

1 Article

# 2 Polyetheroaryl oxadiazole/pyridine-based ligands: a 3 structural tuning for enhancing G-quadruplex 4 binding

5 Filippo Doria,<sup>1</sup> Valentina Pirola,<sup>1</sup> Michele Petenzi,<sup>1</sup> Marie-Paule Teulade-Fichou,<sup>2,3</sup> Daniela  
6 Verga<sup>2,3\*</sup> and Mauro Freccero <sup>1,\*</sup>

7 <sup>1</sup> Dipartimento di Chimica, Università di Pavia, Pavia, 27100, Italy; filippo.doria@unipv.it;  
8 valentina.pirola01@universitadipavia.it, michele.petenz@gmail.com.

9 <sup>2</sup> CNRS UMR9187, INSERM U1196 Institut Curie, PSL Research University, F-91405, Orsay (France);  
10 mp.teulade-fichou@curie.fr

11 <sup>3</sup> CNRS UMR9187, INSERM U1196, Université Paris Sud, Université Paris-Saclay, F-91405, Orsay (France);  
12 mp.teulade-fichou@curie.fr

13 \* Correspondence: daniela.verga@curie.fr (D.V.); mauro.freccero@unipv.it (M.F.);  
14 Tel.: +39-0382-987668 (M.F.)

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16 **Abstract:** Acyclic oligoheteroaryl-based compounds represent a valuable class of ligands for nucleic  
17 acid recognition. In this regard, acyclic pyridyl polyoxazoles and polyoxadiazoles were recently  
18 identified as selective G-quadruplex stabilizing compounds with high cytotoxicity and promising  
19 anticancer activity. Herein, we describe the synthesis of a new family of polyheteroaryl  
20 oxadiazole/pyridine-ligands targeting DNA G-quadruplexes. In order to perform a structure-  
21 activity analysis identifying determinants of activity and selectivity, we followed a convergent  
22 synthetic pathway to modulate the nature and number of the heterocycles (1,3-oxazole *vs* 1,2,4-  
23 oxadiazole and pyridine *vs* benzene). Each ligand was evaluated towards secondary nucleic acid  
24 structures, which have been chosen as a prototype to mimic cancer-associated G-quadruplex  
25 structures (e.g., the human telomeric sequence, c-myc and c-kit promoters). Interesting,  
26 heptapyridyl-oxadiazole compounds showed preferential binding towards the telomeric sequence  
27 (22AG) in competitive conditions *vs* duplex DNA. In addition, G4-FID assays suggest a different  
28 binding mode from the classical stacking on the external G-quartet. Additionally, CD titrations in  
29 the presence of the two most promising compounds for affinity, TOxAzaPy and TOxAzaPhen,  
30 display a structural transition of 22AG in K-rich buffer. This investigation suggests that the pyridyl-  
31 oxadiazole motif is a promising recognition element for G-quadruplexes, combining seven  
32 heteroaryls in a single binding unit.

33 **Keywords:** G-quadruplex; oxadiazole/pyridine polyheteroaryls; G4-ligands; FRET-melting; G4-  
34 FID; circular dichroism

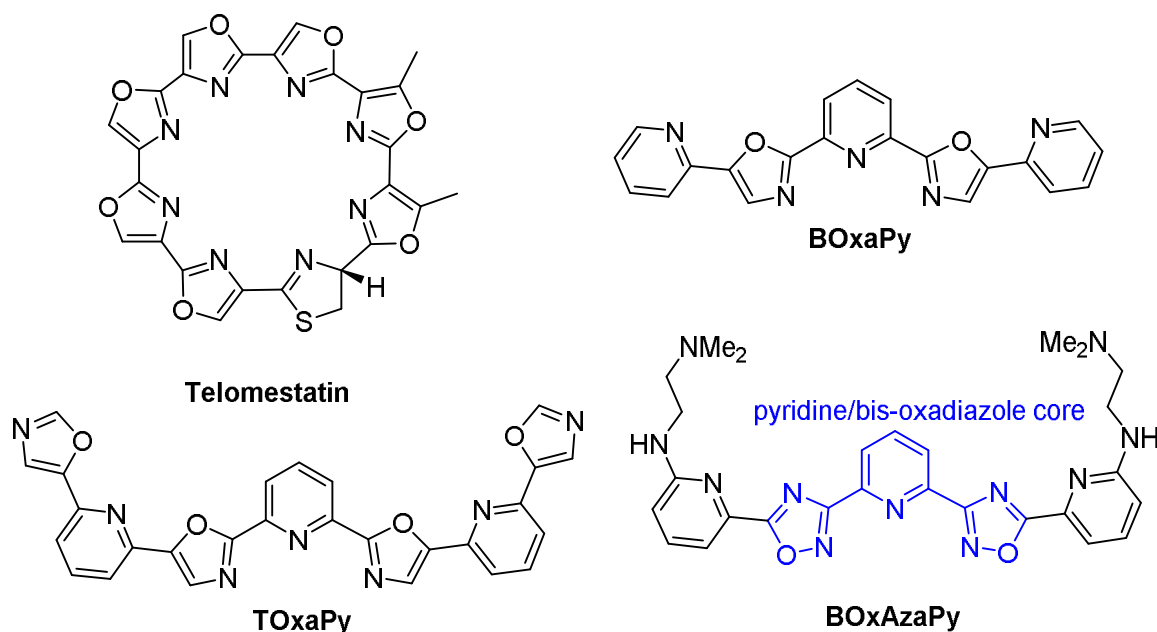
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## 36 1. Introduction

37 G-quadruplex DNAs (G4s) are tetra-helical structures arising in nucleic acid sequences  
38 containing repeats of runs of three or four adjacent guanines that fold over due to self-assembly of  
39 four guanine bases in square co-planar arrays (G-tetrads) through Hoogsteen hydrogen bonding [1].  
40 Several G-tetrads, upon K<sup>+</sup> coordination [2,3] can stack on each other to form a G4, with the  
41 interconnecting sequences extruded as single-strand loops of various length [4,5]. It is well  
42 documented that these structures exist in a conformational equilibrium with the single stranded DNA  
43 domains transiently generated during key biological processes, such as telomere maintenance,  
44 recombination, replication, transcription and epigenetic regulation [6], together with genome

45 instability [7,8]. Therefore, G4s have been the object of intense study with the aim of defining their  
46 potential as regulatory elements and/or therapeutic targets. Thus, G4s are considered druggable  
47 targets that offer the possibility to control these fundamental processes in cells [9]. Organic  
48 compounds capable of recognising and stabilizing G4 structures have attracted great attention as  
49 selective probes and anticancer agents since they are assumed to act selectively at genomic G-rich loci  
50 such as telomeres and oncogene promoters [10], which have higher propensity to fold into G-  
51 quadruplex structures than other putative quadruplex sequences (PQs) within the human genome.  
52 Dual selectivity towards G4 versus *ds* DNA and different G4 architectures is a key aspect for  
53 developing efficient G4 DNA binding drugs and diagnostic tools. Nowadays, we are far from being  
54 able to identify G4 ligands binding selectively a given G4 structure. Nevertheless, small molecules  
55 able to induce down-regulation of transcription of specific oncogenes, due to the targeting of G4  
56 structures localized within gene promoters, are reported in literature. In this context, the  
57 downregulation of c-kit by benzo[*a*]phenoxazines [11], c-myc by the ellipticin analogue GQC-05 [12],  
58 and HSP90 by phenyl bis-oxazole derivatives [13] are paradigmatic examples.

59 A massive synthetic effort has been focused on the development of potent and selective G4  
60 ligands, the main class of compounds being represented by extended planar aromatic compounds  
61 able to interact by  $\pi$ -stacking with the external quartets. Often to increase binding affinity and water  
62 solubility, these ligands are functionalized with cationic moieties or with lateral chains that can be  
63 protonated in physiological conditions [14-17]. However at present, a limited number of binders  
64 simulates the peculiar characteristics of the natural macrocycle telomestatin [18] (Figure 1), which  
65 exhibits excellent binding properties and remarkable antitumor activity. Its unique molecular  
66 features inspired the design and synthesis of a large number of both cyclic [19,20] and acyclic  
67 polyheterocyclic ligands (Scheme 1) [13,21-25]. Among them, the acyclic binders with high rotational  
68 flexibility are rather underrepresented as a class of G-quadruplex ligands. However, the high  
69 rotational flexibility of these oligoaryl systems should arise the number of possible binding modes  
70 occurring between the ligand and the G4 structures, from classical  $\pi$ -stacking to grove binding or  
71 hybrid interactions (Figure 1). The initial structure-activity investigation reported for these  
72 polyaromatic binders focused on the study of the effect of ligand length on target affinity, suggested  
73 that the neutral symmetric (BOxaPy and TOxaPy, Figure 1) and asymmetric pyridyl-oxazole motif,  
74 particularly when consisting of a minimum of five alternating units, serves as a potent binding  
75 element for G-quadruplexes recognition[21,23,24].



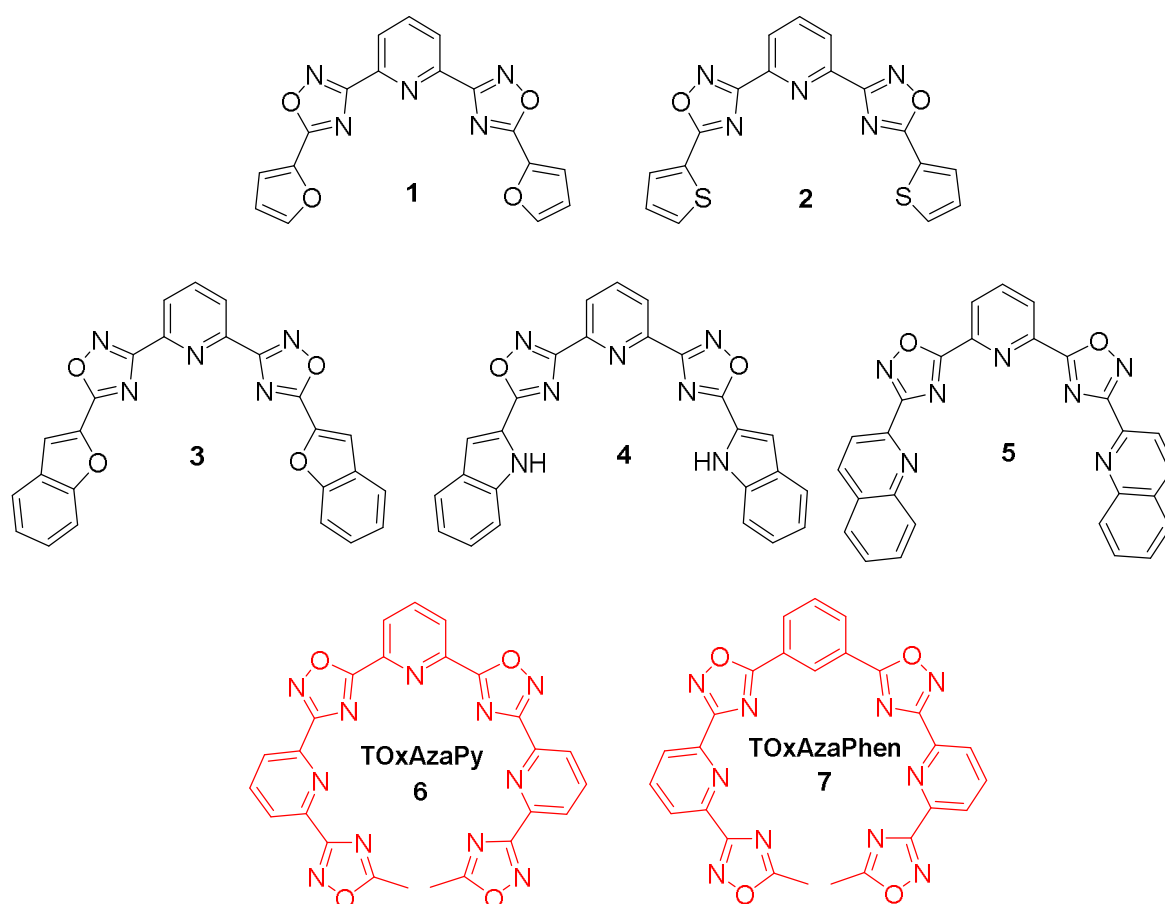
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**Figure 1.** Structure of Telomestatin, together with acyclic and flexible oligoaromatics previously evaluated as G4[21,22].

79 Although these pentaaromatic compounds are modest stabilizers of pre-folded G-quadruplex  
 80 structures, they could be further optimized by exploiting a synthetic strategy that allows the  
 81 functionalization of the structures with amino/cationic side-chains (BOxAzaPy, Figure 1) [22]. These  
 82 promising results prompted us to develop a pool of conformationally flexible ligands (1-7, Figure 2),  
 83 allowing adaptability to the G-quadruplex DNA target. In particular, starting from the “lead”  
 84 pyridyl–oxadiazole central core, we focused our attention on the number of aromatic rings and the  
 85 introduction of different heteroaryl units, with 1,2,4-oxadiazole being the ever present structural  
 86 moiety. Herein, we describe the synthesis of a small library of these oligo-hetero-aromatics also  
 87 outlining the preliminary biophysical data: FRET melting and G4-FID assays collected in the presence  
 88 of prototype G4s chosen to mimic cancer-associated G4s. The effect produced by these structural  
 89 changes on G4 selectivity and stabilization has been evaluated and the most potent ligands were then  
 90 further investigated by circular dichroism (CD) titration to appraise their ability to induce  
 91 conformational changes.



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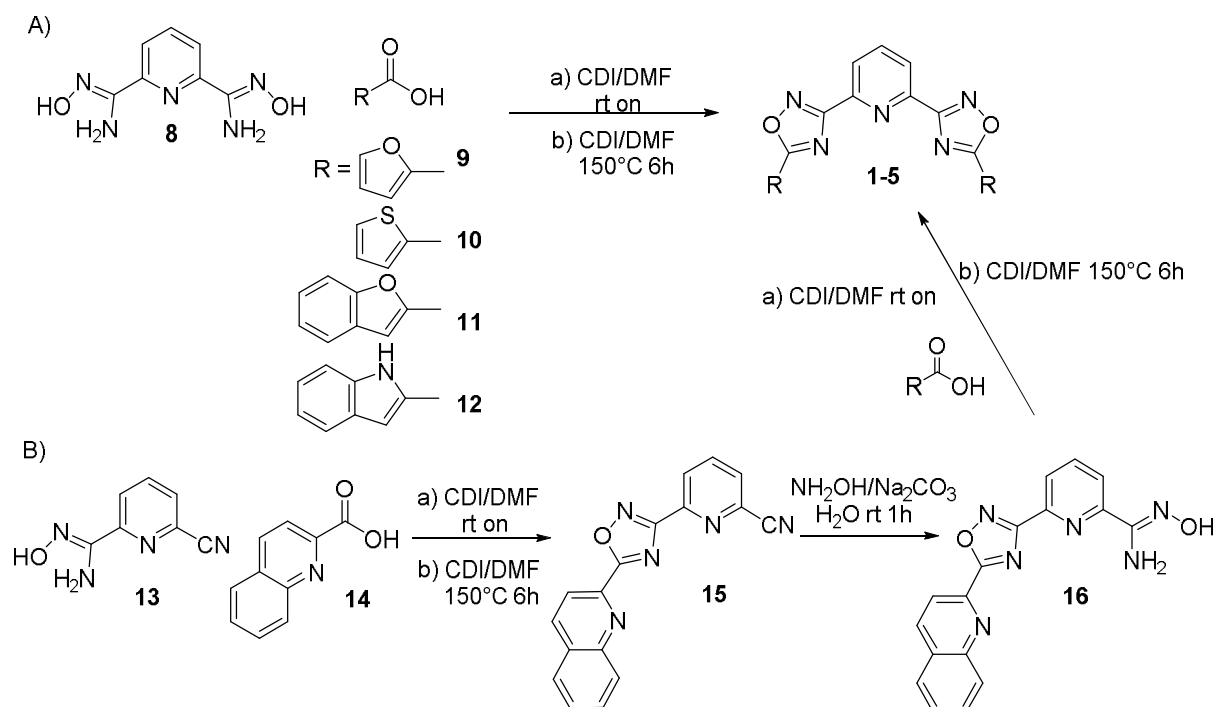
93 **Figure 2.** Structures of the newly synthesized acyclic penta-hetero-aromatics 1-5 and hepta-hetero-  
 94 aromatics TOxazaPy (6) and TOxazaPhen (7).

## 95 2. Results and Discussion

### 96 2.1. Design and Synthesis of the polyheteroaryl oxadiazole/pyridine-ligands

97 In order to implement the potency and selectivity of the BOxazaPy G4 ligand and assess the  
 98 effect of the nature of the aromatic rings composing the ligand on G4 binding, we introduced different  
 99 heterocycles directly bound to the central pyridine/bis-oxadiazole core (Figure 2), performing in a  
 100 single step the synthesis of both oxadiazoles (Pathway A, Scheme 1). The synthesis is a one-pot  
 101 process constituted by an initial O-acylation followed by a cycloaromatization. The condensation  
 102 between pyridine-2,6-bis-amidoxime 8 and a small set of carboxylic acids (9-12, 14, Scheme 1)  
 103 occurred in the presence of 1,1'-carbonyldiimidazole (CDI) as a coupling reagent. The subsequent

104 cycloaromatization was achieved without isolation of the resulting intermediate O-  
 105 acylbenzamidoxime, generating the final BOxAzaPys **1-4** in low yields (Table 1). However, this one-  
 106 pot protocol yielded the quinolinium oxadiazole **5** only as by-product. Therefore, the synthesis of the  
 107 latter was more effectively optimised by a stepwise process, introducing the oxadiazole rings one at  
 108 the time (Pathway B, scheme 1). The (Z)-6-cyano-N'-hydroxypicolinimidamide **13** was initially O-  
 109 acylated by carboxylic acids **14**, and the resulting intermediate cycloaromatized to the cyano-  
 110 oxadiazole **15** (Scheme 1). The cyano group was subsequently converted into amidoxime **16** in the  
 111 presence of hydroxylamine, which yielded the final product **5** after O-acylation and  
 112 cycloaromatization, as described above.



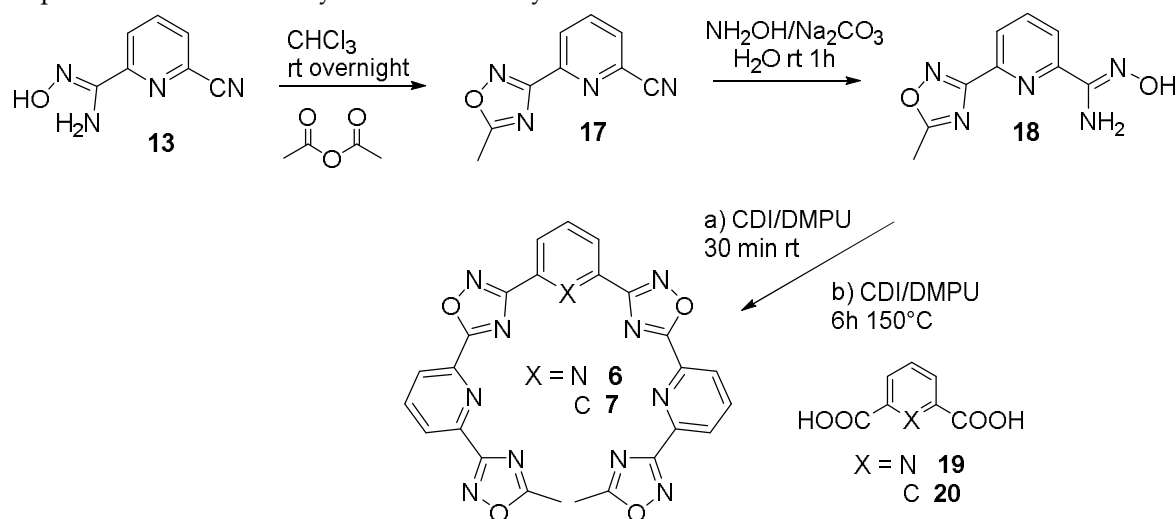
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 114 **Scheme 1.** Synthetic protocol developed for the preparation of the penta-heterocyclic compounds: (A)  
 115 One-step synthetic pathway followed for the preparation of compounds **1-4**, and (B) Multi-step  
 116 synthetic pathway for the preparation of compound **5**.

117 **Table 1.** Penta-heterocyclic compounds.

R	Pathway	Compound	Yield (%)
	A	<b>1</b>	11
	A	<b>2</b>	11
	A	<b>3</b>	28
	A	<b>4</b>	10
	B	<b>5</b>	53

118  
 119 TOxAzaPy (**6**) and TOxAzaPhen (**7**) were synthesized according to the stepwise protocol shown  
 120 in Scheme 2. The picolin-cyano-monoamidoxime **13** was converted into oxadiazole **17** exploiting a

121 one-pot, two-step synthesis. **13** was firstly O-acylated by acetic anhydride and then cycloaromatized  
 122 in acetic acid to give **17**. Hydroxylamine addition in aqueous ethanol, yielded the formation and the  
 123 precipitation of the amidoxime **18** as white crystals, directly employed for the last step. The  
 124 dicarboxylic acid **19** or **20** was activated in order to perform in a single step the O-acylation and the  
 125 cycloaromatization to obtain the final oxadiazoles **6** and **7**. A change in solvent, from DMF to DMPU,  
 126 implemented the reaction yield of the final cyclization reaction.



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129 **Scheme 2.** Multistep synthetic protocol developed for the preparation of the heptaheterocyclic  
 130 compounds TOxAzaPy and TOxAzaPhen (**6-7**).

131 **Table 2.** Overall reaction yields of the heptaheterocyclic compounds

X	Compound	Yield (%)
N	<b>6</b>	19
C	<b>7</b>	38

132

## 133 2.2. Screening of the ligands

134 The ability of the oligoheteroaryl oxadiazole/pyridine-compounds **1-7** (Figure 2) to bind and  
 135 stabilize G-quadruplex structures was evaluated by performing two well-known biophysical assays,  
 136 namely FRET melting and G4-FID assay.

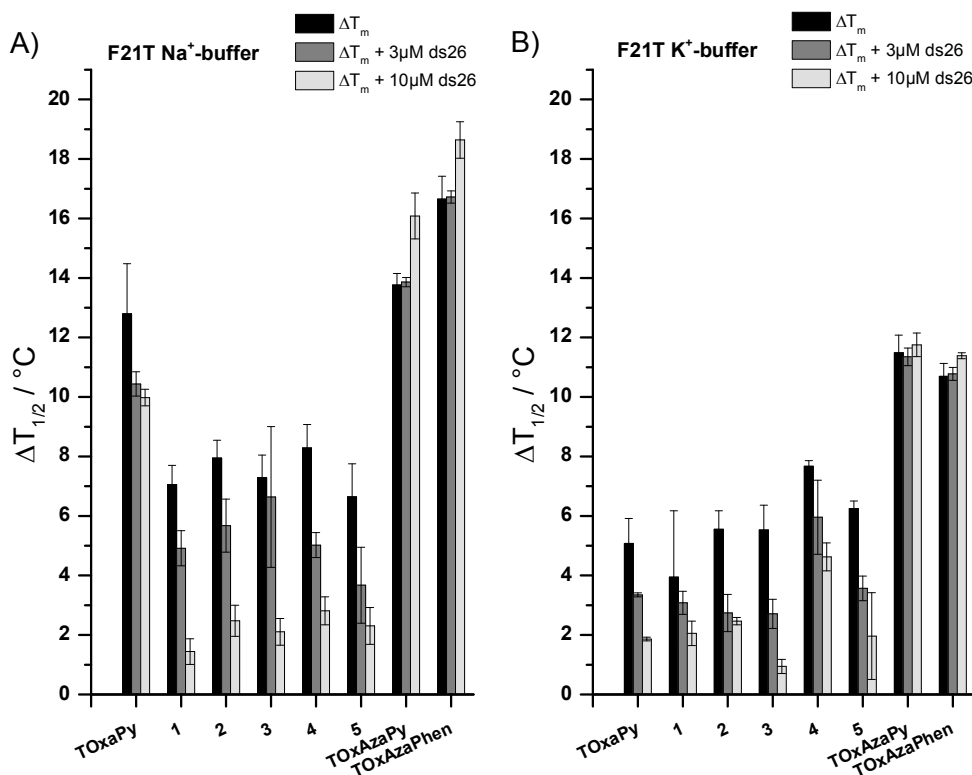
### 137 2.1.1. FRET-melting

138 The G4 stabilizing properties of the compounds **1-7** were evaluated by FRET (Förster resonance  
 139 energy transfer) melting assay [26]. This assay employs fluorescently doubly labelled (FAM and  
 140 TAMRA dyes as FRET partners) single stranded oligonucleotides mimicking G4 structures. FRET  
 141 melting provides an assessment of the stabilization effect produced by the binding of the ligand on a  
 142 G4 structure by measuring the change in the melting temperature ( $\Delta T_m$ ). In detail, the stabilization is  
 143 measure as a difference of melting temperature in the presence and in the absence of ligand at the  
 144 inflection point of the melting curve. First, the experiment was carried out in the presence of F21T  
 145 (0.2  $\mu$ M), a 21-mer oligonucleotide mimicking the human telomeric sequence, in both Na<sup>+</sup>- and K<sup>+</sup>-  
 146 rich buffer and the polyheteroaryl oxadiazole/pyridine-compounds **1-7** at the fixed concentration of  
 147 1  $\mu$ M.

148 All the newly synthesized compounds were able to stabilize F21T and the stabilization was  
 149 higher in Na<sup>+</sup>- than in K<sup>+</sup>-rich buffer. The  $\Delta T_m$  values of F21T for the pentaryl compounds **1-5** ranged  
 150 from 6.6 to 8.3 °C in Na<sup>+</sup>-rich buffer and from 3.9 to 7.7 °C in K<sup>+</sup>-rich buffer (Figure 3). Following the  
 151 same trend, the heptameric ligands TOxAzaPy and TOxAzaPhen showed much higher stabilization

152 ability than the pentameric counterparts, with  $\Delta T_m$  values ranging from 13.8 to 16.6 °C in Na<sup>+</sup>-rich  
 153 buffer (Figure 3A) and from 10.8 to 11.5 °C in K<sup>+</sup>-rich buffer (Figure 3B), respectively. This trend of  
 154 higher stabilization in Na<sup>+</sup>-rich buffer was previously observed for the heptyl TOxPy [21], used as  
 155 a reference compound in this study, although in this specific case the differences in stabilization in  
 156 the two buffer conditions are less pronounced. As suggested for TOxPy, these results point out the  
 157 unusual behavior of these heteroaryl compounds to preferentially recognize and stabilize the  
 158 antiparallel form of the telomeric sequence (predominantly formed in Na<sup>+</sup>-buffer) more than the [3+1]  
 159 hybrid conformation adopted in K<sup>+</sup>-rich buffer. Based on these results, we decided to analyze our  
 160 compounds with additional FRET melting studies in the presence of the highly polymorphic  
 161 telomeric sequence.

162 To check the selectivity of these ligands towards G4 structures *vs* ds DNA, competition FRET  
 163 melting experiments were carried out in the presence of a large excess of a not-fluorescently labeled  
 164 double stranded DNA (ds26). This experiment was performed in the classical conditions for FRET  
 165 melting in the presence of F21T (0.2  $\mu$ M), ligands (1  $\mu$ M), and ds26 (either at 3  $\mu$ M or 10  $\mu$ M) in Na<sup>+</sup>-  
 166 or K<sup>+</sup>-rich buffer respectively (Figure 3). The results showed that penta-aryl compounds 1-5 were  
 167 strongly affected by the presence of the competitor in both buffer conditions. In particular, in the  
 168 presence of 10  $\mu$ M ds26 (50 mol. eq. competitor) the stabilizing effect towards F21T was almost erased.  
 169 Differently, the ability to stabilize G4 structure by heptyl TOxazaPy and TOxazaPhen was not  
 170 affected by the presence of a large excess of ds-competitor, making them highly selective G4 ligands.  
 171 From the point of view of the selectivity, both the newly made neutral ligand TOxazaPy and  
 172 TOxazaPhen appeared to be much superior than TOxPy both in Na<sup>+</sup>- and K<sup>+</sup>-rich buffered  
 173 conditions. These outcomes support the hypothesis that the minimal number of cycles constituting  
 174 the oligoheteroaryl ligands to achieve both high affinity and selectivity toward G4s is seven.  
 175 Additionally, a comparison between TOxPy *vs* TOxazaPy, TOxazaPhen melting data suggests that  
 176 the replacement of the 1,3-oxazoles by the 1,2,4-oxadiazole moiety positively affect both G4 affinity  
 177 and G4 *vs* ds selectivity.



178

179 **Figure 3.** Quantitative analysis of the FRET-melting competition experiments with the telomeric  
 180 sequence F21T (0.2  $\mu$ M). Stabilization in: A) Na<sup>+</sup>-rich buffer, or B) K<sup>+</sup>-rich buffer is indicated for the  
 181 reference compound TOxPy, in the absence (black bars) or in the presence of double-stranded DNA

182 (ds26) at 3  $\mu\text{M}$  (dark grey bars) or 10  $\mu\text{M}$  (fair grey bars). Analysed compounds: 1-5, TOxAzaPy (6)  
 183 and TOxazaPhen (7) (1  $\mu\text{M}$ ). Error bars corresponds to SD of three independent experiments.  
 184 Experiments were performed in A) 10 mM lithium cacodylate buffer (pH 7.2), 100 mM NaCl and B)  
 185 10 mM lithium cacodylate buffer (pH 7.2), 90 mM LiCl, and 10 mM KCl.

186 The stabilization effect of the selective heptaryls TOxAzaPy and TOxazaPhen was challenged  
 187 towards other G4-forming sequences able to form G-quadruplex structures with different topologies  
 188 and among the parallel conformations several sequences with different central loop sizes were  
 189 analyzed. Besides F21T, we selected six additional sequences suitable for FRET melting experiments:  
 190 parallel forming sequences such as c-myc oncogene promoter (FMycT) [5], c-kit2 oncogene promoter  
 191 (Fkit2T) [27], human minisatellite repeats – native sequence CEB25wt (FCEB25wtT) [28], modified  
 192 sequence CEB25L111T (FCEB25L111TT) [29], Bcl2 proto-oncogene promoter (FBcl2T) [30] and an  
 193 antiparallel forming sequence from the human telomeric sequence variant 22CTA (F21CTAT) [31].  
 194 FMycT sequence was chosen based on previous experiments performed on oxazol-heteroaryl [23,24].  
 195 The idea was to evaluate the effect of the nature of the heterocycle (1,3-oxazole *vs* 1,2,4-oxadiazole)  
 196 on the stabilization of this sequence. TOxAzaPy and TOxazaPhen show very low stabilization effect  
 197 towards this sequence,  $\Delta T_m$  values 1.9 and 3.9  $^{\circ}\text{C}$  respectively (Table 3, Figure 4). Similar values were  
 198 observed with the oxazole family [23,24] suggesting that these oligo-heterocycles are poor ligands for  
 199 c-myc. By observing the  $\Delta T_m$  values measured in the presence of the two heptaryl compounds (Table  
 200 3, Figure 4) it is possible to remark the striking preferential binding towards the human telomeric  
 201 sequence both  $\text{Na}^+$ - and  $\text{K}^+$ -buffer, the moderate stabilization of c-kit2 and 22CTA, and the total lack  
 202 of stabilization for all the other sequences. The striking difference of the binding toward F21T in  $\text{Na}^+$   
 203 *vs* F21CTAT, which share the same topology, by both TOxAzaPy and TOxazaPhen suggest that  
 204 TOxazaPhen and to lower extent TOxAzaPy are structure selective ligands for the antiparallel wild  
 205 type human telomeric sequence.

206 **Table 3.**  $\Delta T_m$  values for 6 and 7 (1  $\mu\text{M}$ ) towards several G4 structures.

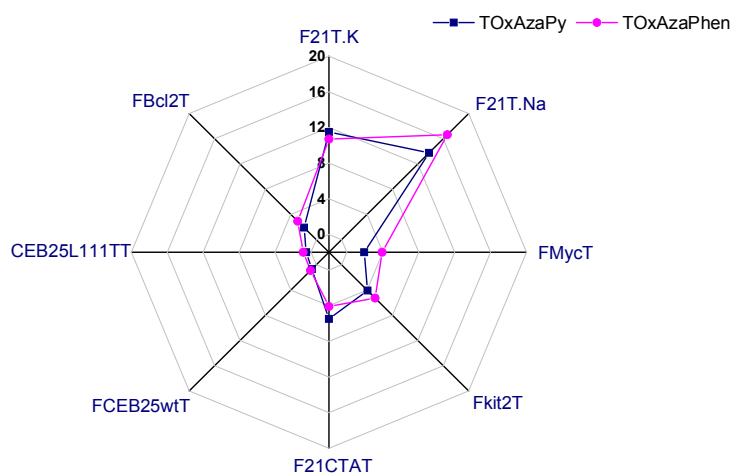
G4 sequences	Topology	TOxAzaPy $\Delta T_m$ ( $^{\circ}\text{C}$ )	TOxazaPhen $\Delta T_m$ ( $^{\circ}\text{C}$ )
F21T.Na <sup>+</sup>	Antiparallel [32]	13.8 $\pm$ 0.4	16.6 $\pm$ 0.8
F21T.K <sup>+</sup>	Hybrid/mix [32-35]	11.5 $\pm$ 0.6	10.8 $\pm$ 0.4
FMycT	parallel	1.9 $\pm$ 0.1	3.9 $\pm$ 0.9 <sup>1</sup>
Fkit2T	parallel	4.1 $\pm$ 0.3	5.3 $\pm$ 0.4 <sup>2</sup>
FCEB25wtT	parallel	0.7 $\pm$ 0.9	0.9 $\pm$ 0.7 <sup>1</sup>
FCEB25L111TT	parallel	0.5 $\pm$ 0.3	0.9 $\pm$ 0.8 <sup>1</sup>
FBcl2T	parallel	1.9 $\pm$ 0.7	2.9 $\pm$ 0.7 <sup>2</sup>
F21CTAT	antiparallel	5.5 $\pm$ 0.2	4.1 $\pm$ 0.1 <sup>2</sup>

<sup>1</sup> 10 mM lithium cacodylate buffer (pH 7.2), 99 mM LiCl, and 1 mM KCl.

<sup>2</sup> 10 mM lithium cacodylate buffer (pH 7.2), 90 mM LiCl, and 10 mM KCl.

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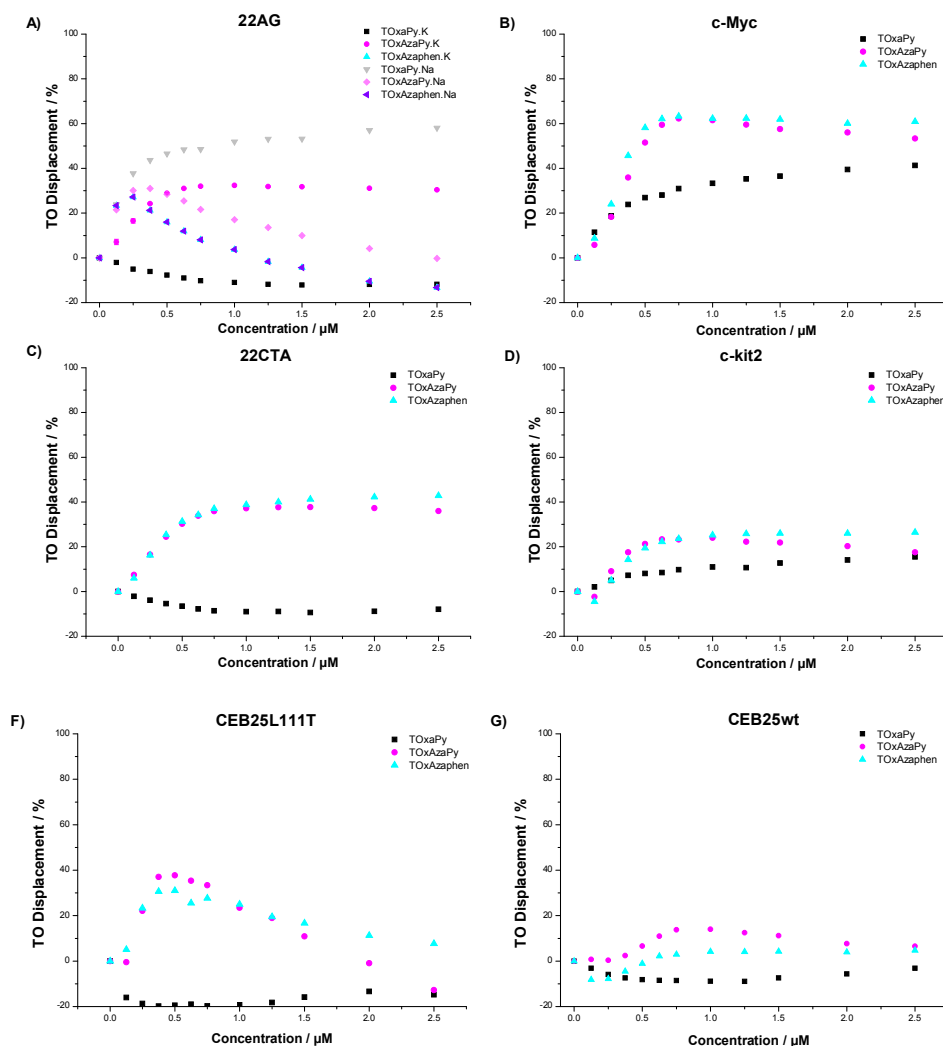
212 **Figure 4.** Stabilization of compounds TOxazaPy and TOxazaPhen towards several G4 structures.  
 213  $\Delta T_m$  values are plotted for each sequence in the buffer conditions mentioned in Table 3.

#### 214 2.1.2. G4-FID

215 According to literature data, BOxazaPy (Figure 1) and other pentaryl compounds [22] bind the  
 216 G4 through a combination of groove binding and  $\pi$ -stacking interactions independently of the  
 217 structure of the target. The hepta-aryl TOxazaPy showing a more elongated structure have been  
 218 suggested to bind according to a groove binding mode [21]. Although, it is likely to expect a similar  
 219 binding mode for both TOxazaPy and TOxazaPhen, the structural selectivity for the antiparallel  
 220 human telomeric sequence suggest an additional investigation on the binding modality.

221 To gain understanding on the binding modality of TOxazaPy and TOxazaPhen, G4-FID  
 222 titrations were performed. Concisely, this assay is based on the displacement of the fluoresce  
 223 probe TO (thiazole orange), which binds with no specificity several oligonucleotide structures. In this  
 224 context the assay was performed in the presence of six G4 sequences (22AG both in Na<sup>+</sup>- and K<sup>+</sup>-  
 225 buffer, called 22AG.N and 22AG.K, c-myc, c-kit2, 22CTA, CEB25wt, and CEB25L111T) and a duplex  
 226 DNA (ds26) as a control. We measured TO displacement by TOxazaPy, TOxazaPhen and TOxazaPy  
 227 in the presence of the G4 sequences previously mentioned, using the latter as reference compound  
 228 (Figure 5). The curves of both compounds never reach 100% TO displacement; either they reach  
 229 saturation a low percentage values (the probe is displaced by up to 40-60 % in the best cases with  
 230 22CTA and c-myc), or they display complex shape curves (22AG.Na, 22AG.K, c-kit2, and  
 231 CEB25L111T) suggesting complex equilibrium processes, or they exhibit no TO displacement  
 232 (CEB25wt). As previously mentioned for this class of compounds [21,22], this kind of curves may  
 233 reflect a low affinity of the heteroaryl compounds for the TO binding sites or/and an indirect  
 234 competition between the ligands and TO as a result of binding to different sites of the G4 structure.  
 235 The indirect competition is caused by the morphological changes onto the G4 structure triggered by  
 236 the ligand-binding. This competition may cause partial TO displacement or TO re-localization in a  
 237 site where its fluorescence is brighter than the original one. The re-localization might explain the  
 238 decreasing in the displacement curves [21]. At last, FID curves in the presence of ds26 (data not  
 239 shown) show no binding to duplex DNA. Interestingly, the two new compounds 6 and 7 behave  
 240 differently in comparison to TOxazaPy. Due to the similar hepta-heteroaromatic structure of the  
 241 ligands, we expected a similar behavior in the TO displacement, but surprisingly we obtained very  
 242 different results showing that the nature of the penta-aromatic ring has an effect on the binding mode  
 243 of the ligand towards the selected G4 structures.





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**Figure 5.** G4-FID plots of TOxAzaPy and TOxAzaPhen in the presence of: A) 22AG.Na and 22AG.K, B) c-Myc, C) 22CTA, D) c-kit2, F) CEB25wt, and G) CEB25L111T. TOxPy is used as a reference compound.

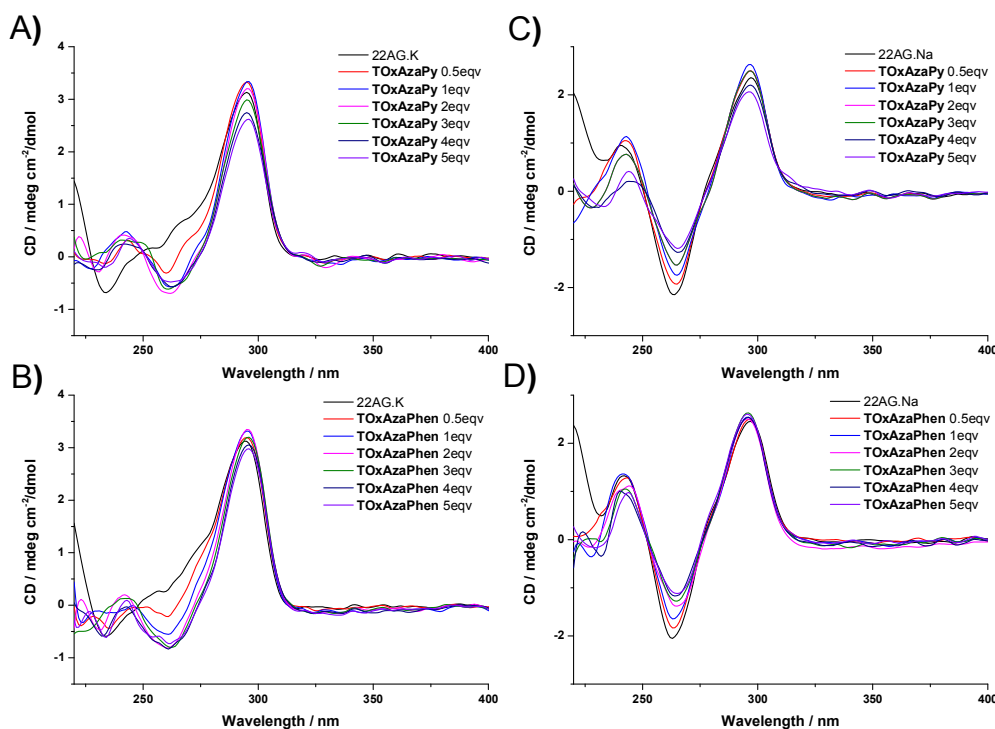
249 The flexible nature of these compounds makes them suitable for loop and groove interactions,  
 250 as previously described for the structural related TOxPy [21] and BOxAzaPy derivatives [22], or a  
 251 mixed binding mode as reported from structural studies obtained for distamycin A [36]. Remarkably,  
 252 by analyzing FID data we were able to determine interactions between G4s and both TOxAzaPy and  
 253 TOxAzaPhen that were not detected by FRET melting assay. A striking example is represented by  
 254 the c-myc binding, where no stabilization was measured by FRTE melting in the presence of both  
 255 TOxAzaPy and TOxAzaPhen, whereas TO was displaced up to 60%. These results support the idea  
 256 that FRET melting and G4 FID assays are two complementary biophysical methods that should be  
 257 used in concertation while analyzing the binding and the affinity of G4 ligands towards G4 structures.  
 258 Their combination should allow to gain much more understanding about the binding mode of these  
 259 family of ligands.

### 260 2.3. CD Titrations

261 To gain insights into the binding event, highlighting the morphological changes induced by ligand  
 262 binding to the chiral G4 structures, circular dichroism (CD) experiments were conducted. Due to the  
 263 higher stabilization observed by FRET melting assay in the presence of TOxAzaPy and TOxAzaPhen,  
 264 we selected the human telomeric sequence 22AG and we performed CD titrations in both  $\text{Na}^+$ - and  
 265  $\text{K}^+$ -buffer. The CD spectrum of the pre-folded 22AG alone in  $\text{K}^+$ -rich buffer shows the characteristic

signature reported in literature for this sequence, displaying a maximum at 290 nm, a shoulder around 265 nm, and a minimum at 235 nm [32-35], corresponding to a 3+1 hybrid structure [32]. After addition of TOxAzaPy an increase of the band at 290 nm together with a conversion of the positive band a 265 nm to negative was observed, suggesting a TOxAzaPy:22AG.K complex formation. This signature is characteristic of an antiparallel structure induced by the presence of the ligand. This trend is observed up to 1 eq. of TOxAzaPy, with the generation of two isoelliptical points at 285 nm and 250 nm suggesting a single equilibrium converting the free DNA into a bound species (Figure 6A). Increasing concentrations of ligand produced a dropping of the intensity of the band a 290 nm, a stabilization of the intensity of the band a 265 nm, and the disappearance of the two isoelliptical points suggesting multiple equilibriums. A very similar behavior was observed during the CD titration performed in the presence of TOxAzaPhen (Figure 6B) with the only difference that the higher molar ellipticity value at 290 nm and the lower value at 265 nm were reached at 2 eq. of ligand.

The same experiments were carried out in the presence of 22AG in Na<sup>+</sup>-rich buffer. As reported in literature, the main conformation generated by the telomeric sequence in this condition is an antiparallel structure characterized by positive bands at 295 nm and 245 nm and a negative band at 265 nm [32,37]. Upon addition of TOxAzaPy an increase in intensity of the bands at 295 nm, 265 nm, and 245 nm together with a 3 nm hypochromic shift at 295 nm and a 3 nm bathochromic shift at 265 nm was recorded. The first equivalent of ligand induced an increase of the intensity of the bands at 295 nm and 245 nm, differently higher ligand concentrations reduced the band intensities to lower values as compare to the initial ellipticity together with a 6 nm bathochromic shift at 245 nm. Increasing concentration of ligand produced a continuous increasing of the intensity of the band at 265 nm (Figure 6C). Two neat isoelliptical points are visible at 275 nm and 285 nm and a not clear isoelliptical point around 250 nm. TOxAzaPhen showed a similar behavior (Figure 6D) with a mild increase in intensity of the band at 295 nm, a strong increase at 265 nm, and a strong decrease at 245 nm. Two clear isoelliptical points are visible at 275 nm and at 285 nm and a not clear isoelliptical point around 255 nm. The titration experiments carried out in the presence of TOxAzaPy and TOxAzaPhen and the telomeric sequence both in K<sup>+</sup>- and Na<sup>+</sup>-rich buffer show the generation of heptaryl:22AG complexes. In the presence of K<sup>+</sup> the ligand binding causes a modification of the spectra shape due to a shift of the equilibrium towards the antiparallel topology. In the presence of Na<sup>+</sup> the dichroic changes suggest a strong interaction without substantial structural transitions. Summing up, both the ligands TOxAzaPy and TOxAzaPhen exhibit a remarkable preference for the human telomeric sequence in the antiparallel conformation.



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**Figure 6.** CD spectra of 22AG 5  $\mu$ M alone and upon addition of increasing concentration of A) TOxAzaPy in 10 mM lithium cacodylate buffer (pH 7.2), 100 mM KCl; B) TOxAzaPhen in 10 mM lithium cacodylate buffer (pH 7.2), 100 mM KCl; C) TOxAzaPy in 10 mM lithium cacodylate buffer (pH 7.2), 100 mM NaCl; and D) TOxAzaPhen in 10 mM lithium cacodylate buffer (pH 7.2), 100 mM NaCl.

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### 3. Materials and Methods

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#### 3.1. General Information

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All chemicals and solvents were purchased from Sigma Aldrich and used without further purification. Melting points were taken on a Olympus Bx41 polarization microscope apparatus and are uncorrected. TLC analysis was carried out on silica gel (Merck 60F 254) with visualization at 254 and 366 nm. Flash chromatography was performed with silica gel 60 (40–63  $\mu$ m, Merck). Biotage (MPLC) Isolera ONE, an automated flash purification system with variable wavelength detector, was performed with SNAP 10 g, 50 g, and 100 g Column, KP-SIL Pk 20, Biotage. All anhydrous reactions were carried out under positive pressure of nitrogen or argon. Elemental analysis were provided by Carlo Erba CHN analyzer. All  $^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra were recorded on a Bruker Advance 300 MHz spectrometer using deuterated solvents and TMS as internal standard. The spectra are reported in ppm and referenced to deuterated DMSO (2.49 ppm for  $^1\text{H}$ , 39.5 ppm for  $^{13}\text{C}$ ) or deuterated chloroform (7.26 ppm for  $^1\text{H}$ , 77 ppm for  $^{13}\text{C}$ ). The following abbreviations are used: singlet (s), doublet (d), triplet (t) and multiplet (m). Naked and modified oligonucleotide sequences were purchased from Eurogentec. FRET-melting assay were performed in 96-well plates on real time PCR apparatus 7900HT Fast Real-Time PCR System. G4-FID assays were performed in a fluorescence Agilent Cary Eclipse spectrophotometer, and a temperature of 20  $^\circ\text{C}$  was kept constant with thermostated cell holders (1 mL reaction volume). CD experiments were carried out at 20  $^\circ\text{C}$  with a JASCO J-710 spectropolarimeter equipped with a Peltier temperature controller (Jasco PTC-348WI) interfaced to a PC, by using 1 cm path rectangular quartz cells (1 mL reaction volume).

## 326 3.2. Synthetic Methods

## 327 3.2.1. General procedure for the synthesis of 1-4

328 The correspondent carboxylic acid (3.4 mmol) and 1,1'-carbonyldiimidazole (CDI, 3.4 mmol)  
329 were dissolved in DMF (15 mL) and stirred at room temperature for 30 min. (2Z, 6Z)-N'2,N'6-  
330 Dihydroxypyridine-2,6 bis(carboximidamide) (1.53 mmol) was added and the reaction mixture was  
331 stirred at room temperature overnight. CDI (3.4 mmol) was further added and the reaction mixture  
332 was heated at 150 °C for 6 h. After cooling down, the reaction mixture was poured into water to  
333 induce the precipitation of a solid, purified by Isolera ONE (CHCl<sub>3</sub>:MeOH gradient 0 to 20%), which  
334 was filtered and characterised as pure product.

335 *2,6-bis(5-(furan-2-yl)-1,2,4-oxadiazol-3-yl)pyridine (1)*. White solid, Yield: 11 %, m.p.>300 °C (dec.); <sup>1</sup>H  
336 NMR (300 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 8.37 (d, 3J(H,H) = 7.8 Hz, 2 H), 8.10 (t, 3J(H,H) = 8 Hz, 1 H),  
337 7.77 (d, 3J(H,H) = 1.2 Hz, 2 H), 7.51 (d, 3J(H,H) = 3.6 Hz, 2 H), 6,7 (dd, 3J(H,H) = 3.5 Hz; 3J(H,H) = 1.6  
338 Hz, 2 H). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 168.3, 167.8, 146.9, 146.8, 139.8, 138.3, 125.1, 117.3,  
339 112.6; elemental analysis calcd (%) for C<sub>17</sub>H<sub>9</sub>N<sub>5</sub>O<sub>4</sub>: C, 58.79; H, 2.61; N, 20.17; found C, 57.98; H, 2.70;  
340 N, 20.07.

341 *2,6-bis(5-(thiophen-2-yl)-1,2,4-oxadiazol-3-yl)pyridine (2)*. White solid, Yield: 11 %, m.p.>300 °C; <sup>1</sup>H NMR  
342 (300 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 8.46 (d, 3J(H,H) = 7.8 Hz, 2 H), 8.19 (t, 3J(H,H) = 7.8 Hz, 2 H), 8.17  
343 (dd, 3J(H,H) = 2.4 Hz, 3J(H,H) = 1.3 Hz 2 H), 7.82 (dd, 3J(H,H) = 5 Hz, 3J(H,H) = 1.05 Hz, 2 H), 7.36  
344 (dd, 3J(H,H) = 4.9 Hz, 3J(H,H) = 3.8 Hz, 2 H). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 172.2, 167.9,  
345 146.9, 138.2, 132.5, 132.4, 128.5, 125.3, 124.9; elemental analysis calcd (%) for C<sub>17</sub>H<sub>9</sub>N<sub>5</sub>O<sub>2</sub>S<sub>2</sub>: C, 53.81; H,  
346 2.39; N, 18.46; found C, 53.12; H, 2.51; N, 19.02.

347 *2,6-bis(5-(benzofuran-2-yl)-1,2,4-oxadiazol-3-yl)pyridine (3)*. White solid, Yield: 28 %, m.p.>300 °C; <sup>1</sup>H  
348 NMR (300 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 8.44 (d, 3J(H,H) = 7.8 Hz, 2 H), 8.16 (d, 3J(H,H) = 7.8 Hz, 2  
349 H), 7.88 (s, 2 H), 7.80 (d, 3J(H,H) = 7.7 Hz, 2 H), 7.72 (d, 3J(H,H) = 8.4 Hz, 2 H), 7.55 (t, 3J(H,H) = 7.5  
350 Hz, 2 H), 7.41 (t, 3J(H,H) = 7.5 Hz, 2 H). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 168.5, 167.7, 146.4,  
351 140.3, 138.0, 127.6, 126.6, 126.1, 124.9, 123.9, 122.4, 113.1, 111.9; elemental analysis calcd (%) for  
352 C<sub>25</sub>H<sub>13</sub>N<sub>5</sub>O<sub>4</sub>: C, 67.11; H, 2.93; N, 15.65; found C, 57.01; H, 2.77; N, 15.02.

353 *2,6-bis(5-(1H-indol-2-yl)-1,2,4-oxadiazol-3-yl)pyridine (4)*. White solid, Yield: 3 %, m.p.>300 °C; <sup>1</sup>H NMR  
354 (300 MHz, CDCl<sub>3</sub>, 25°C, TMS): δ = 9.49 (s, 2 H), 8.38 (d, 3J(H,H) = 7.7 Hz, 2 H), 8.10 (t, 3J(H,H) = 7.3  
355 Hz, 1 H), 7.77 (d, 3J(H,H) = 7.7 Hz, 2 H), 7.52 (s, 2 H), 7.40 (t, 3J(H,H) = 7.2 Hz, 2 H), 7.31 (m, 4 H).  
356 elemental analysis calcd (%) for C<sub>25</sub>H<sub>15</sub>N<sub>7</sub>O<sub>2</sub>: C, 67.41; H, 3.39; N, 22.01; found C, 67.12; H, 3.51; N,  
357 21.82.

## 358 3.2.2. Stepwise synthesis for (5)

359 *6-(5-(quinolin-2-yl)-1,2,4-oxadiazol-3-yl)picolinonitrile 15*. Quinoline-2-carboxylic acid (0.12 g, 0.68  
360 mmol) and 1,1'-carbonyldiimidazole (CDI, 0.11 g, 0.68 mmol) were dissolved in DMF (10 mL) and  
361 stirred at room temperature for 30 min. (Z)-6-cyano-N'-hydroxypicolinimidamide (0.1 g, 0.62 mmol)  
362 was added and the reaction mixture was stirred at room temperature overnight. CDI (0.11 g, 0.68  
363 mmol) was further added and the reaction mixture was heated at 150 °C for 6 h. After cooling down,  
364 the reaction mixture was poured into water to induce the precipitation of a pale yellow solid, purified  
365 by Isolera ONE (CHCl<sub>3</sub>:MeOH gradient) obtaining a white solid (0.11 g; yield 60%), which was  
366 filtered and characterised as pure product. M.p.>300 °C (dec.); <sup>1</sup>H NMR (300 MHz, [D<sub>6</sub>]DMSO, 25°C,  
367 TMS): δ = 8.73 (d, 3J(H,H) = 8.5 Hz, 1 H), 8.53 (d, 3J(H,H) = 7.8 Hz, 1 H), 8.43 (d, 3J(H,H) = 7.8 Hz, 1  
368 H), 8.36 (t, 3J(H,H) = 7.8, 1 H), 8.27 (d, 3J(H,H) = 8.4 Hz, 2 H), 8.17 (d, 3J(H,H) = 7.9 Hz, 1 H), 7.96 (t,  
369 3J(H,H) = 7.2, 3J(H,H) = 8.1 Hz, 1 H), 7.82 (t, 3J(H,H) = 7.4, 3J(H,H) = 7.6 Hz, 1 H). <sup>13</sup>C NMR (75 MHz,

370 [D6]DMSO, 25°C, TMS):  $\delta$  = 175.2, 167.5, 147.4, 147.0, 142.6, 139.9, 138.4, 133.4, 131.2, 130.9, 129.7,  
371 128.9, 128.2, 127.1, 120.6, 116.8.

372 (*Z*)-*N'*-hydroxy-6-(5-(quinolin-2-yl)-1,2,4-oxadiazol-3-yl)-picolinimidamide **16**. A mixture of  
373 hydroxylamine hydrochloride (0.047 g, 0.68 mmol) and of Na<sub>2</sub>CO<sub>3</sub> (0.036 g, 0.34 mmol) in water (10  
374 mL) was added dropwise in 45 minutes to a solution containing 6-(5-(quinolin-2-yl)-1,2,4-oxadiazol-  
375 3-yl)picolinonitrile (0.1 g, 0.33 mmol) in EtOH (30 mL) and the reaction mixture was stirred at r.t. for  
376 1 hour. The precipitated white solid (0.095 g, yield 92%) was filtered and dried. M.p. = 225-230°C; <sup>1</sup>H  
377 NMR (300 MHz, [D6]DMSO, 25°C, TMS):  $\delta$  = 10.09 (s, 1 H), 8.74 (d, 3J(H,H) = 8.5 Hz, 1 H), 8.45 (d,  
378 3J(H,H) = 8.5 Hz, 1 H), 8.27 (m, 2 H), 8.18 (d, 3J(H,H) = 8.2, 1 H), 8.11 (d, 3J(H,H) = 4.1 Hz, 2 H), 7.96  
379 (dt, 3J(H,H) = 1.4 Hz, 3J(H,H) = 8.3, 1 H), 7.82 (t, 3J(H,H) = 1.2, 3J(H,H) = 8.1 Hz, 1 H), 5.89 (s, 2 H). <sup>13</sup>C  
380 NMR (75 MHz, [D6]DMSO, 25°C, TMS):  $\delta$  = 174.8, 168.3, 150.8, 148.9, 147.3, 144.3, 142.8, 138.4, 138.3,  
381 131.2, 129.7, 129.1, 128.9, 128.3, 123.7, 121.8, 120.6.

382 2,6-bis(5-(quinolin-2-yl)-1,2,4-oxadiazol-3-yl)pyridine (**5**). Quinoline-2-carboxylic acid (0.126 g, 0.72  
383 mmol) and 1,1'-carbonyldiimidazole (CDI, 0.12 g, 0.72 mmol) were dissolved in DMF (10 mL) and  
384 stirred at room temperature for 30 min. (2*Z*, 6*Z*)-*N'*,*N'*6-Dihydroxypyridine-2,6-  
385 bis(carboximidamide) (0.2 g, 0.62 mmol) was added and the reaction mixture was stirred at room  
386 temperature overnight. CDI (0.12 g, 0.72 mmol) was further added and the reaction mixture was  
387 heated at 150 °C for 6 h. After being cooled, the reaction mixture was poured into water to induce the  
388 precipitation of a pale white solid, purified by Isolera ONE (CHCl<sub>3</sub>:MeOH gradient) obtaining a white  
389 solid (0.151 g; yield 53%), which was filtered and characterised as pure product; m.p. >300 °C (dec.).  
390 <sup>1</sup>H NMR (300 MHz, [D6]DMSO, 25°C, TMS):  $\delta$  = 8.73 (d, 3J(H,H) = 8.4 Hz, 2 H), 8.47 (m, 4 H), 8.40 (t,  
391 3J(H,H) = 7.1 Hz, 1 H), 8.29 (d, 3J(H,H) = 8.3, 2 H), 8.17 (d, 3J(H,H) = 7.95 Hz, 2 H), 7.96 (t, 3J(H,H) =  
392 7.1 Hz, 3J(H,H) = 7.5 Hz, 2 H), 7.82 (t, 3J(H,H) = 6.9, 3J(H,H) = 7.3 Hz, 1 H). HRMS (ESI-MS): 470.1356  
393 (calculated: 470.1365 C<sub>27</sub>H<sub>17</sub>N<sub>7</sub>O<sub>4</sub><sup>+</sup>). ; elemental analysis calcd (%) for C<sub>27</sub>H<sub>15</sub>N<sub>7</sub>O<sub>2</sub>: C, 69.08; H, 3.22; N,  
394 20.89; found C, 69.18; H, 3.10; N, 20.13.

### 395 3.2.3. Stepwise synthesis of TOxAzaPy and TOxAzaPhen

#### 396 Synthesis of (*Z*)-*N'*-hydroxy-3-(5-methyl-1,2,4-oxadiazol-3-yl)benzimidamide **18**

397 6-(5-methyl-1,2,4-oxadiazol-3-yl)picolinonitrile **17**. (*Z*)-6-cyano-*N'*-hydroxypicolinimidamide (0.3 g, 1.86  
398 mmol), was dissolved in 15 mL of chloroform and treated with acetic anhydride (0.21 mL, 2.22 mmol)  
399 and Et<sub>3</sub>N (0.3 mL, ) and the mixture was stirred overnight at room temperature. The solvent was  
400 removed under reduce pressure and the crude product was dissolved in acetic acid and refluxed for  
401 4 hours. The reaction mixture was neutralized with a saturated solution of NaHCO<sub>3</sub>. A white solid  
402 precipitated was filtered and washed with water (0.23 g, yield 65%). <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>, 25°C,  
403 TMS):  $\delta$  = 8.35 (d, 3J(H,H) = 7.2 Hz, 1 H), 8.05 (t, 3J(H,H) = 7.8 Hz, 1 H), 7.86 (d, 3J(H,H) = 7.8 Hz, 1 H),  
404 2.75 (s, 1 H). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>, 25°C, TMS):  $\delta$  = 179.8, 168.7, 149.7, 140.1, 136.3, 131.7, 127.8,  
405 118.1, 14.2.

406 (*Z*)-*N'*-hydroxy-6-(5-methyl-1,2,4-oxadiazol-3-yl)picolinimidamide **18**. An aqueous solution (30 mL)  
407 containing hydroxylamine hydrochloride (0.150 g, 1.1 mmol), and sodium carbonate (0.116 g, 1.08  
408 mmol) was added to a solution of 6-(5-methyl-1,2,4-oxadiazol-3-yl)picolinonitrile **17** (0.2 g, 1.08  
409 mmol) in ethanol (50 mL). The hydroxylamine was added dropwise in 30 minutes and then the  
410 mixture was stirred for further 1 hour. The precipitated white solid (0.21 g, yield 89%) was filtered,  
411 dried, and used without any further purification. <sup>1</sup>H NMR (300 MHz, [D6]DMSO, 25°C, TMS):  $\delta$  =  
412 10.13 (s, 1 H), 8.04 (m, 3 H), 5.87 (s, 2 H), 2.71 (s, 3 H). <sup>13</sup>C NMR (75 MHz, [D6]DMSO, 25°C, TMS):  $\delta$   
413 = 177.9, 167.3, 150.5, 148.8, 144.5, 138.2, 123.3, 121.5, 12.07.

#### 414 General procedure for the synthesis of TOxAzaPy-Me (**6**) and TOxAzaPhen (**7**)

415 The corresponding dicarboxylic acid (0.21 mmol) and 1,1'-carbonyldiimidazole (CDI, 0.46 mmol)  
 416 were dissolved in DMPU (2 mL) and stirred at room temperature for 30 min. (Z)-N'-hydroxy-6-(5-  
 417 methyl-1,2,4-oxadiazol-3-yl)picolinimidamide **18** (0.46 mmol) was added and the reaction mixture  
 418 was stirred at room temperature overnight. CDI (0.46 mmol) was further added and the reaction  
 419 mixture was heated at 150 °C for 6 h. After cooling down, the reaction mixture was poured into water  
 420 to induce the precipitation of a white solid, purified by Isolera ONE (CHCl<sub>3</sub>:MeOH 0 to 5%) obtaining  
 421 a solid, which was filtered and characterised as pure product.

422 *2,6-bis(5-(6-(5-methyl-1,2,4-oxadiazol-3-yl)pyridin-2-yl)-1,2,4-oxadiazol-3-yl)pyridine* TOxAzaPy (**6**).  
 423 White solid, Yield: 19 %, m.p.>300 °C (dec.); <sup>1</sup>H NMR (300 MHz, [D<sub>6</sub>]DMSO, 25 °C, TMS): δ = 8.68 (d,  
 424 3J(H,H) = 7.8 Hz, 2 H), 8.49 (t, 3J(H,H) = 8.1 Hz, 1 H), 8.41 (dd, 3J(H,H) = 6.2 Hz, 3J(H,H) = 2.7 Hz 2  
 425 H), 8.31 (d, 3J(H,H) = 6.2 Hz, 2 H), 2.74 (s, 3 H). <sup>13</sup>C NMR (75 MHz, [D<sub>6</sub>]DMSO, 25 °C, TMS): δ = 177.9,  
 426 174.1, 168.5, 167.4, 146.8, 146.2, 143.6, 140.4, 139.4, 127.5, 125.4, 11.9; elemental analysis calcd (%) for  
 427 C<sub>25</sub>H<sub>15</sub>N<sub>11</sub>O<sub>4</sub>: C, 56.29; H, 2.83; N, 28.88; found C, 56.93; H, 2.92; N, 29.02.

428 *1,3-bis(5-(6-(5-methyl-1,2,4-oxadiazol-3-yl)pyridin-2-yl)-1,2,4-oxadiazol-3-yl)benzene* TOxAzaPhen (**7**).  
 429 White solid, Yield: 38 %, m.p.>300 °C (dec.); <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>, 25 °C, TMS): δ = 9.22 (s, 1 H),  
 430 8.56 (d, 3J(H,H) = 7.5 Hz, 2 H), 8.39 (d, 3J(H,H) = 7.8, 2 H), 8.29 (d, 3J(H,H) = 7.7 Hz, 2 H), 8.09 (t,  
 431 3J(H,H) = 7.8 Hz, 2 H), 7.82 (t, 3J(H,H) = 7.8 Hz, 1 H), 2.75 (s, 3 H). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>, 25 °C,  
 432 TMS): δ = 177.9, 174.1, 168.5, 167.4, 146.8, 146.2, 143.6, 140.4, 139.4, 127.5, 125.4, 11.9; elemental analysis  
 433 calcd (%) for C<sub>26</sub>H<sub>16</sub>N<sub>10</sub>O<sub>4</sub>: C, 58.65; H, 3.03; N, 26.31; found C, 58.12; H, 3.21; N, 26.08.

### 434 3.3. Biophysical Assays

#### 435 3.3.1. FRET Melting Assay

436 Stabilization of compounds with quadruplex-structure was monitored via FRET-melting assay  
 437 performed in 96-well plates on real time PCR apparatus 7900HT Fast Real-Time PCR System as  
 438 follow: 5 min at 25 °C, then increase of 0.5 °C every minute until 95 °C. Each experimental condition  
 439 was tested in duplicated in a volume of 25 μL for each sample. FRET-melting assay was performed  
 440 with oligonucleotides that mimic the human telomeric sequence, as well as other quadruplex-  
 441 forming oligonucleotides, equipped with FRET partners (here: FAM and TAMRA) at each extremity  
 442 in the presence of the G4 ligand. To determine ligand G4 *vs* duplex selectivity, an unlabelled DNA  
 443 competitor ds26 was used. The oligonucleotides were prepared at 0.2 μM, the ligands at 1 μM and  
 444 competitors at 0, 3, and 10 μM final concentration. Measurements were made with excitation at 492  
 445 nm and detection at 516 nm in a buffer of lithium cacodylated (10 mM, pH 7.2), NaCl (100 mM) or  
 446 KCl (10 mM, completed by 90 mM LiCl for F21T, and 1 mM, completed by 99 mM LiCl for all the  
 447 others G-quadruplex sequences). To identify if the ligands are able to stabilize the G-quadruplex  
 448 structure, we determined the temperature at half denaturation of the G4 in the absence and in the  
 449 presence of ligand. The experiments were carried out on prefolded G-quadruplex structures: the  
 450 sequences were heated at 90 °C for 5 min and left to cool down at 4 °C overnight. At last, in order to  
 451 identify a preferential binding towards a particular G-quadruplex conformation, a series of labelled  
 452 oligonucleotides covering a range of possible G4 conformations was used:

453 **Table 4.** DNA sequences employed for the FRET melting assay

Sequence Name	Sequence (5'-3')
F21T	Fam-G <sub>3</sub> TTAG <sub>3</sub> TTAG <sub>3</sub> TTA G <sub>3</sub> -Tamra
FMycT	Fam-TTGAG <sub>3</sub> TG <sub>3</sub> TAG <sub>3</sub> TG <sub>3</sub> TAA-Tamra
F21CTAT	Fam-G <sub>3</sub> CTAG <sub>3</sub> CTAG <sub>3</sub> CTAG <sub>3</sub> -Tamra
FKit2T	Fam-G <sub>3</sub> CG <sub>3</sub> CGCGAG <sub>3</sub> AG <sub>4</sub> -Tamra
FCEB25wtT	Fam-AAG <sub>3</sub> TG <sub>3</sub> TGTAAGTGTG <sub>3</sub> TG <sub>3</sub> T-Tamra
FCEB25-L111TT	Fam-AAG <sub>3</sub> TG <sub>3</sub> TG <sub>3</sub> TG <sub>3</sub> T-Tamra
FCEB25-L121(AA)TT	Fam-AAG <sub>3</sub> TG <sub>3</sub> AAAG <sub>3</sub> TG <sub>3</sub> T-Tamra

FCEB25-L121TT	Fam-AAG <sub>3</sub> TG <sub>3</sub> TTG <sub>3</sub> TG <sub>3</sub> T-Tamra
FBcl2T	Fam-AG <sub>4</sub> CG <sub>3</sub> CGCG <sub>3</sub> AG <sub>2</sub> AAG <sub>5</sub> CG <sub>3</sub> AGCG <sub>4</sub> CTG-Tamra
ds26	CAATCGGATCGAATTCGATCCGATTG

### 454 3.3.2. G4-FID Assay

455 Cell G4-FID assays were performed in a fluorescence Agilent Cary Eclipse spectrophotometer.  
 456 A temperature of 20 °C was kept constant with thermostated cell holders. Each experiment was  
 457 performed in a 1mL cell, in 10 mM lithium cacodylate buffer (pH 7.4) with 100 mM KCl or NaCl  
 458 depending on the experiment, in a total volume of 1 mL. The G4-FID assay was designed as follows:  
 459 0.25 μM pre-folded DNA target was mixed with thiazole orange (0.50 μM for G4-DNA, 0.75 μM for  
 460 ds26). Each ligand addition (from 0.5 to 10 equivalents) was followed by a 3 min equilibration time,  
 461 after which the fluorescence spectrum was recorded. The percentage of displacement was calculated  
 462 as follows: TO displacement (%) = 100 - [(FA / FA<sub>0</sub>) × 100], where FA and FA<sub>0</sub> stand for the  
 463 fluorescence emission area of TO bound to DNA after each ligand addition and FA<sub>0</sub> before ligand  
 464 addition (area measured from 510 to 750 nm, λ<sub>exc</sub> = 495 nm, slit = 10nm). The percentage of  
 465 displacement was then plotted as a function of the concentration of added ligand. As for the FRET  
 466 melting assay a series of oligonucleotides covering a range of possible G4 conformations was used:

467 **Table 5.** DNA sequences employed for the G4-FID assay

Sequence Name	Sequence (5'-3')
22AG	AG <sub>3</sub> TTAG <sub>3</sub> TTAG <sub>3</sub> TTAG <sub>3</sub>
Myc22	TGAG <sub>3</sub> TG <sub>3</sub> TAG <sub>3</sub> TG <sub>3</sub> TAA
22CTA	AG <sub>3</sub> CTAG <sub>3</sub> CTAG <sub>3</sub> CTAG <sub>3</sub>
c-kit2	G <sub>3</sub> CG <sub>3</sub> CGCGAG <sub>3</sub> AG <sub>4</sub>
CEB25wt	AAG <sub>3</sub> TG <sub>3</sub> TGTAAGTG <sub>3</sub> TG <sub>3</sub> T
CEB25-L111T	AAG <sub>3</sub> TG <sub>3</sub> TG <sub>3</sub> TG <sub>3</sub> T
Bcl2	AG <sub>4</sub> CG <sub>3</sub> CGCG <sub>3</sub> AG <sub>2</sub> AAG <sub>5</sub> CG <sub>3</sub> AGCG <sub>4</sub> CTG
ds26	CAATCGGATCGAATTCGATCCGATTG

### 468 3.3.3. Circular Dichroism (CD) Titration

469 CD experiments were carried out at 20 °C with a JASCO J-710 spectropolarimeter equipped with  
 470 a Peltier temperature controller (Jasco PTC-348WI) interfaced to a PC, by using 1 cm path rectangular  
 471 quartz cells (1 mL reaction volume). Scans were recorded from 220 nm to 400 nm with the next  
 472 parameters: 100 mdeg sensitivity, 1 nm data pitch, 200 nm min<sup>-1</sup> scan speed, 1 s response, 1 nm band  
 473 width, and 4 accumulations. Solutions containing 5 μM 22AG were titrated with increasing  
 474 concentration of ligands (0 to 5 eq, 10 min stabilization time) in lithium cacodylate (10 mM)  
 475 supplemented with potassium or sodium chloride (100 mM) at pH 7.2. The scan of the buffer was  
 476 subtracted from the average of the scan of the samples. The signal was further smoothed with a  
 477 Savitzky-Golay method (2 order, 20 points window).

## 478 4. Conclusions

479 In conclusion, a new series of acyclic oligoheteroaryls (**1-7**) containing 1,2,4-oxadiazoles and  
 480 carbo- or hero-aromatics have been synthesized according to a one-pot, two-step synthesis or  
 481 following a more time consuming, but more efficient stepwise protocol. **1-7** have been evaluated as  
 482 ligands towards secondary nucleic acid structures mimicking cancer-associated G-quadruplexes  
 483 (e.g.: 22AG for human telomeric sequence, c-myc, and c-kit promoters). Their binding properties have  
 484 been compared to the structurally related reference compound TOxAPy, by FRET melting, G4-FID  
 485 assays and circular dichroism. Pentaheteroaryls showed to be very poor G-quadruplex ligands,  
 486 corroborating previous results on cationic analogues (BOxAzaPy), which binding properties were  
 487 mainly conferred by the charged appendages. On the contrary, more extended heptapyridyl-

488 oxadiazole compounds, such as TOxAzaPy and TOxAzaPhen, showed preferential binding towards  
489 the antiparallel telomeric sequence (22AG) with remarkable selectivity *vs* duplex DNA. From this  
490 point of view, and for potency as well, they compare favorably to TOxaPy as G-quadruplex ligands.  
491 Our evidences, suggest that 1,2,4-oxadiazole moieties introduced in the G4-ligand as structural  
492 modification, implement and ameliorate the G-quadruplex binding properties of the  
493 heptaheteroaryls, which mainly act as groove ligands.

494 **Author Contributions:**

495

496 Methodology, F.D.; Investigation, V.P. and M.P.; Resources, M.F. and M.-P. T. F; Data Curation,  
497 D.V.; Writing-Original Draft Preparation, F.D., and D.V.; Writing-Review & Editing, M.F.;  
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505 **Sample Availability:** Samples of the compounds 1-7 are available from the authors.

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