

Supplementary Data

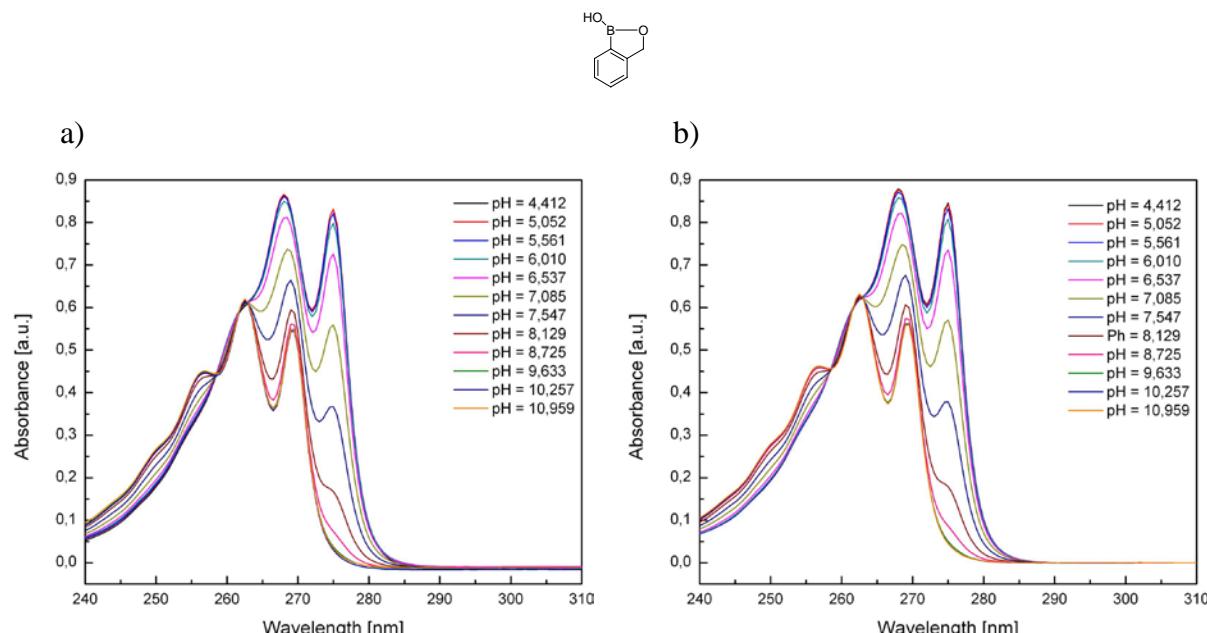
Lewis acidity and sugar receptor activity of 3-aminosubstituted benzoxaboroles and their *ortho*-aminomethylphenylboronic acids analogues

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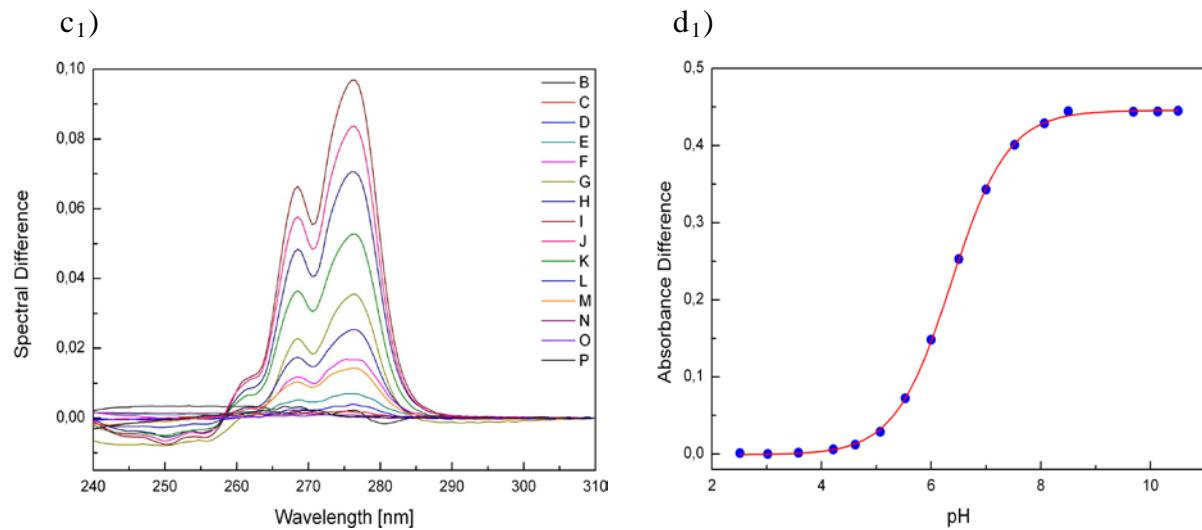
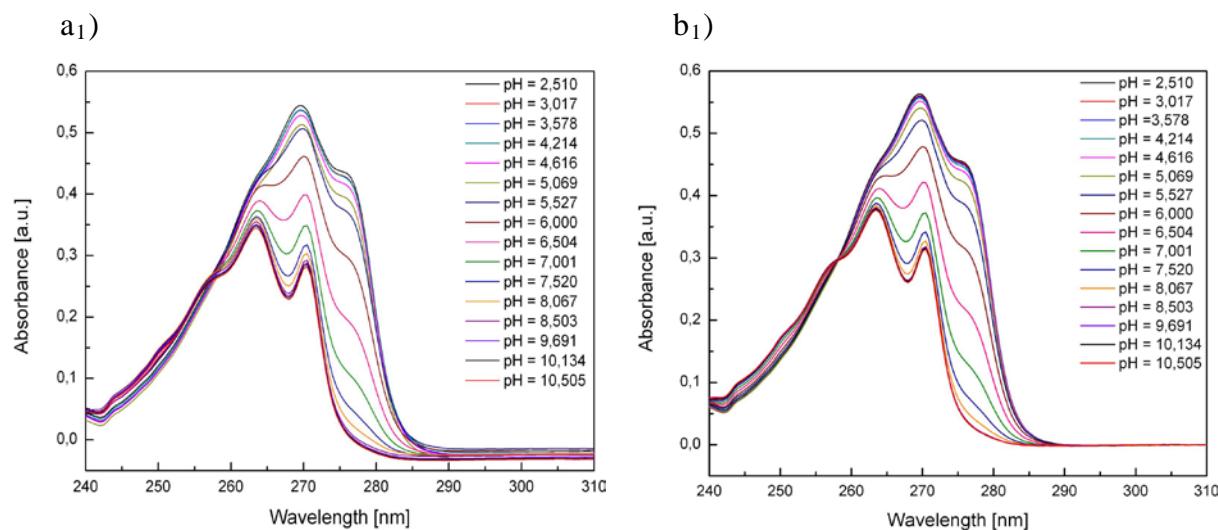
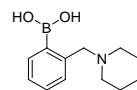
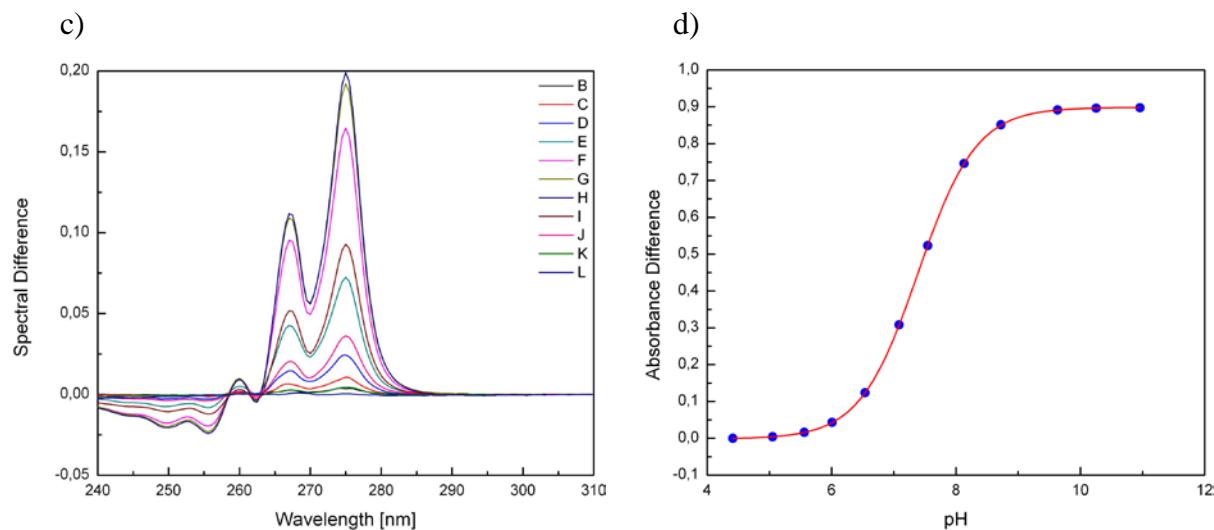
Table 1. pK_a values of **1**, **2a-c** and **3a-c** determined in three independent experiments

	1	2a	2b	2c	3a	3b	3c
pK_{a1}	7,391	7,484	7,485	7,469	4,895	5,212	6,298
pK_{a2}	7,387	7,480	7,326	7,376	4,943	5,134	6,371
pK_{a3}	----	7,219	7,324	7,335	4,965	5,252	6,350
pK_a	7,389	7,394	7,378	7,393	4,934	5,199	6,340
+ / -	0,002	0,124	0,075	0,056	0,029	0,049	0,031

Figure 1. Spectral data and pK_a determination; a) UV/vis spectral scans for compounds **1**, **3c** and phenylboronic acid (PBA) in solutions at various pH values; b) Normalized spectral scans ($\text{Absorbance}_{310\text{nm}} = 0$); c) Spectral differences between solutions of the compounds under study; d) pK_a plots



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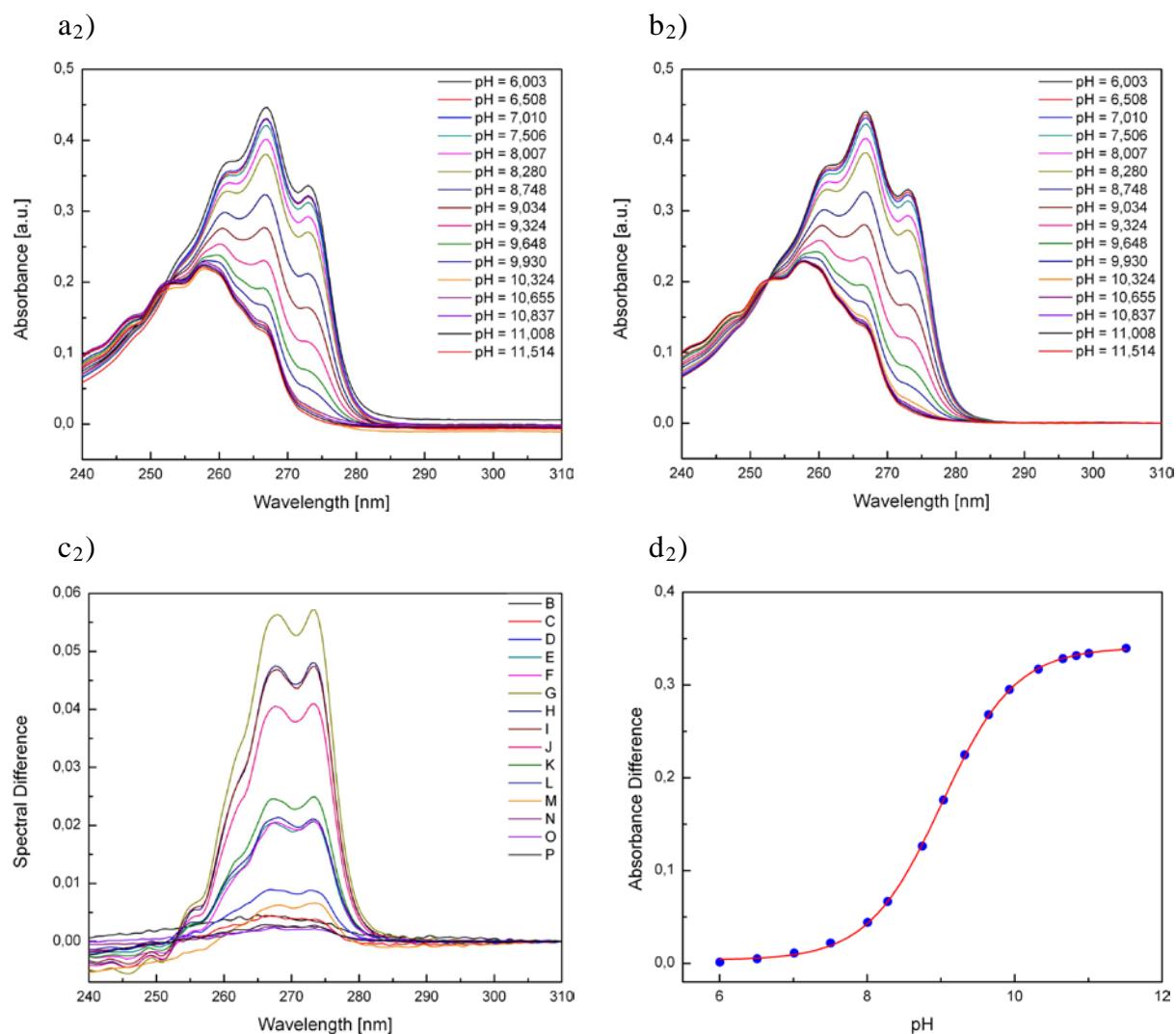
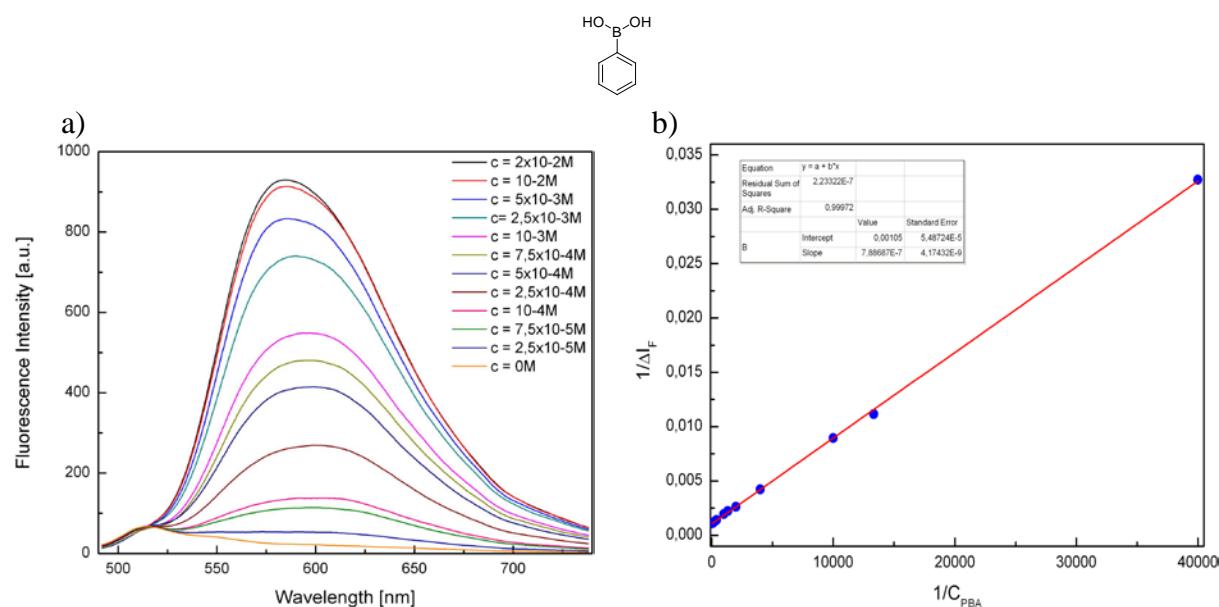


Table 2. The apparent binding constants ($K [\text{M}^{-1}]$) of **1**, **2a-c**, **3a-c** with ARS, fructose, galactose and glucose.

	1	2a	2b	2c	3a¹	3b¹	3c
K_{ARS}	659	1750 ± 105	2556	2315 ± 124	2900 ± 120	720 ± 40	3209 ± 159
K_{Fru}	50	395.8 ± 31.7	263	930.1 ± 8.9	120 ± 30	55 ± 4	449 ± 12.7
K_{Gal}	19.1 ± 0.6	78.4 ± 12.1	17.46	58.7 ± 0.2	100 ± 20	5.7 ± 0.8	48.5 ± 15.5
K_{Glu}	3.9	10.05 ± 0.17	14.22	33.05 ± 1.65	13 ± 5	2.4 ± 0.9	32.10 ± 9.02

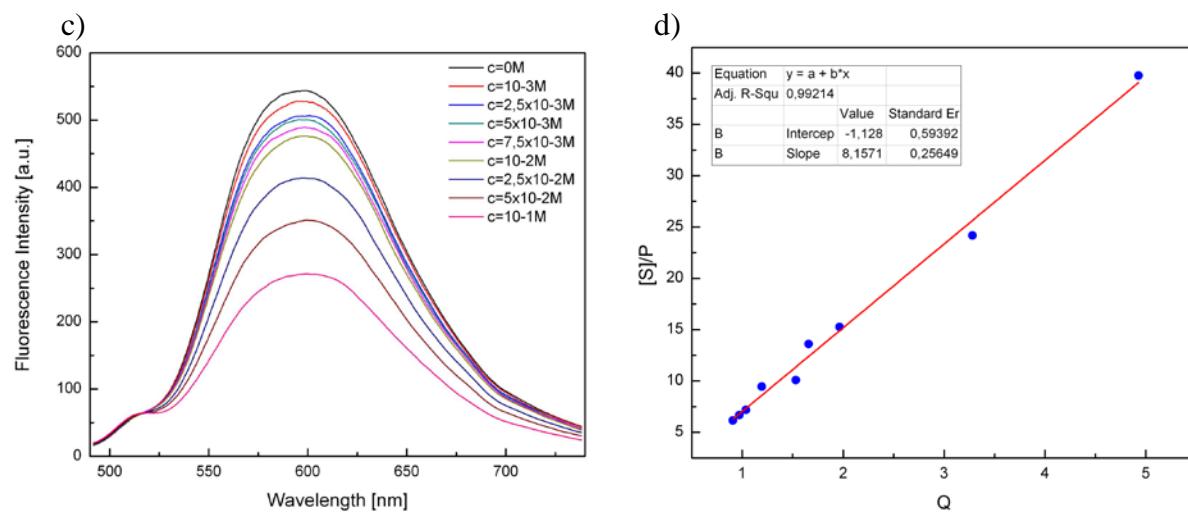
Figure 2) The apparent binding constants determination by ARS method; a) Fluorescence spectrum of ARS at different concentrations of boronic acid; b) $1/\Delta I_F$ versus $1/C_{\text{boronic acid}}$; c) Titration of fructose into a solution of ARS and boronic acid where 20% of ARS is in a free form; d) $[S]/P$ versus Q



Equations for binding constant determination:

$$\frac{1}{\Delta I_F} = (\Delta k p_0 K_{eq1})^{-1} \frac{1}{[C_{\text{boronicacid}}]} + (\Delta k p_0 I_0)^{-1}$$

The binding constant of ARS-phenylboronic acid (K_{ARS}) is the quotient of the intercept and the slope in a plot of $1/\Delta I_F$ versus $1/C_{\text{phenylboronic acid}}$. We compared value of $K_{\text{ARS}} = 1331 \text{ M}^{-1}$ (obtained in our lab) to the previously reported^{2,3} ($K_{\text{ARS}} = 1300 \text{ M}^{-1}$) to validate our method.



The binding constant of fructose-phenylboronic acid (K_{FRU}) is determined by plotting $[S]/P$ versus Q where:

$$P = [L_0] - \frac{1}{QK_{\text{ARS}}} - \frac{[I_0]}{Q+1}$$

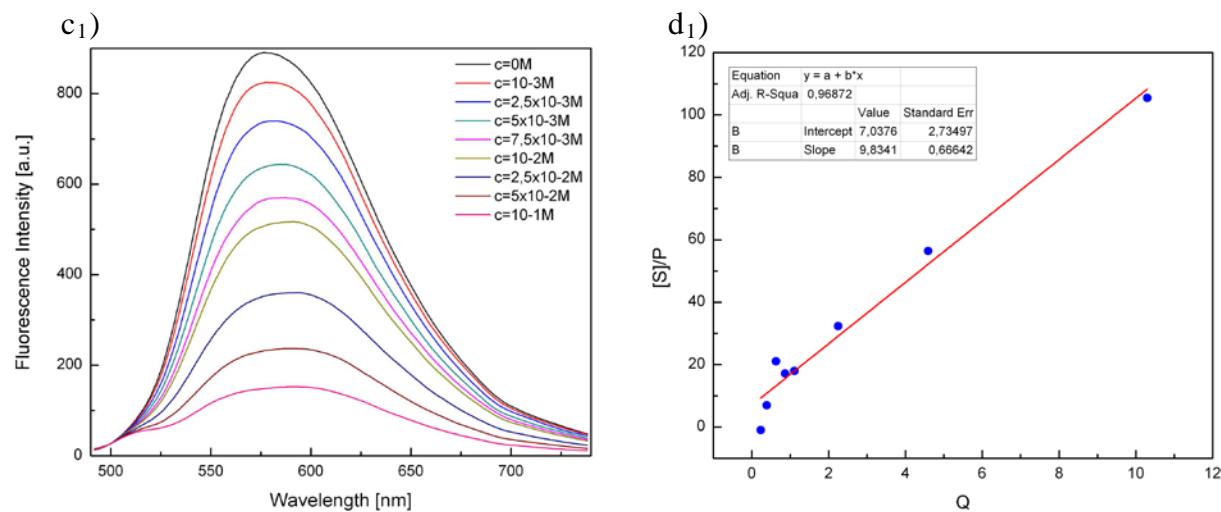
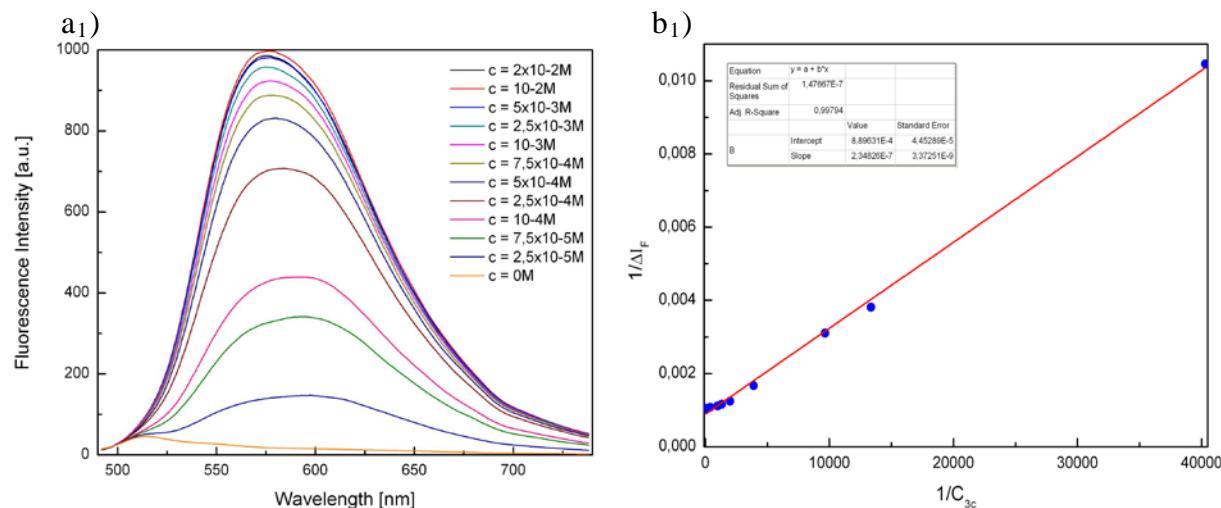
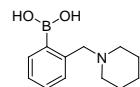
L_0 – Total concentration of phenylboronic acid

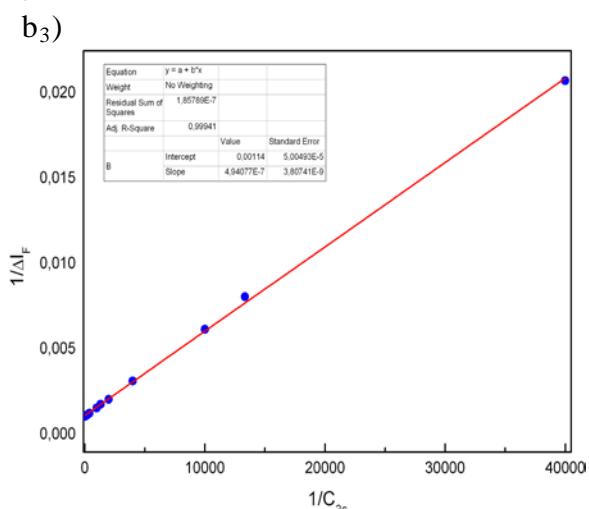
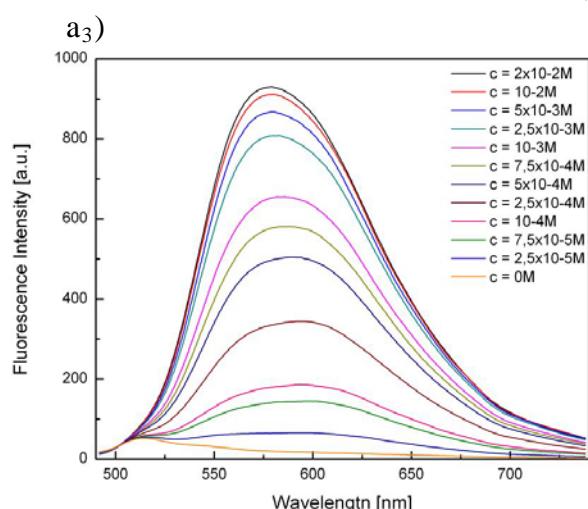
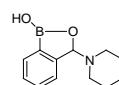
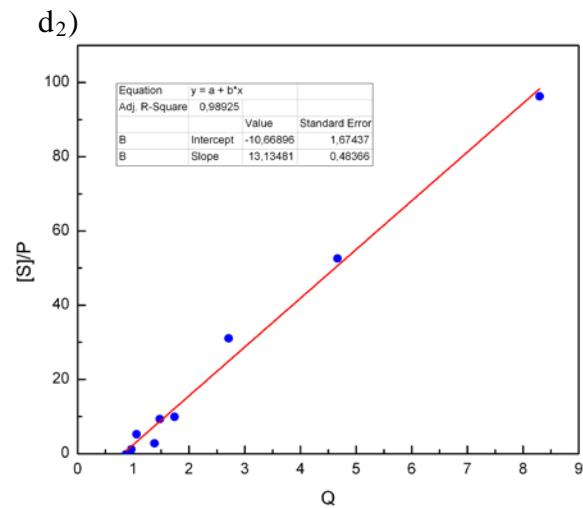
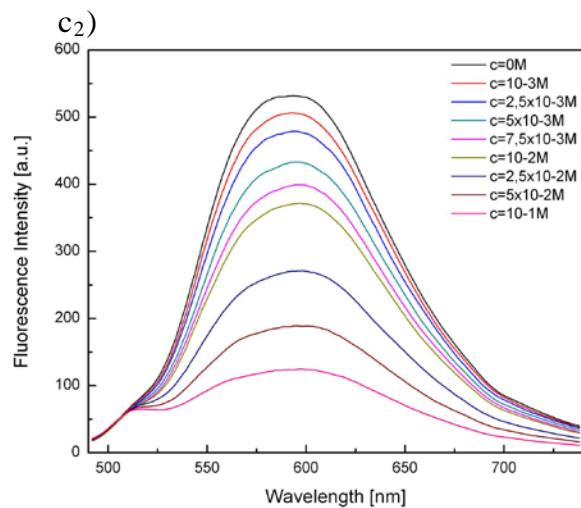
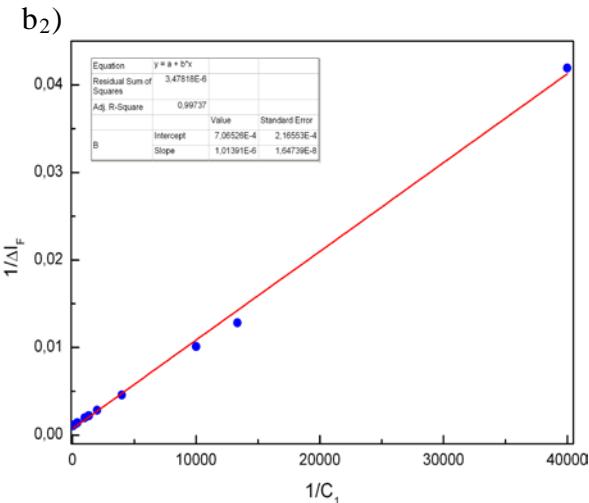
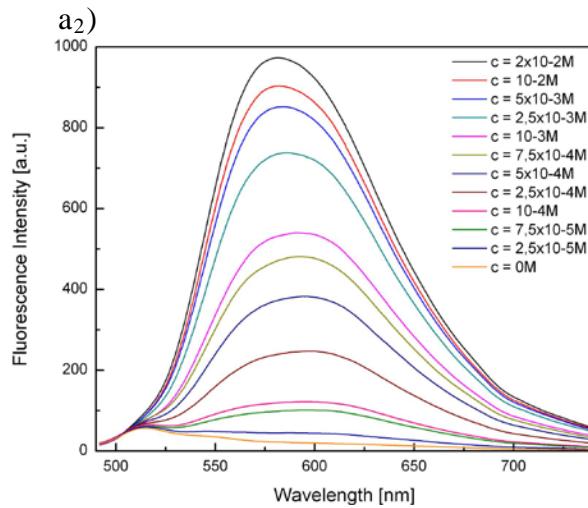
I_0 – Total concentration of ARS

Q – ratio of concentration of free ARS to complexed ARS

The binding constant of fructose-phenylboronic acid (K_{FRU}) can be calculated by dividing K_{ARS} by the slope of the plot equation:

$$\frac{[S_0]}{P} = \frac{K_{ARS}}{K_{FRU}} Q + 1 \text{ where } [S_0] - \text{Total concentration of fructose}$$





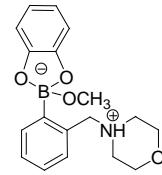
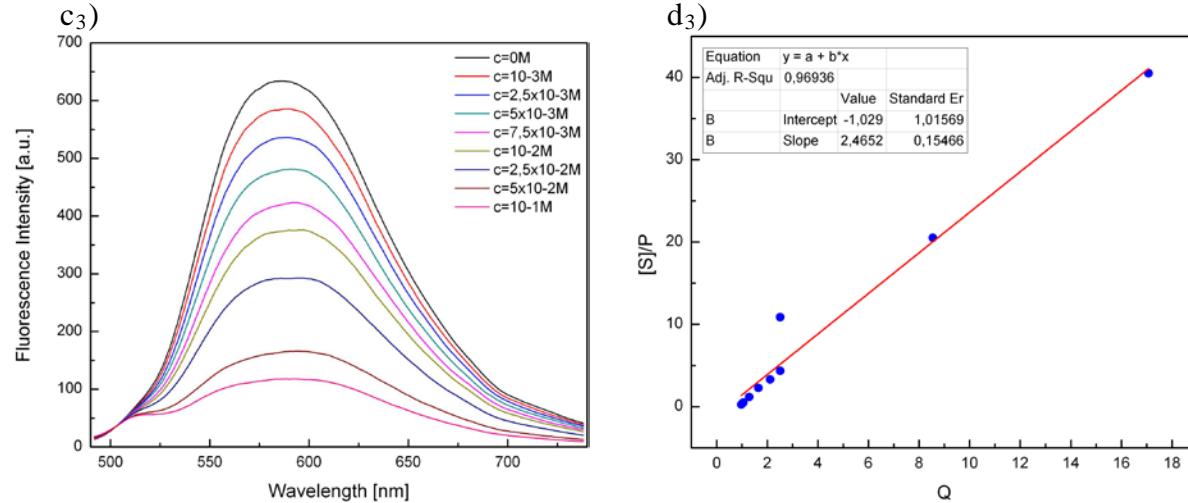


Table 3. Selected geometrical parameters for **5a**.

bond lengths, Å		bond angles, °	
O1–C12	1.3541(15)	C11–N1–C7	112.16(9)
C1–B1	1.6230(17)	C11–N1–C8	109.83(9)
O1–B1	1.5152(15)	C8–N1–C7	112.16(9)
O3–C10	1.4259(15)	C6–C1–B1	120.08(10)
O3–C9	1.4243(15)	C2–C1–B1	122.68(10)
O4–B1	1.4580(15)	O1–B1–C1	109.74(9)
O4–C18	1.4306(15)	O4–B1–O1	111.77(10)
O2–C13	1.3572(15)	O4–B1–O2	110.21(9)
O2–B1	1.5102(15)	O4–B1–C1	109.55(9)
N1–C11	1.4940(15)	O2–B1–O1	103.20(9)
N1–C7	1.5094(15)	O2–B1–C1	112.27(9)
N1–C8	1.4969(15)	N1–C7–C2	110.97(9)
torsion angles, °			
C2–C1–B1–O4	-59.21(14)	C2–C1–B1–O2	178.00(10)
C1–C2–C7–N1	70.77(13)	C11–N1–C7–C2	68.01(12)
C11–N1–C8–C9	-55.25(13)	C7–N1–C8–C9	179.30(9)

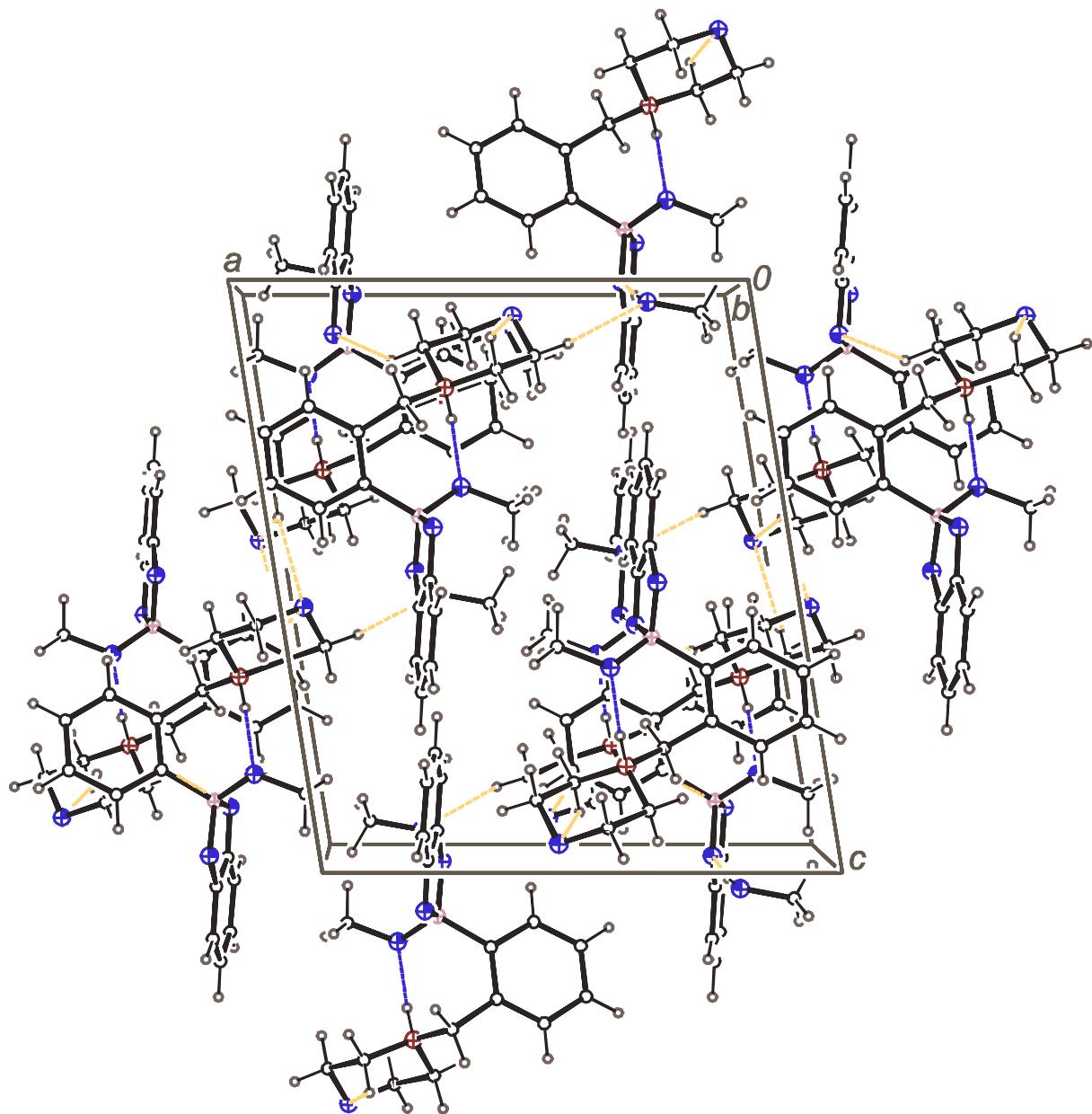
Table 4. Geometry of intra- and intermolecular interactions in **5a** (\AA , $^\circ$).

	H...A	D...A	D–H...A
N1–H1...O4	1.75(2)	2.6250(13)	157(2)
O5–H5A...O2 ⁱ	1.97(2)	2.7940(14)	171(2)
C8–H8B...O3 ⁱⁱ	2.46	3.3787(15)	154
C11–H11B...O1 ⁱⁱⁱ	2.53	3.3267(15)	138
C17–H17...O5	2.56	3.3863(17)	145
C9–H9A...O5 ^{iv}	2.57	3.2996(17)	131
C10–H10A...Cg1 ^{iii a}	2.55	3.3643(13)	139
C18–H18B...Cg1	2.62	2.8736(14)	95
C18–H18A...Cg2 ^v	2.80	3.4091(14)	121
C5–H5...Cg2 ^{vi}	2.96	3.8364(13)	154

Symmetry codes: (i) $x, -1+y, z$; (ii) $1-x, 2-y, 2-z$; (iii) $1/2-x, 1/2+y, 3/2-z$; (iv) $1/2+x, 3/2-y, 1/2+z$; (v) $1-x, 2-y, 1-z$; (vi) $-x, 2-y, 1-z$.

^aCg1 and Cg2 denote the gravity centers of the phenyl and catechol ring, respectively.

Figure 3. Packing diagram⁴ of **5a** (view along [010] direction). The intermolecular hydrogen bonds are denoted with dashed lines.



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