

## Electronic Supplementary Information

### Dual Responsive Salen-type Schiff Bases for the Effective Detection of L-Arginine *via* Static Quenching Mechanism.

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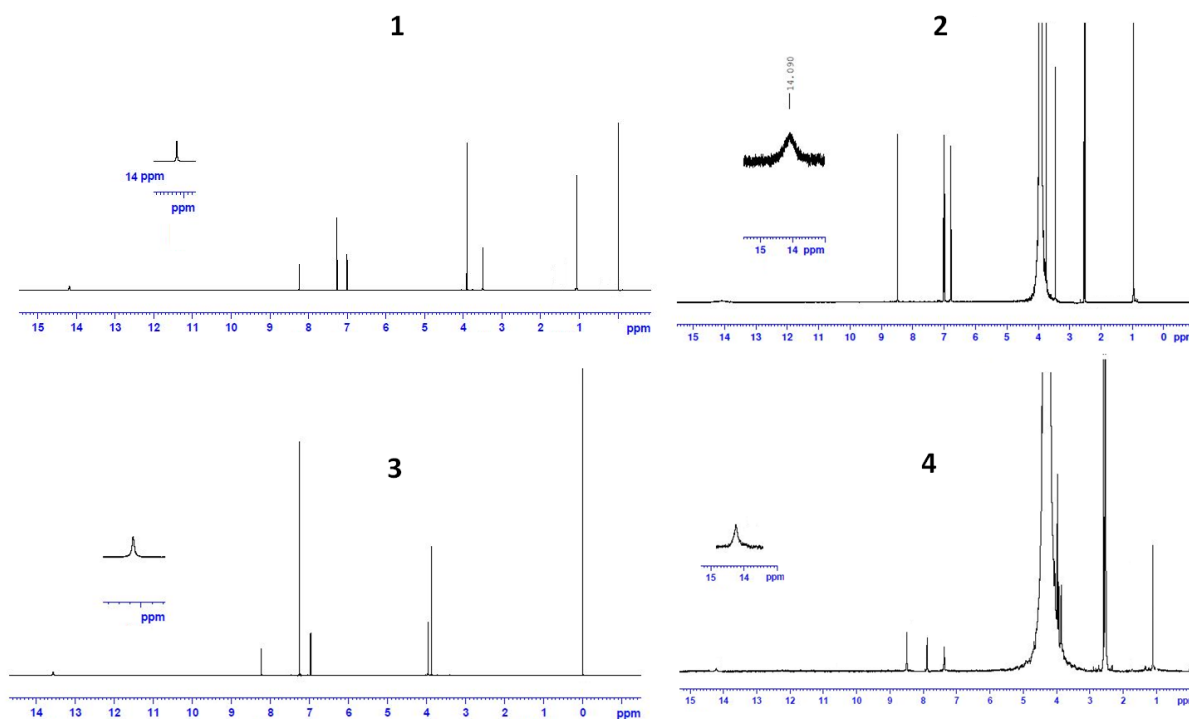
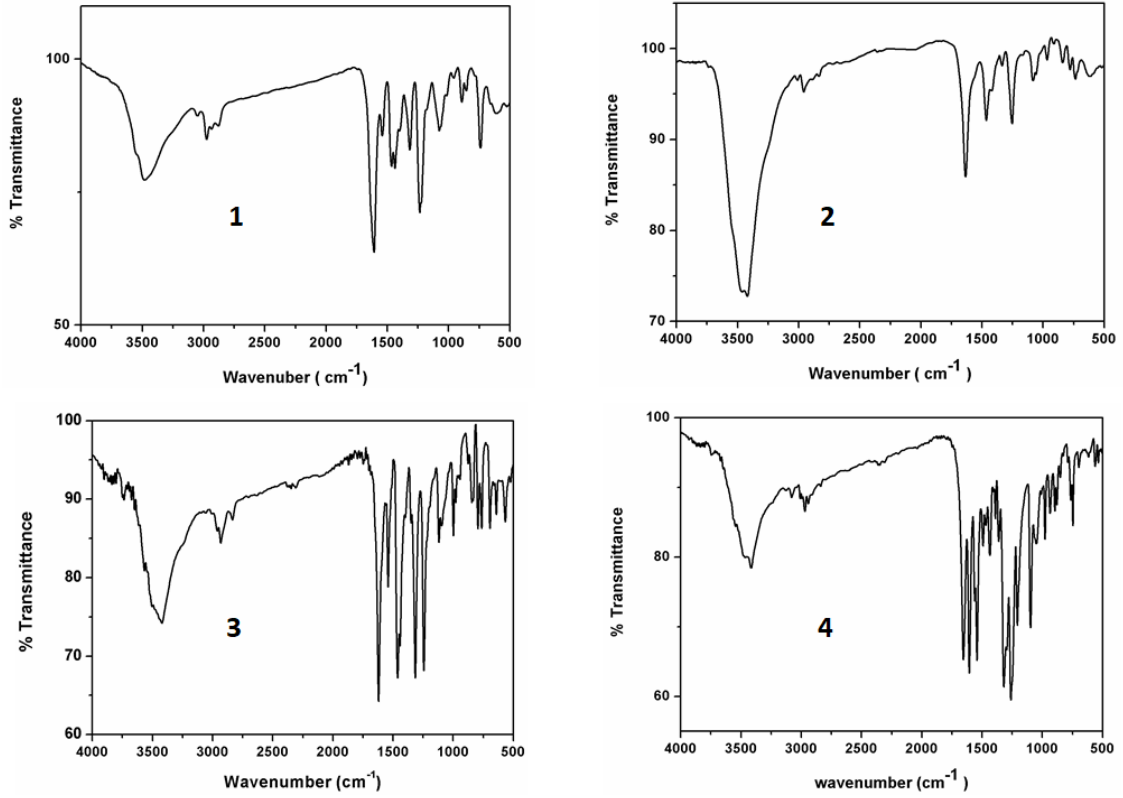
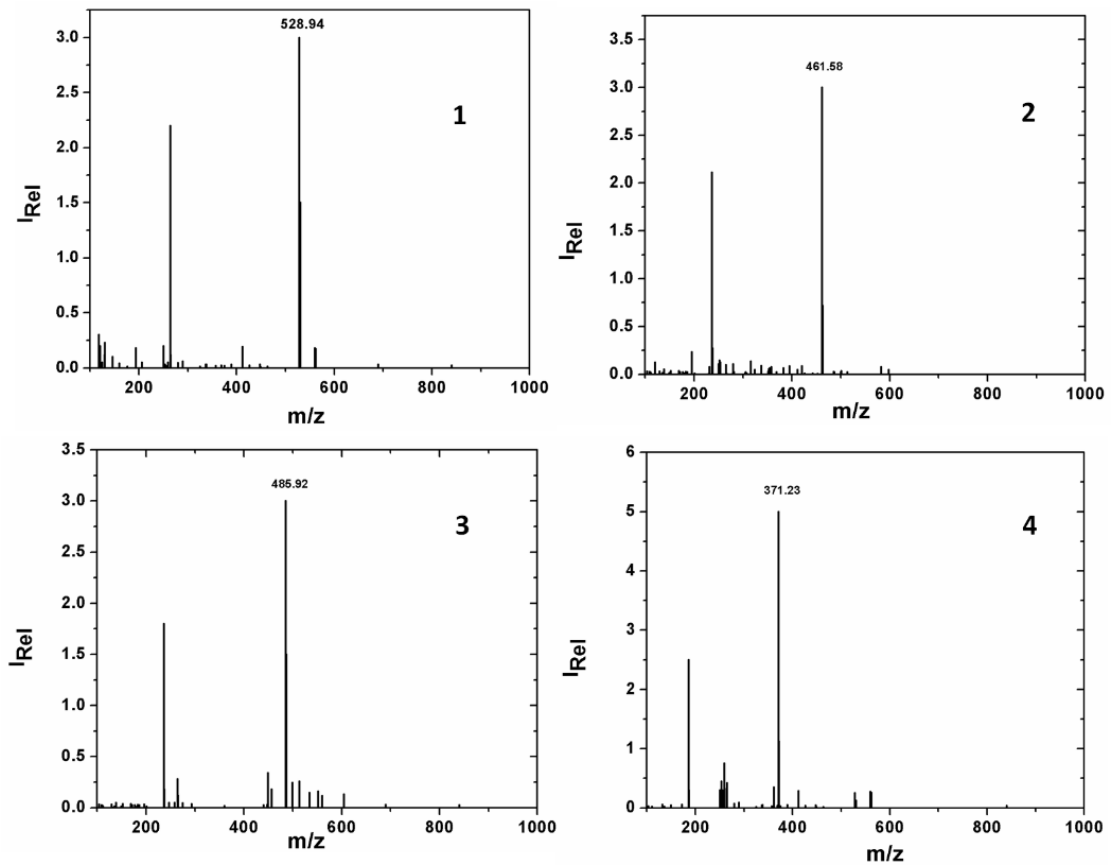


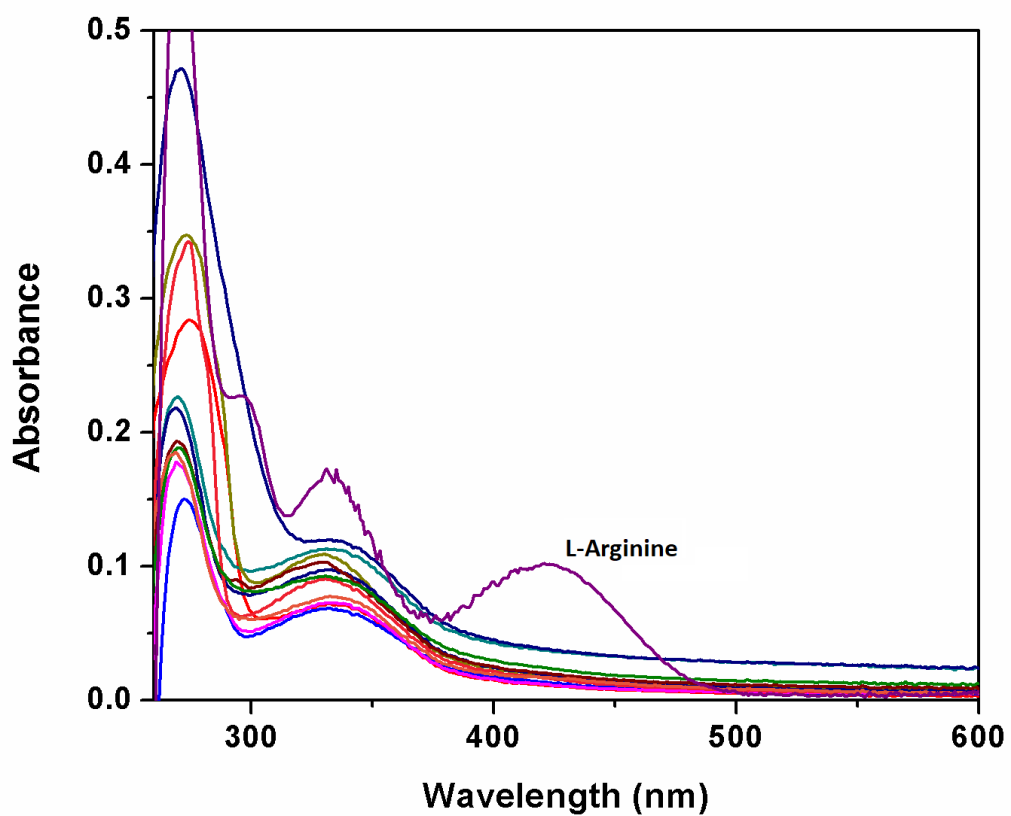
Figure S1. <sup>1</sup>H NMR spectra of sensors 1 to 4.



**Figure S2.** FT-IR spectra of sensors **1** to **4**.



**Figure S3.** GC-MS data of sensors **1** to **4**.



**Figure S4.** Absorption spectra of sensor **1** with five equivalents of various amino acids.

**Table S1**Crystal data and structural refinement parameters of **2**.

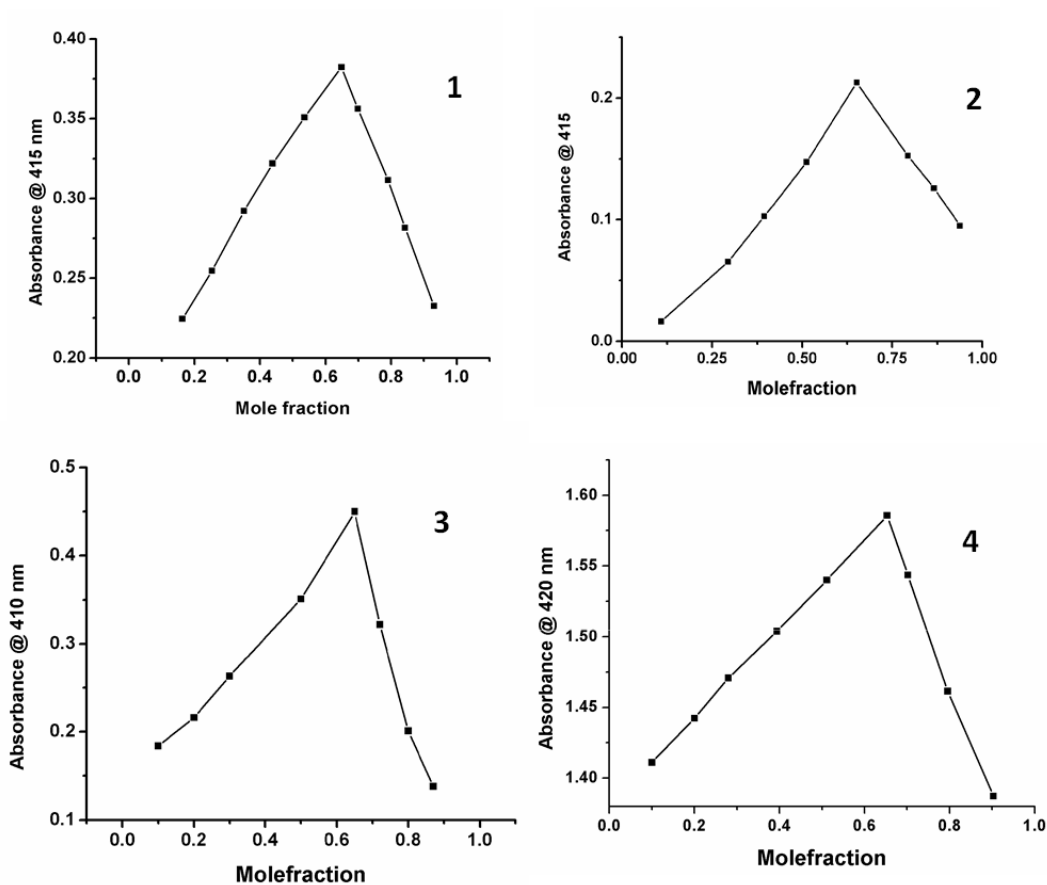
Compound	<b>2</b>
Empirical formula	C <sub>21</sub> H <sub>26</sub> N <sub>2</sub> O <sub>4</sub>
Formula weight	370.44
Color	Yellow
Crystal system	Monoclinic
Space group	<i>P</i> <sub>2</sub> <sub>1</sub> / <i>n</i>
Cell parameters	
a (Å)	15.1698(10)
b (Å)	6.9547(4)
c (Å)	19.5022(12)
α (°)	90
β (°)	93.802(3)
γ (°)	90
Volume V (Å <sup>3</sup> )	2053.0(2)
Z	4
Calculated density (ρ) (Mg m <sup>-3</sup> )	1.198
Absorption coefficient, μ (mm <sup>-1</sup> )	0.083
F(000)	792
Crystal size mm <sup>3</sup>	0.30 x 0.20 x 0.20
θ(°) range for data collection	2.69 to 28.35
Limiting indices	-20 ≤ h ≤ 20, -9 ≤ k ≤ 9, -26 ≤ l ≤ 24
Reflections collected	24051
Unique Reflections (R <sub>int</sub> )	5084(0.0485)
Absorption correction	Semi-empirical from equivalents
Maximum and minimum transmission	0.995 and 0.982
Refinement method	Full-matrix least- squares on F <sup>2</sup>
Data / restraints / parameters	5084/ 3 / 245
Goodness-of-fit on F <sup>2</sup>	0.879
Final R indices [I > 2σ (I)]	R <sub>1</sub> = 0.0604, wR <sub>2</sub> = 0.1659
R indices (all data)	R <sub>1</sub> = 0.1547, wR <sub>2</sub> = 0.2384
Largest difference peak and hole (e Å <sup>-3</sup> )	0.264 and -0.206

$$R_1 = \frac{\sum ||F_o| - |F_c||}{\sum |F_o|} \quad wR_2 = \left[ \frac{\sum w(F_o^2 - F_c^2)^2}{\sum w(F_o^2)^2} \right]^{1/2}$$

**Table S2**Intramolecular hydrogen bonding interactions in **2**.

<b>Hydrogen bonding interactions</b>				
D–H···A	D–H (Å)	H···A (Å)	D···A (Å)	∠D–H···A (°)
O(2)–H(2)···N(1) <sup>a</sup>	0.82	1.86	2.59	147
O(3)–H(3)···N(2) <sup>a</sup>	0.82	1.87	2.59	147
C(11)–H(11A)···N(2) <sup>b</sup>	0.96	2.58	2.90	115

Equivalent position codes : a = 1/2-x, -1/2+y, 1/2-z; b = -1+x, -1+y, z.  
D, Donor; A, acceptor.

**Figure S5.** Jobs plot of sensors **1** to **4** showing 1:2 complex formation between the sensor and the analyte.

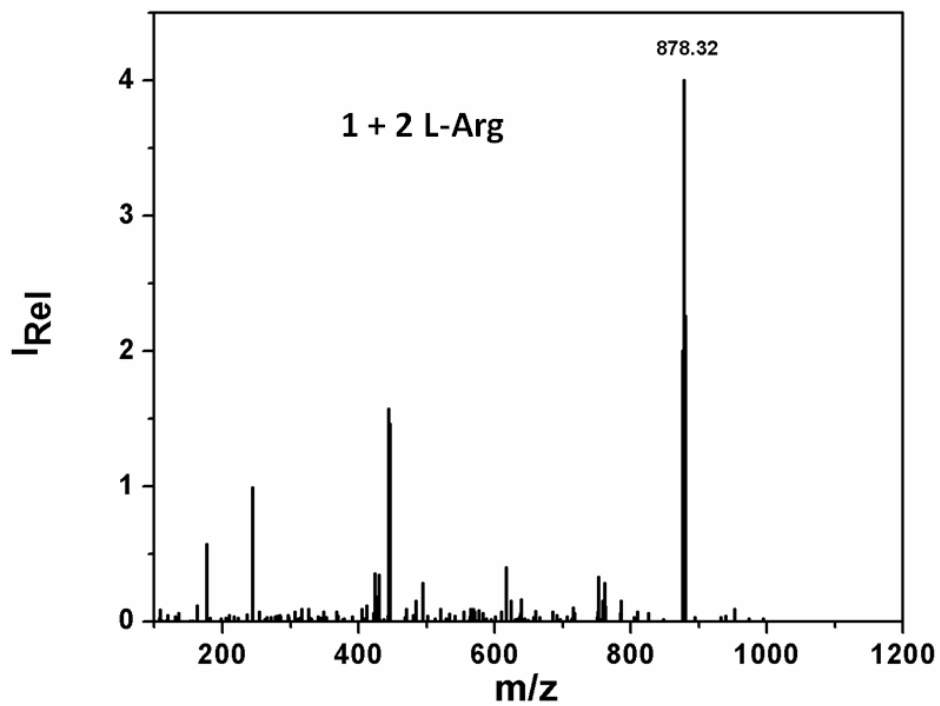


Figure S6. Mass spectrum of 1 + 2 L-Arginine.

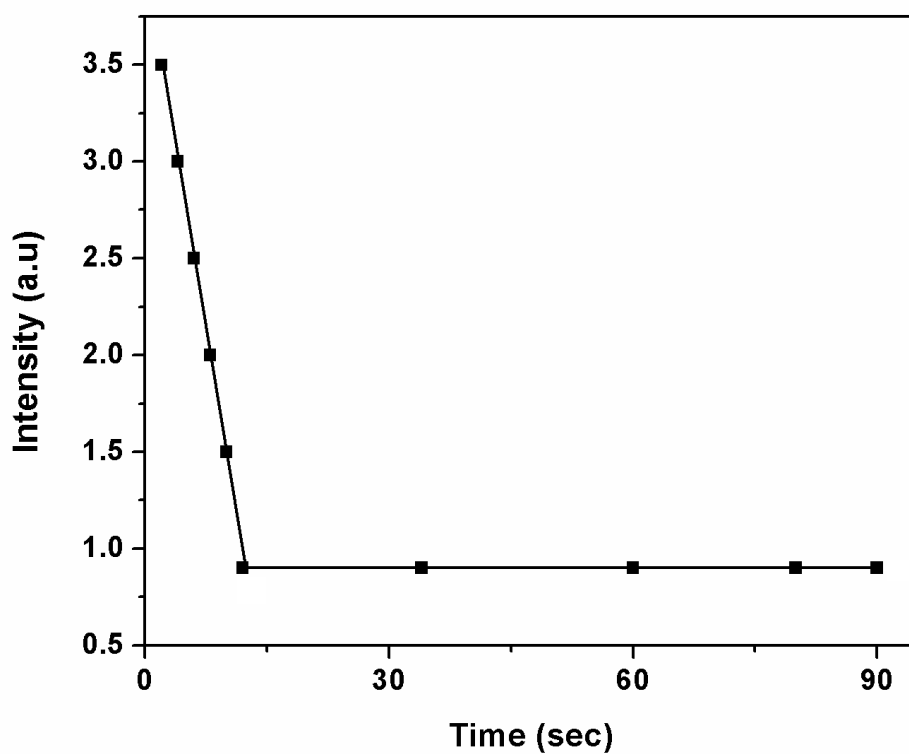


Figure S7. Fluorometric response time of the sensor 4 towards L-Arginine.

**Table S3**

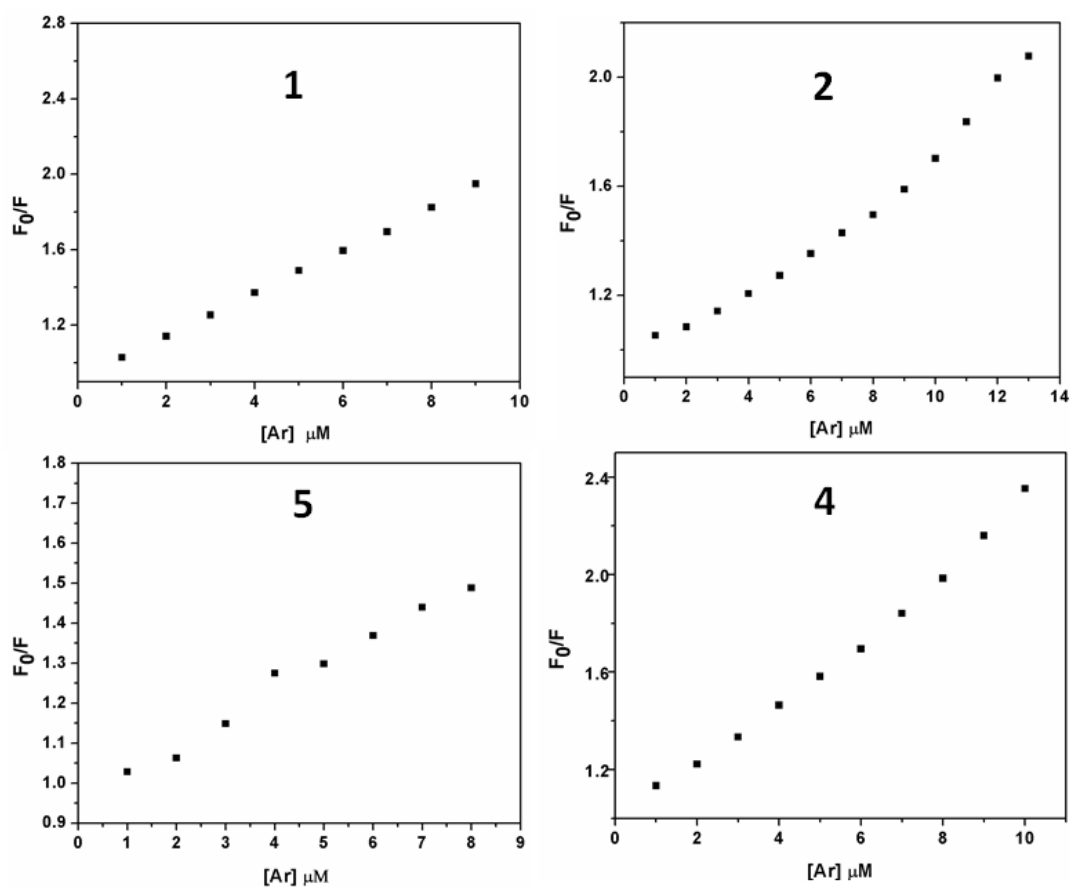
Life time of sensors with and without analyte showing static quenching.

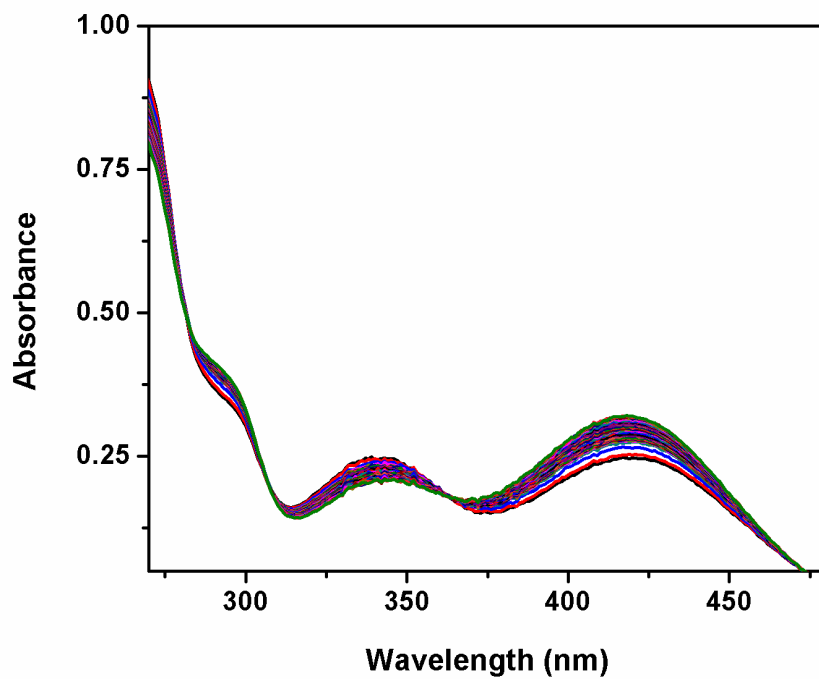
Compounds	Average Lifetime (ns)
Sensor 1	1.40
Sensor 1 + L-Ar	1.52
Sensor 2	0.95
Sensor 2 + L-Ar	0.91
Sensor 3	0.62
Sensor 3 + L-Ar	0.68
Sensor 4	1.25
Sensor 4 + L-Ar	1.49

**Table S3**

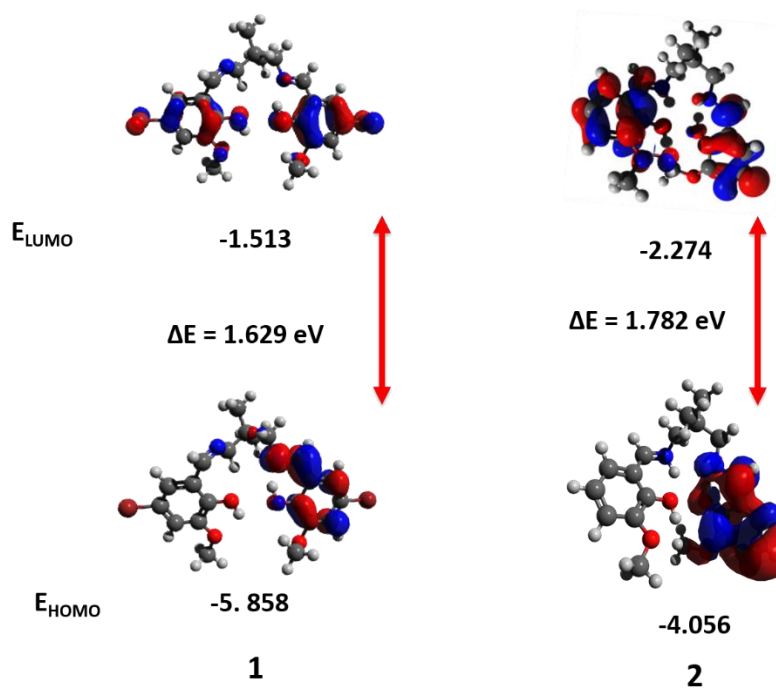
$K_{SV}$  values as obtained from Stern-Volmer plots and quantum yield measurements against standard quinine sulphate for 1-4.

Sensor	1	2	3	4
$K_{SV}(10^4)$	11.43	6.42	6.89	13.85
$\Phi$	0.107	0.274	0.110	0.386

**Figure S8.** Stern-Volmer plots for sensors 1 to 4.

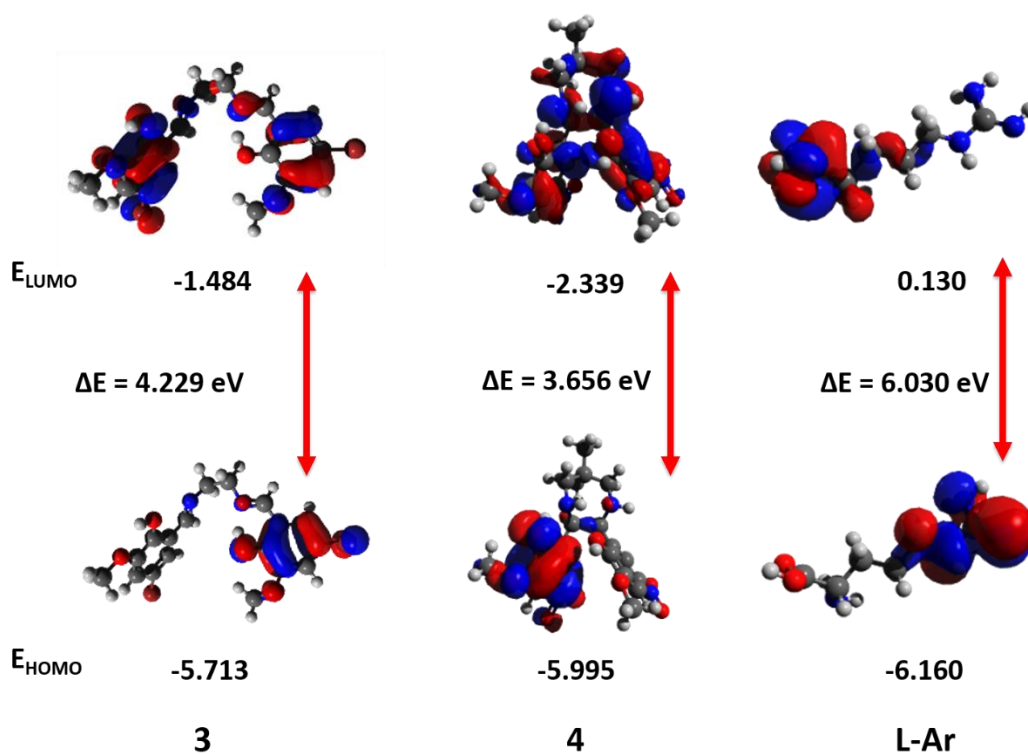


**Figure S9.** Absorption titration of **1** with guanidine.



**Figure S10.** HOMO-LUMO plots of DFT optimized structures of **1** and **2** with the associated energy gaps.





**Figure S11.** HOMO-LUMO plots of DFT optimized structures of **3**, **4** and **L-Ar** with the associated energy gaps.

### Frontier orbital compositional analysis

Frontier orbital calculations were performed using Mulliken method as employed in multiwfn program version 3.3.9.<sup>1</sup> In the case of **1**, HOMO is on one of the benzene rings 73.15% with a little contribution from the azomethine group of the same side (19.13%). But the LUMO is evenly distributed among the two sides with phenolic groups having the highest contribution of about 86.65%. For **2**, HOMO is on the benzene ring of one side with a contribution of about 79.37% and on the methoxy oxygen group (15.44%). In the case of **3**, the trend is similar with the HOMO being concentrated on one of the benzene ring (75.68%) and azomethine nitrogen atom (19.15%) and the LUMO on phenolic moieties (82.27%) on either side with a slight contribution from bromine atoms (9.19%). In **4**, HOMO is mainly concentrated on one of the benzene rings with a cumulative contribution of about 88.96% and there is some share from the oxygen atoms of the nitro group (7.21%) while LUMO is concentrated on phenolic moieties on either side with a total contribution of about 76.58%. In the case of L-Arginine, the HOMO is concentrated on the nitrogen atoms of the amine side with a cumulative contribution of about 85.51% and the LUMO is on the acidic part oxygen atoms with a total of about 79.10%.

### Reference

1. T. Lu and F. Chen, *J. Comput. Chem.*, 2012, **33**, 580-592.