

SYNTHESIS, STRUCTURE, AND ANTIMICROBAL ACTIVITY OF A SERIES OF NOVEL
HETEROCYCLIC AZO-COLORANTS DERIVED FROM α -DICARBONYL COMPOUNDS

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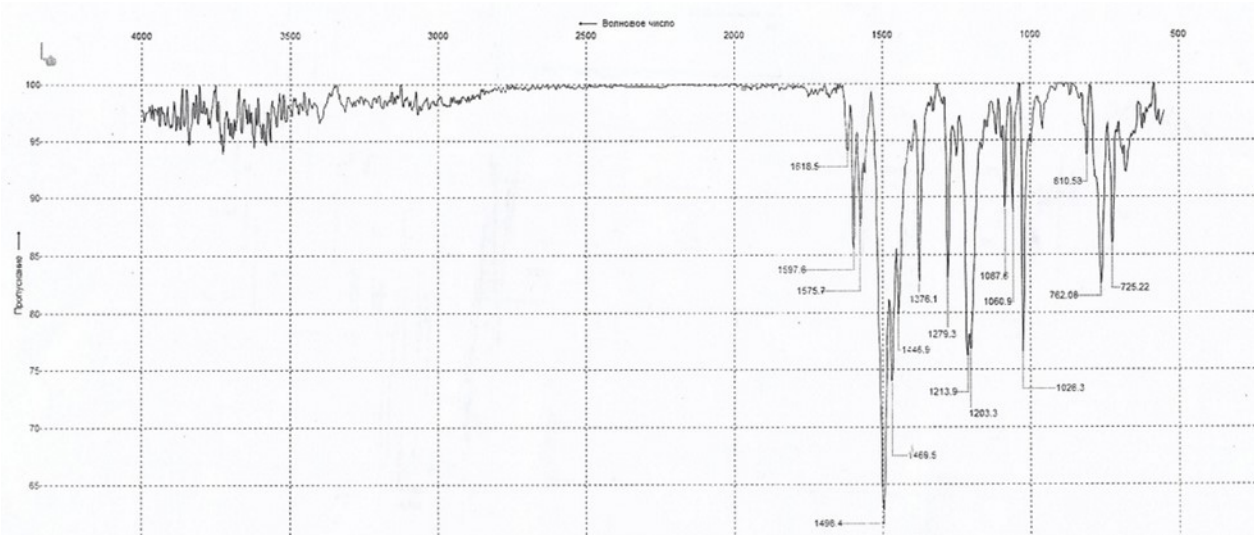


Fig 1S. IR Spectrum of L¹

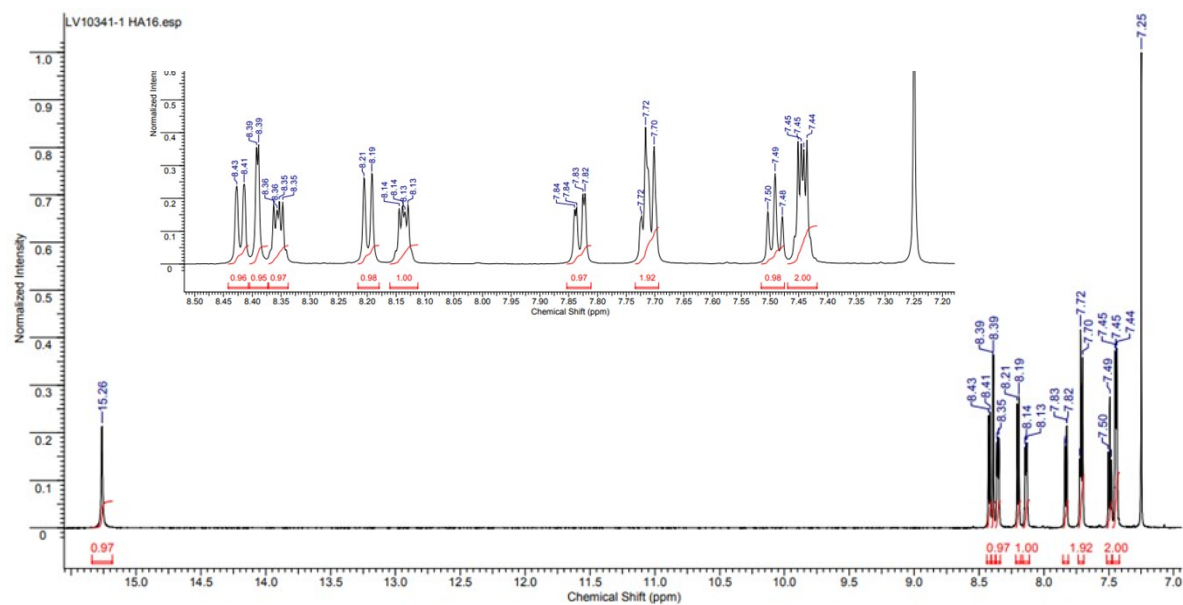


Fig 2S. ¹H NMR spectrum of L¹

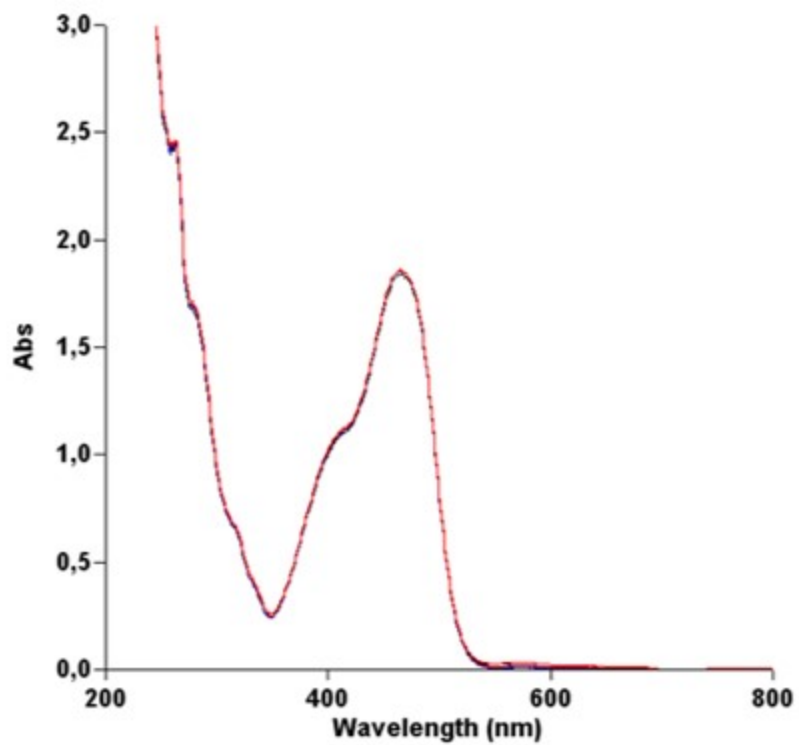


Fig 3S. UV-Vis spectrum of L¹ in ethanol

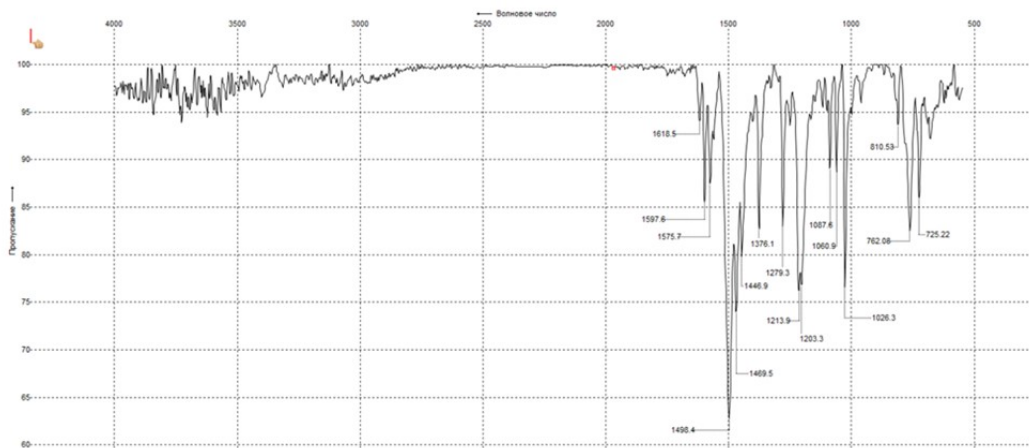


Fig 4. IR spectrum of L³

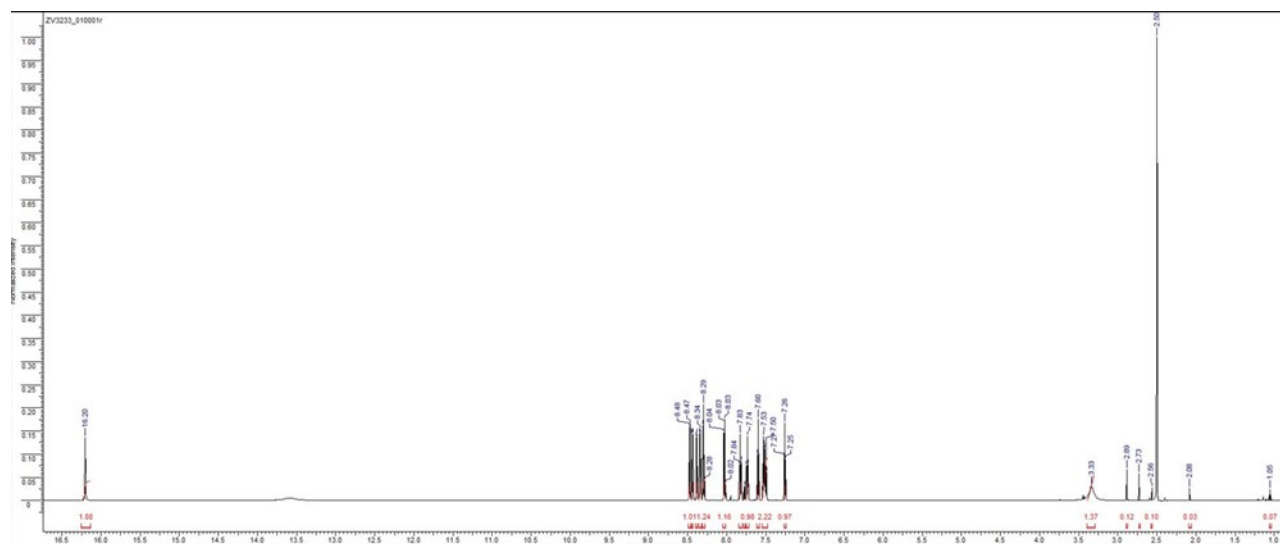


Fig 5. ^1H NMR spectrum of L^4

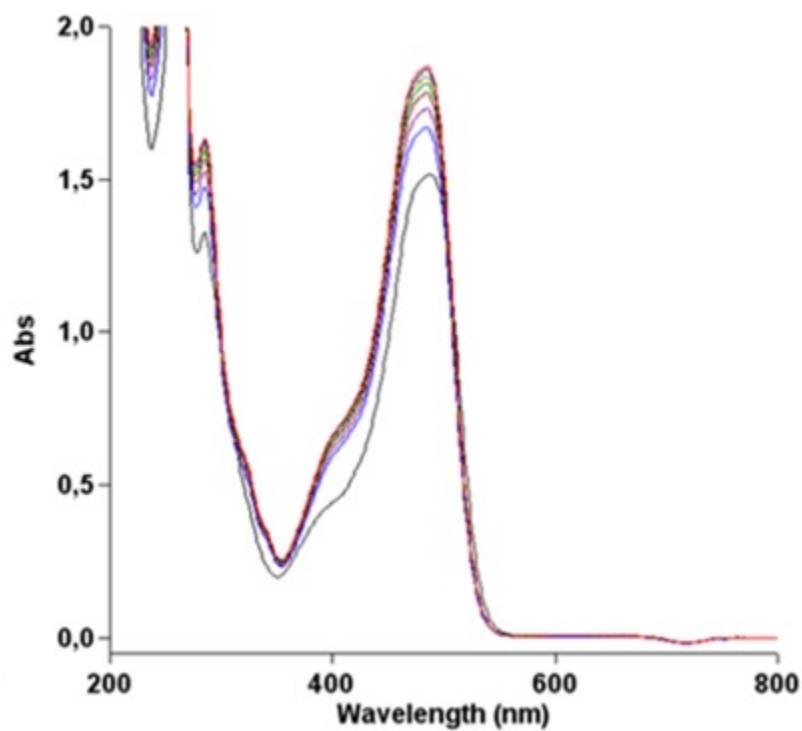


Fig 6: UV-Vis spectrum of L^3

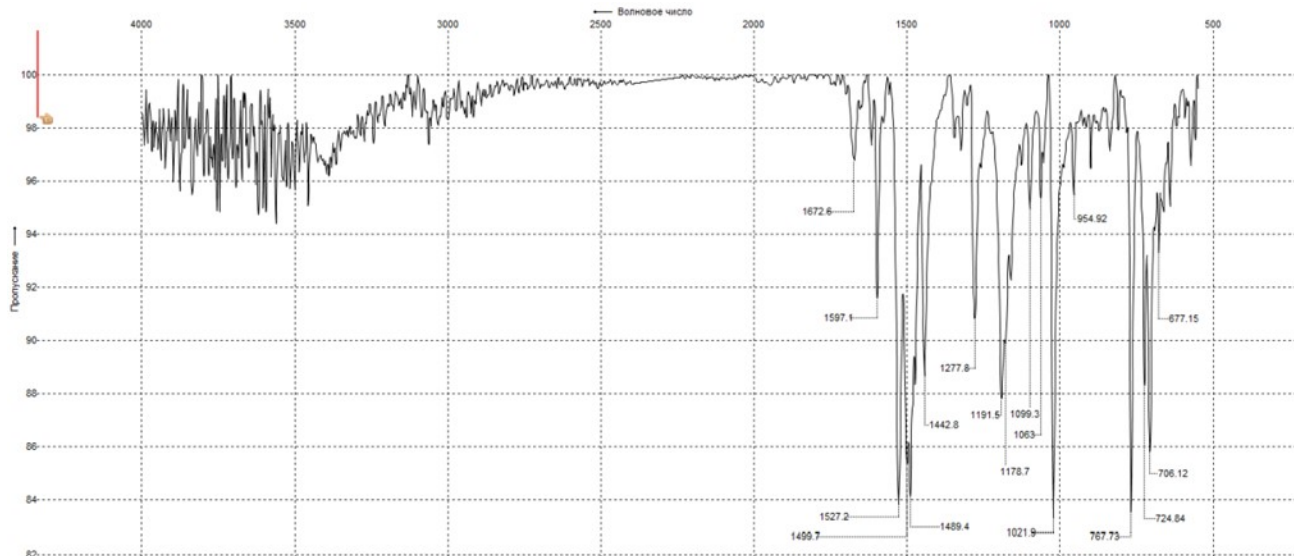


Fig 7S. IR spectrum of L⁴

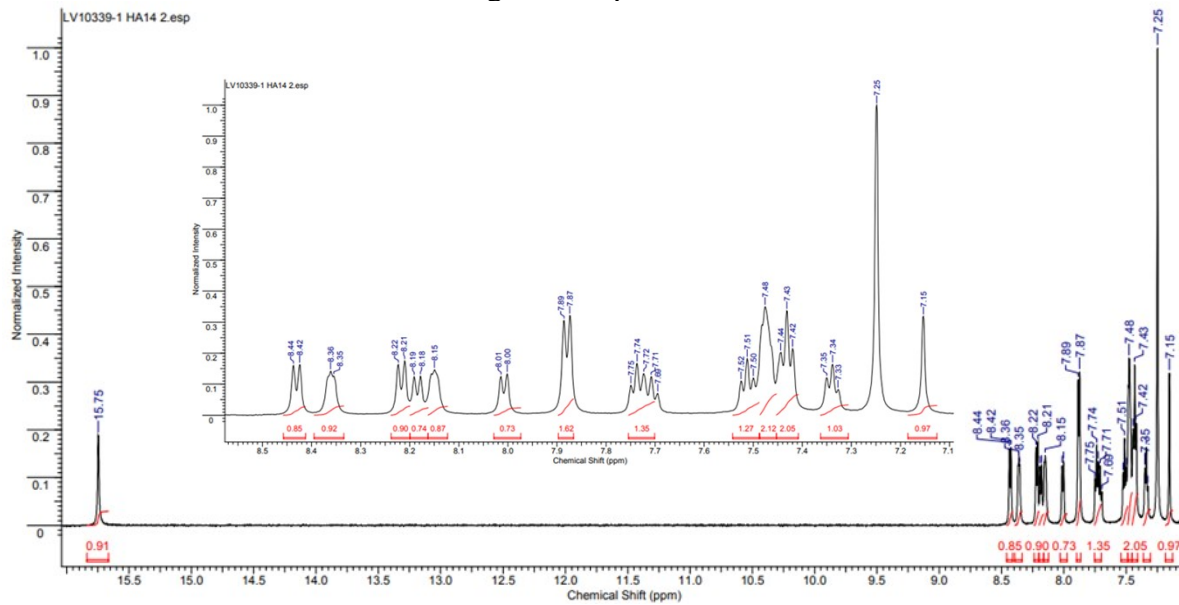


Fig 8S. ¹H NMR spectrum of L⁴

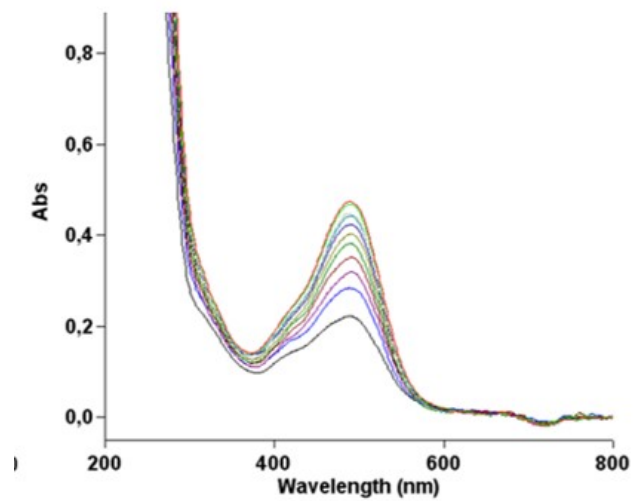


Fig 9S. UV-Vis spectrum of L⁴

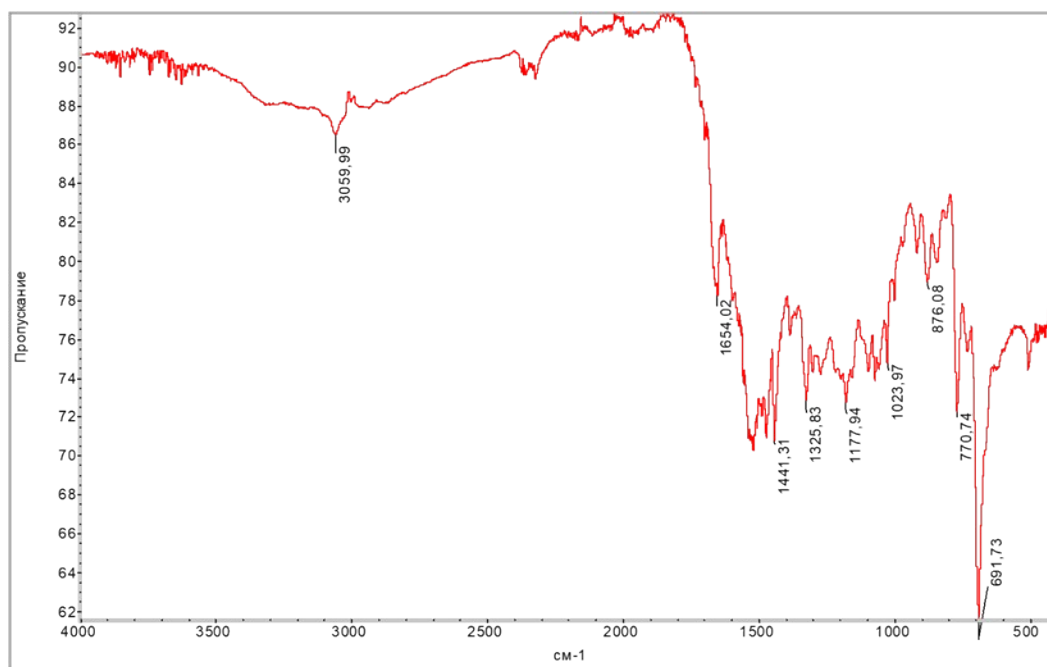


Fig 10S. IR spectrum of L⁵

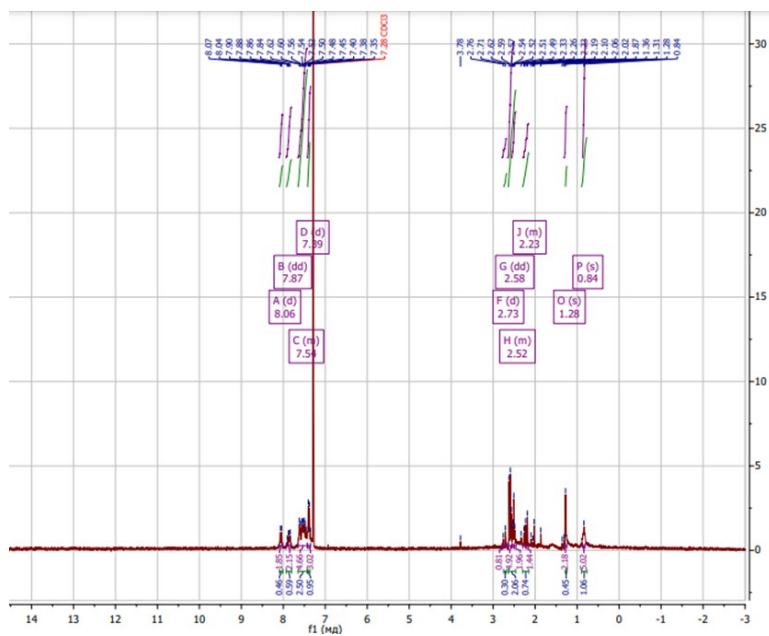


Fig 11S. ¹H NMR spectrum of L⁵

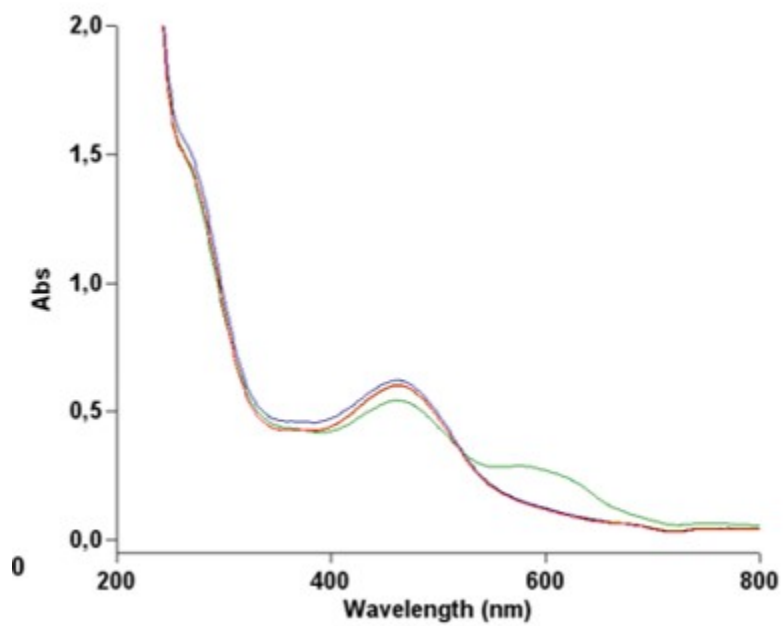


Fig 12S. UV-Vis spectrum of L⁵

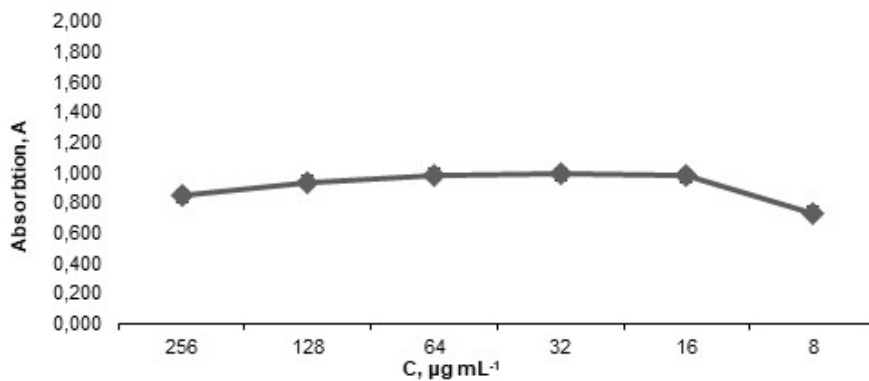


Figure 13S. Change in absorption when L¹ is exposed to the *Escherichia coli* C1 strain at concentrations of 8-256 µg mL⁻¹ (example of an inactive compound).

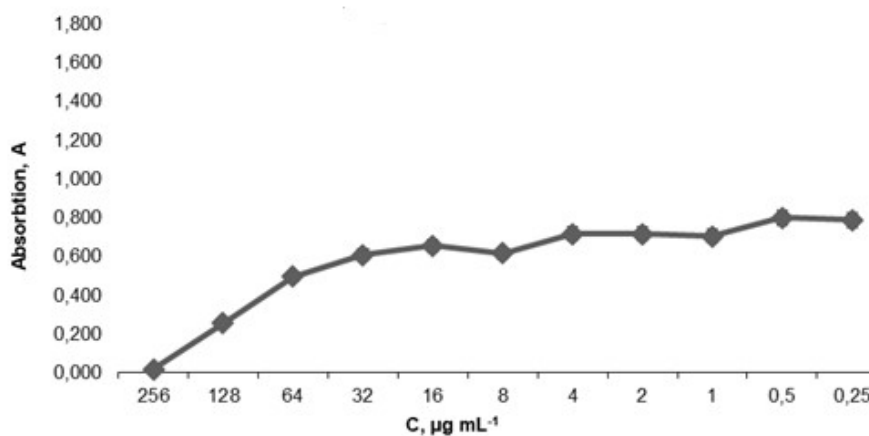


Figure 14S. Change in absorption when L³ is exposed to the *Escherichia coli* C1 strain at concentrations of 0,25-256 µg mL⁻¹ (example of an active compound).

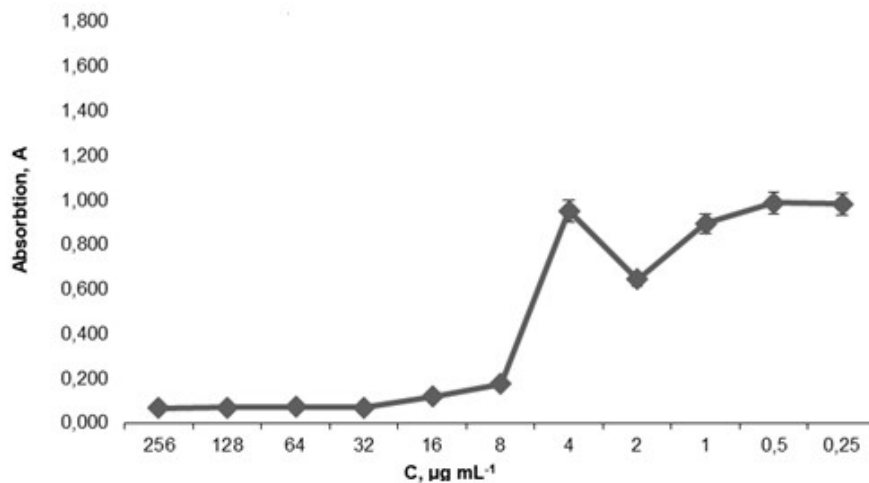


Figure 15S. Change in absorption when L⁴ is exposed to the *Staphylococcus aureus* ATCC-25923 strain at concentrations of 0,25-256 µg mL⁻¹ (example of an active compound).

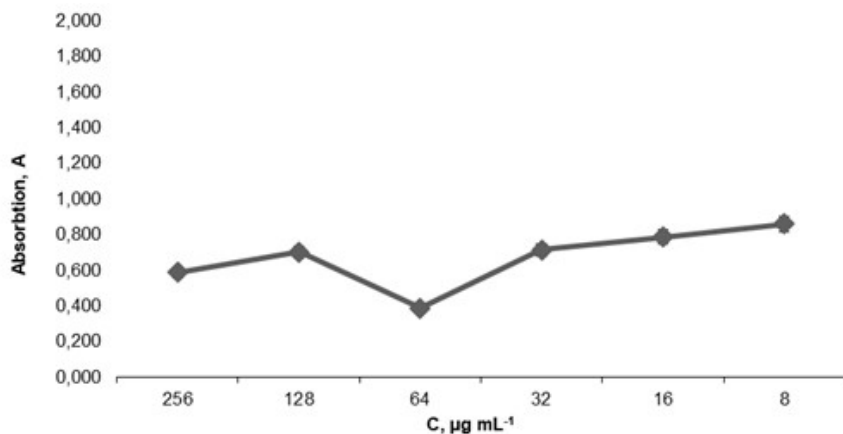
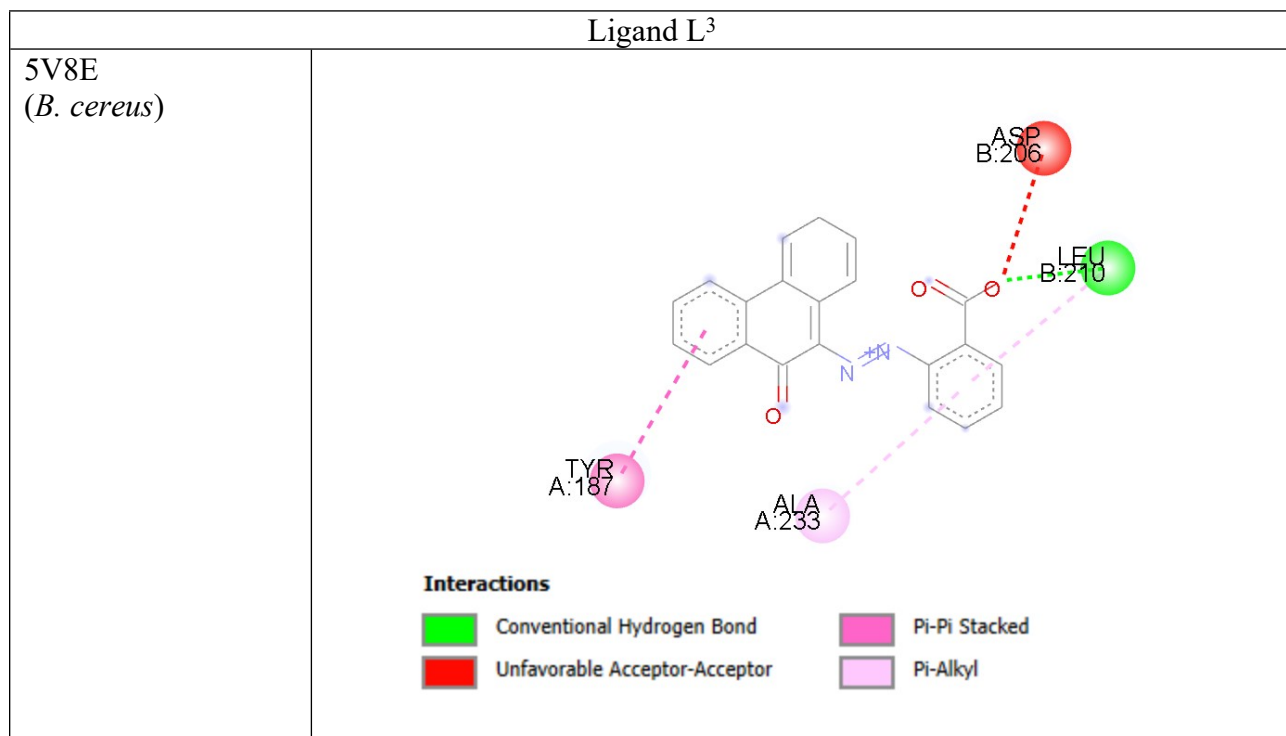
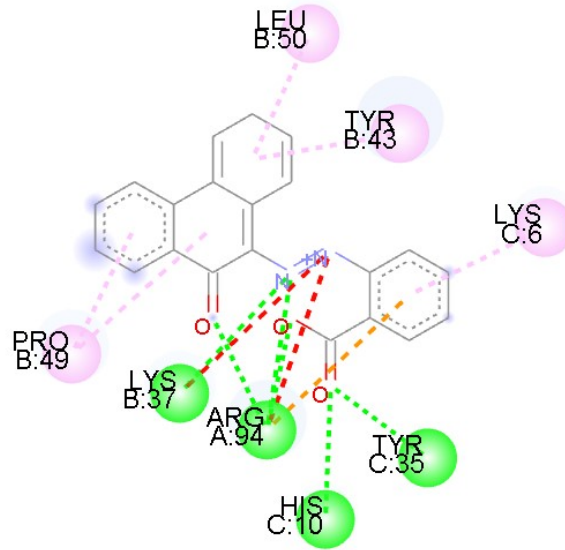


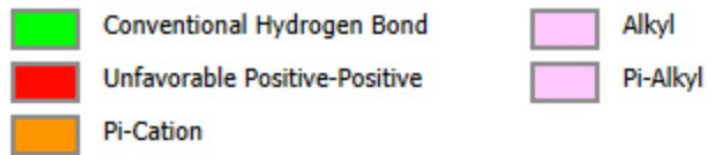
Figure 16S. Change in absorption when L¹ is exposed to the *Staphylococcus aureus* ATCC-25923 strain at concentrations of 8-256 µg mL⁻¹ (example of an inactive compound).



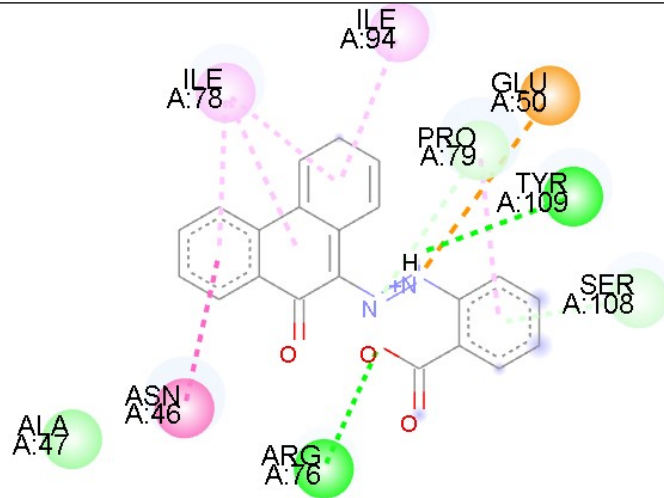
2BL8
(*E. faecium*)



Interactions



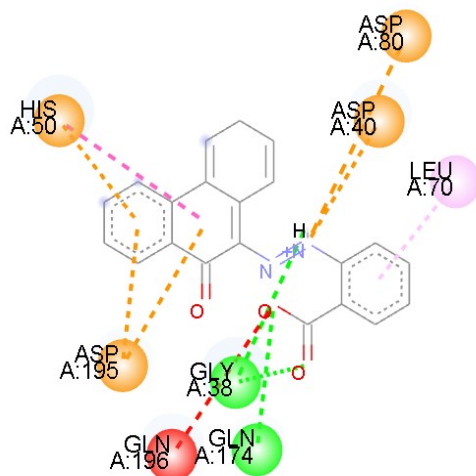
4PRV
(*E. coli*)










Interactions



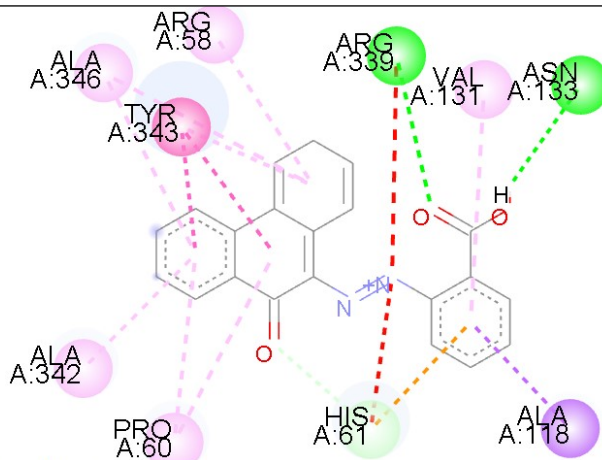
1JIL
(*S. aureus*)











Interactions

- | | | | |
|---|-------------------------------|--|---------------|
|  | Attractive Charge |  | Pi-Anion |
|  | Conventional Hydrogen Bond |  | Pi-Pi Stacked |
|  | Unfavorable Acceptor-Acceptor |  | Pi-Alkyl |
|  | Pi-Cation | | |

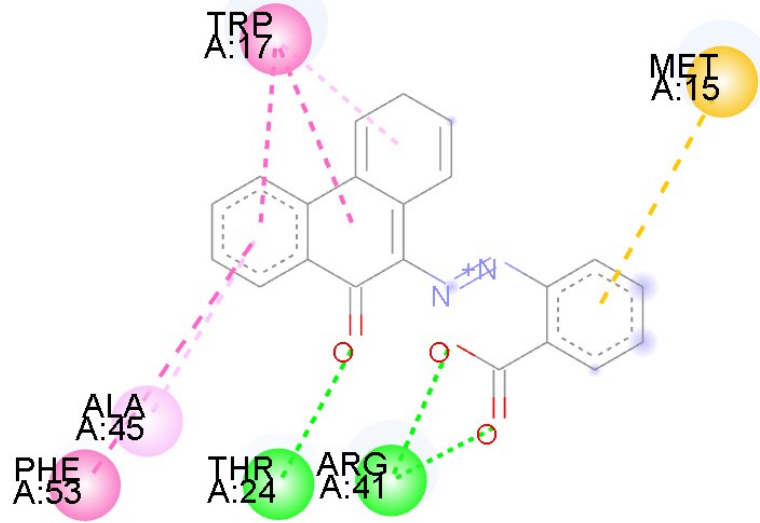
1GWE
(*M. luteus*)



Interactions

- | | | | |
|---|-------------------------------|--|---------------|
|  | Conventional Hydrogen Bond |  | Pi-Sigma |
|  | Carbon Hydrogen Bond |  | Pi-Pi Stacked |
|  | Unfavorable Positive-Positive |  | Alkyl |
|  | Pi-Cation |  | Pi-Alkyl |

5V00
(*P. fluorescens*)

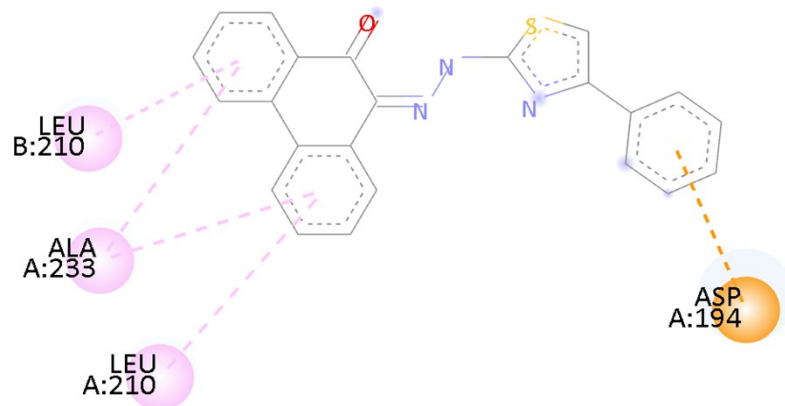


Interactions

- Conventional Hydrogen Bond
- Pi-Sulfur
- Pi-Pi Stacked
- Pi-Pi T-shaped
- Pi-Alkyl

L⁴

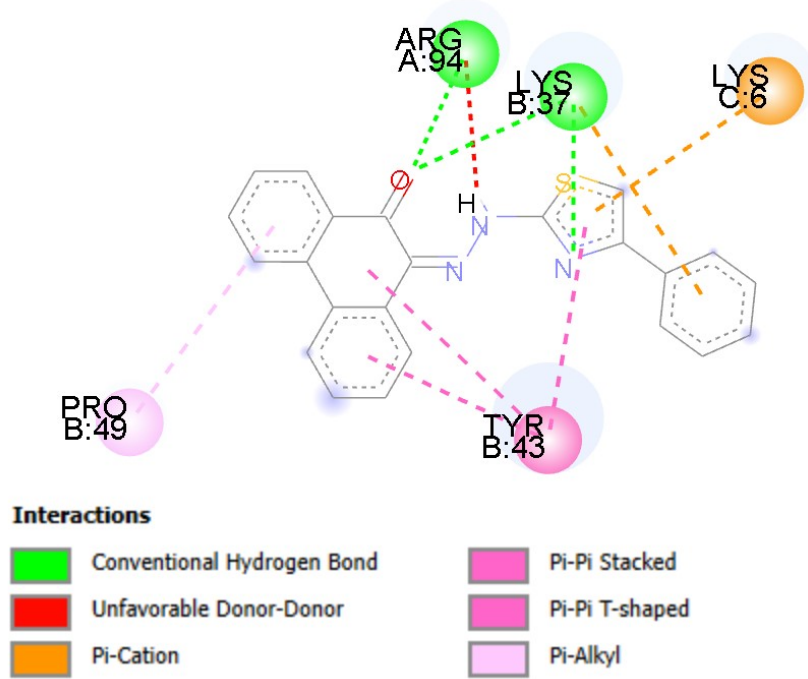
5V8E
(*B. cereus*)



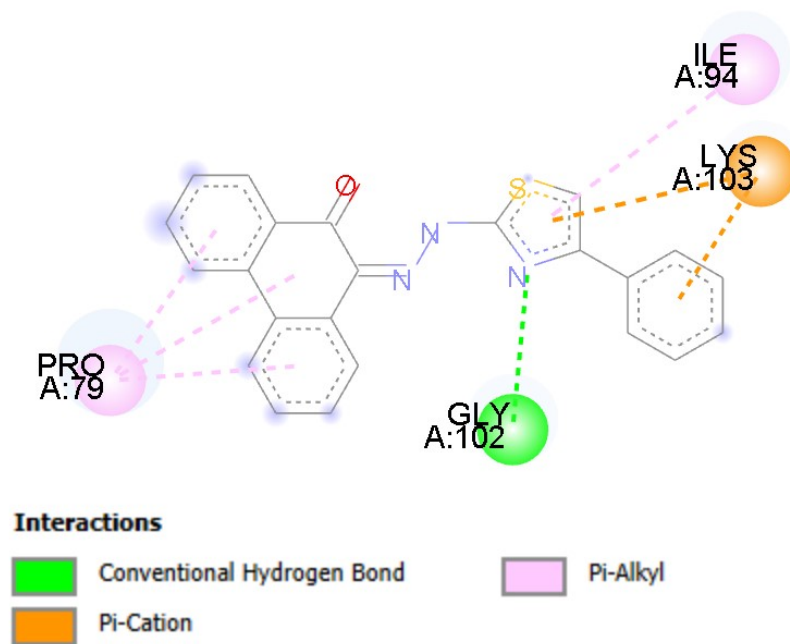
Interactions

- Pi-Anion
- Pi-Alkyl

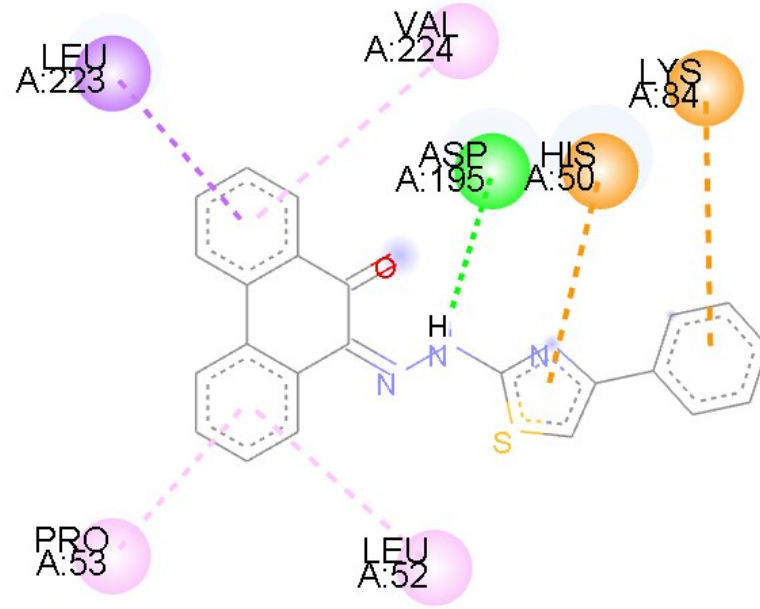
2BL8
(*E. faecium*)








4PRV
(*E. coli*)



1JIL
(*S. aureus*)



Interactions

- | | |
|--|--|
|  Conventional Hydrogen Bond |  Pi-Pi T-shaped |
|  Pi-Cation |  Pi-Alkyl |
|  Pi-Sigma | |

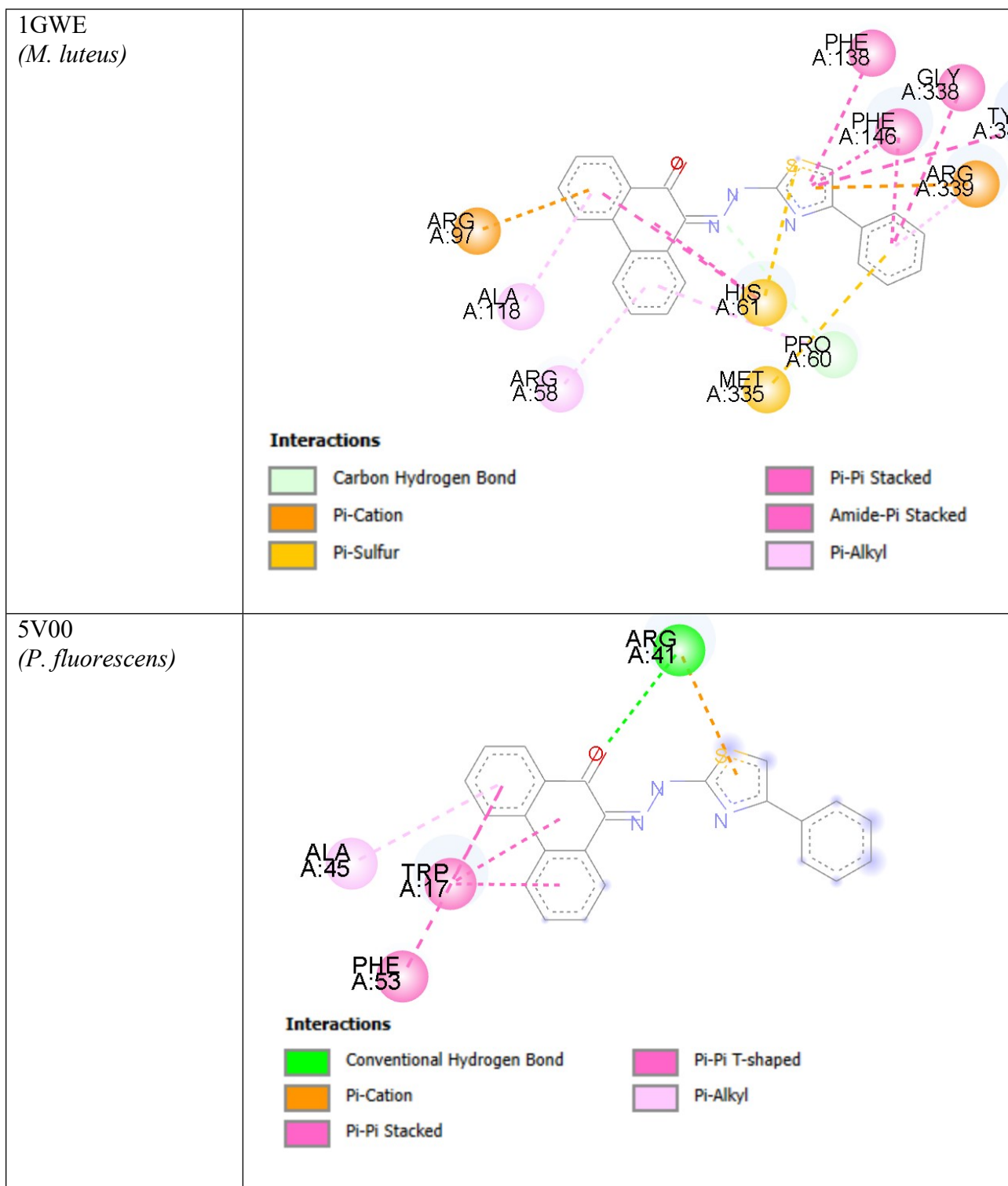


Figure 17S. 2D Chemical Structures representing protein-ligand interactions of L³ and L⁴

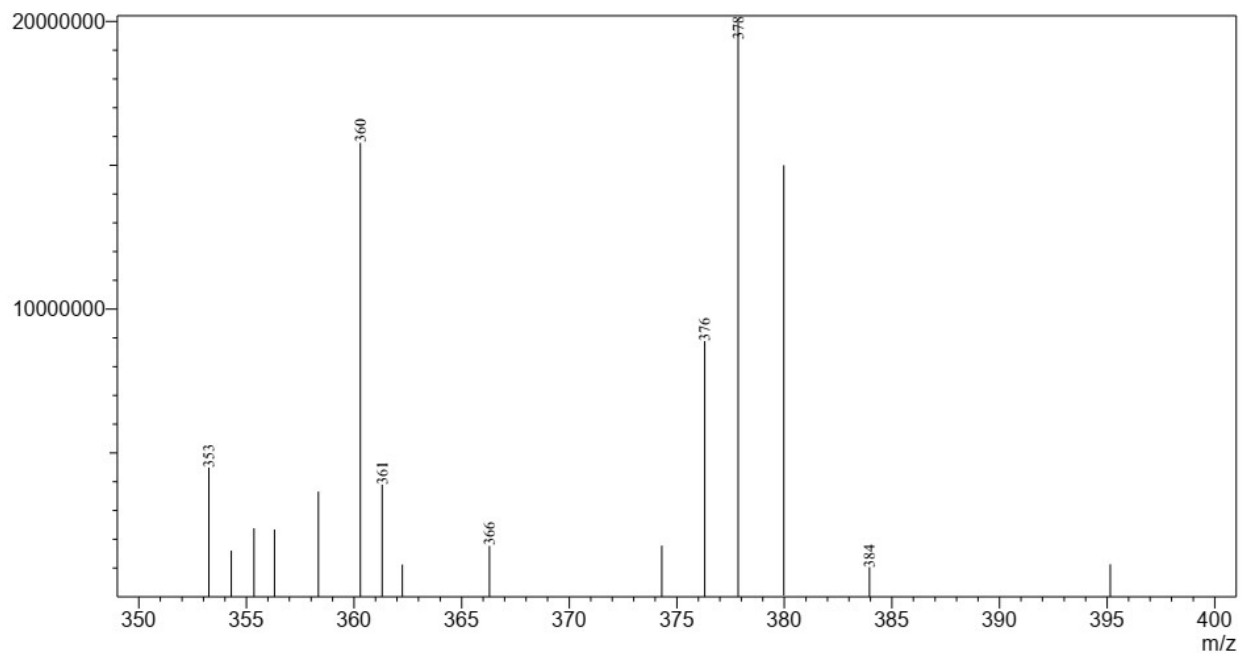


Fig 18S. Mass spectrum of L¹

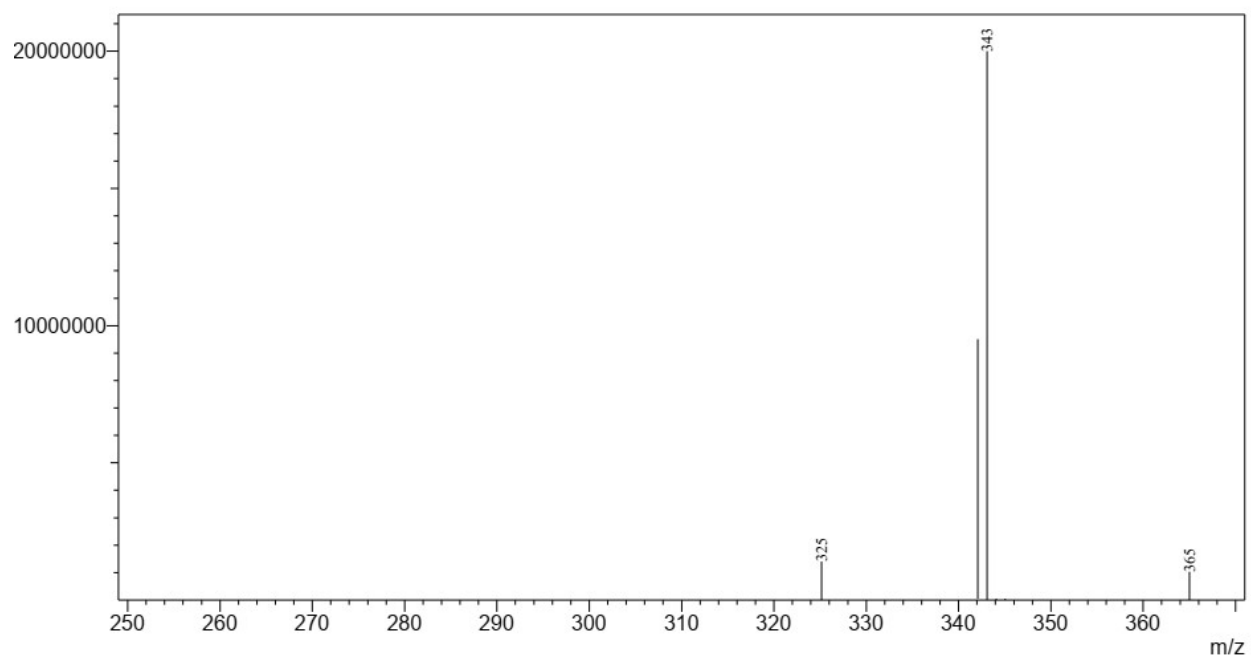


Fig 19S. Mass spectrum of L³

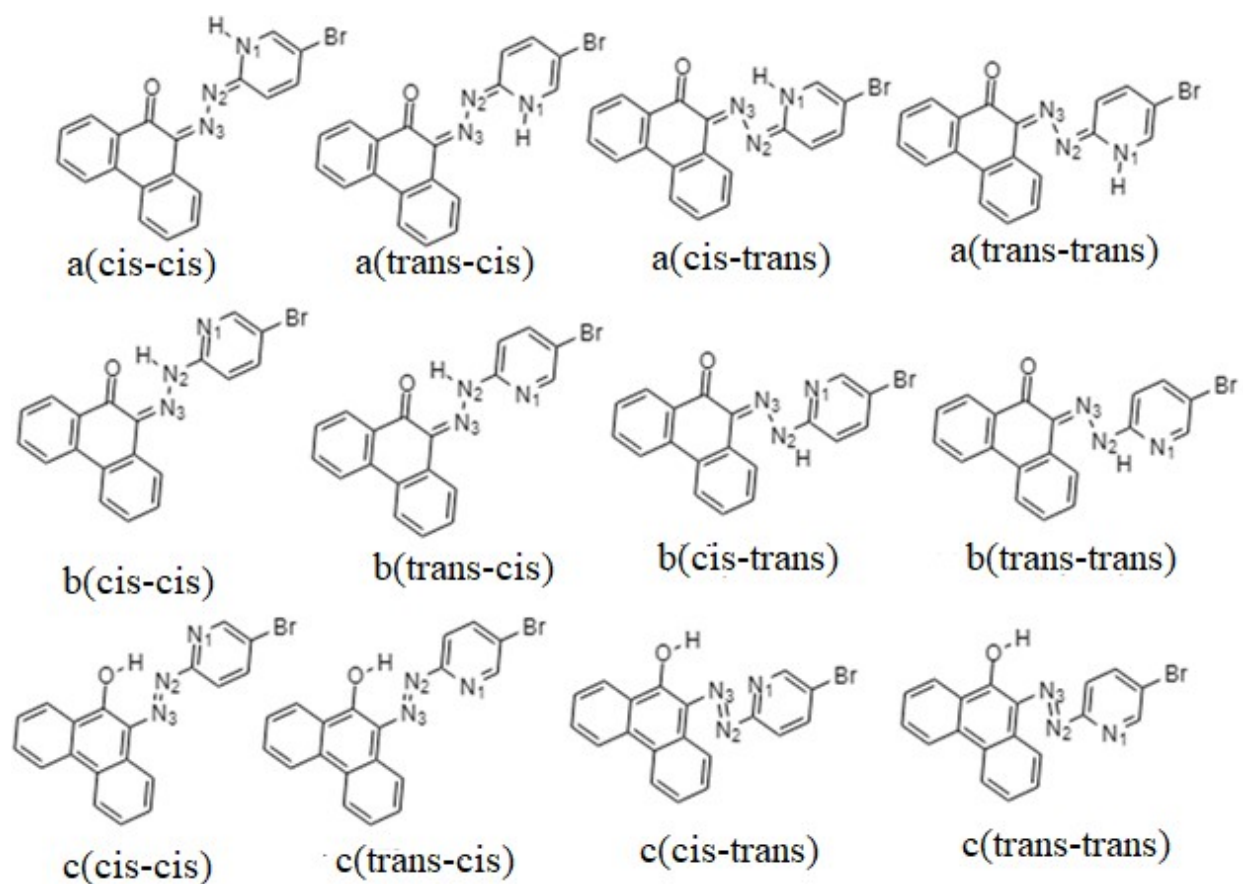


Figure 20S. Possible tautomeric forms of L¹

Table 1S. Relative energies (kJ/mole) of cis-isomers of zwitter-ionic (a), hydrazo (b) and azo (c) forms of L¹-L⁶ according to DFT calculations.

Tautomer	Compound	L ¹	L ²	L ³	L ⁴	L ⁵	L ⁶
a		96.5	86.34	—	70.86	47.47	221.01
b		0	0	0	0	9.04	0
c		33.8	0	31.44	24.14	0	12.51

Table 2S. Hemolytic activity of L³ and L⁴.

Compound	HC ₅₀ , µg mL ⁻¹ RBC
L ³	>32
L ⁴	>32

Table 3S. The crystal data and structure refinement of L² and L⁴.

Compound	L ²	L ⁴
Empirical formula	C ₁₈ H ₁₁ ClN ₄ O	C ₂₃ H ₁₅ N ₃ OS
Formula weight	334.76	381.44
T(K)	100(2)	100(2)
λ (Å)	1.54184	1.54184
Crystal system	orthorhombic	monoclinic
Space group	<i>Pna</i> 2 ₁	<i>P</i> 2 ₁ / <i>c</i>
Unit cell dimensions:		
a, Å	16.4130(11)	20.2762(5)
b, Å	22.3174(14)	5.30357(13)
c, Å	3.9518(2)	16.5308(4)
α, deg.	90	90
β, deg.	90	104.649(2)
γ, deg.	90	90
V, Å ³	1447.53(15)	1719.87(7)
Z	4	4
D _{calc} (g cm ⁻³)	1.536	1.473
Absorption coefficient, µ, mm ⁻¹	2.449	1.830
F(000)	688	792
Crystal shape	needle	plate
Crystal color	light-yellow	dark-red
Crystal size, mm	0.53×0.04×0.02	0.12×0.12×0.02
Theta range for data collection	3.343–79.337	2.252–80.128
Index ranges	-20 ≤ h ≤ 19;	-25 ≤ h ≤ 24;

	-28 ≤ k ≤ 24;	-6 ≤ k ≤ 6;
	-4 ≤ l ≤ 5	-21 ≤ l ≤ 20
Reflections collected	10280	41624
Independent reflections	2709	3593
Reflections collected with I > 2σ(I)	2249	3412
Absorption correction	Gaussian	Multi-scan
T_{\min} / T_{\max}	0.392 / 1.000	0.575 / 1.000
Data / restraints / parameters	2709 / 1 / 220	3593 / 0 / 258
Goodness-of-fit (GOF) on F^2	1.033	1.020
R ₁ [I > 2σ(I)]	0.0836	0.1020
wR ₂ [all data]	0.2133	0.2271
Largest dif. peak and hole	1.195 / -0.707	0.914 / -0.693
CCDC No	2482127	2482128
