Thiacalixarene "knot" effect on protein binding by oligolactic acid particles

Olga A Mostovaya^a, Vladimir V. Gorbachuk^a, Olga B. Bazanova^b, Alexander V. Gerasimov^a, Vladimir G. Evtugyn^c, Yury N. Osin^c, Viktor D. Myakushev^d, Ildar Kh. Rizvanov^b, Ivan I. Stoikov^a

^{a.} Kazan Federal University, A.M. Butlerov Chemistry Institute, 420008, Kremlevskaya Street, 18, Kazan, Russian Federation.E-mail: Ivan.Stoikov@mail.ru; Fax: +7-8432-752253; Tel: +7-8432-337463

b. Arbuzov Institute of Organic and Physical Chemistry, FRC Kazan Scientific Center of RAS

^{c.} Interdisciplinary Center for Analytical Microscopy of Kazan Federal University, 420008 Kazan, Russian Federation

^{d.} Enikolopov Institute of Synthetic Polymeric Materials, Russian Academy of Sciences, Moscow 117393, Russian Federation

Electronic Supplementary Information

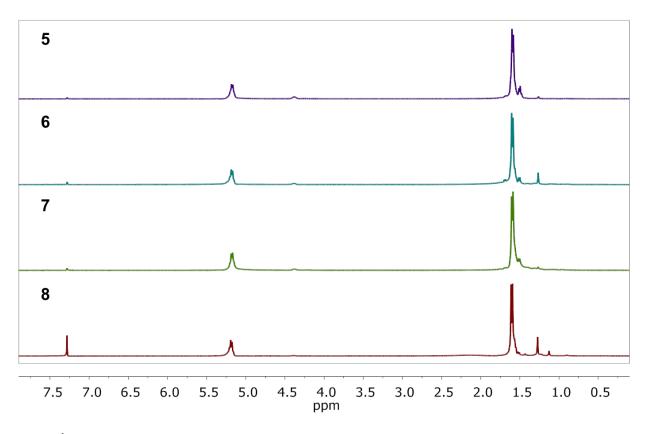


Fig.S1. ¹H NMR (CDCl₃) spectra of products 5-8.

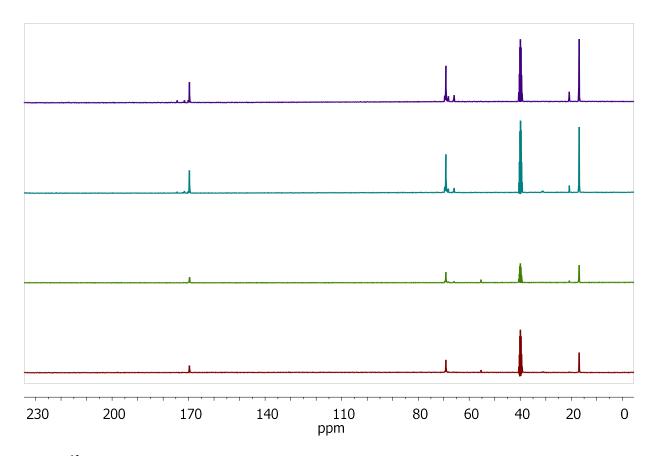


Fig.S2. ¹³C{H} NMR (DMSO D_6) spectra of products 5-8.

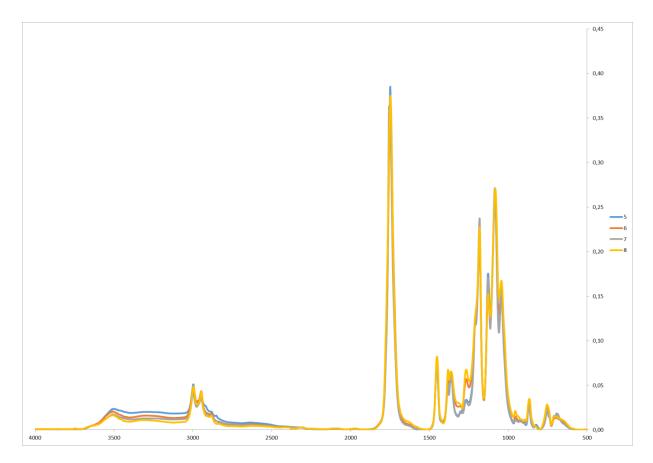


Fig.S3. FTIR-ATR spectra of products 5-8.

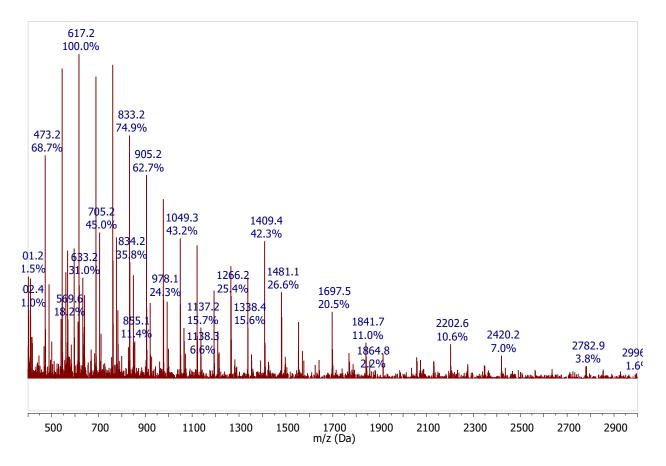


Fig.S4. MALDI mass-spectrum of product 5, matrix - 2,5-dihydroxybenzoic acid

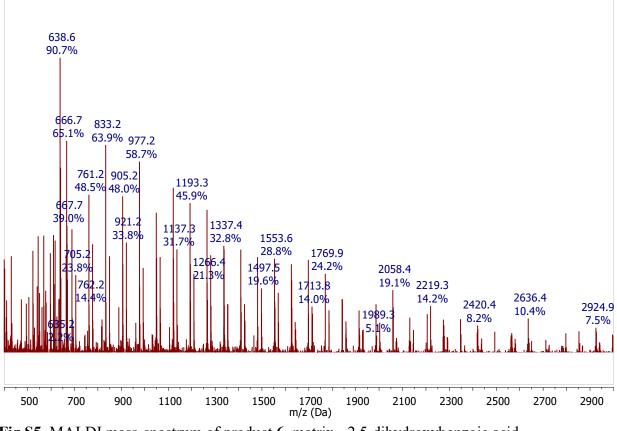


Fig.S5. MALDI mass-spectrum of product 6, matrix - 2,5-dihydroxybenzoic acid

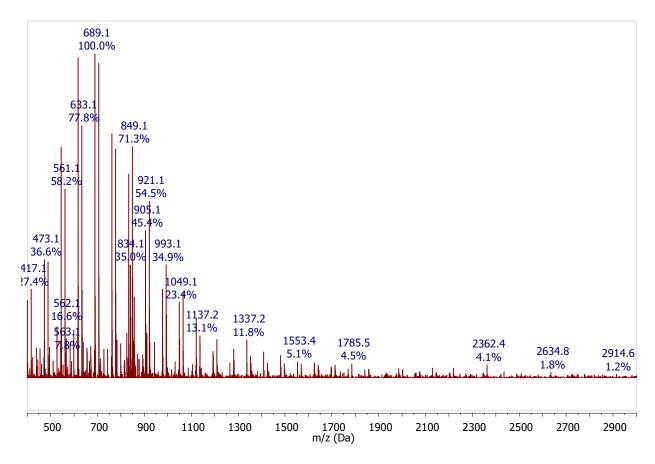


Fig.S6. MALDI mass-spectrum of product 7, matrix - 2,5-dihydroxybenzoic acid

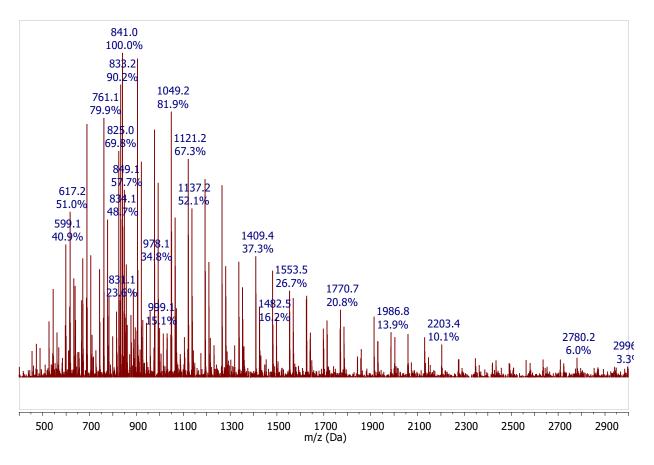


Fig.S7. MALDI mass-spectrum of product 8, matrix - 2,5-dihydroxybenzoic acid

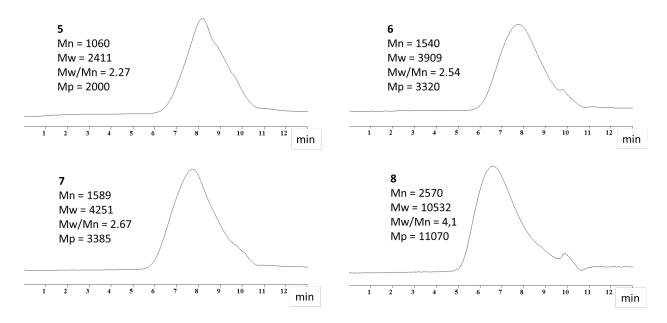


Fig.S8. GPC curves of products 5-8 (eluent-THF, calibrated by PS standards)

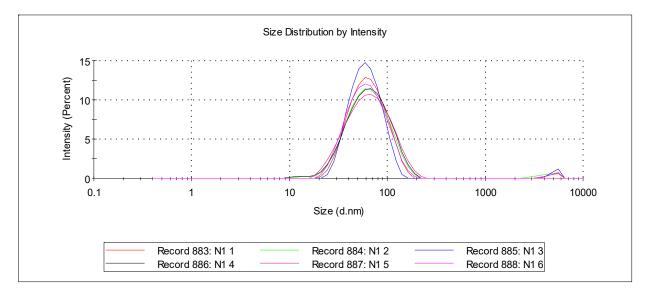


Fig.S9. Size distribution by intensity (DLS) of solution of the macrocycle **6**(*cone*, 0.19 mg/ml) in the 50 mM phosphate buffer at pH 7.4.

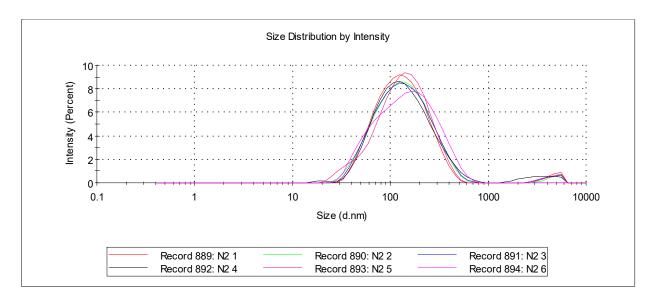


Fig.S10. Size distribution by intensity (DLS)of solution of the macrocycle 7(*partial cone*, 0.19 mg/ml) in the 50 mM phosphate buffer at pH 7.4.

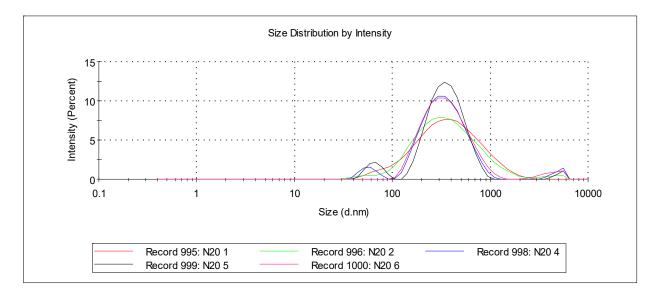


Fig.S11. Size distribution by intensity (DLS) of solution of the macrocycle **8**(*1,3-alternate*, 0.19 mg/ml) in the 50 mM phosphate buffer at pH 7.4.

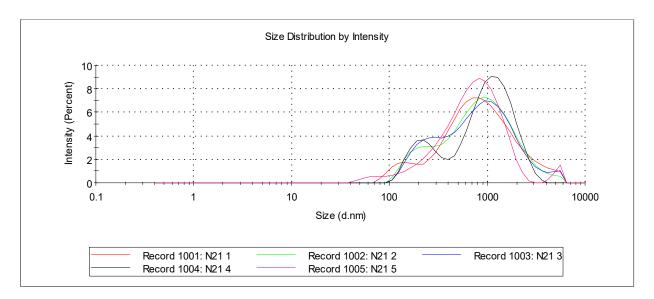


Fig.S12. Size distribution by intensity (DLS) of solution of the OLA-**5** (0.19 mg/ml) in the 50 mM phosphate buffer at pH 7.4.

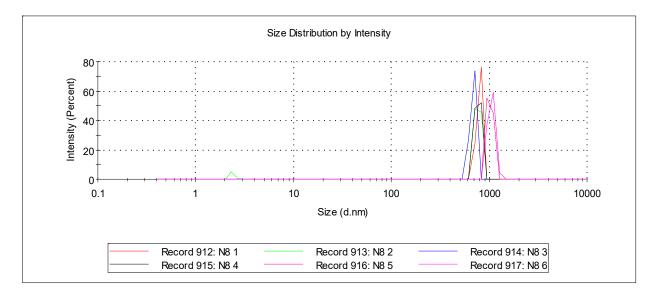


Fig.S14. Size distribution by intensity (DLS) of solution of the macrocycle 6(cone, 0.19 mg/ml) and lysozyme (10 μ M)in the 50 mM phosphate buffer at pH 7.4.

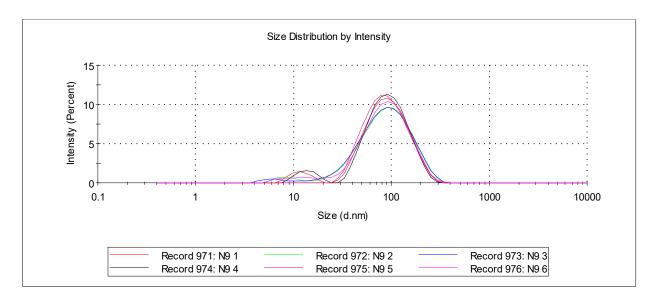


Fig.S15. Size distribution by intensity (DLS) of solution of the macrocycle **6** (*cone*, 0.19 mg/ml) and BSA (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

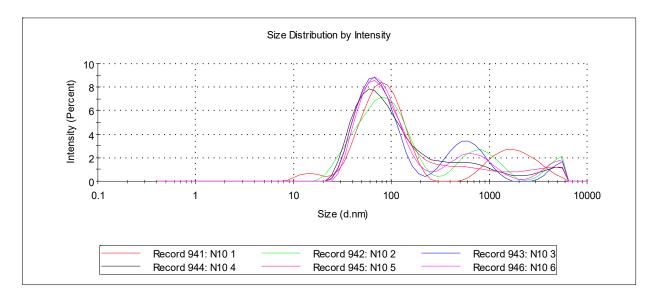


Fig.S16. Size distribution by intensity (DLS) of solution of the macrocycle **6** (*cone*, 0.19 mg/ml) and hemoglobin (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

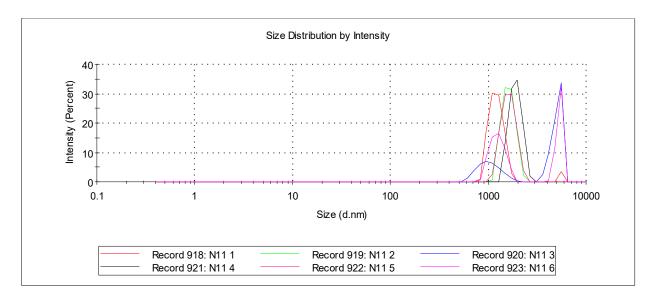


Fig.S17. Size distribution by intensity (DLS) of solution of the macrocycle 7 (*partial cone*, 0.19 mg/ml) and lysozyme (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

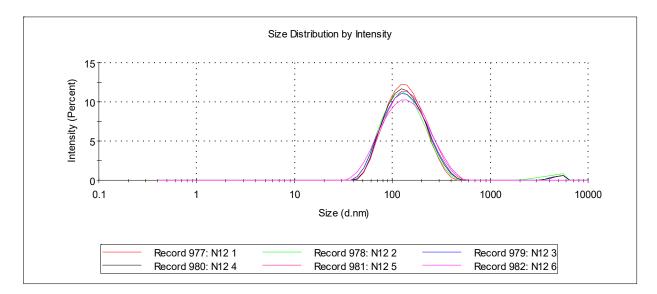


Fig.S18. Size distribution by intensity (DLS) of solution of the macrocycle 7 (*partial cone*, 0.19 mg/ml) and BSA (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

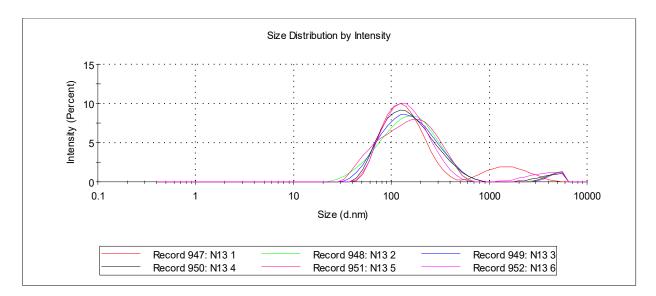


Fig.S19. Size distribution by intensity (DLS) of solution of the macrocycle 7 (*partial cone*, 0.19 mg/ml) and hemoglobin (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

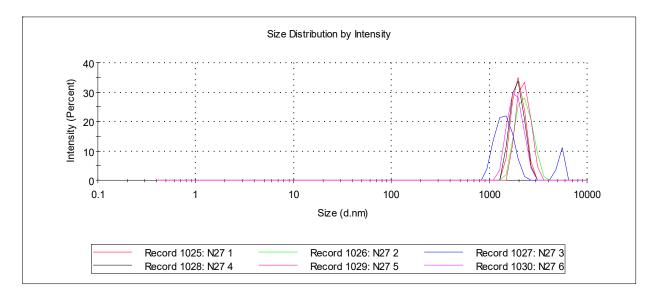


Fig.S20. Size distribution by intensity (DLS) of solution of the macrocycle **8** (*1,3-alternate*, 0.19 mg/ml) and lysozyme (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

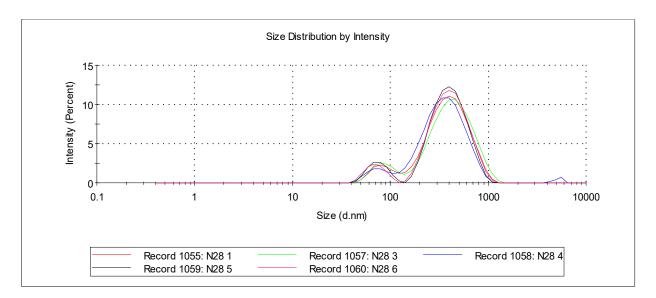


Fig.S21. Size distribution by intensity (DLS) of solution of the macrocycle **8** (*1,3-alternate*, 0.19 mg/ml) and BSA (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

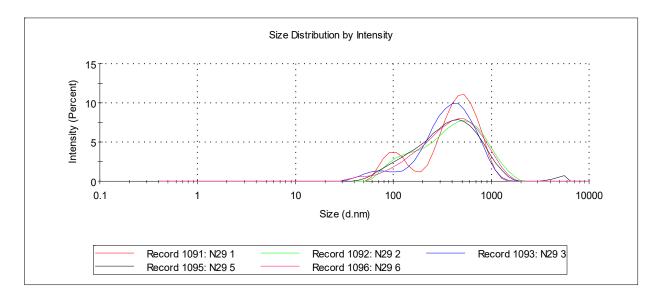


Fig.S22. Size distribution by intensity (DLS) of solution of the macrocycle **8** (*1*,*3-alternate*, 0.19 mg/ml) and hemoglobin (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

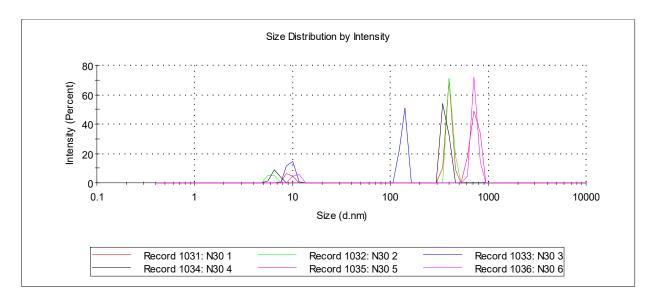


Fig.S23. Size distribution by intensity (DLS) of solution of the OLA-5 (0.19 mg/ml) and lysozyme (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

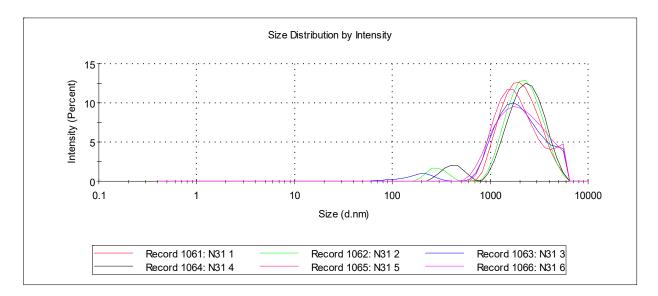


Fig.S24. Size distribution by intensity (DLS) of solution of the OLA-5 (0.19 mg/ml) and BSA (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

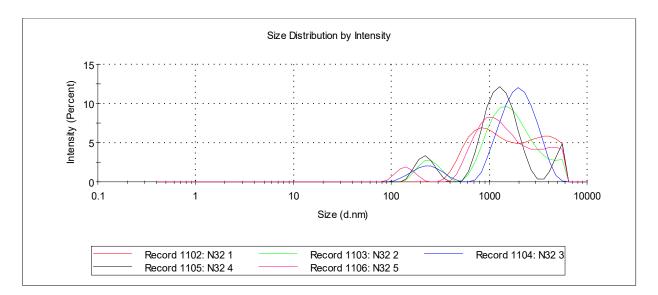


Fig.S25. Size distribution by intensity (DLS) of solution of the OLA-5 (0.19 mg/ml) and hemoglobin (10 μ M) in the 50 mM phosphate buffer at pH 7.4.

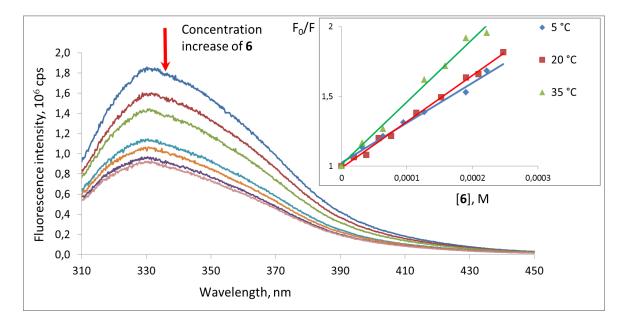


Fig.S26. Fluorescence spectra of BSA (5 μ M) at different concentrations of the compound **6** (0-0.5 mg/ml) in phosphate buffer at pH 7.4. The inset shows the plots of F₀/F vs. concentration of the compound **6** at different temperatures (5, 20 and 35 °C). λ_{ex} =285 nm; λ_{em} =330 nm.

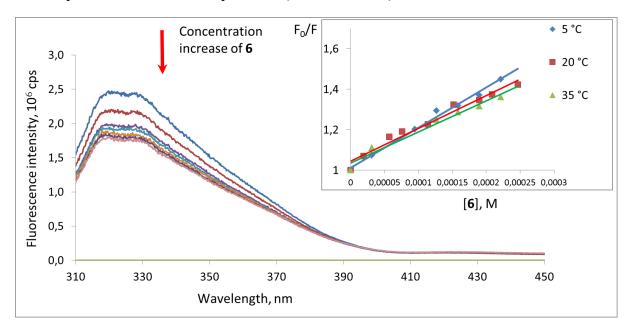


Fig.S27. Fluorescence spectra of Hb (5 μ M) at different concentrations of the compound **6** (0-0.5 mg/ml) in phosphate buffer at pH 7.4. The inset shows the plots of F₀/F vs. concentration of the compound **6** at different temperatures (5, 20 and 35 °C). λ_{ex} =285 nm; λ_{em} =330 nm.

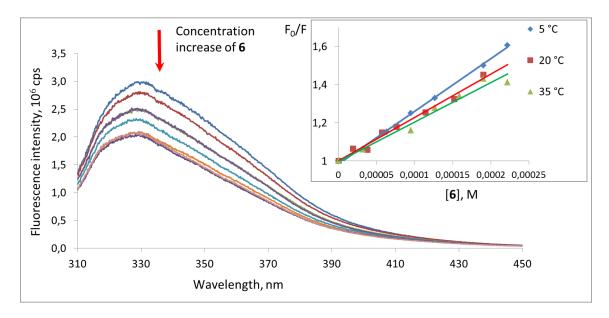


Fig.S28. Fluorescence spectra of Lys (5 μ M) at different concentrations of the compound **6** (0-0.5 mg/ml) in phosphate buffer at pH 7.4. The inset shows the plots of F₀/F vs. concentration of the compound **6** at different temperatures (5, 20 and 35 °C). λ_{ex} =285 nm; λ_{em} =330 nm.

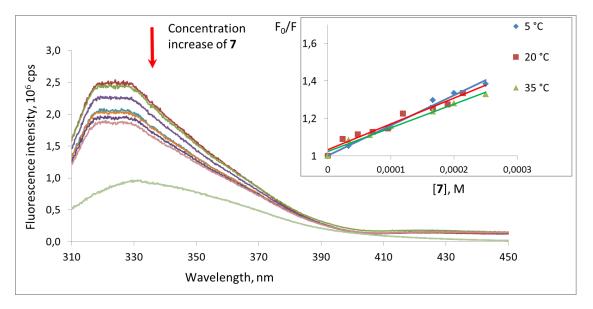


Fig.S.29. Fluorescence spectra of Hb (5 μ M) at different concentrations of the compound 7 (0-0.5 mg/ml) in phosphate buffer at pH 7.4. The inset shows the plots of F₀/F vs. concentration of the compound 7 at different temperatures (5, 20 and 35 °C). λ_{ex} =285 nm; λ_{em} =330 nm.

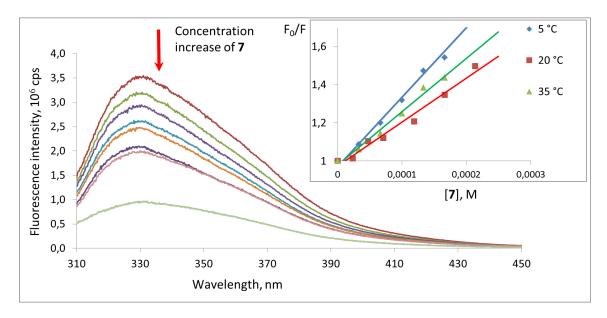


Fig.S30. Fluorescence spectra of Lys (5 μ M) at different concentrations of the compound 7 (0-0.5 mg/ml) in phosphate buffer at pH 7.4. The inset shows the plots of F₀/F vs. concentration of the compound 7 at different temperatures (5, 20 and 35 °C). λ_{ex} =285 nm; λ_{em} =330 nm.

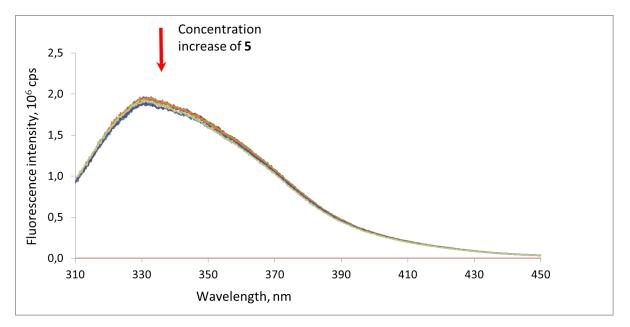


Fig.S31. Fluorescence spectra of BSA (5 μ M) at different concentrations of the compound **5** (0-0.5 mg/ml) in phosphate buffer at pH 7.4.

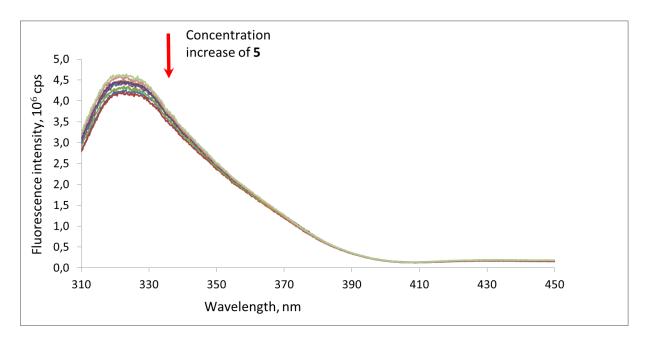


Fig.S32. Fluorescence spectra of Hb (5 μ M) at different concentrations of the compound **5** (0-0.5 mg/ml) in phosphate buffer at pH 7.4.

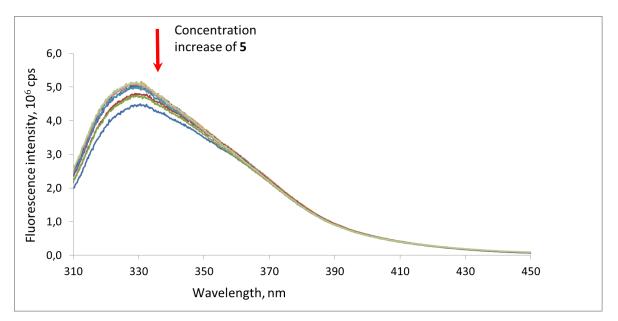


Fig.S33. Fluorescence spectra of Lys (5 μ M) at different concentrations of the compound **5** (0-0.5 mg/ml) in phosphate buffer at pH 7.4.

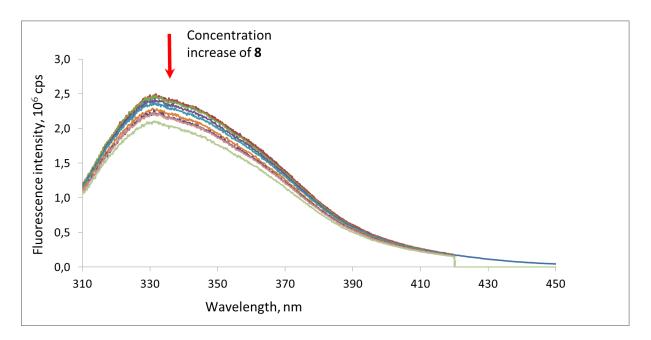


Fig.S34. Fluorescence spectra of BSA (5 μ M) at different concentrations of the compound **8** (0-0.5 mg/ml) in phosphate buffer at pH 7.4.

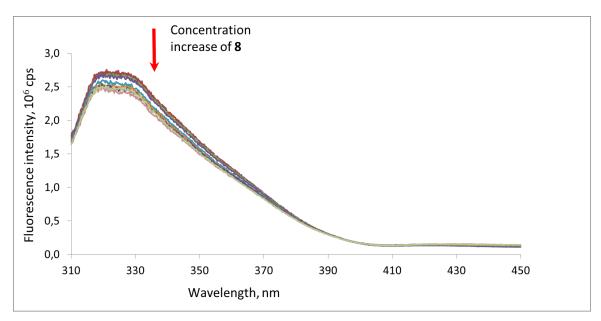


Fig.S35. Fluorescence spectra of Hb (5 μ M) at different concentrations of the compound **8** (0-0.5 mg/ml) in phosphate buffer at pH 7.4.

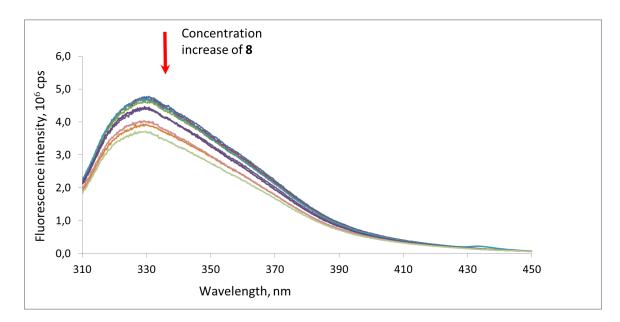


Fig.S36. Fluorescence spectra of Lys (5 μ M) at different concentrations of the compound 8 (0-0.5 mg/ml) in phosphate buffer at pH 7.4.

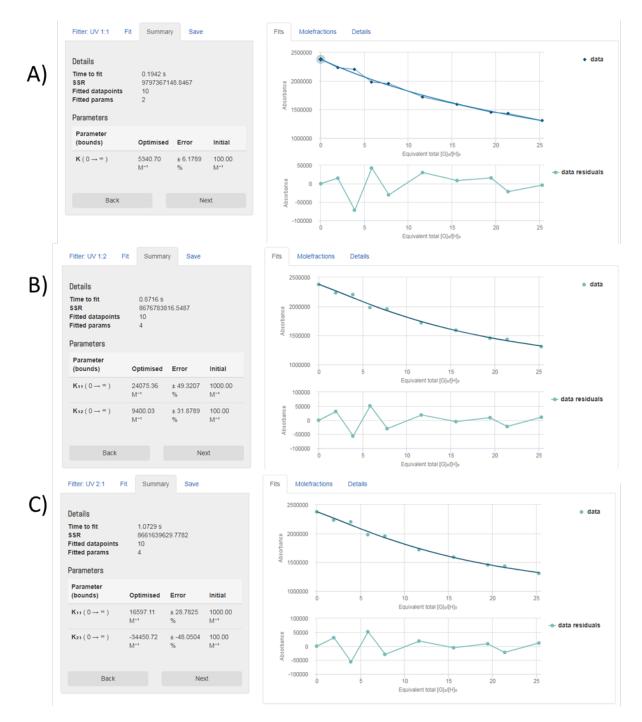


Fig.S37. Screenshots taken from the summary window of the website supramolecular.org. These screenshots show the raw data for titration of **6** with BSA, the data fitted to 1:1 (A), 1:2 (B) and 2:1 (C) binding models.

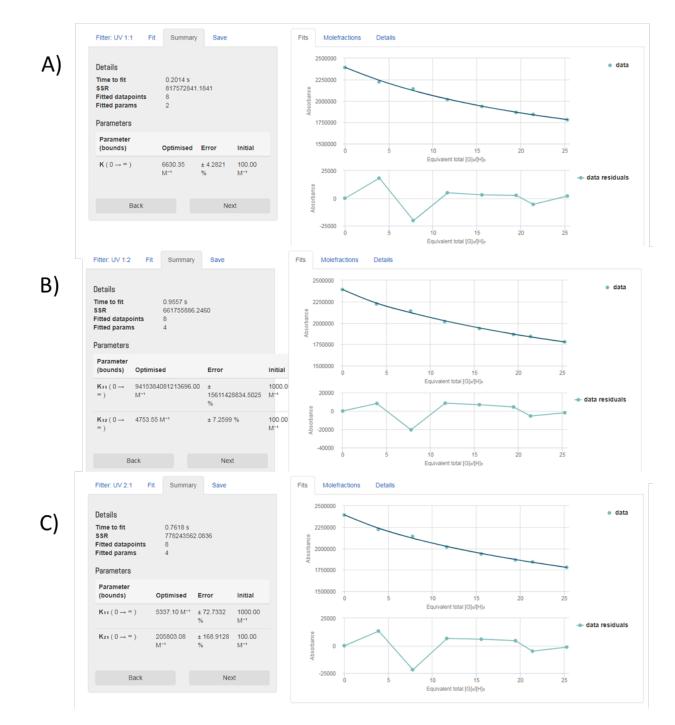


Fig.S38. Screenshots taken from the summary window of the website supramolecular.org. These screenshots show the raw data for titration of **6** with Hb, the data fitted to 1:1 (A), 1:2 (B) and 2:1 (C) binding models..

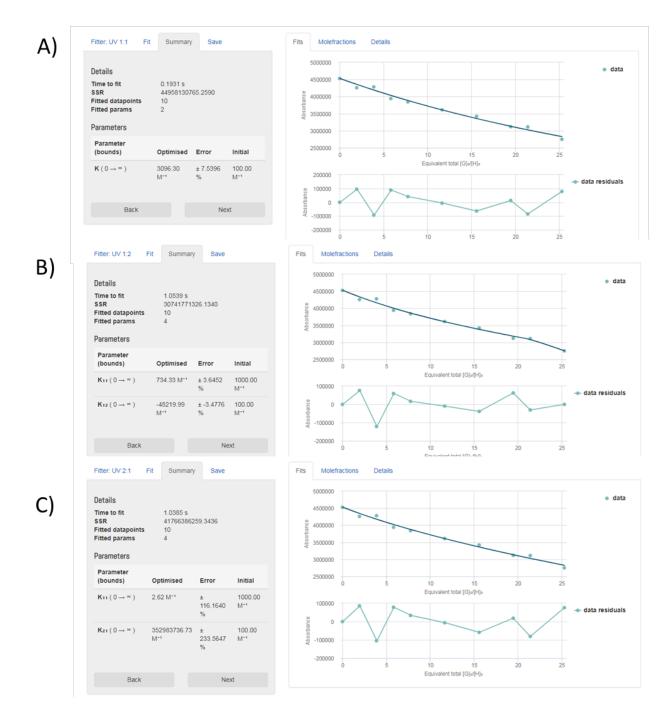


Fig.S39. Screenshots taken from the summary window of the website supramolecular.org. These screenshots show the raw data for titration of **6** with Lys, the data fitted to 1:1 (A), 1:2 (B) and 2:1 (C) binding models.

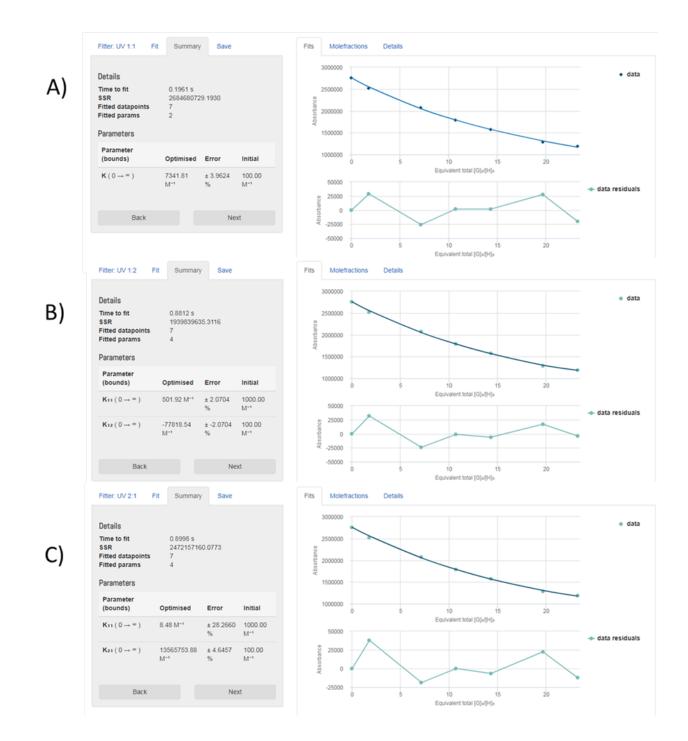


Fig.S40. Screenshots taken from the summary window of the website supramolecular.org. These screenshots show the raw data for titration of 7 with BSA, the data fitted to 1:1 (A), 1:2 (B) and 2:1 (C) binding models.

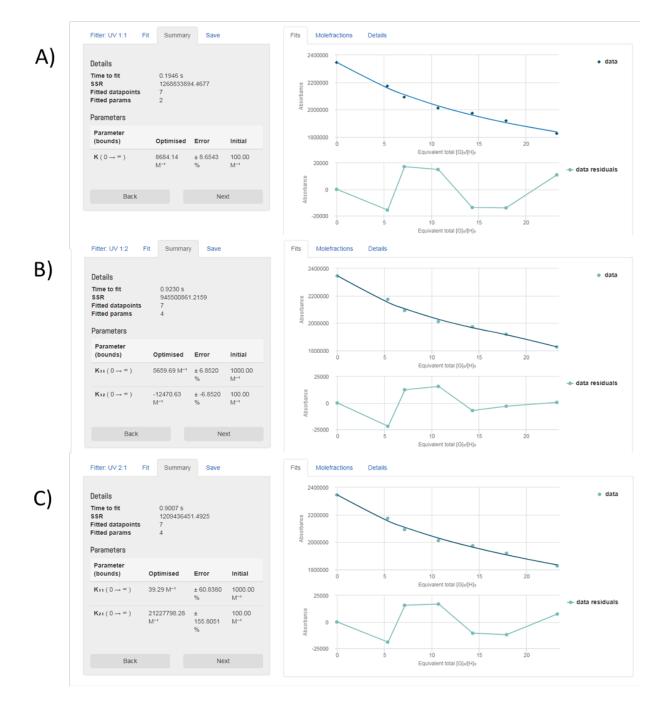


Fig.S41. Screenshots taken from the summary window of the website supramolecular.org. These screenshots show the raw data for titration of **7** with Hb, the data fitted to 1:1 (A), 1:2 (B) and 2:1 (C) binding models.

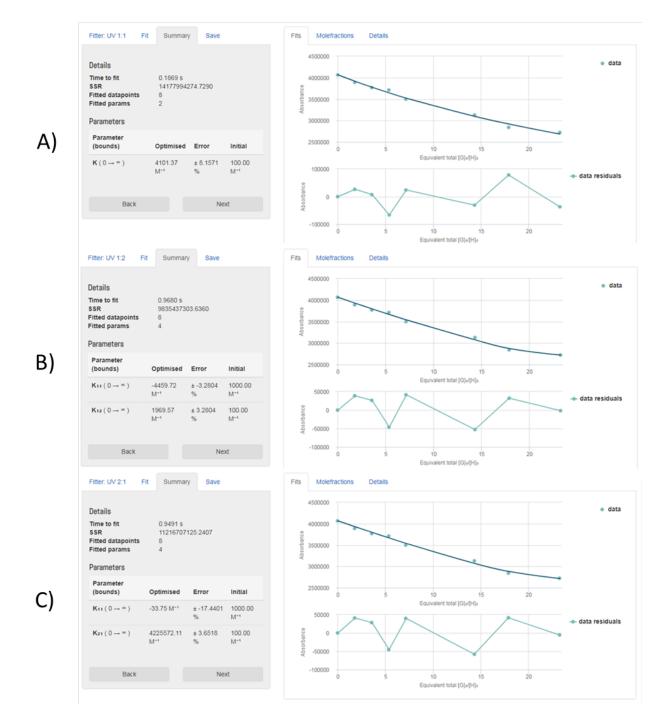


Fig.S42. Screenshots taken from the summary window of the website supramolecular.org. These screenshots show the raw data for titration of **7** with Lys, the data fitted to 1:1 (A), 1:2 (B) and 2:1 (C) binding models.