

## Supporting Information

### **AIE active sulfonyldianiline derived Schiff base for the detection of folic acid, hemoglobin and glucose**

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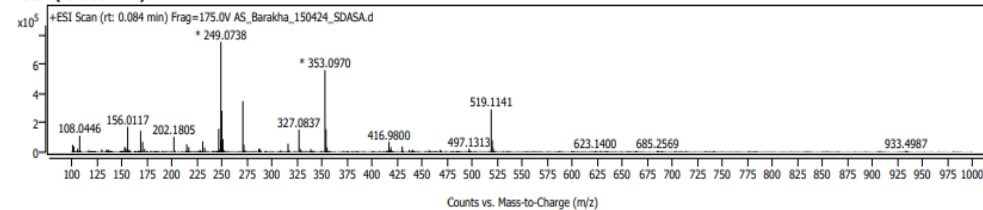
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## Sample Information

<b>Name</b>	AS_Barakha_150424_SDASA	<b>Data File Path</b>	D:\Projects\SEP_2023\Data\Apr_2024\AS_Barakha_150424_SDASA.d
<b>Sample ID</b>		<b>Acq. Time (Local)</b>	15-04-2024 15:22:12 (UTC+05:30)
<b>Instrument</b>	LCQTOF	<b>Method Path (Acq)</b>	D:\Projects\SEP_2023\Methods\POSITIVE.m
<b>MS Type</b>	QTOF	<b>Version (Acq SW)</b>	6200 series TOF/6500 series Q-TOF (11.0.203.0)
<b>Inj. Vol. (ul)</b>	2	<b>IRM Status</b>	Success
<b>Position</b>	P1-B2	<b>Method Path (DA)</b>	D:\Projects\INSTALLATION\Methods\IIT Roorkee.m
<b>Plate Pos.</b>		<b>Target Source Path</b>	
<b>Operator</b>	SYSTEM (SYSTEM)	<b>Result Summary</b>	1 qualified (1 targets)

## Sample Spectra

## + Scan (rt: 0.084 min)

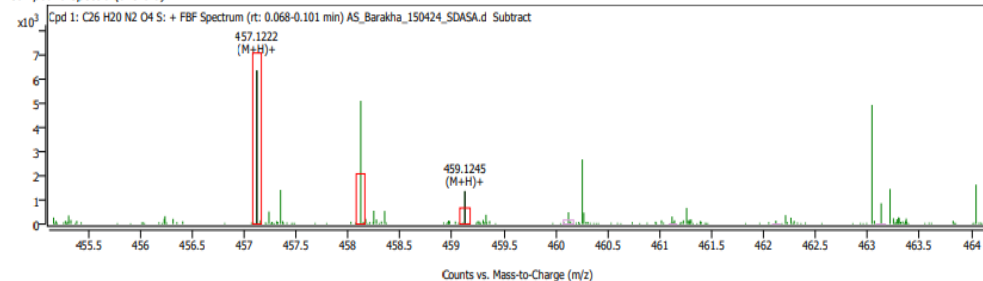


## Compound Details

Cpd. 1: C<sub>26</sub>H<sub>20</sub>N<sub>2</sub>O<sub>4</sub>S

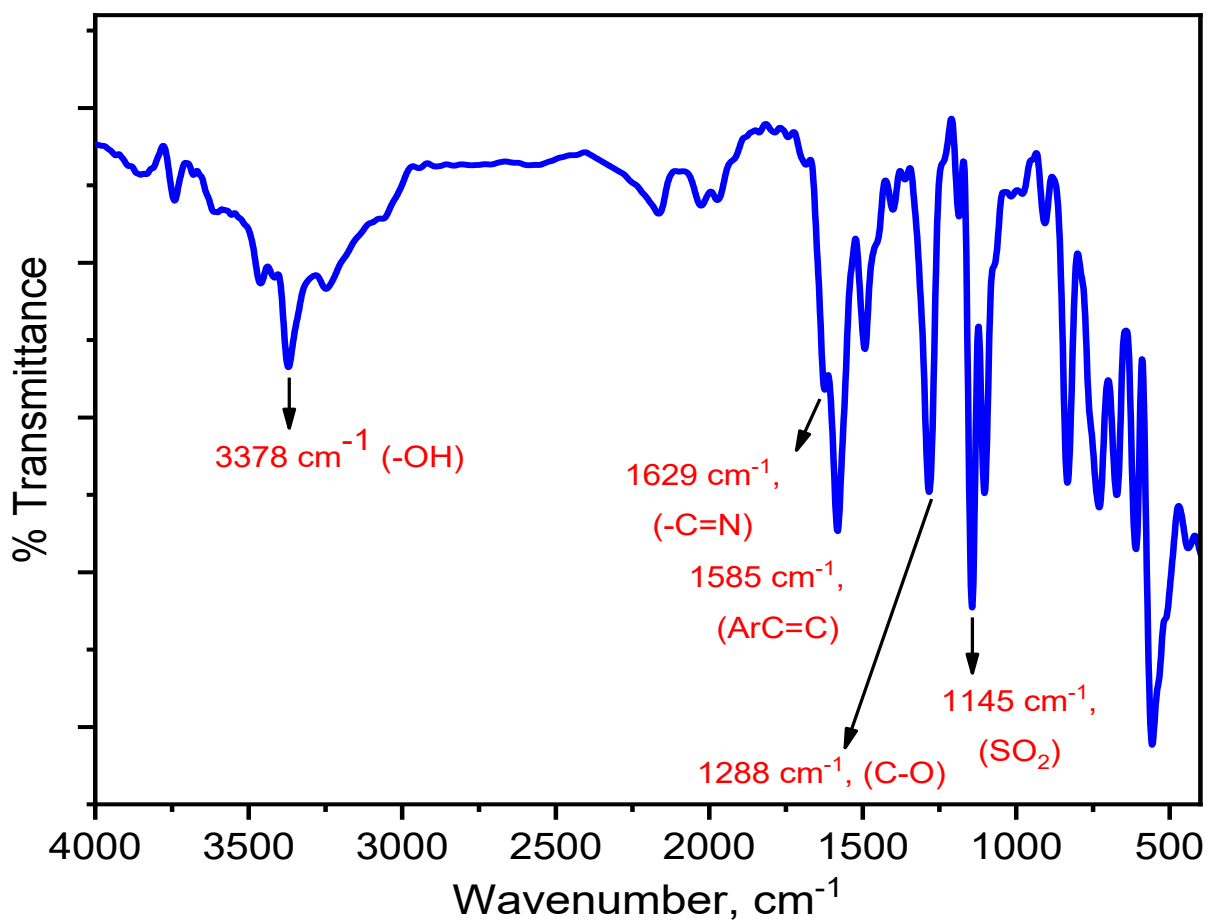
Formula	Mass	Score	Algorithm	Diff (Tgt, ppm)	Polarity
C <sub>26</sub> H <sub>20</sub> N <sub>2</sub> O <sub>4</sub> S	456.1151	57.79	FBF	1.64768286951381	Positive

## Compound Spectra (overlay)

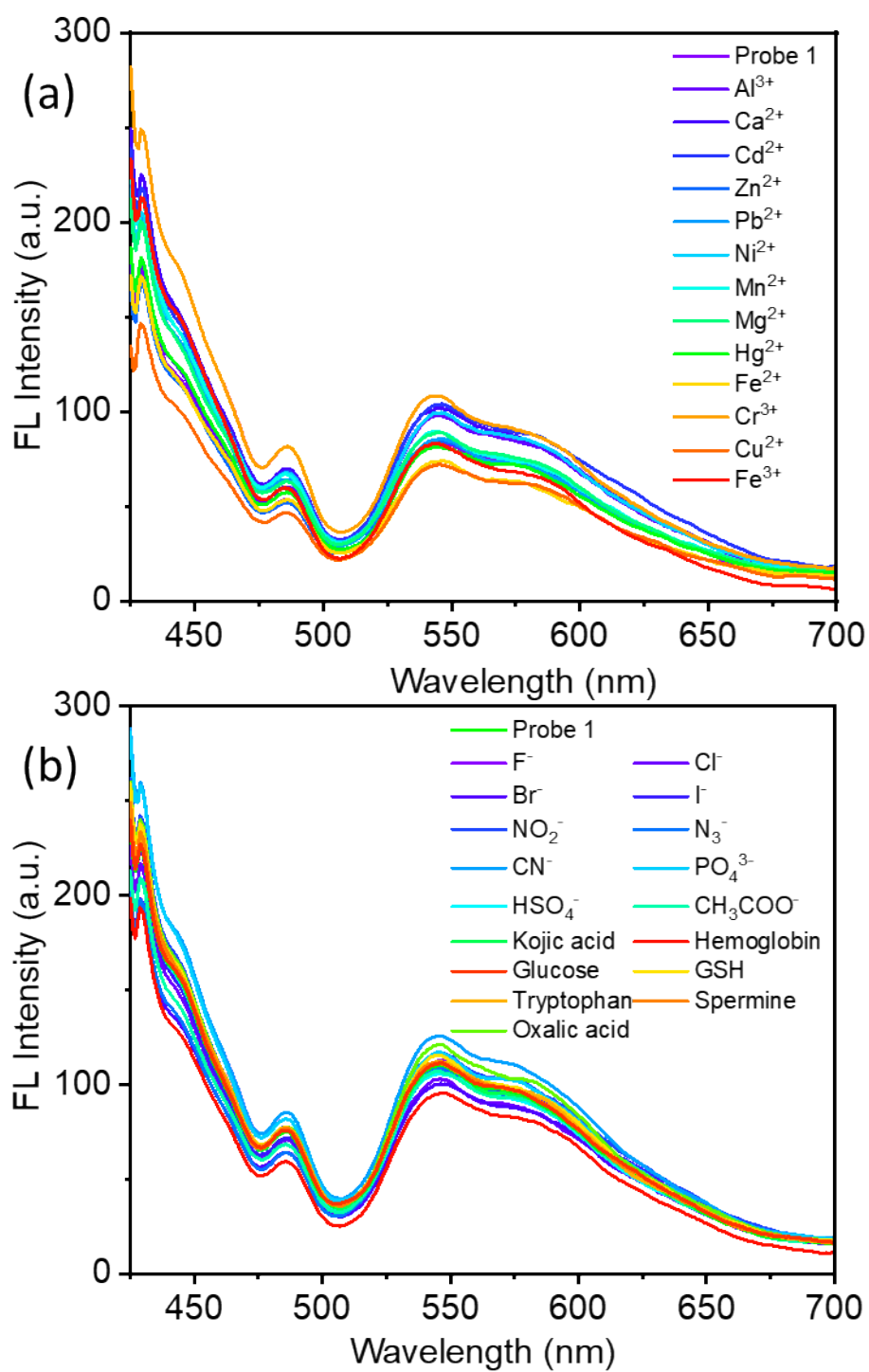


MassHunter Qual 10.0  
(End of Report)

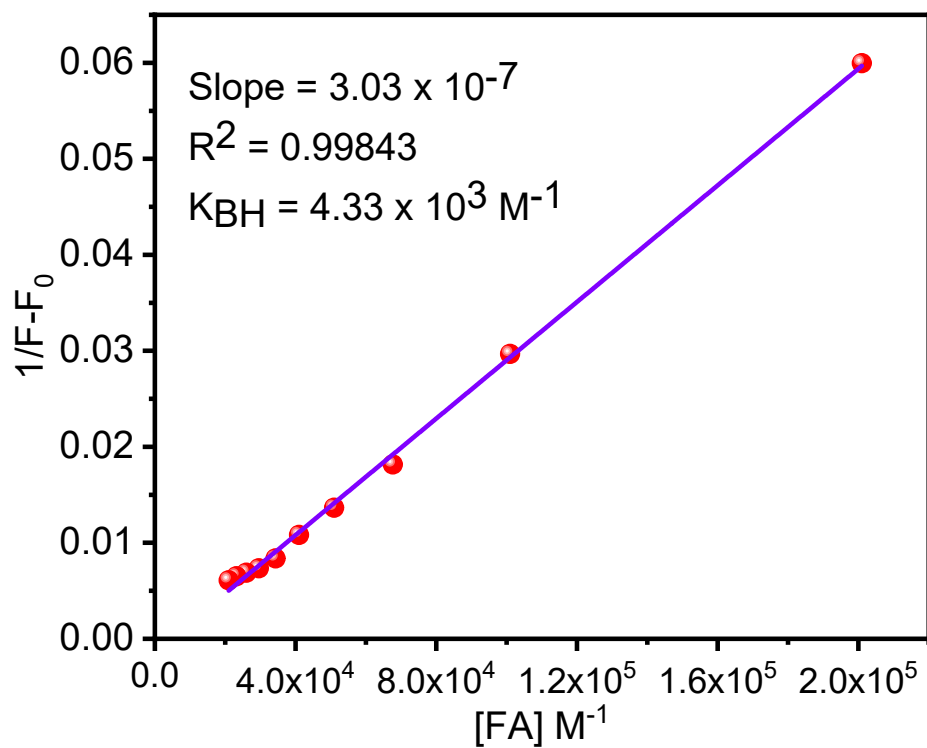
**Fig. S1.** Mass spectrum of SDASA.



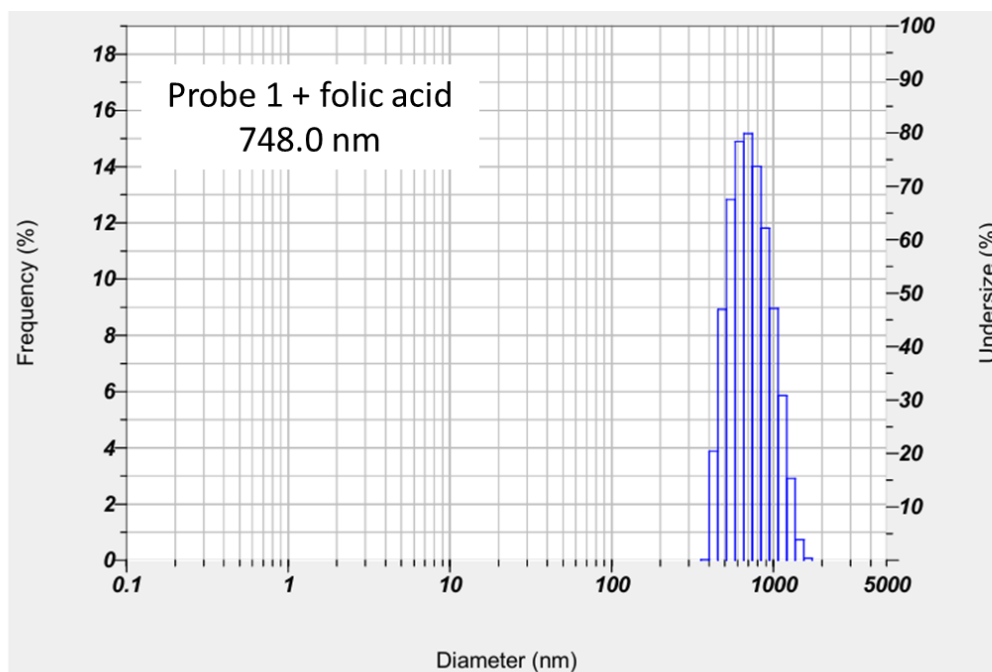
**Fig. S2.** FT-IR spectra of SDASA.



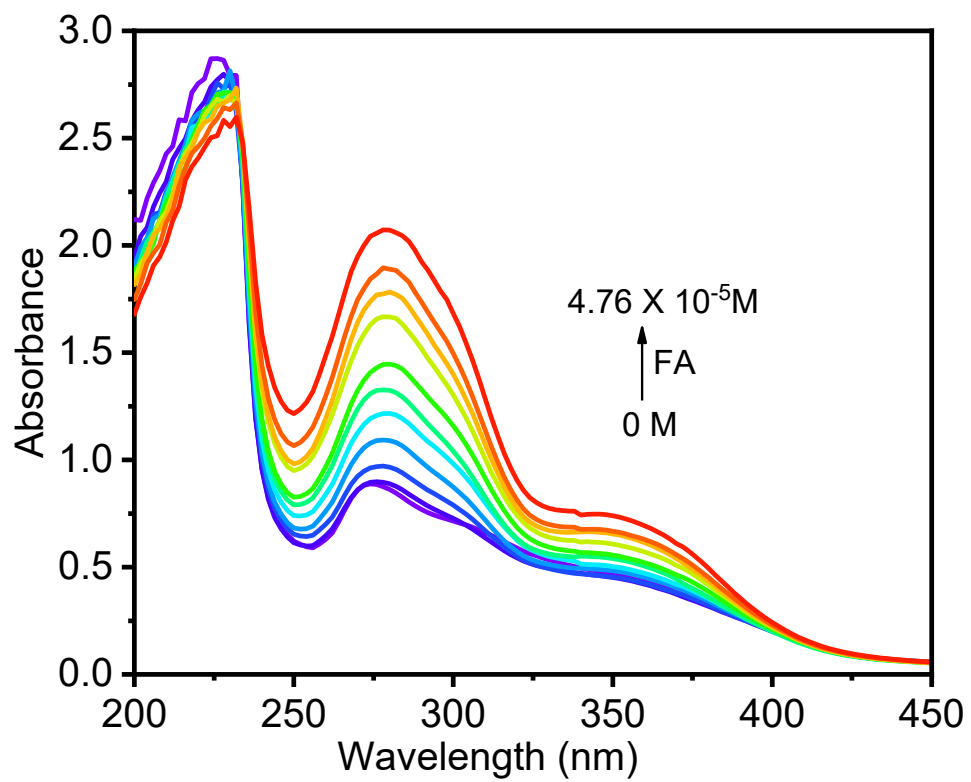
**Fig. S3.** The fluorescence spectral changes of Probe 1 ( $5 \times 10^{-5}$  M) in the absence and presence of different analytes ( $5 \times 10^{-5}$  M).



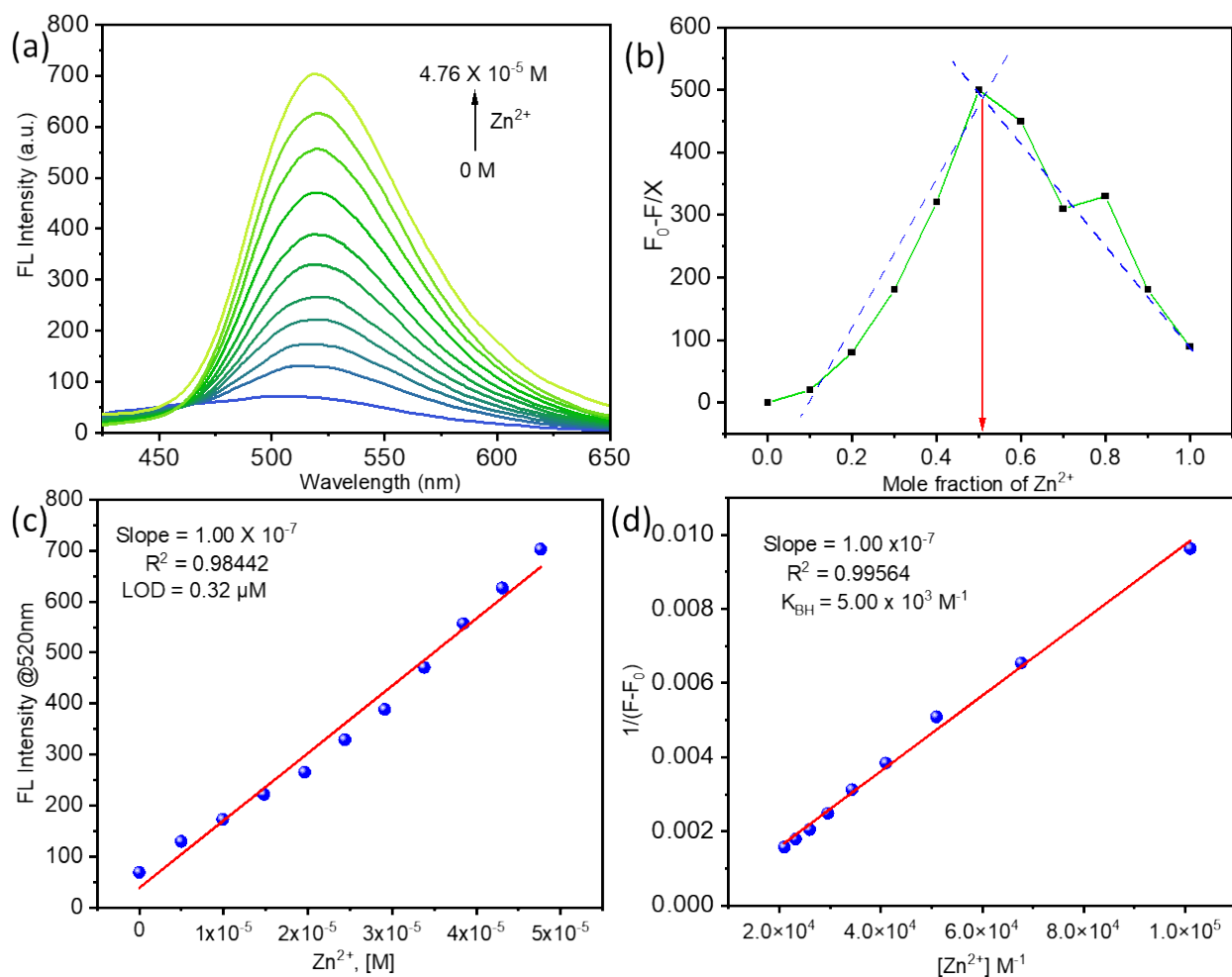
**Fig. S4.** The B-H plot of probe 1 with FA.



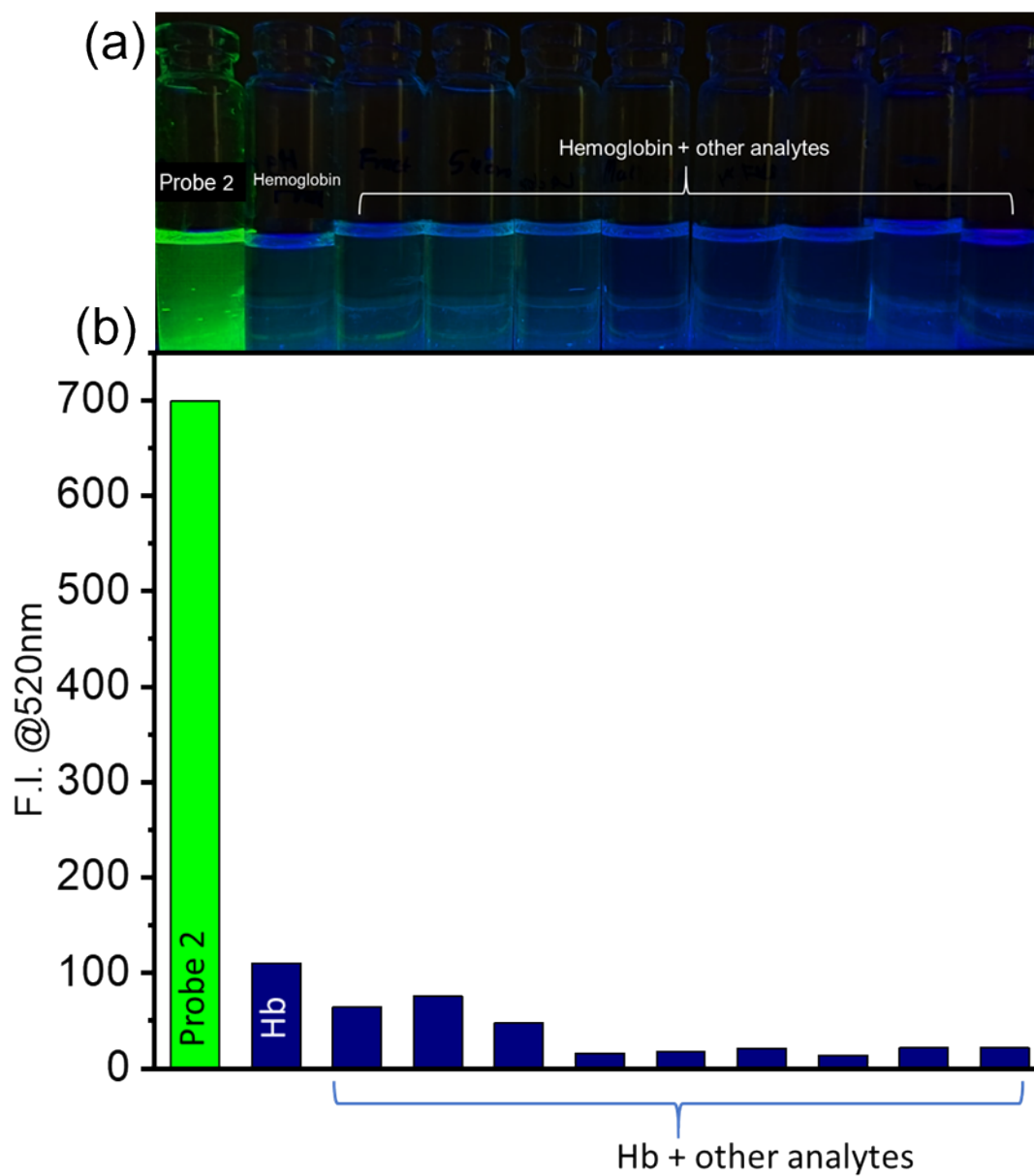
**Fig. S5.** DLS of probe 1 in the presence of FA.



**Fig. S6.** UV-Vis spectral changes of probe 1 upon successive incremental addition of FA.

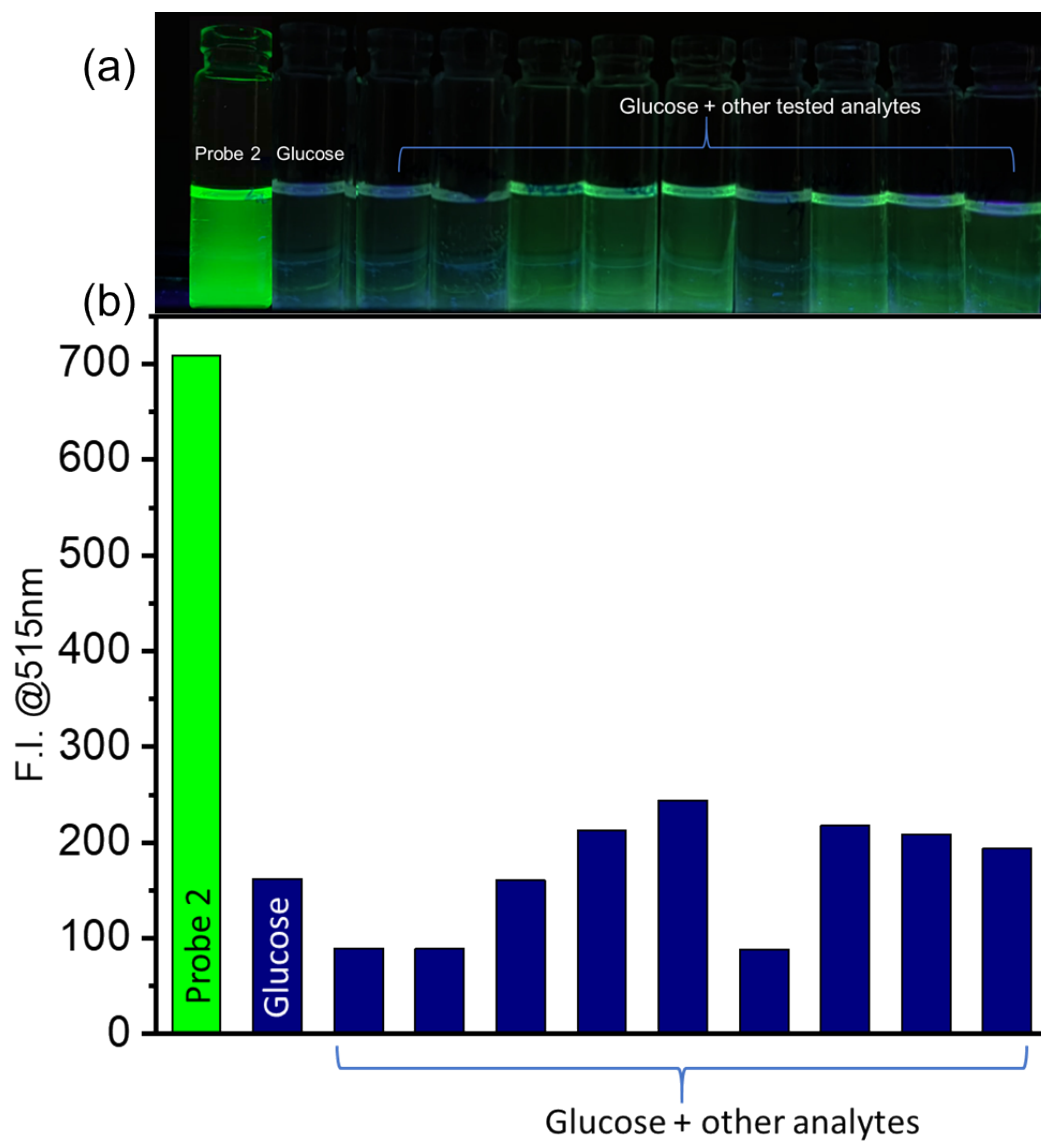


**Fig. S7.** (a) Fluorescence spectral changes of SDASA ( $5 \times 10^{-5}$  M, DMSO) upon successive incremental addition of  $\text{Zn}^{2+}$ . (b) The calibration plot for estimation the LOD for  $\text{Zn}^{2+}$ . (c) The Job's plot of SDASA- $\text{Zn}^{2+}$  complex at 520 nm, plotted against the mole fraction of  $\text{Zn}^{2+}$ . (d) The B-H plot for SDASA- $\text{Zn}^{2+}$  complex.

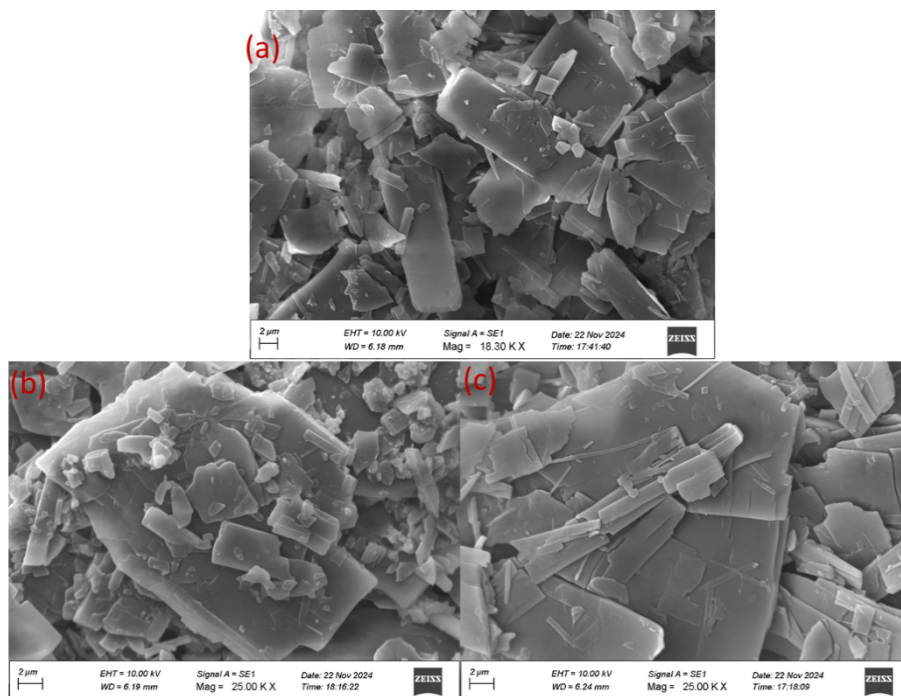


**Fig. S8.** Fluorescence colour (a) and spectral (b) changes of probe 2 ( $5 \times 10^{-5}$  M) in the presence of Hb ( $2.5 \times 10^{-5}$  M) and an equimolar amount of other analytes.

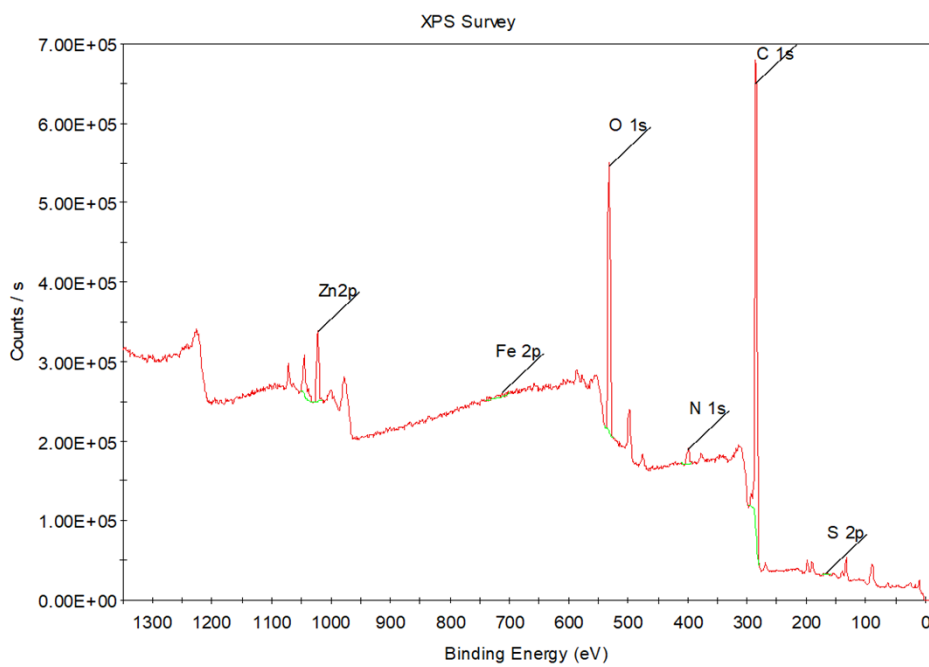




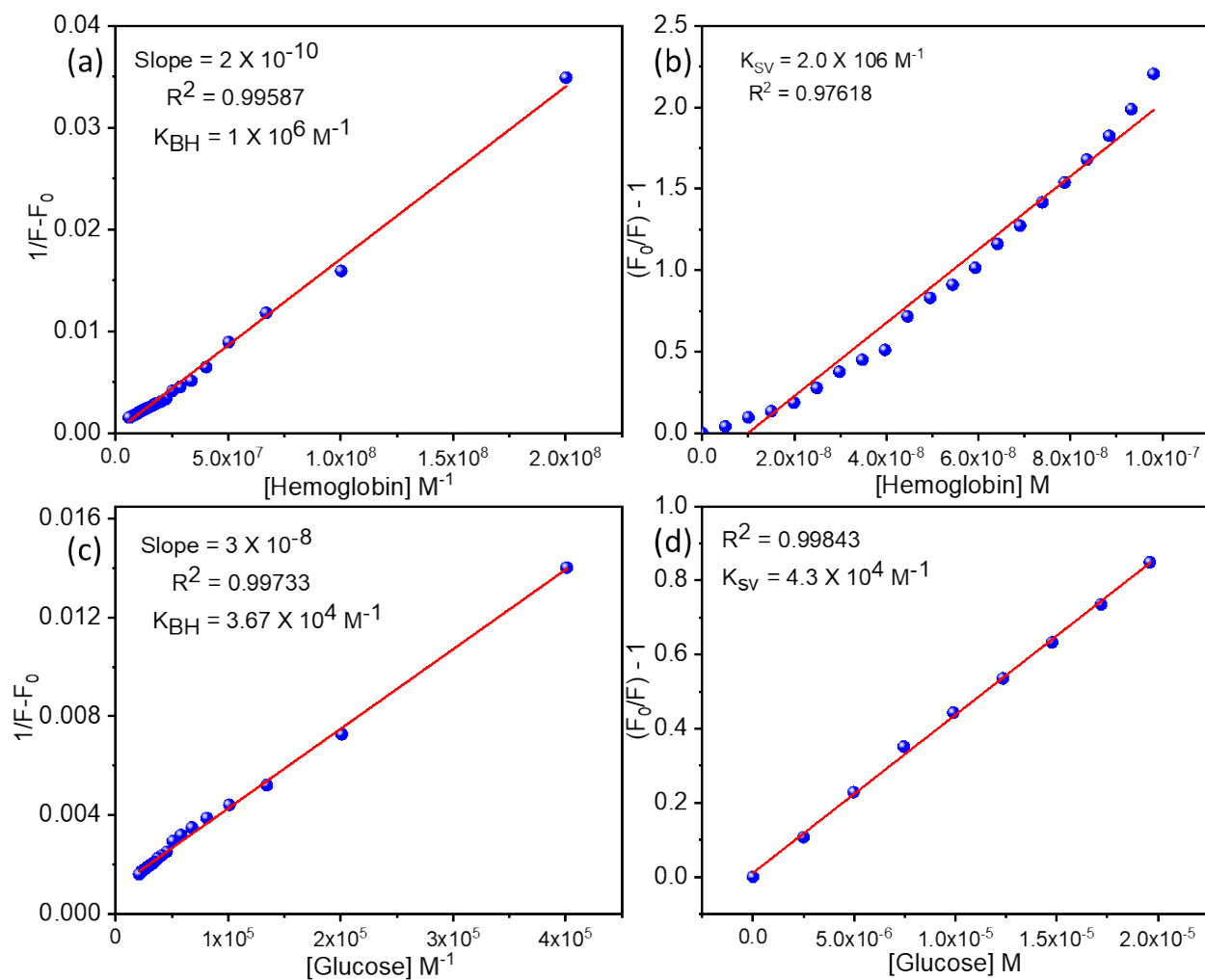
**Fig. S9.** Fluorescence colour (a) and spectral (b) changes of probe 2 ( $5 \times 10^{-5}$  M) in the presence of glucose ( $2.5 \times 10^{-5}$  M) and an equimolar amount of other analytes.



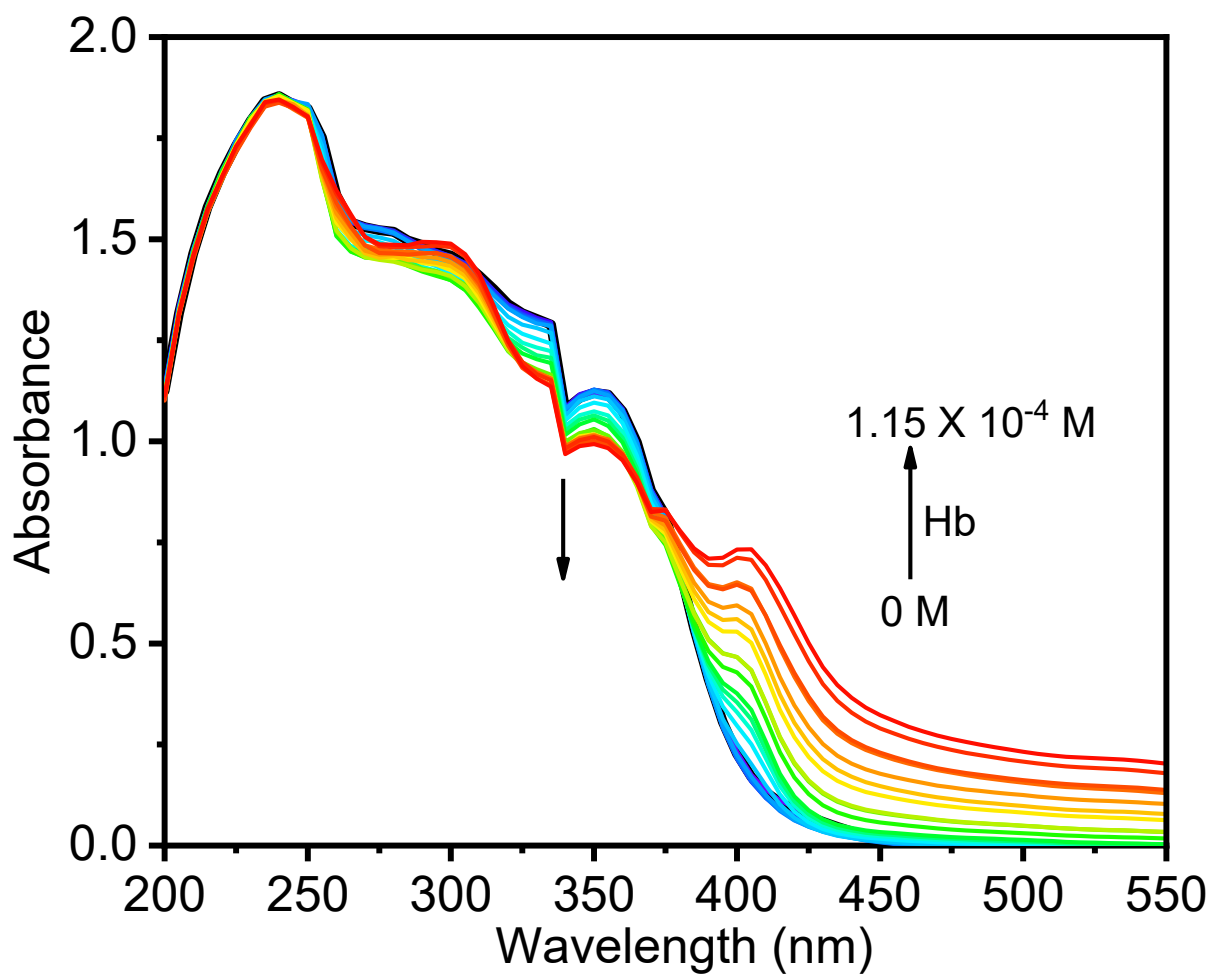
**Fig. S10.** The SEM images of probe 2 in the absence (a) and presence of hemoglobin (b) and glucose (c).



**Fig. S11.** XPS survey scan of probe 2 with hemoglobin.



**Fig. S12.** The B-H plot of probe 2 with Hb (a) and glucose (c). The Stern-Volmer plot for the fluorescence quenching of probe 2 by Hb (b) and glucose (d).



**Fig. S13.** UV-visible spectral changes of probe 2 with the gradual incremental addition of Hb.

**Table S1.** Fluorescence lifetime decay parameters of probe 1 and probe 1 with FA.

	$\tau^1$ (ns)	$\tau^2$ (ns)	$\tau^3$ (ns)	Average $\tau$ (ns)
Probe 1	0.20 ns	0.50 ns	2.62 ns	1.10 ns
Probe 1+ FA	1.30 ns	0.40 ns	2.30 ns	1.33 ns

**Table S2.** Calculated atomic percentage of different elements of probe 2 in the presence of hemoglobin from the XPS scan.

Name	Start BE	Peak BE	End BE	Height CPS	FWHM eV	Area (P) CPS.eV	Area (N) TPP-2M	Atomic %
S 2p	175	166.61	157	2034.56	7.157	18532.78	0.01	0.31
C 1s	298	284.7	279.3	574852.32	3.559	2321461.02	3.69	76.69
N 1s	410	399.22	392	18238.74	4.866	98829.92	0.1	2.07
O 1s	539.34	531.87	525	334040.45	4.202	1478233.23	0.94	19.43
Fe 2p	740	712.02	700	4466.61	1.46	68909.89	0.01	0.21
Zn2p	1052	1022.29	1015	88234.94	4.049	710086.12	0.06	1.3

**Table S3.** Fluorescence lifetime decay parameters of probe 2 and probe 2 with glucose.

	$\tau^1$ (ns)	$\tau^2$ (ns)	$\tau^3$ (ns)	Average $\tau$ (ns)
Probe 2	0.67 ns	1.34 ns	2.68 ns	1.56 ns
Probe 2+ Glucose	0.80 ns	1.61 ns	3.22 ns	1.87 ns

**Table S4.** Real sample analysis of folic acid using probe 1.

Sample	Folic acid			
	Added, M	Found, M	Recovery, %	RSD, %
Serum	$4.97 \times 10^{-6}$	$4.50 \times 10^{-6}$	90.54 %	1.20
	$1.23 \times 10^{-5}$	$1.15 \times 10^{-5}$	93.49 %	2.16
	$1.47 \times 10^{-5}$	$1.42 \times 10^{-5}$	97.93 %	0.68

**Table S5.** Real sample analysis of hemoglobin using probe 2.

Sample	Hemoglobin			
	Added, M	Found, M	Recovery, %	RSD, %
Serum	$7.38 \times 10^{-8}$	$7.34 \times 10^{-8}$	99.45%	0.13

	$7.87 \times 10^{-8}$	$7.52 \times 10^{-8}$	95.55 %	0.42
	$8.35 \times 10^{-8}$	$7.93 \times 10^{-8}$	94.97 %	0.20

**Table S6.** Real sample analysis of glucose using probe 2.

Sample	Glucose			
	Added, M	Found, M	Recovery, %	RSD, %
Serum	$3.14 \times 10^{-5}$	$3.05 \times 10^{-5}$	97.13 %	1.91
	$3.38 \times 10^{-5}$	$3.15 \times 10^{-5}$	93.19 %	0.69
	$3.61 \times 10^{-5}$	$3.33 \times 10^{-5}$	92.24 %	0.64

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