

Discovery of new G-quadruplex binding chemotypes†

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We report here on the discovery and preliminary evaluation of a novel non-macrocyclic low molecular weight quadruplex-stabilizing chemotype. The lead compounds, based on a furan core, show high G-quadruplex stabilisation and selectivity as well as potent *in vitro* anti-proliferative activity.

Quadruplexes (G4s) are higher-order nucleic acid arrangements involving a core of π - π stacked guanine-quartets (G-quartets) rather than the Watson-Crick base pairs of double-helical nucleic acids.¹ G4-forming sequences are widely prevalent in eukaryotic telomeric sequences as well as being over-represented in other genomes,² notably promoter and 5'-UTR sequences of genes involved in cellular proliferation.³ The recent demonstration of the presence of G4s in human cells⁴ has added credence to the concept that G4s can be targets for therapeutic intervention, at the single gene or poly-gene levels.⁵ Appropriate small molecules can serve to stabilise G4s and the resulting complexes can then act as impediments to telomere maintenance, transcription or translation, depending on the nature of the quadruplex target site.⁶ These effects have been shown in several target genes of relevance to human cancer such as *c-MYC*⁷ and *c-KIT*.⁸

A large number of small molecule chemotypes have been reported as G4-binding ligands.⁶ The overwhelming majority are heteroaromatic with large flat surfaces, designed to complement the surface characteristics of a terminal G-quartet in a typical quadruplex structure. A second class of ligand is represented by the cyclic polyoxazole natural product telomestatin.⁹ A number of cyclic and acyclic analogues have been reported, some of which show potent biological activity.¹⁰ The acyclic compounds tend to be characterised by a crescent shape. For example, pyridostatin¹¹ and several series of phenyl- and pyridyl-bis-oxazoles^{12–14} all selectively target G4s (Fig. 1). A more general

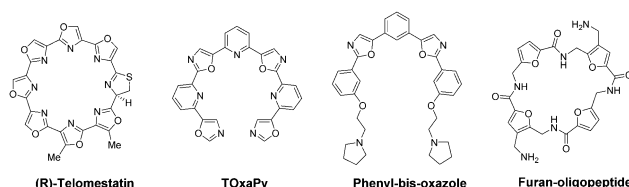


Fig. 1 Structures of various non-polycyclic G-quadruplex ligands.

requirement of most G4-binding ligands is the possession of side-chains terminating in cationic charge.⁶

Few G4-binding small molecules have proceeded to *in vivo* evaluation in models of human cancer, and to date only one compound, Quarfloxin, has been evaluated in clinical trials.¹⁵ The perceived lack of drug-like characteristics in many G4-binding compounds may have hindered progress to the clinic. We report here on a study to discover novel ligands with MW < 400 Da that could be suitable starting-points for future drug discovery efforts.

Thirty-eight representative members of a large chemical library from the anti-parasitic drug discovery programme at Georgia State University^{16,17} (several hundred compounds), with highly diverse scaffolds and functional groups, were screened using a high-throughput 96-well FRET (Fluorescence Resonance Energy Transfer) assay.¹⁸ G4 stabilisation was initially evaluated using dual-labelled F21T (human telomeric 21-mer) and *c-KIT*2 (a tyrosine kinase oncogene) G4s, as well as a duplex DNA sequence (T-loop). The ten most active compounds were subsequently screened against an expanded panel of fluorescently-labelled promoter G4-forming sequences, with HSP90A, HSP90B (heat shock protein 90 promoter sequences),¹² *k-RAS*21 (in the promoter of the *k-RAS* oncogene)¹⁹ and AR, a G4 recently identified in the promoter of the androgen receptor (involved in prostate cancer development).²⁰

Six acyclic furan- and thiophene-based compounds (Fig. 2: 1–6), representing two distinctive chemotypes were identified with high (>15 °C) ΔT_m values. A competition assay using unlabelled calf thymus duplex DNA and compounds 1–6 examined the ability of these compounds to differentiate between duplex and the F21T G4 DNA at high duplex : G4 ratios. The compounds were also examined

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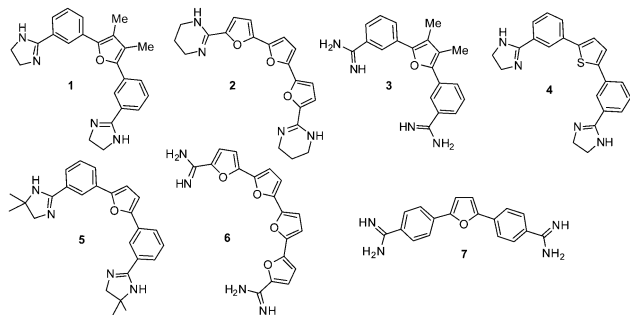


Fig. 2 Structures of the lead compounds **1–6** identified in this study, together with a control compound **7**.

in a 96 h short-term sulforhodamine B (SRB) assay, to determine their ability to inhibit cancer cell growth (Tables 1 and 2). All six compounds showed potent G4 stabilising abilities, as judged by the large changes in ΔT_m values for the selected G4s. In particular the bis-phenyl-mono-furan compounds **1** and **5** had especially high ΔT_m values, broadly comparable to those for established high-affinity G4-binding compounds such as tetra-substituted naphthalene diimides.²¹ The tri-furan compound **2** is consistently more effective in stabilising the G4s than the polyfuran compound **6**. Compound **2** and **6**, representatives of a tetra-furan second chemotype, were less selective at high duplex ratios and were inactive in the SRB assay, possibly because of aqueous solubility and cellular uptake issues. Switching from a furan (**1**, **5**) to a thiophene (**4**), does slightly affect G4 stabilisation and selectivity *vs.* duplex DNA, though not *in vitro* potency, which at least in the cell lines examined, is comparable to that of compounds **1** and **5**.

Overall, F21T and the two HSP90 G4s have been most stabilised by compounds **1–6**. Comparison with the behaviour of a tetra-substituted naphthalenediimide compound previously

examined by us,²⁰ shows that **1–6** exhibit only moderate ΔT_m values with the AR G4, which are generally lower than with other G4s. Compounds **1–6** produced slightly reduced but still significant stabilisation with the *c-KIT2* and *k-RAS21* G4s, suggesting that these compounds have the ability to act simultaneously on multiple G4 targets (G4 poly-targeting). The stabilisation of a duplex DNA sequence (T-loop) was not significantly affected by any of the compounds at the biologically relevant concentration employed here (1 μ M). Compounds **1** and **5** in particular are highly selective for G4 *versus* duplex DNA, as found in a series of competition assays, where the G4 stabilisation ability of both compounds is undiminished by adding calf thymus DNA in excess, at ratios up to 1:100/300. A control compound, **7**, an established duplex DNA minor groove binder, also showed significant G4 stabilisation, albeit with greater effects on the duplex DNA used (with a sub-optimal sequence for this compound).

Circular dichroism (CD) was employed to qualitatively evaluate the binding mode of the lead compounds and to examine induced structural transitions in the telomeric G4. The CD spectra (Fig. 3 and ESI†) show that compounds **1** and **3–6** produce very small induced CD signals in a human telomeric quadruplex sequence. Such weak induced CD signals are characteristic of quadruplex end-stacking compounds and the small differences in the CD signal patterns for different compounds indicate minor differences in the stacking geometries of the ligands at the terminal G-quartets.

All the compounds exhibiting weak induced CD signals that show a binding preference for an anti-parallel type quadruplex conformation as observed by decreases in the CD signal around 260 nm. Compound **5** however behaves differently, inducing a much larger conformational transition in the telomeric quadruplex upon complex formation. Upon titrating **5**, the CD intensity around 290 nm decreases with a subsequent increase in the CD signal around 260 nm. This is most likely due to an induced

Table 1 FRET G4 stabilization (ΔT_m at 1 μ M in °C) and calf thymus DNA (CT) competition data (the latter showing % retention of the F21T ΔT_m at 1 μ M). (E_{sd}) \pm 0.5 °C, from triplicate measurements. n/a: indicates unsuccessful curve fitting to the melting data. Compound **7**, a negative control, is a *para* analogue of the mono-furan compounds. Compound **8** is a tetra-substituted naphthalene diimide derivative²¹ used here as a G4 control

Compound	Mol. Wt	F21T	<i>c-KIT2</i>	HSP-90A	HSP-90B	<i>k-RAS21R</i>	AR	T-loop	G4:CT 1:1	G4:CT 1:10	G4:CT 1:100	G4:CT 1:300
1	384.5	22.3	16.8	27.7	23.7	19.6	15.4	<2	100	100	100	100
2	364.4	20.4	16.3	26.0	22.0	18.5	13.8	<2	100	100	92.7	63.3
3	330.5	17.6	17.8	21.2	18.1	15.1	11.9	2.2	100	100	65.9	43.9
4	372.5	18.0	13.9	22.5	19.1	15.9	10.2	<2	100	100	60.5	39.9
5	412.5	22.6	18.5	26.6	22.7	18.3	14.7	<2	100	100	100	n/a
6	350.3	18.6	n/a	20.6	16.2	13.4	10.1	<2	100	100	53.1	35.4
7	302.5	14.4	12.2	19.0	15.7	16.6	9.9	3.4	100	100	24.5	2.5
8	830.6	26.6	22.0	33.1	28.6	n/a	15.9	4.9	100	100	n/a	27.2

Table 2 IC₅₀ values of compounds **1–8** determined by a 96 h SRB assay (see the ESI for further details). (E_{sd}) \pm 0.3 μ M for compounds **1–6**

	A549	MCF7	ALT	Mia-PaCa2	Panc1	RCC4	786-O	WI38
1	0.3	0.9	1.0	0.3	0.5	4.9	3.0	1.3
2	>25	>25	>25	>25	>25	>25	>25	>25
3	0.8	0.9	6.6	2.0	1.7	>25	>25	>25
4	0.4	1.4	1.9	0.7	1.0	3.3	5.6	2.0
5	0.5	1.2	1.4	0.5	1.0	3.9	4.8	2.0
6	4.2	15.6	18.8	7.0	>25	>25	12.8	>25
7	3.7	11.7	>25	8.90	10.9	>25	>25	>25
8	0.019	0.070	0.063	0.011	0.003	0.560	0.320	0.230



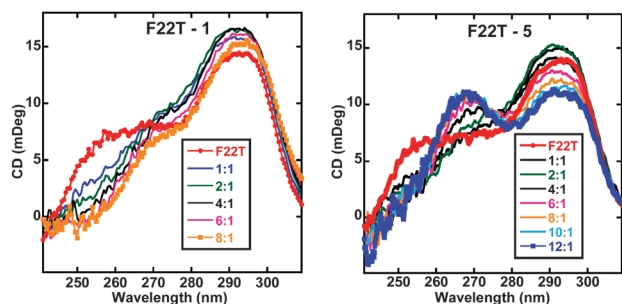


Fig. 3 Circular dichroism (CD) spectra for compounds **1** and **5** at differing ligand : G4 ratios. F22T is a 22-mer analogue of the F21T sequence.

conformational transition from a hybrid to a more parallel-type G4 form.

Analysis of the equilibrium conformation for compound **1** was undertaken with classical molecular mechanics (MM2 and AMBER) force fields and by *ab initio* (STO-3G) calculations. The MM2 analysis suggested that overall arrangement involved coplanar rings. This somewhat implausible conclusion was at variance with results from the AMBER force field and the *ab initio* calculations, which concurred in suggesting a twisted conformation due to the repulsive effects of the two furanoid methyl groups on the attached phenyl rings. The barrier between planar and twisted conformations is likely to be low so a qualitative analysis of plausible G4-bound conformations for compounds **1** and **2** has been undertaken. The NMR structure of an anti-parallel human telomeric G4 complexed with a telomestatin derivative has been used as a starting-point.²² Compounds **1** and **2** have similarities to the overall shape and curvature of telomestatin (Fig. 4), albeit in an acyclic manner. The NMR structure shows that the telomestatin derivative is non-planar and its out-of-plane distortions complement the non-planarity of the G-quartet to which it is bound. The qualitative low-energy structures proposed for compounds **1** and **2** incorporate such distortions, which still enable effective π - π stacking onto the terminal G-quartet of this G4, as well as some additional stabilisation from a thymine. The modelling show that the *meta* position of the dihydroimidazole substituents in **1** and the tetrahydropyrimidine substituents in **2**, is crucial for shape similarity to telomestatin. The *para* substitution in the control compound **7** enables it to effectively bind to the minor groove of duplex DNA, which is not possible for any of the six compounds arising from the screen.

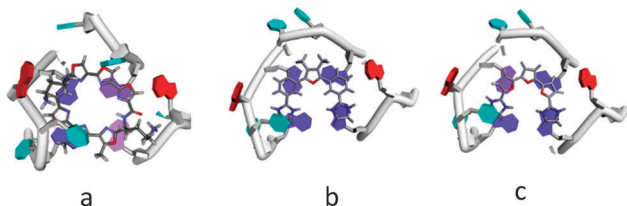


Fig. 4 Plots showing the terminal G-quartet of anti-parallel G4 complexes, with guanines shown in dark blue, thymine in cyan and adenines in red. Ligands are shown in stick representation. (a) The telomestatin derivative complex,²² (b) results of qualitative modelling with compound **1** and (c) with compound **2**.

Several of the compounds showed low μ M anti-proliferative activity (Table 2) in a cancer cell line panel [A549 (lung cancer), MCF7 (breast cancer), RCC4 and 786-O (renal cancer), Panc1 and Mia-PaCa2 (pancreatic cancer), ALT (transformed WI38 lung fibroblast cells characterised by Alternative Maintenance of Telomeres) and WI38 (non-transformed lung fibroblast cells)].

The two poly-furan compounds **2** and **6** have low anti-proliferative activity, even though both have G4-stabilising activity comparable to the other four compounds in the group. This may be due to cell uptake and nuclear localisation problems as well as limited aqueous solubility; the lack of observed precipitation during the SRB assay supports the former suggestions. Compounds from the mono-furan and mono-thiophene series on the other hand show activity in several cell lines in the low- μ M range. The lung cancer cell line A549 is slightly more sensitive to these compounds, although we do not currently have a molecular explanation for this.

We cannot exclude the possibility of non-G4 targets being involved in these cellular effects: experiments are currently underway to examine links with G4 affinity for these compounds. G4 selectivity appears to be limited even though for the two lead compounds at least, duplex DNA is less likely to be a target. This suggests that they may be acting as poly-targeting agents, affecting a number of genes and oncogenes involved in cellular proliferation and also that the compounds do not have identical cellular targets. It is notable that compounds **3** and **6** have high selectivity for several of the cancer lines compared to the normal WI38 line and have some activity in the ALT line, which maintains telomere length by non-telomerase mechanisms. This suggests that telomere maintenance rather than telomerase *per se*, is being targeted.

We report here that screening putative ligands using a HTS-FRET assay against a panel of G4s with a duplex control sequence, has resulted in the discovery of *meta*-substituted bisphenyl-monofurans as a novel G4 stabilizing chemotype. A similar chemotype, with a urea group replacing the furan ring, has been reported²³ as having high G4 affinity. These compounds are structurally-simple, conformationally flexible and chemically readily accessible with MW < 400 Da. They have G4 stabilisation ability comparable to those previously observed with polycyclic heteroaromatic compounds²¹ (*cf.* compound **8** (4,9-bis((3-(4-methyl-piperazin-1-yl)propyl)amino)-2,7-bis(3-morpholinopropyl)benzo-*[lmn]*[3,8]phenanthroline-1,3,6,8(2*H*,7*H*)-tetraone) in Table 1), but with low duplex DNA affinity. They inhibit cancer cell growth at low μ M/high nM levels, suggesting that these or related compounds may have potential as drug-like poly-quadruplex targeting agents.

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