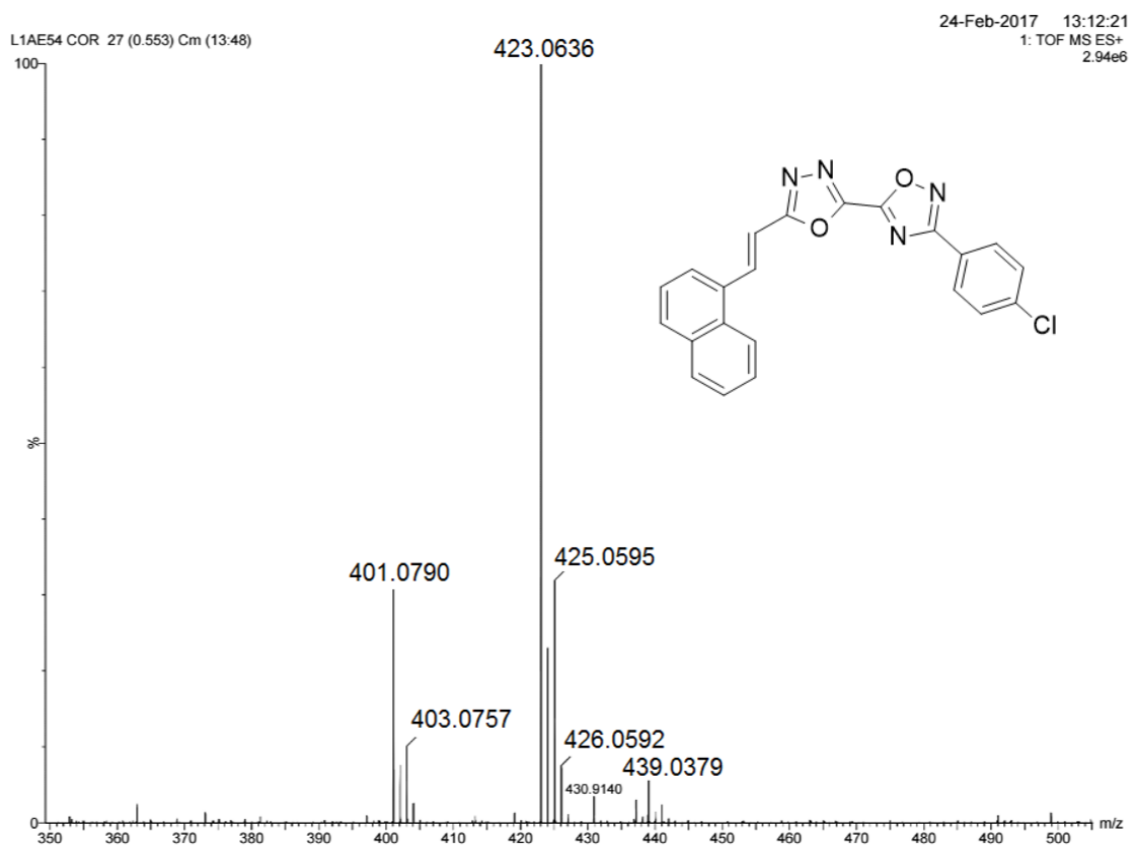


Figure S29. HRMS Spectrum of compound **7bd** (ESI+).

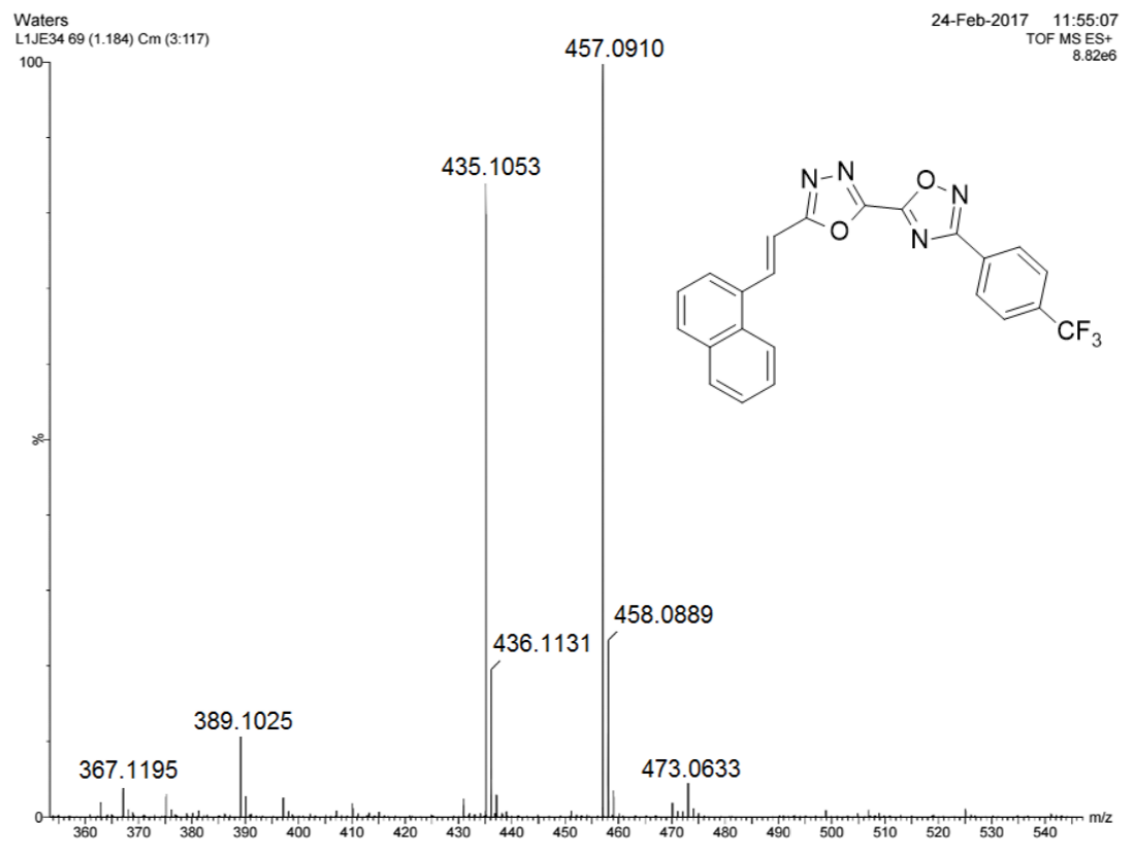


Figure S30. HRMS Spectrum of compound **7be** (ESI+).

Table S1. Crystal data and details of the refinement of the crystal structure of **7aa**.

Compound	7aa
Empirical formula	C ₁₈ H ₁₂ N ₄ O ₂
Formula weight	316.32
Temperature (K)	100(2)
Wavelength	0.71073
Crystal system	Monoclinic
Space group	<i>P</i> 2 ₁ / <i>c</i>
<i>a</i> (Å)	15.8581(13)
<i>b</i> (Å)	4.2689(3)
<i>c</i> (Å)	22.0230(18)
α (°)	90
β (°)	94.541(3)
γ (°)	90
Volume (Å ³)	1486.2(2)
<i>Z</i>	4
Calculated density(mg·m ⁻³)	1.414
Absorp. coefficient (mm ⁻¹)	0.096
<i>F</i> (000)	656
Crystal size (mm)	0.373 x 0.104 x 0.080
Theta range for data collection (°)	2.34 to 27.13
Limiting indices	-20 ≤ <i>h</i> ≤ 20. -5 ≤ <i>k</i> ≤ 5. -28 ≤ <i>l</i> ≤ 28
Reflections collected / unique	12694 / 3290
Completeness to theta	99.8%
Absorption correction	Semi-empirical from equivalents
Max. and min. trans.	0.8620/0.6741
Data / restraints / parameters	3290 / 0 / 217
Goodness-of-fit on <i>F</i> ²	1.015
Índice <i>R</i> _{int}	0.0768
Final R indices <i>R</i> ₁ and <i>wR</i> ₂ [<i>I</i> > 2σ(<i>I</i>)]	0.0572 / 0.01129
<i>R</i> indices (all data)	0.01129 / 0.01344
Largest diff. peak and hole	0.243 and -0.333 e·Å ⁻³

Table S2. Bond lengths [Å] and angles [deg] for **7aa**.

O(12)-C(13)	1.364(2)
O(12)-C(16)	1.374(2)
O(1)-C(5)	1.346(2)
O(1)-N(2)	1.413(2)
N(15)-C(16)	1.302(3)
N(15)-N(14)	1.402(2)
N(14)-C(13)	1.286(3)
N(2)-C(3)	1.310(3)
N(4)-C(5)	1.290(3)
N(4)-C(3)	1.392(2)
C(13)-C(5)	1.444(3)
C(6)-C(11)	1.384(3)
C(6)-C(7)	1.402(3)
C(6)-C(3)	1.460(3)
C(19)-C(20)	1.388(3)
C(19)-C(24)	1.397(3)
C(19)-C(18)	1.472(3)
C(17)-C(18)	1.339(3)
C(17)-C(16)	1.437(3)
C(8)-C(9)	1.378(3)
C(8)-C(7)	1.382(3)
C(10)-C(11)	1.384(3)
C(10)-C(9)	1.392(3)
C(20)-C(21)	1.389(3)
C(21)-C(22)	1.375(4)
C(22)-C(23)	1.382(4)
C(24)-C(23)	1.382(3)
C(13)-O(12)-C(16)	101.45(16)
C(5)-O(1)-N(2)	104.81(16)
C(16)-N(15)-N(14)	106.42(18)
C(13)-N(14)-N(15)	105.77(17)
C(3)-N(2)-O(1)	104.41(16)
C(5)-N(4)-C(3)	101.83(17)
N(14)-C(13)-O(12)	113.99(18)
N(14)-C(13)-C(5)	126.46(19)
O(12)-C(13)-C(5)	119.49(19)
C(11)-C(6)-C(7)	119.6(2)
C(11)-C(6)-C(3)	121.18(18)
C(7)-C(6)-C(3)	119.2(2)
C(20)-C(19)-C(24)	118.7(2)
C(20)-C(19)-C(18)	118.7(2)
C(24)-C(19)-C(18)	122.6(2)
N(4)-C(5)-O(1)	115.14(18)
N(4)-C(5)-C(13)	129.97(19)
O(1)-C(5)-C(13)	114.87(19)
C(18)-C(17)-C(16)	124.0(2)

C(17)-C(18)-C(19)	125.4(2)
N(2)-C(3)-N(4)	113.8(2)
N(2)-C(3)-C(6)	121.88(18)
N(4)-C(3)-C(6)	124.29(19)
N(15)-C(16)-O(12)	112.36(17)
N(15)-C(16)-C(17)	126.9(2)
O(12)-C(16)-C(17)	120.75(19)
C(9)-C(8)-C(7)	119.98(19)
C(8)-C(7)-C(6)	120.1(2)
C(11)-C(10)-C(9)	120.0(2)
C(19)-C(20)-C(21)	121.1(2)
C(8)-C(9)-C(10)	120.3(2)
C(10)-C(11)-C(6)	120.09(18)
C(22)-C(21)-C(20)	119.6(2)
C(21)-C(22)-C(23)	120.1(2)
C(23)-C(24)-C(19)	120.0(2)
C(22)-C(23)-C(24)	120.6(2)

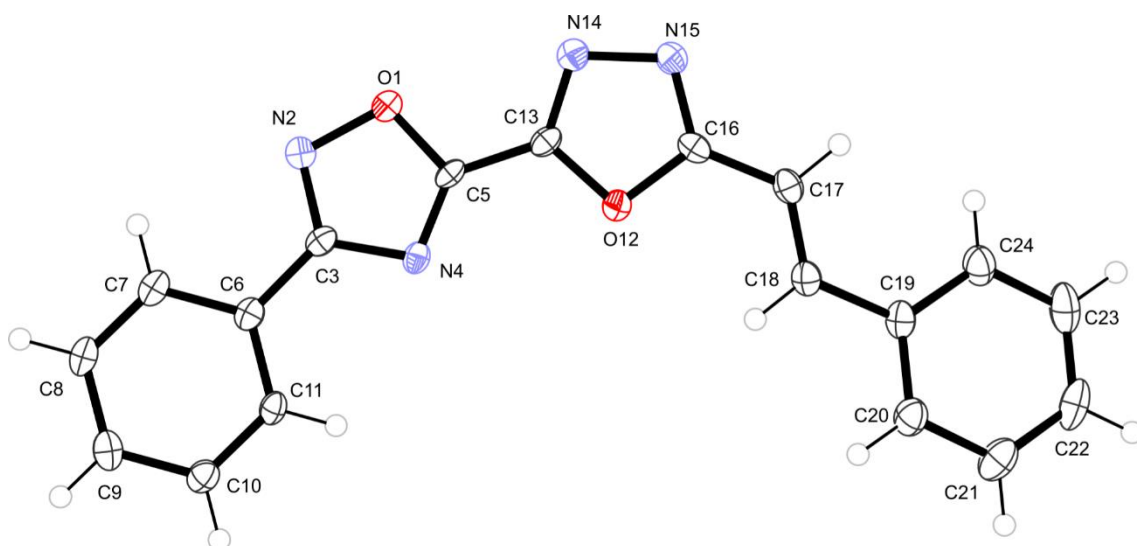


Figure S31. Projection of the molecular structure of the **7aa** ligand. Thermal ellipsoids with 50% probability level.

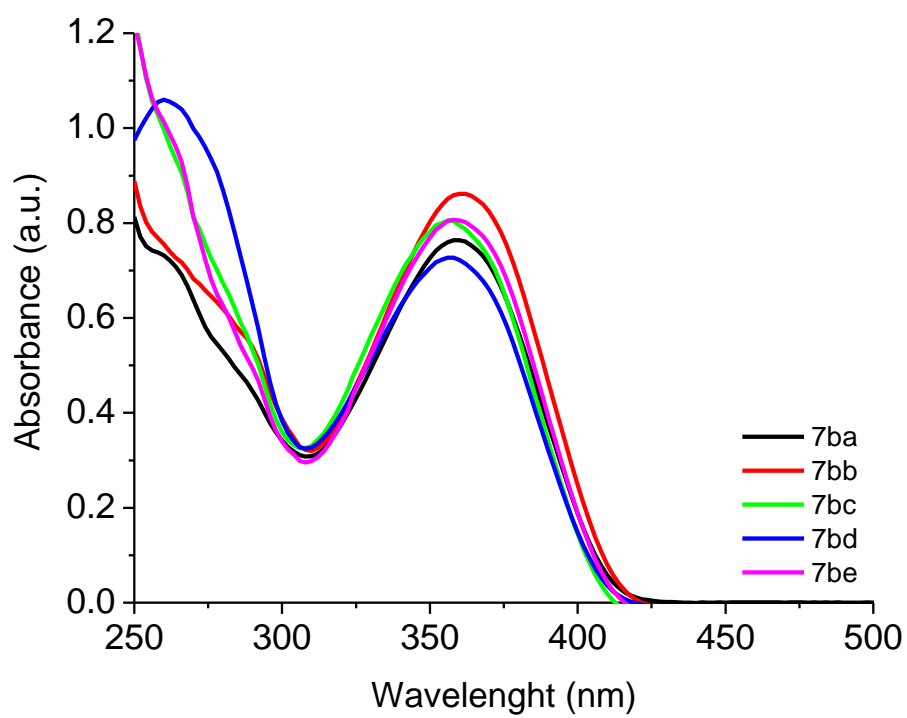


Figure S32. UV-Vis absorption spectra of derivatives **7ba-be** in CHCl_3 solution ($[\] = 2.00 \times 10^{-5} \text{ M}$).

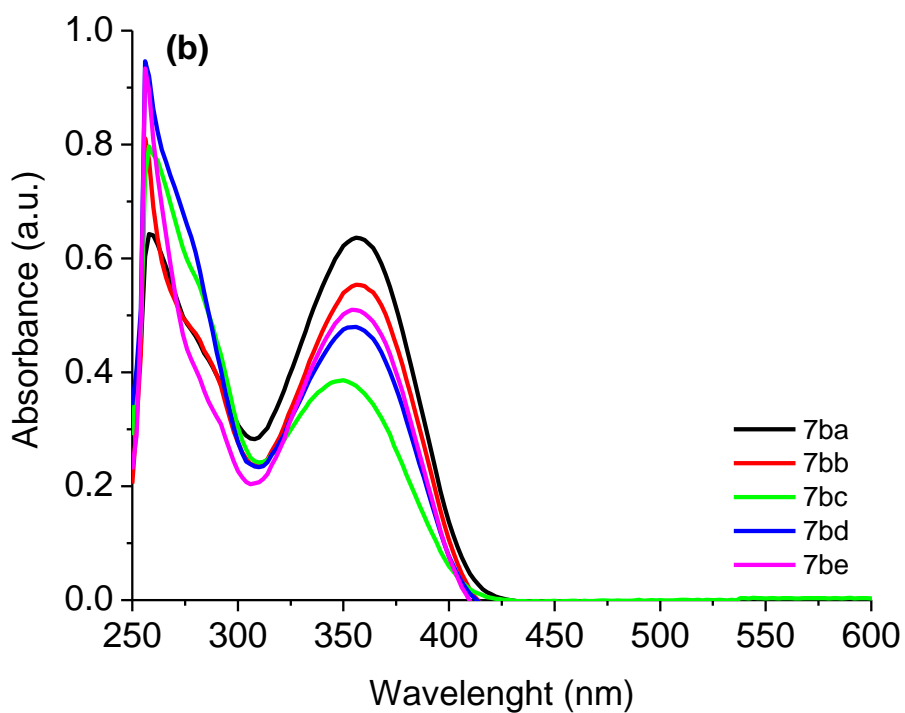
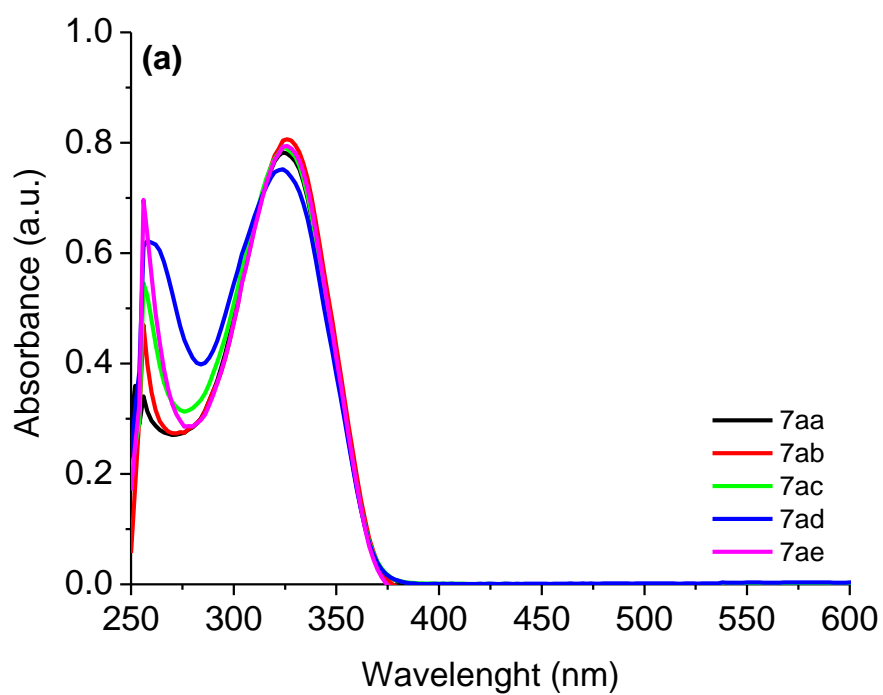


Figure S33. UV-Vis absorption spectra of derivatives (a) **7aa-ae** and (b) **7ba-be** in DMSO solution ($[] = 2.00 \times 10^{-5} \text{ M}$).

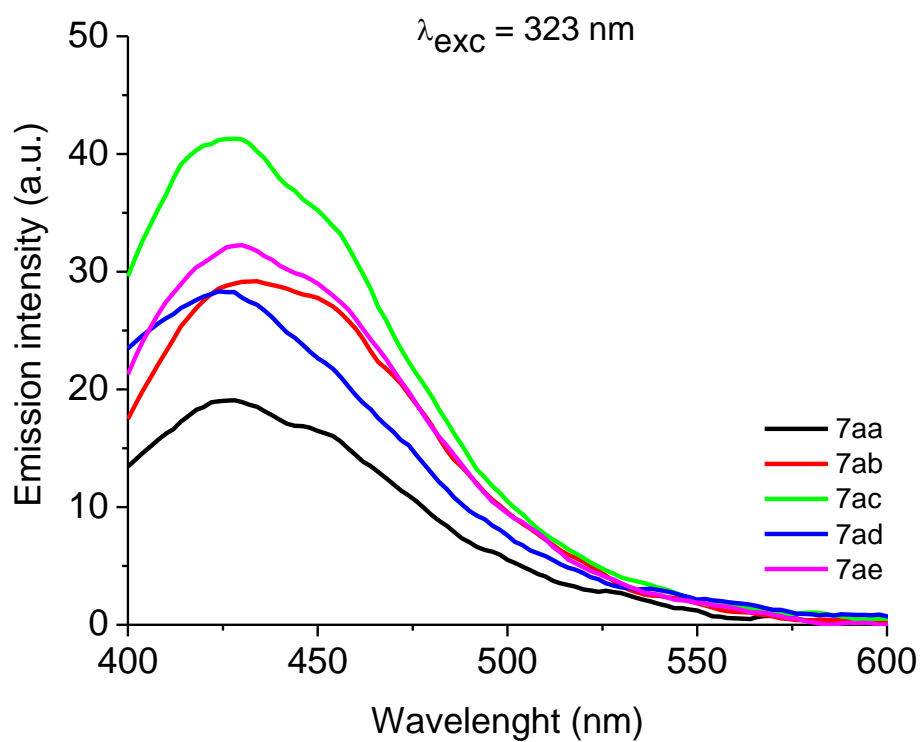


Figure S34. Steady-state emission spectra of derivatives **7aa-ae** ($\lambda_{\text{exc}} = 323 \text{ nm}$) in saturated argon chloroform solution ($[] = 1.00 \times 10^{-6} \text{ M}$).

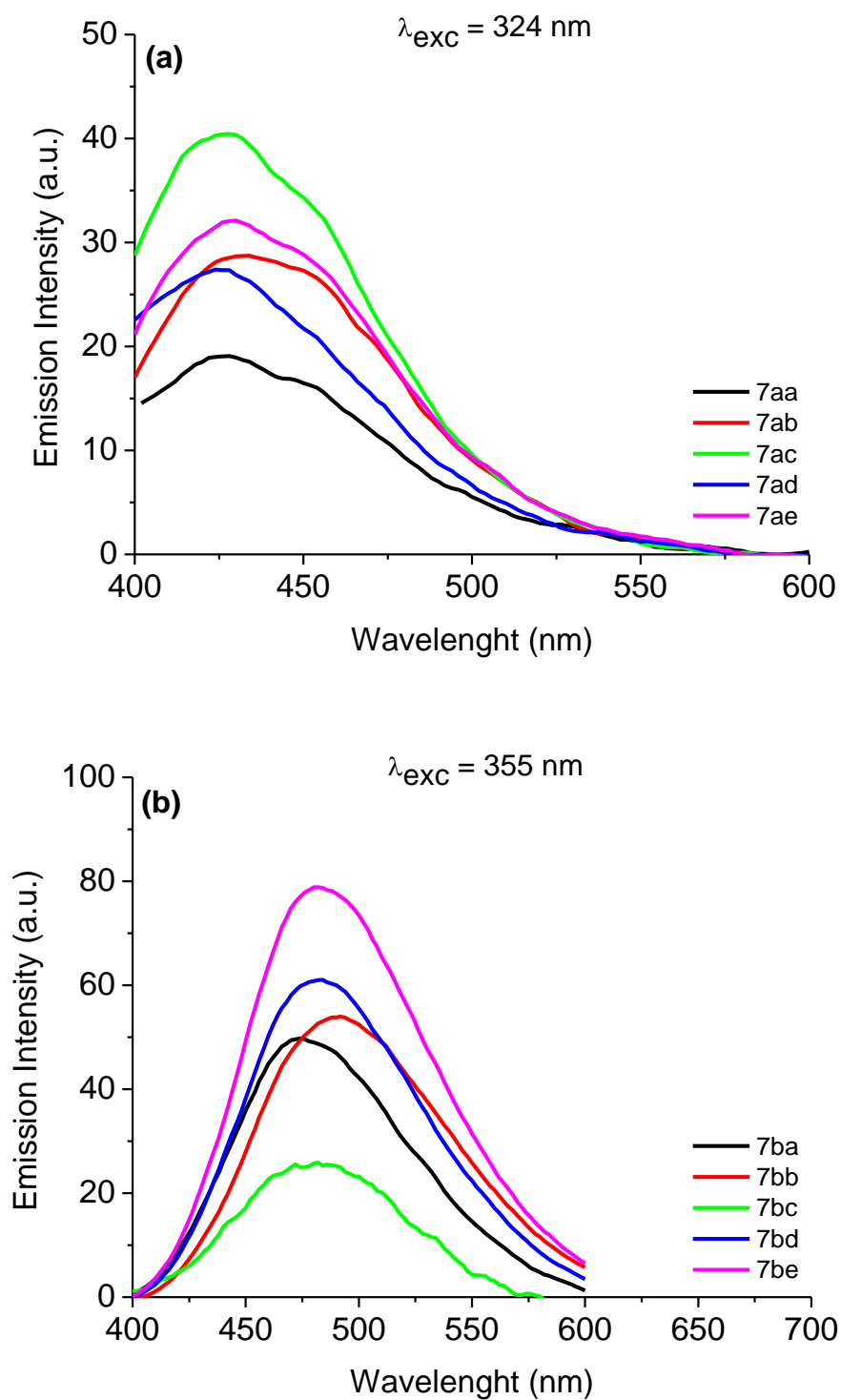


Figure S35. Steady-state emission spectra of derivatives **(a) 7aa-ae** ($\lambda_{\text{exc}} = 324 \text{ nm}$) and **(b) 7ba-be** ($\lambda_{\text{exc}} = 355 \text{ nm}$) in saturated argon DMSO solution ($[] = 1.00 \times 10^{-6} \text{ M}$).

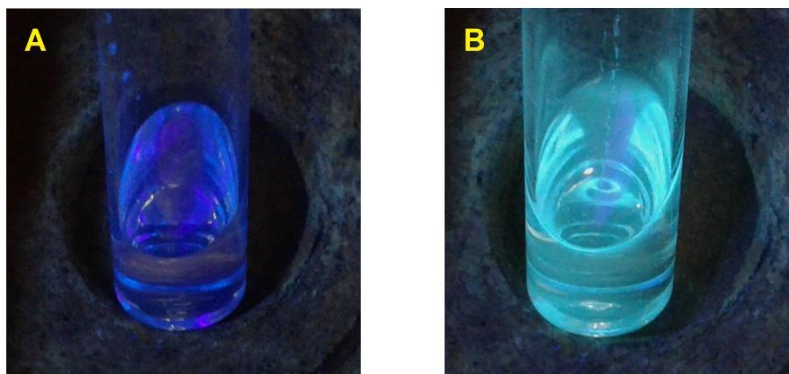


Figure S36. Representative samples of compound **7aa** at 1.0 mg/mL (1.05×10^{-3} M) solution in DMSO (A), and compound **7ba** at 1.0 mg/mL (9.10×10^{-4} M) solution in DMSO (B), both under UV illumination (365 nm).

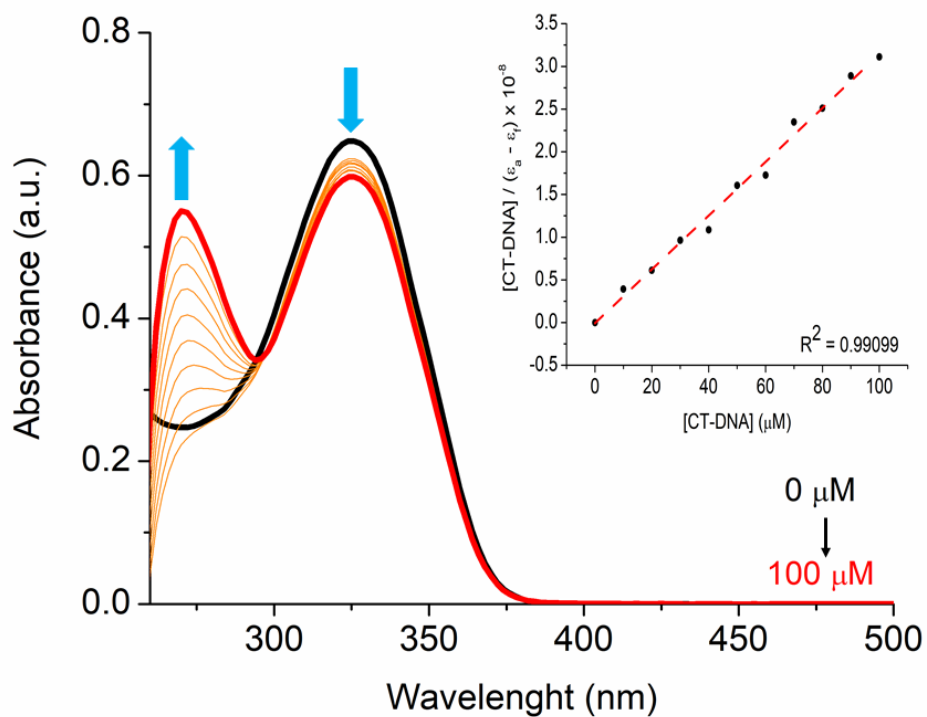


Figure S37. UV-Vis titration absorption spectra of derivative **7ab**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}] / (\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

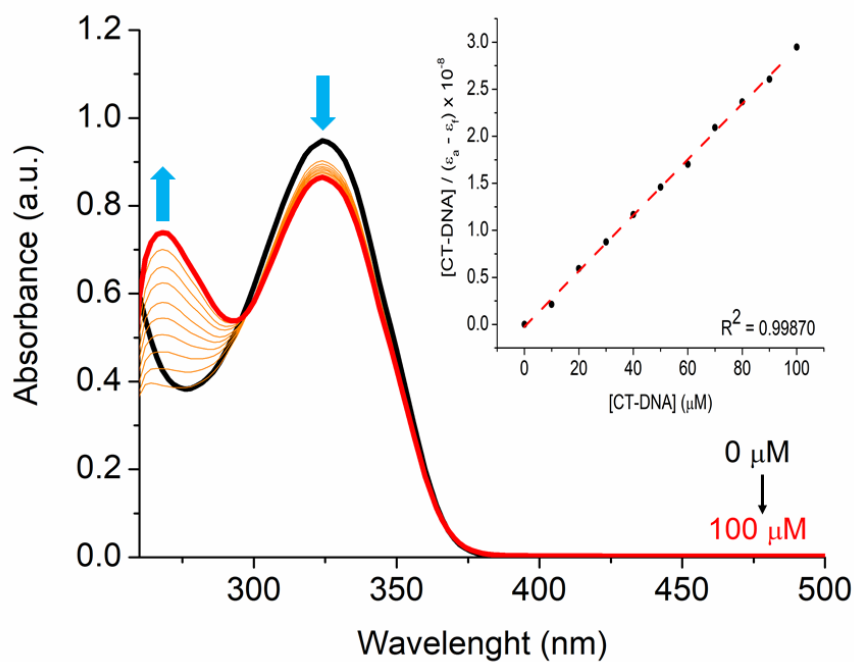


Figure S38. UV-Vis titration absorption spectra of derivative **7ac**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

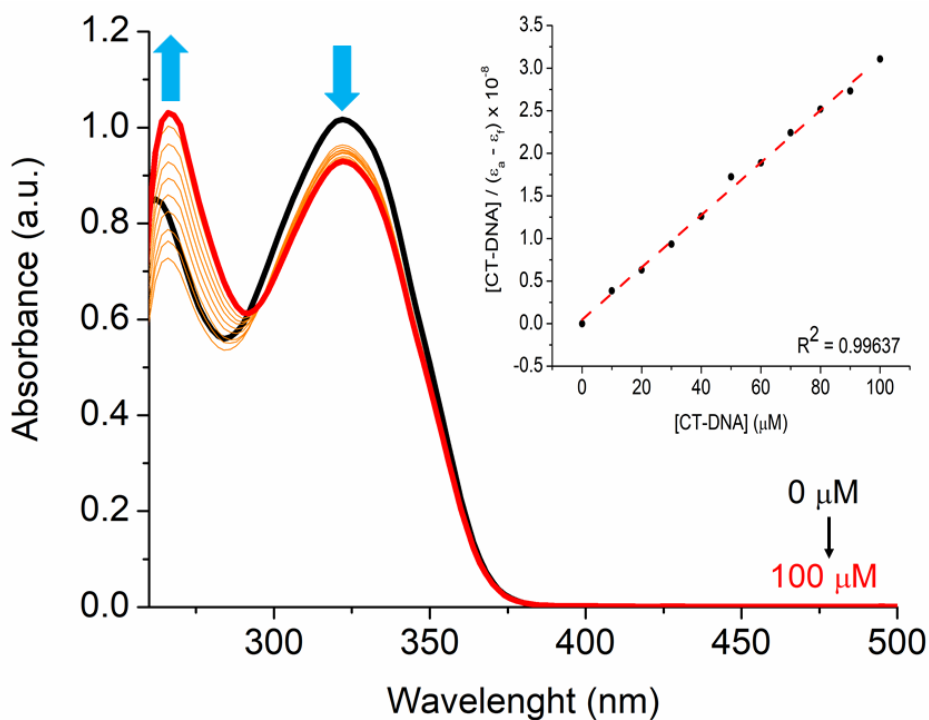


Figure S39. UV-Vis titration absorption spectra of derivative **7ad**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

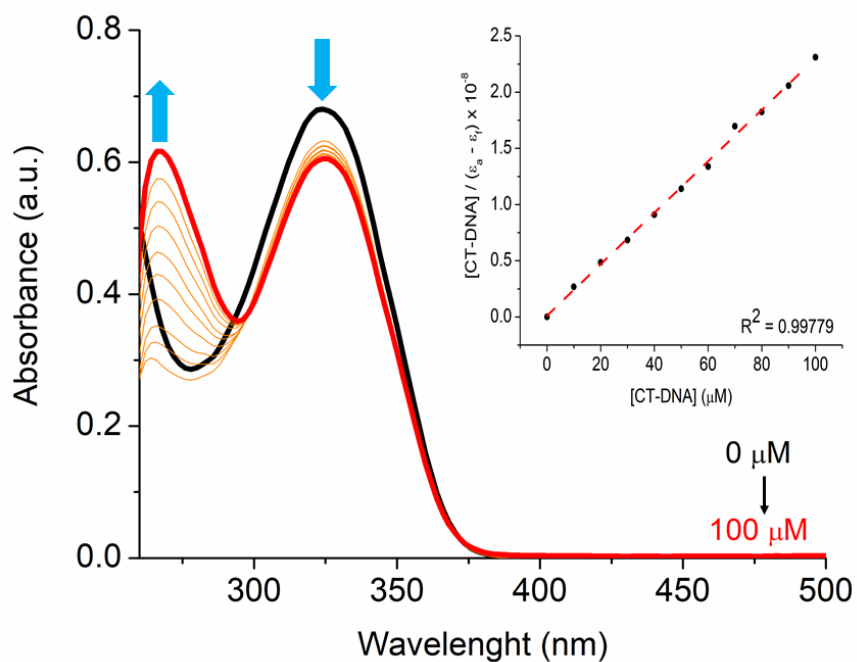


Figure S40. UV-Vis titration absorption spectra of derivative **7ae**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

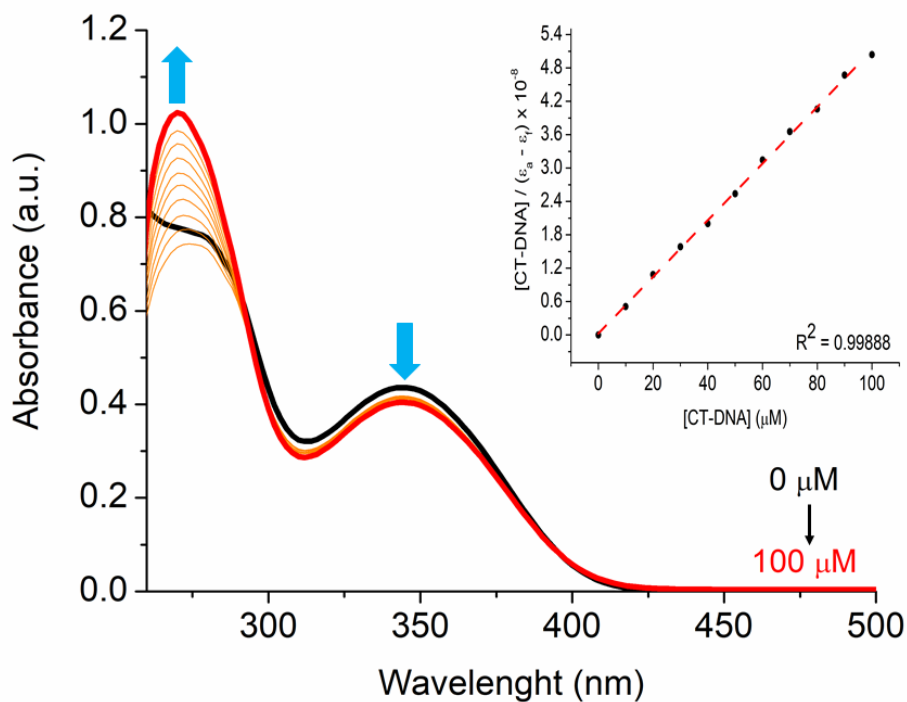


Figure S41. UV-Vis titration absorption spectra of derivative **7bb**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

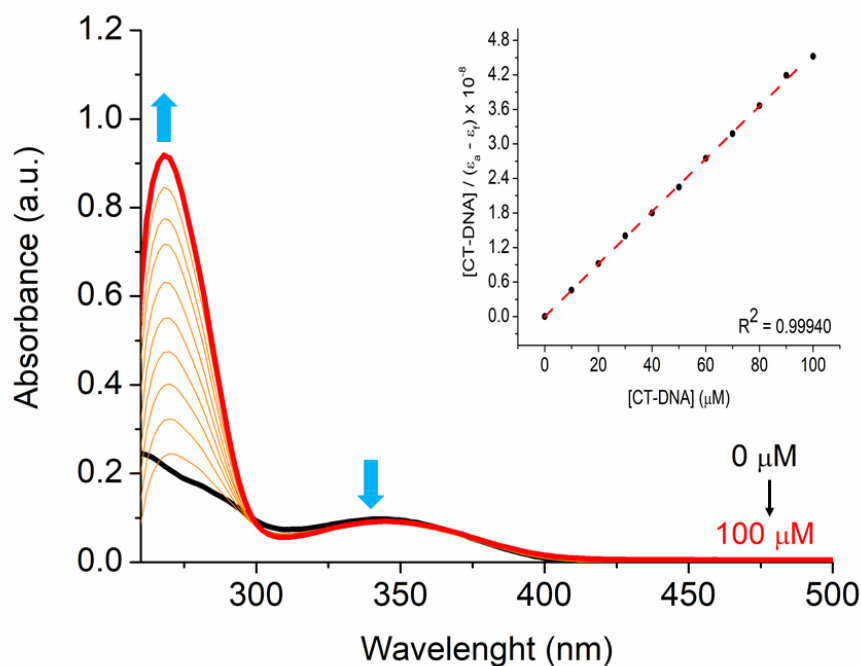


Figure S42. UV-Vis titration absorption spectra of derivative **7bc**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

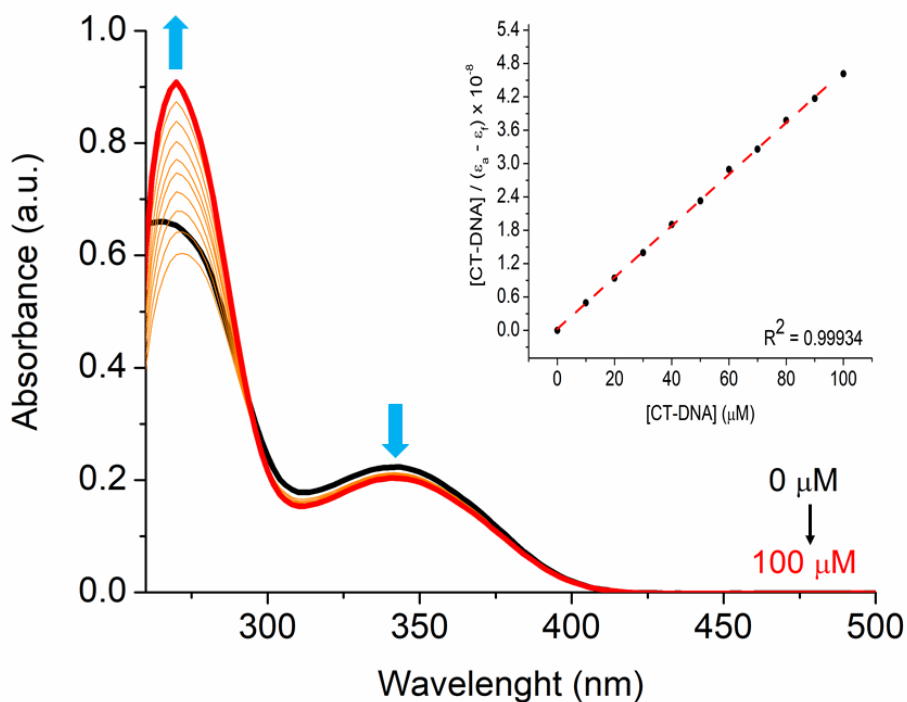


Figure S43. UV-Vis titration absorption spectra of derivative **7bd**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

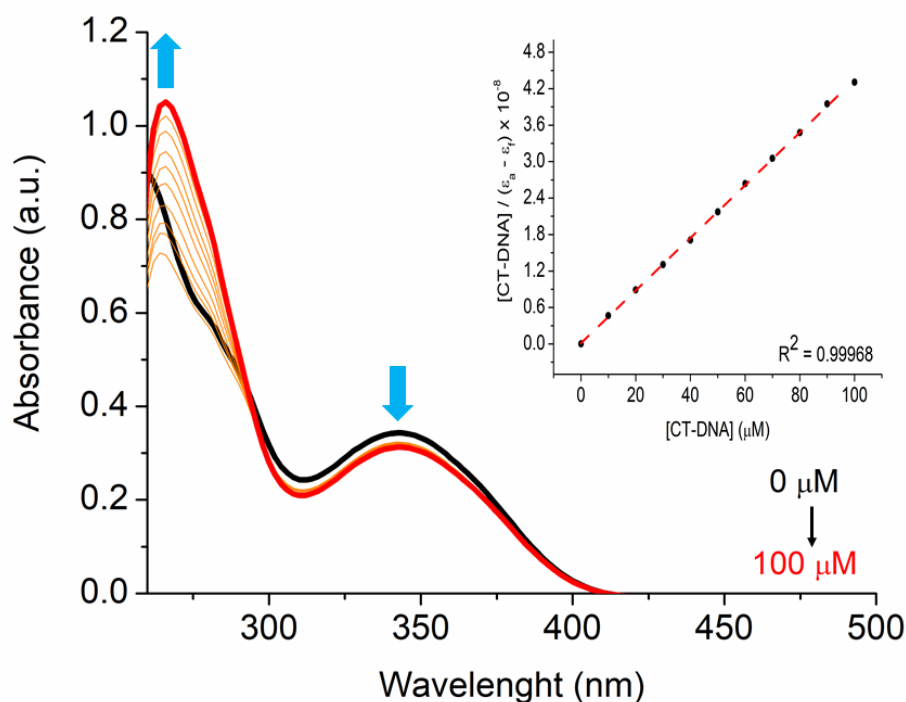


Figure S44. UV-Vis titration absorption spectra of derivative **7be**, in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of CT-DNA ranged from 0 to 100 μM . Insert graph shows the plot of $[\text{DNA}]/(\epsilon_a - \epsilon_f)$ versus $[\text{DNA}]$.

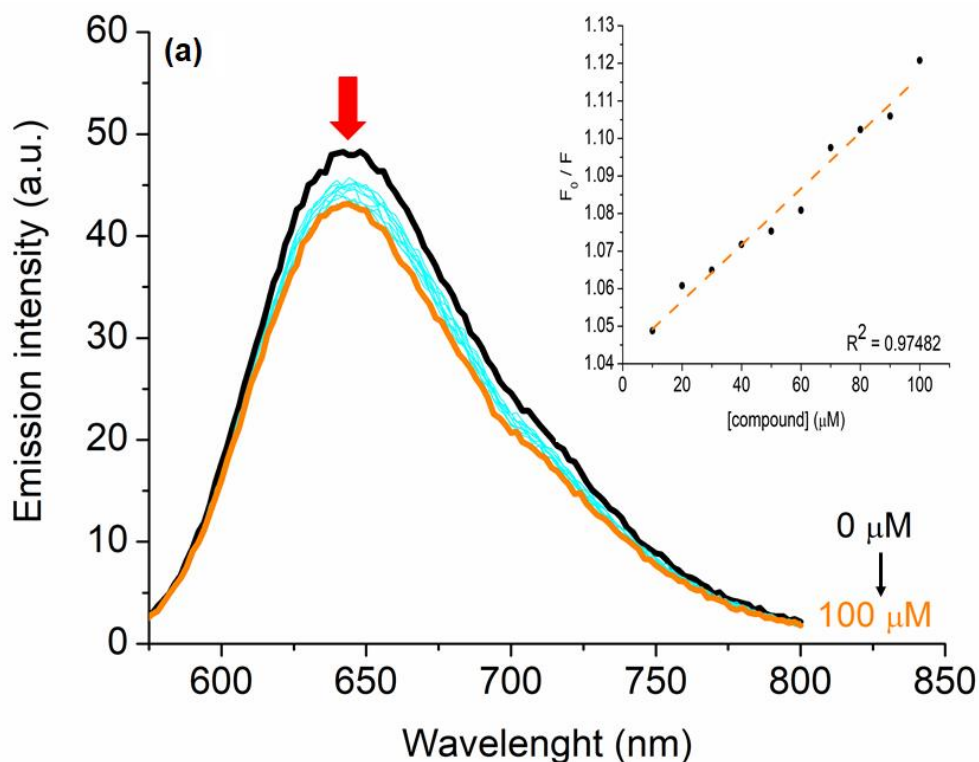


Figure S45. Competitive emission spectra of EB-DNA with derivative **7aa** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus $[\text{compound}]$.

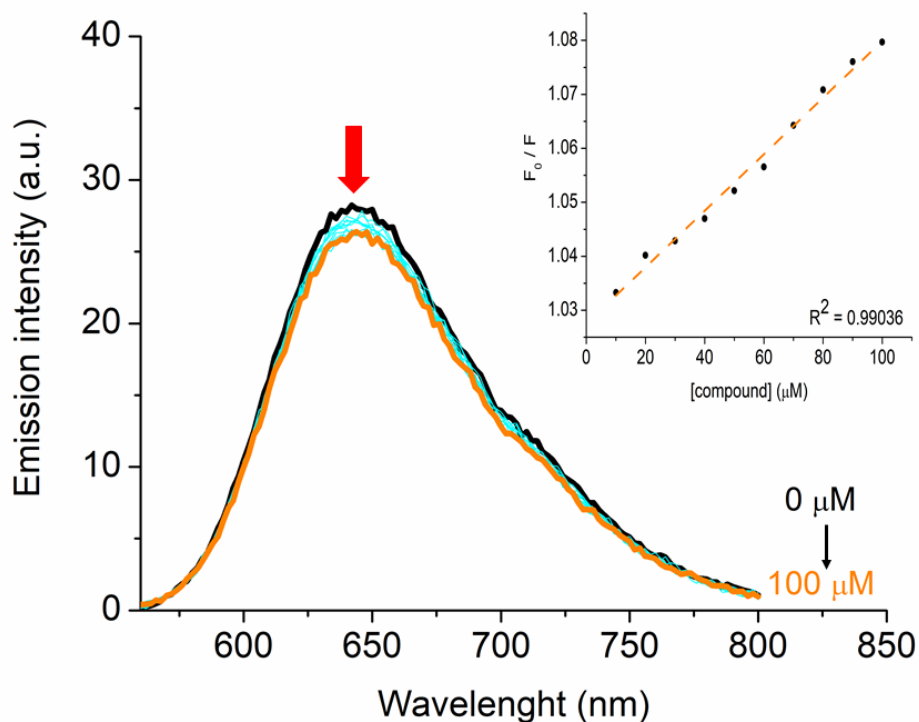


Figure S46. Competitive emission spectra of EB-DNA with derivative **7ab** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

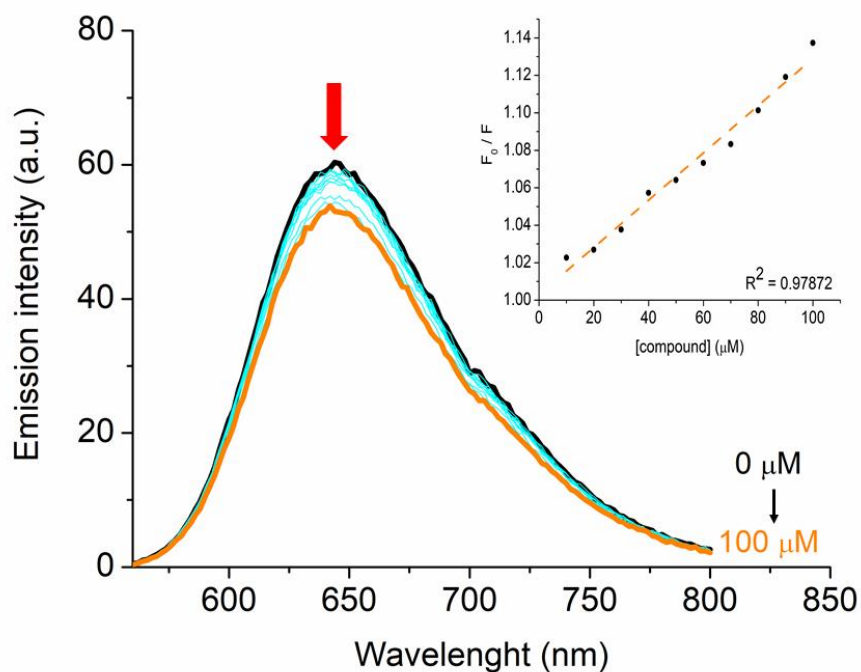


Figure S47. Competitive emission spectra of EB-DNA with derivative **7ac** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

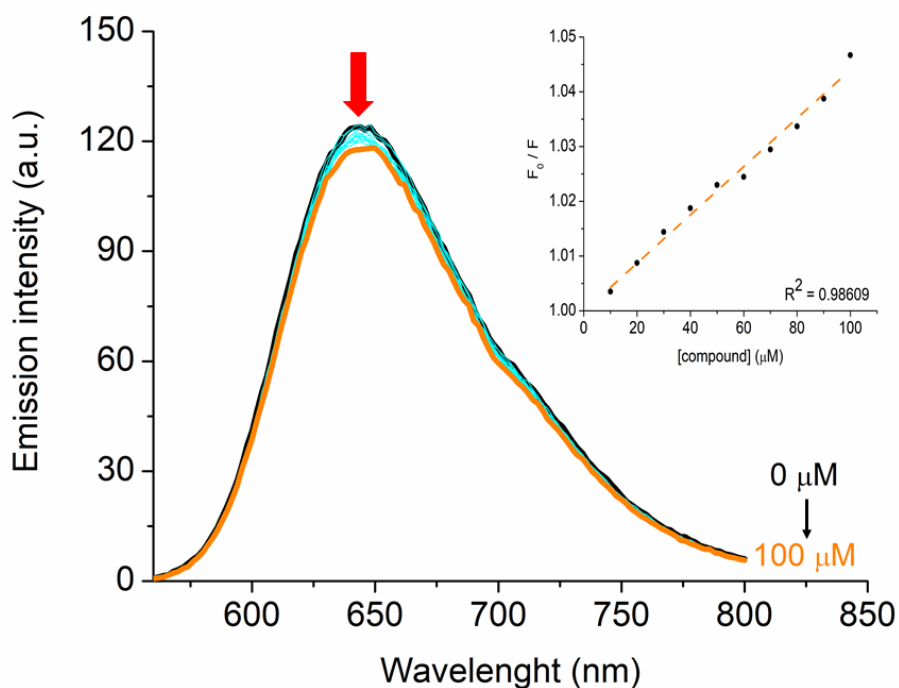


Figure S48. Competitive emission spectra of EB-DNA with derivative **7ad** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

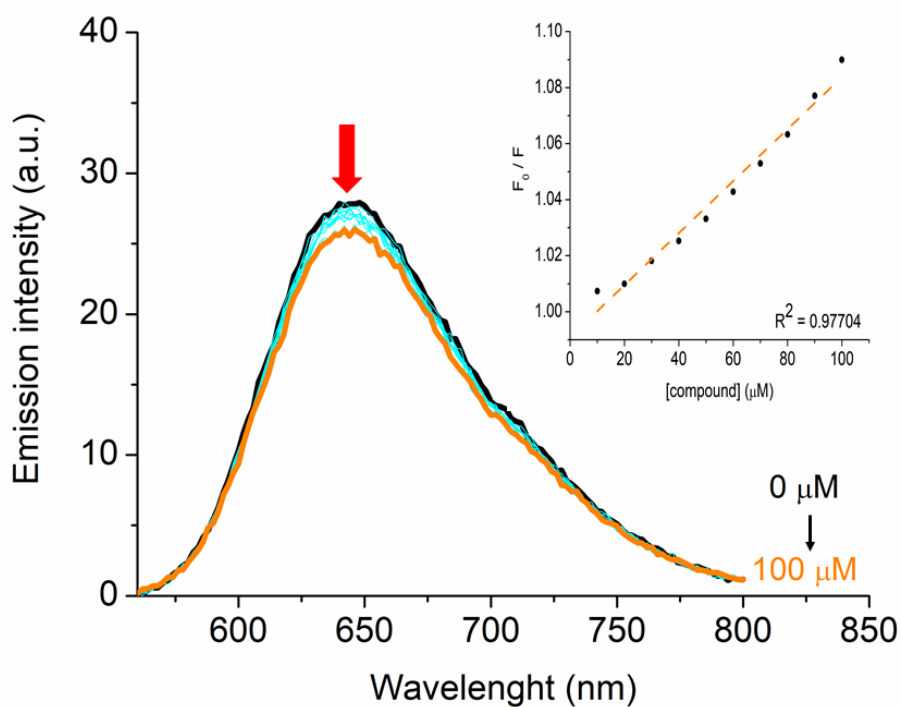


Figure S49. Competitive emission spectra of EB-DNA with derivative **7ae** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

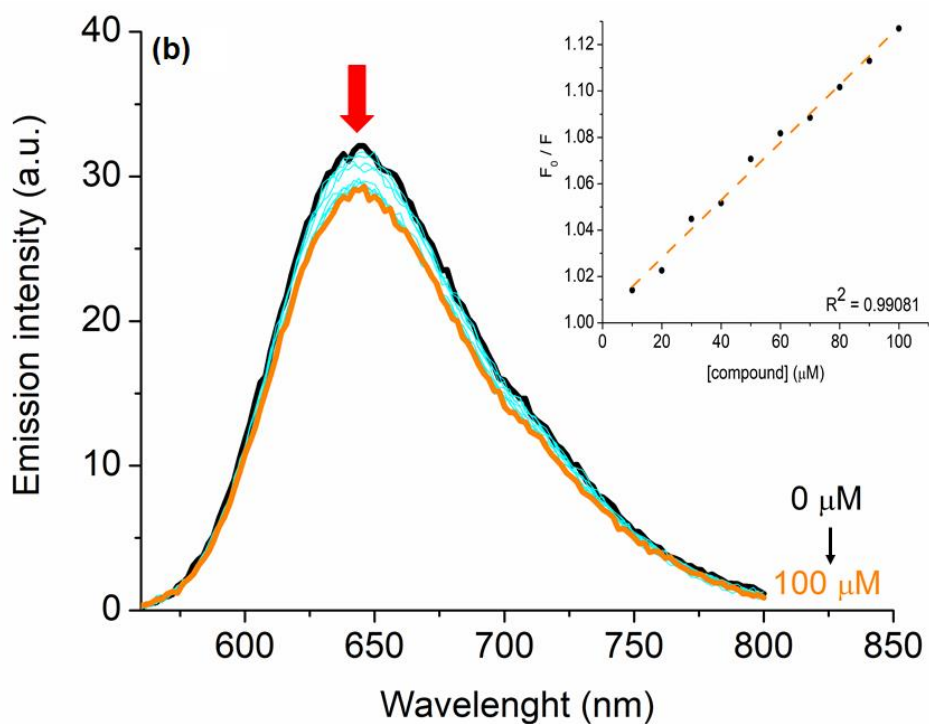


Figure S50. Competitive emission spectra of EB-DNA with derivative **7ba** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

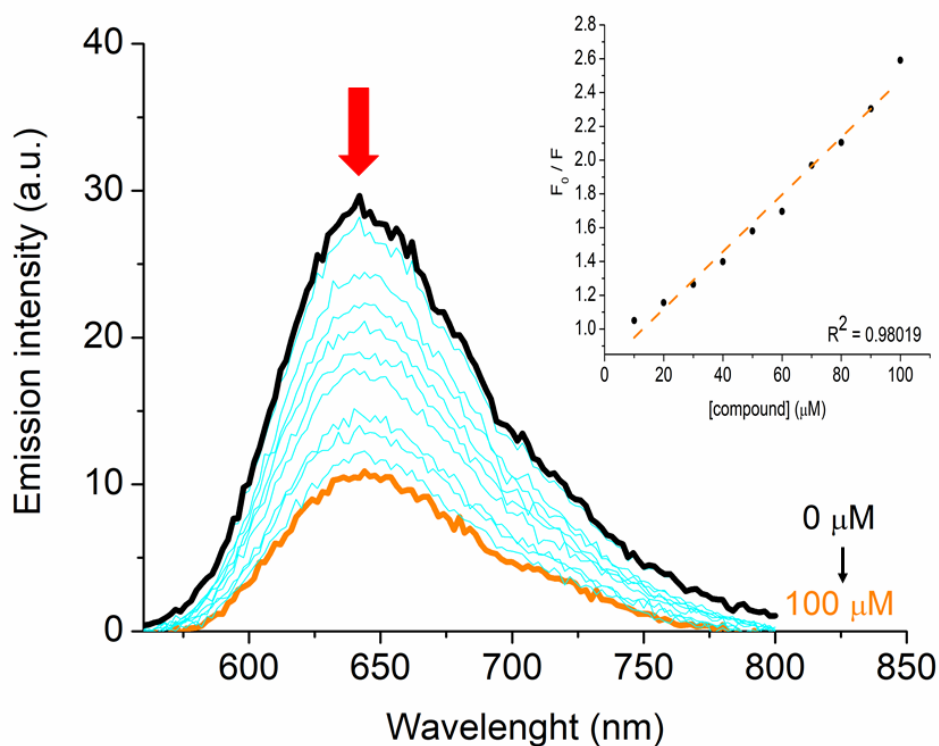


Figure S51. Competitive emission spectra of EB-DNA with derivative **7bb** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

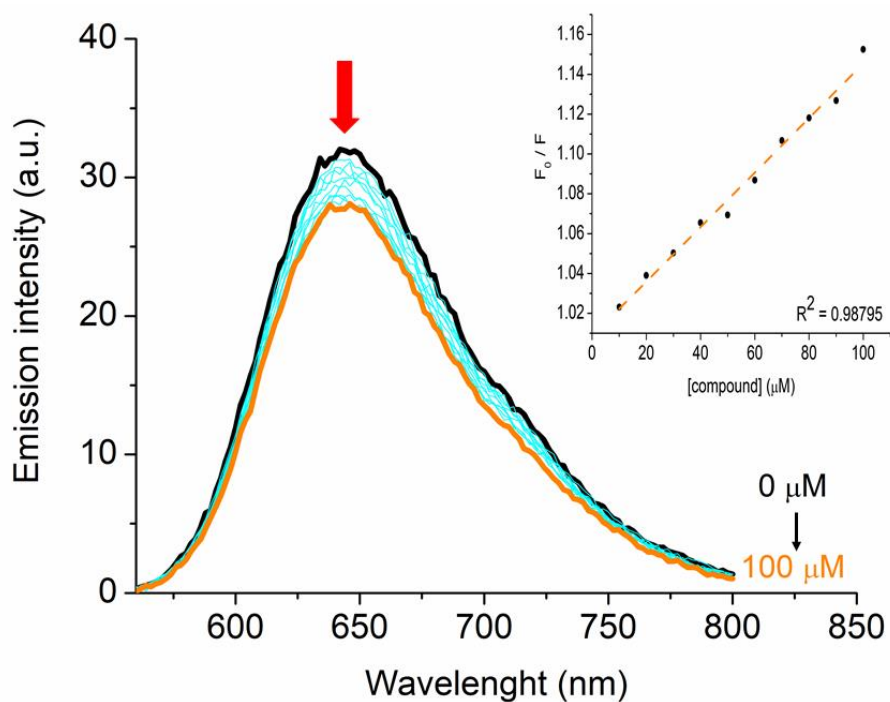


Figure S52. Competitive emission spectra of EB-DNA with derivative **7bc** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

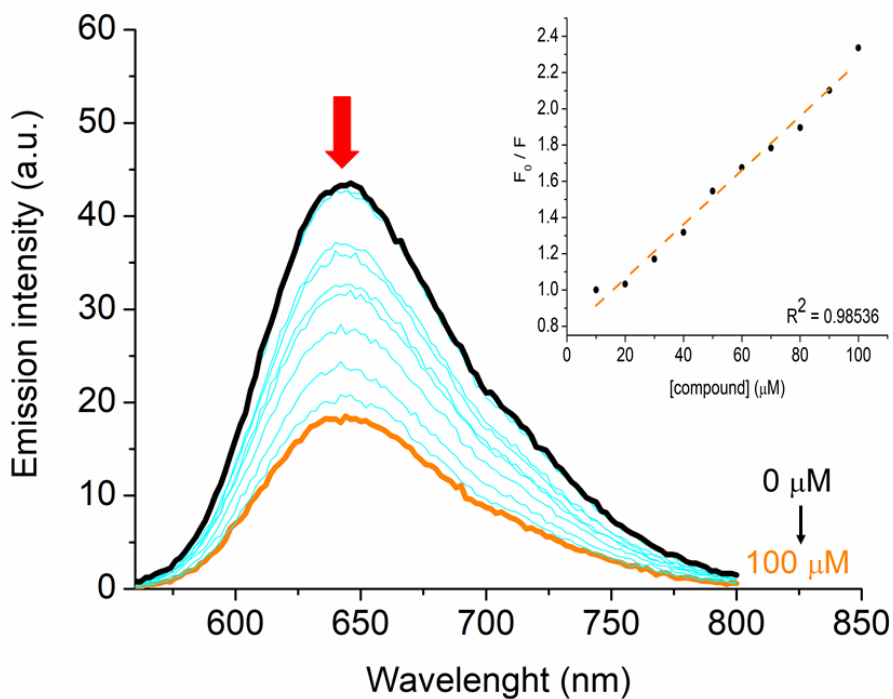


Figure S53. Competitive emission spectra of EB-DNA with derivative **7bd** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

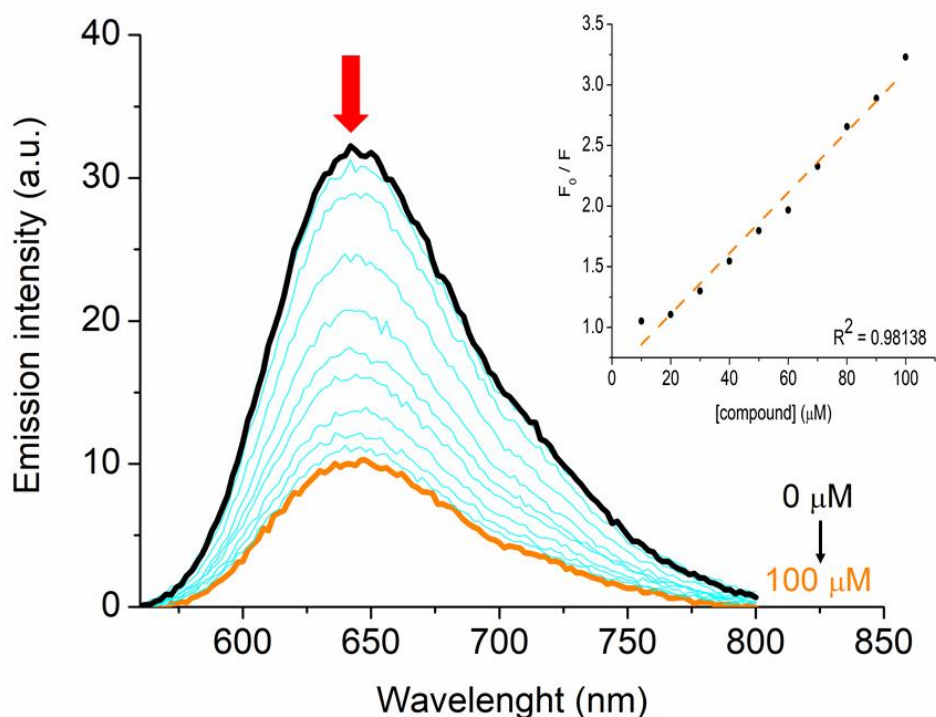


Figure S54. Competitive emission spectra of EB-DNA with derivative **7be** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

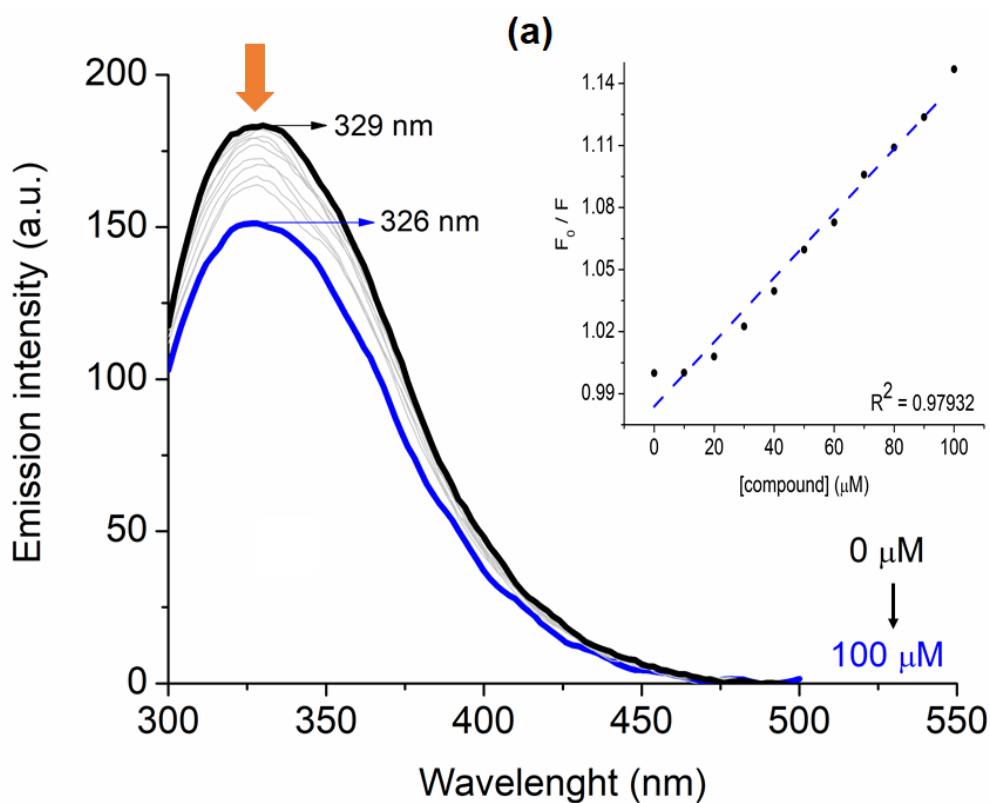


Figure S55. HSA-binding emission spectra with derivative **7aa** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . Insert graph shows the plot of F_0/F versus [compound].

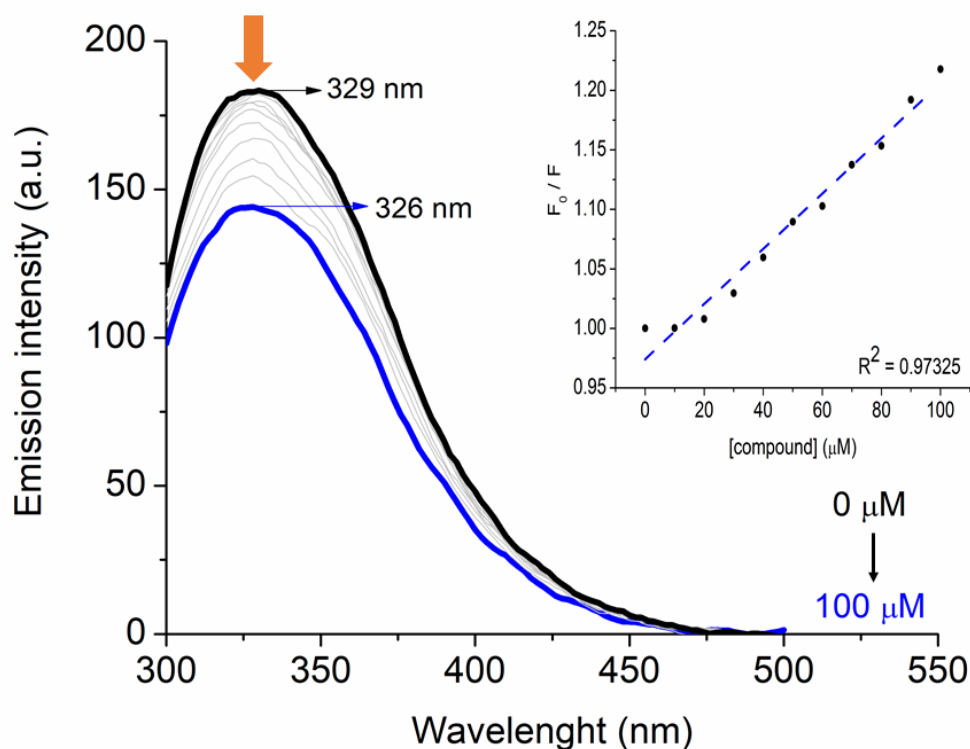


Figure S56. HSA-binding emission spectra with derivative **7ab** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph shows the plot of F_0/F versus [compound].*

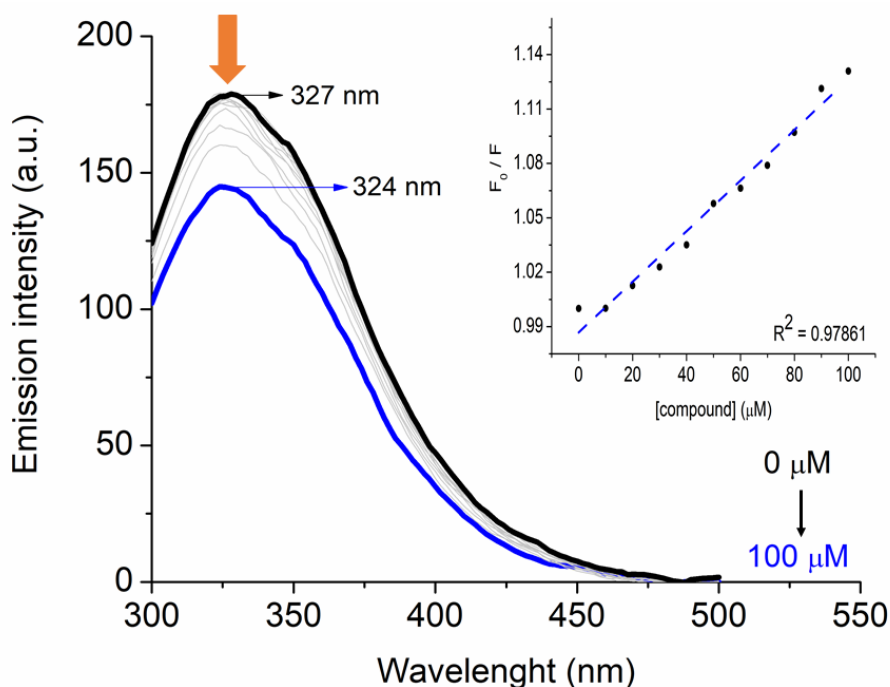


Figure S57. HSA-binding emission spectra with derivative **7ac** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph shows the plot of F_0/F versus [compound].*

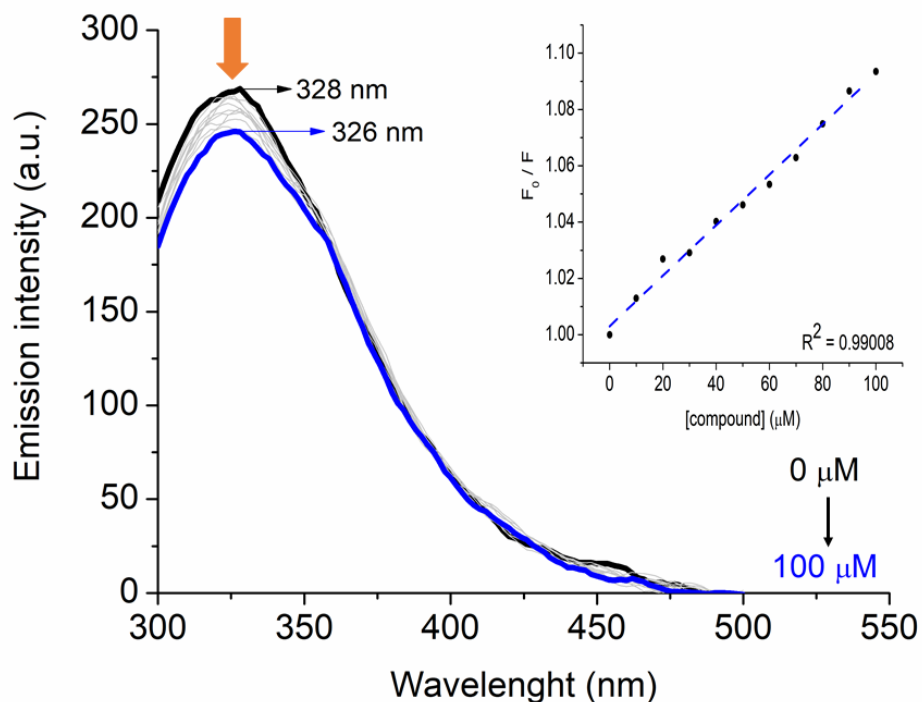


Figure S58. HSA-binding emission spectra with derivative **7ad** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph* shows the plot of F_0/F versus [compound].

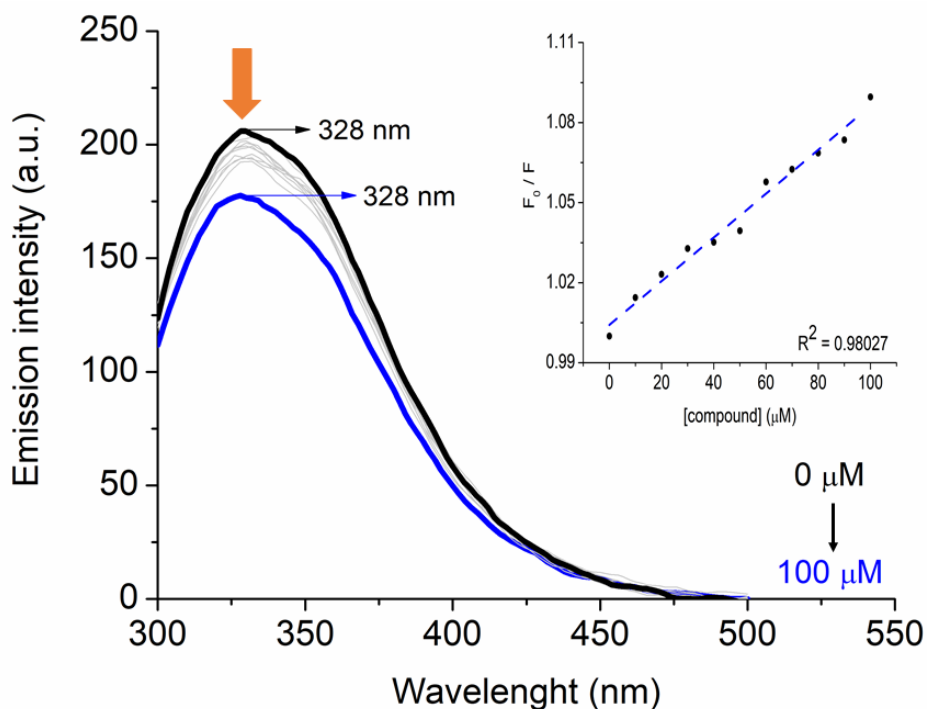


Figure S59. HSA-binding emission spectra with derivative **7ae** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph* shows the plot of F_0/F versus [compound].

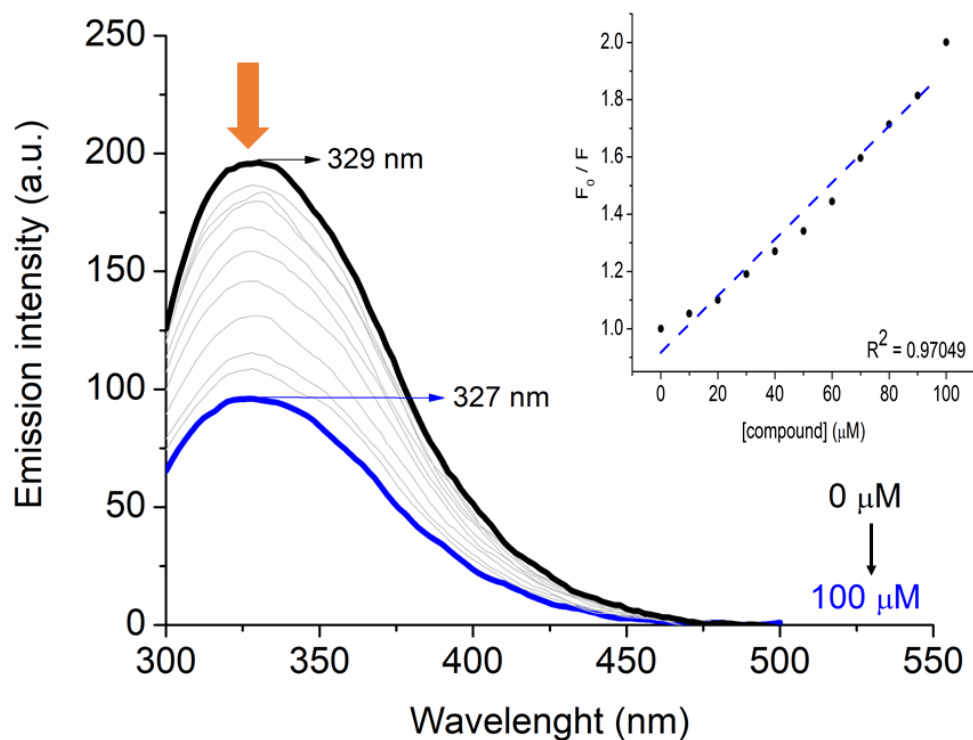


Figure S60. HSA-binding emission spectra with derivative **7bb** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph shows the plot of F_0/F versus [compound].*

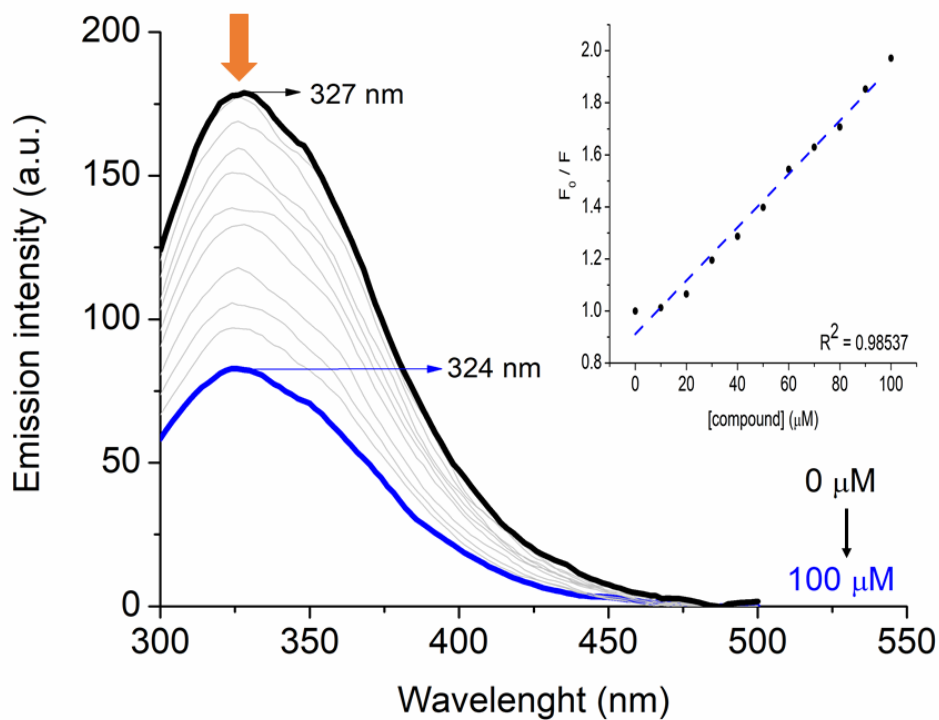


Figure S61. HSA-binding emission spectra with derivative **7bc** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph shows the plot of F_0/F versus [compound].*

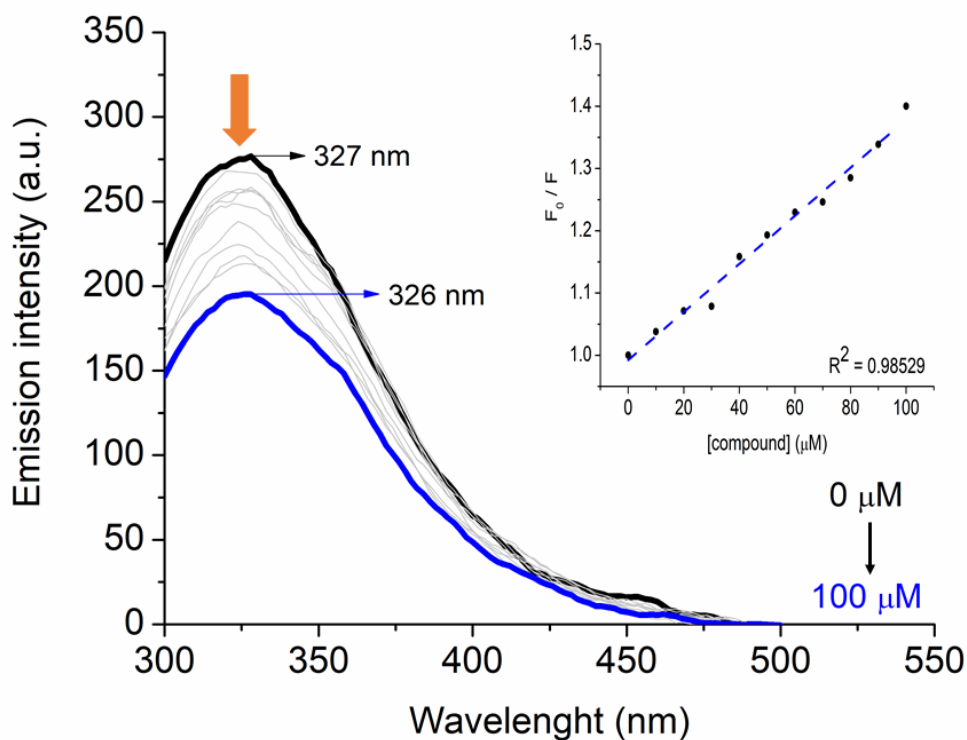


Figure S62. HSA-binding emission spectra with derivative **7bd** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph shows the plot of F_0/F versus [compound].*

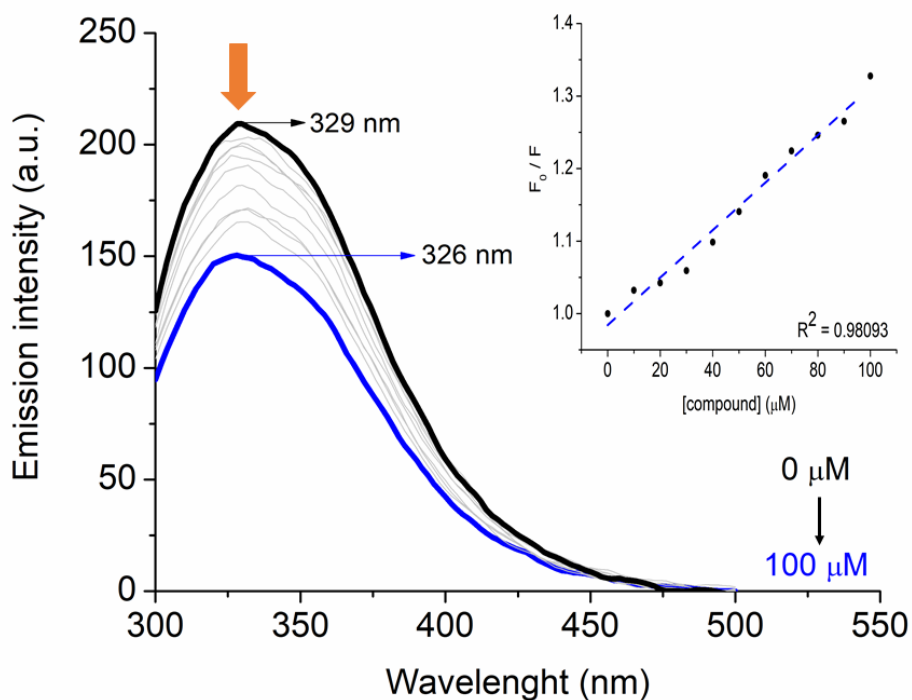


Figure S63. HSA-binding emission spectra with derivative **7be** in a DMSO (2%)/Tris-HCl buffer (pH 7.2) solution. The concentration of compound ranged from 0 to 100 μM . *Insert graph shows the plot of F_0/F versus [compound].*

Table S3. The main nucleobases and interactions involved in the association DNA:**7aa-ae** and DNA:**7ba-be** in the minor groove.

Compound	Nucleobase	Interaction	Distance (Å)
7aa-ac, ae	DG-04	Van der Waals	2.30
	DA-05	Van der Waals	2.10
	DA-06	Van der Waals	2.70
	DT-07	Van der Waals	2.40
	DT-08	Van der Waals	2.30
	DA-17	Van der Waals	2.80
	DA-18	Van der Waals	1.70
	DT-19	Van der Waals	2.90
	DT-20	Van der Waals	3.30
	DC-21	Van der Waals	3.40
7ad	DA-06	Van der Waals	2.00
	DT-07	Van der Waals	2.70
	DT-08	Van der Waals	2.40
	DA-18	Van der Waals	2.50
	DT-19	Van der Waals	2.40
	DT-20	Van der Waals	3.00
7ba, bc	DA-05	Van der Waals	3.40
	DA-06	Van der Waals	2.20
	DT-07	Van der Waals	2.40
	DT-08	Van der Waals	2.30
	DC-09	Van der Waals	2.50
	DG-16	Van der Waals	2.80
	DA-17	Van der Waals	2.80
	DA-18	Van der Waals	2.70
	DT-19	Van der Waals	3.00
	DT-20	Van der Waals	2.50
7bb	DA-06	Van der Waals	3.10
	DT-07	Van der Waals	2.60
	DC-09	Van der Waals	2.50
	DG-16	Van der Waals	1.70
	DA-17	Van der Waals	3.00
	DA-18	Van der Waals	3.40
	DT-19	Van der Waals	2.50
	DT-19	Van der Waals	2.50
7bd, be	DC-03	Van der Waals	3.10
	DG-04	Van der Waals	2.10
	DA-05	Van der Waals	2.60
	DA-06	Van der Waals	1.50
	DT-07	Van der Waals	2.30
	DT-08	Van der Waals	3.00
	DA-18	Van der Waals	2.50
	DT-19	Van der Waals	2.80

	DT-20	Van der Waals	2.90
	DC-21	Van der Waals	3.60
	DG-22	Van der Waals	2.00
EB	DA-06	Van der Waals	3.60
	DT-07	Van der Waals	2.00
	DT-08	Hydrogen bonding	2.90
	DA-17	Van der Waals	3.60
	DA-18	Van der Waals	1.40
	DT-19	Van der Waals	2.40

Table S4. The main amino acid residues and interactions involved in the association HSA:**7aa-ae** and HSA:**7ba-be** in the site II (subdomain IIIA).

Compound	Amino acid residue	Interaction	Distance (Å)
7aa-ac	Ile-388	Van der Waals	1.30
	Tyr-411	Hydrogen bonding	2.10
	Lys-414	Van der Waals	2.80
	Val-415	Van der Waals	2.30
	Val-433	Van der Waals	3.30
	Cys-437	Van der Waals	2.50
	Leu-453	Van der Waals	2.40
	Leu-457	Van der Waals	3.00
	Leu-460	Van der Waals	2.80
	Phe-488	Van der Waals	3.60
	Ser-489	Hydrogen bonding	1.90
7ad/ae	Ile-388	Van der Waals	1.50
	Tyr-411	Hydrogen bonding	2.10
	Leu-430	Van der Waals	2.10
	Val-433	Van der Waals	2.80
	Cys-437	Van der Waals	2.10
	Leu-460	Van der Waals	2.50
	Phe-488	Van der Waals	3.60
	Ser-489	Van der Waals	2.00
7ba-be	Leu-387	Van der Waals	2.90
	Ile-388	Van der Waals	1.30
	Tyr-411	Hydrogen bonding	2.10
	Val-415	Van der Waals	1.90
	Val-418	Van der Waals	1.30
	Val-426	Van der Waals	1.30
	Leu-430	Van der Waals	2.10
	Val-433	Van der Waals	1.70
	Gly-434	Van der Waals	1.90
	Arg-485	Hydrogen bonding	2.70

