

***Supporting Information for***

**On the selective detection of duplex deoxyribonucleic acids by 2,1,3-benzothiadiazoles fluorophores**

Brenno A. D. Neto,<sup>\*a,c</sup> Alexandre A. M. Lapis,<sup>b,c</sup> Fabiana S. Mancilha,<sup>c</sup> Eraldo Luiz Batista Jr.,<sup>d</sup> Paulo A. Netz,<sup>c</sup> Frank Rominger,<sup>e</sup> Luiz A. Basso,<sup>d</sup> Diógenes S. Santos<sup>d</sup> and Jairton Dupont<sup>\*c</sup>

jairton.dupont@ufrgs.br, brenno.ipi@gmail.com

<sup>a</sup> Laboratory of Medicinal and Technological Chemistry, University of Brasília (IQ-UnB), Brasilia, DF, Brazil. <sup>b</sup> Universidade Federal do Pampa, Unipampa, Bagé, RS, Brazil <sup>c</sup> Laboratory of Molecular Catalysis, IQ-UFRGS, Porto Alegre, RS, Brazil. <sup>d</sup> Centro de Pesquisas em Biologia Molecular e Funcional (CPBMF), Tecnopuc, PUCRS, Brazil. <sup>e</sup> Organisch-Chemisches Institut der Ruprecht-Karls-Universität Heidelberg, Im Neuenheimer Feld 270, D-69120 Heidelberg, Germany.

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Cyclic voltammograms of compounds **3a-c**.

**Table S1.** Selected bond angles and distances from x-ray data of compounds **3a** and **3b**.

selected atoms	d (Å) <b>3a</b>	d (Å) <b>3b</b> <sup>1</sup>	selected atoms	angle (deg) <b>3a</b>	angle (deg) <b>3b</b> <sup>1</sup>
C21-C1	1.439(2)	1.431(3)	N1-S1-N2		
C1-C2	1.197(2)	1.200(3)	C7-N1-S1	101.20(7)	
C2-C3	1.427(2)	1.424(3)	C8-N2-S1	106.61(10)	101.03(9)
C3-C4	1.379(2)	1.371(3)	N1-C7-C8	106.00(11)	107.25(15)
C4-C5	1.414(2)	1.414(3)	N2-C8-C7	112.65(13)	106.26(13)
C5-C6	1.381(2)	1.377(3)	C21-C1-C2	113.53(14)	111.55(17)
C6-C7	1.438(2)	1.440(3)	C1-C2-C3	172.16(17)	113.90(18)
C7-C8	1.437(2)	1.438(3)		176.85(18)	174.90(2)
C8-C3	1.430(2)	1.424(3)			179.40(3)
N1-C7	1.347(2)	1.356(2)			
N2-C8	1.347(2)	1.348(3)			
S1-N1	1.612(1)	1.609(2)			
S1-N2	1.615(1)	1.607(2)			

<sup>1</sup> mean values of three independent molecules

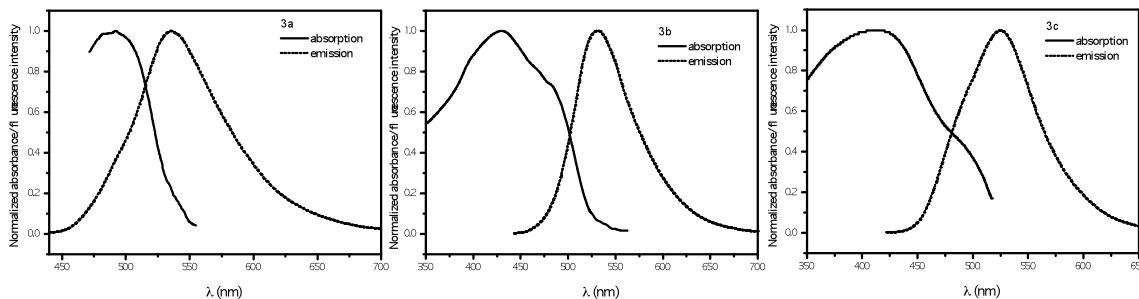
**Table S2.** Summary of the crystal data and structure refinement for **3a** and **3b**.

Parameter	<b>3a</b>	<b>3b</b>
Empirical formula	C <sub>20</sub> H <sub>13</sub> N <sub>3</sub> OS	C <sub>20</sub> H <sub>13</sub> N <sub>3</sub> OS
Formula weight	343.39	343.39
Temperature	200(2) K	200(2) K
Wavelength	0.71073 Å	0.71073 Å
Crystal system	triclinic	triclinic
Space group	p $\bar{1}$	p $\bar{1}$
Z	2	6
Unit cell dimensions		
a (Å)		
b (Å)	8.963(1)	7.4910(1)
c (Å)	9.048(1)	16.4275(3)
$\alpha$ (deg)	10.465(1)	21.0326(3)
	111.391(2)	107.378(1)
$\beta$ (deg)	90.220(2)	96.119(1)
	91.569(2)	97.804(1)
$\gamma$ (deg)		
Volume	789.76(16) Å <sup>3</sup>	2417.40(6) Å <sup>3</sup>
Density (calculated)	1.44 g/cm <sup>3</sup>	1.41 g/cm <sup>3</sup>
Absorption coefficient	0.22 mm <sup>-1</sup>	0.21 mm <sup>-1</sup>
Crystal shape	polyhedron	polyhedron
Crystal size	0.36 x 0.29 x 0.24 mm <sup>3</sup>	0.48 x 0.38 x 0.10 mm <sup>3</sup>
Crystal colour	yellow	yellow
Theta range	2.1 to 28.3 deg.	1.9 to 27.5 deg.
	-11 ≤ h ≤ 11	-9 ≤ h ≤ 9
Index ranges	-12 ≤ k ≤ 12	-21 ≤ k ≤ 21
	-13 ≤ l ≤ 13	-27 ≤ l ≤ 27
Reflections collected	8197	24828
Independent reflections	3875 (R(int) = 0.0193)	10995 (R(int) = 0.0685)
Observed reflections	3307 (I > 2σ(I))	6623 (I > 2σ(I))
Absorption correction	Semi-empirical from equivalents	Semi-empirical from equivalents
Max./min. transmission	0.95 / 0.93	0.98 / 0.90
Data/restraints/parameters	3875 / 0 / 227	10995 / 0 / 679
Goodness-of-fit on F <sup>2</sup>	1.05	1.01
Final R indices (I > 2σ(I))	R1 = 0.042, wR2 = 0.106	R1 = 0.051, wR2 = 0.120
Largest diff. peak and hole	0.34 and -0.26 eÅ <sup>-3</sup>	0.25 and -0.39 eÅ <sup>-3</sup>

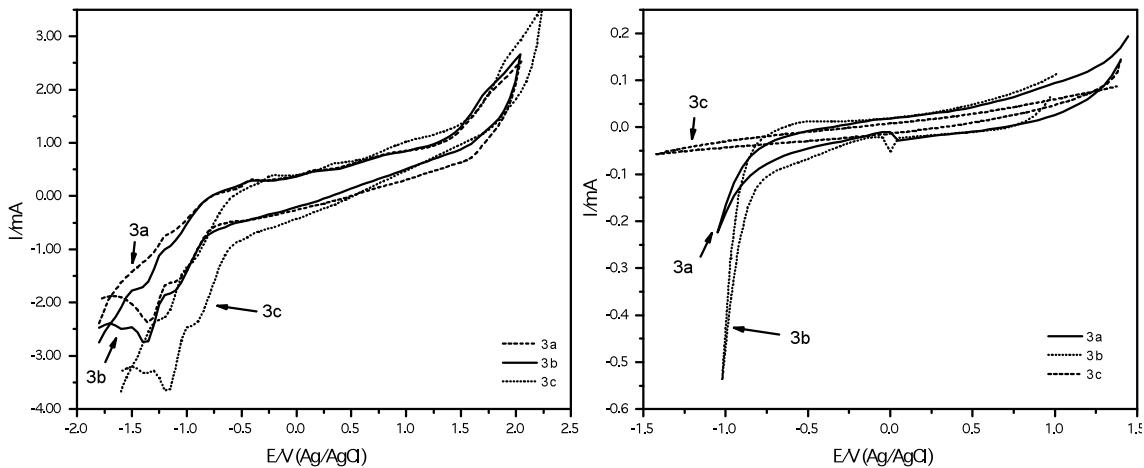
**Table S3.** Spectrometric data of all tested compounds **1a-c**, **2a-b**, **3a-c**.

Dye	Log $\epsilon$ ( $\epsilon$ )	$\lambda_{\text{abs}}^{\text{max}}$ (nm) <sup>a</sup>	$\lambda_{\text{em}}^{\text{max}}$ (nm) <sup>a</sup>	Stokes Shift (nm)	$\lambda_{\text{abs}}^{\text{max}}$ (nm) <sup>b</sup>	$\lambda_{\text{em}}^{\text{max}}$ (nm) <sup>b</sup>	Stokes Shift (nm) <sup>b</sup>	$\Phi_f$	$E_{\text{gap}}^{\text{op}}$ (eV) <sup>d</sup>
<b>1a</b>	4.35 (22342)	365	471	106	452	543	91	0.86	2.81
<b>1b</b>	3.76 (5790)	429	563	134	522	590	68	0.29	2.29
<b>1c</b>	3.81 (6482)	438	552	114	424	543	119	0.37	1.90
<b>2a</b>	3.22 (1644)	411	535	124	418	526	108	0.51	1.85
<b>2b</b>	3.33 (2150)	367	506	139	426	493	67	0.80	2.28
<b>3a</b>	3.97 (9152)	444	544	100	491	535	44	0.40	2.21
<b>3b</b>	3.78 (6076)	401	547	146	430	531	101	0.44	2.20
<b>3c</b>	4.01 (10208)	426	525	99	413	525	112	0.47	2.22

<sup>a</sup> In solution (phosphate buffer 10  $\mu$ M, pH = 7.0). <sup>b</sup> Solid state. <sup>c</sup> Quantum yield of fluorescence in MeCN (quinine sulfate (Riedel) in 1 M H<sub>2</sub>SO<sub>4</sub>,  $\Phi_f$  = 0.55, as standard. <sup>d</sup> Determined in phosphate buffer 100 mM (pH = 7.0).



**Figure S1.** Solid-state (normalized) absorption and emission of dyes **3a-c**.



**Figure S2.** CV of compounds **3a-c** (1.00 mM) dissolved in a 0.10 M solution TBAPF<sub>6</sub> in MeCN recorded at a scan rate of 200 mV/s (left) and CV (thin film coated onto a Pt wire electrode) of compounds **3a-c** recorded at a scan rate of 40 mV/s in a 0.10 M solution of TBAPF<sub>6</sub> in MeCN (right).

All compounds exhibited a very similar electrochemical behavior and it was possible to observe quasi-reversible processes. Compounds **3a-c** presented a large electrochemical window, *i.e.*, from -2.0 V to 2.0 V (Ag/AgCl) for the anodic and cathodic potential sweeps, respectively. For **3a**, the related values were 0.89 V (and -0.34 V), 0.18 V (and -1.09 V) and -0.41 V (related to -1.35 V). For **3b**, we observed 0.76 V (and -0.35 V), 0.16 V (and -1.12 V) and -0.41 V (related to -1.38 V), as for **3b** at 0.76 V (and -0.35 V), 0.16 V (and -1.12 V) and -0.41 V (related to -1.38 V). In the case of **3c**, an anodic shift in the CV and the related observed values were 0.98 V (and -0.27 V), 0.38 V (and -0.91 V), and -0.20 V (related to -1.19 V). In the solid state, the dyes presented a good electrochemical stability. In the electrochemical window tested (-1.5-1.5 V), a well-defined reduction or oxidation peak was not observed, indicating a larger electrochemical window and stability than the limits of the solvent oxidation and reduction windows.

**Table S4.** *Ab initio* calculation of the HOMO-LUMO values for tested dyes and bases.

Structure		Hartree	eV	kJ/mol
<b>3a</b>	HOMO	-0.279	-7.59198	-732.515
	LUMO	0.03545	0.964644	93.07398
	GAP	0.31445	8.556625	825.5885
<b>3b</b>	HOMO	-0.27971	-7.6113	-734.379
	LUMO	0.03292	0.895799	86.43146
	GAP	0.31263	8.5071	820.8101
<b>3c</b>	HOMO	-0.27582	-7.50545	-724.165
	LUMO	0.03656	0.994849	95.98828
	GAP	0.31238	8.500297	820.1537
<b>3a'</b>	HOMO	-0.28316	-7.70518	-743.437
	LUMO	0.04817	1.310773	126.4703
	GAP	0.33133	9.015953	869.9069
<b>3b'</b>	HOMO	-0.28439	-7.73865	-746.666
	LUMO	0.04011	1.091449	105.3088
	GAP	0.32450	8.830099	851.9748
<b>3c'</b>	HOMO	-0.28057	-7.6347	-736.637
	LUMO	0.04514	1.228323	118.5151
	GAP	0.32571	8.863025	855.1516
<b>Adenine</b>	HOMO	-0.29852	-8.12315	-783.764
	LUMO	0.14434	3.927693	378.9647
	GAP	0.44286	12.05084	1162.729
<b>Cytosine</b>	HOMO	-0.32146	-8.74738	-843.993
	LUMO	0.13235	3.601429	347.4849
	GAP	0.45381	12.34881	1191.478
<b>Guanine</b>	HOMO	-0.28788	-7.83362	-755.829
	LUMO	0.15706	4.273822	412.361
	GAP	0.44494	12.10744	1168.19
<b>Thymine</b>	HOMO	-0.33245	-9.04643	-872.847
	LUMO	0.12834	3.492311	336.9567
	GAP	0.46079	12.53874	1209.804