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Introduction

The G-quadruplex structure formed by tandem guanine (G) nucleotides, which are incorporated in nucleic acid sequences, has illuminated a new role for DNA in biology.¹ G-quadruplexes as one of the higher-order DNA and RNA structures have been identified in G-rich human telomeres and gene promoter regions. G-quadruplex as a therapeutic target has attracted a growing interest, due to its profound roles in a wide spectrum of diseases, such as cancer, diabetes and cardiovascular disorders.² Several organic probes have been reported to recognize DNA G-quadruplex via diverse interaction modes.³ Stabilization of G-quadruplex structures in the promoter region of certain oncogenes is an emerging field in anticancer drug design.⁴ Human c-myc gene is one of these oncogenes, and Gquadruplexes have been proven to be the transcriptional controller of this gene.

Human c-myc gene or its product is a central regulator of cellular proliferation and cell growth, and over-expression of cmyc gene is related to various cancers.⁵ C-myc is the first and most extensively studied system of promoter G-quadruplex formation. The major G-quadruplex formed in the c-myc

Selective recognition of c-myc promoter G-quadruplex and down-regulation of oncogene c-myc transcription in human cancer cells by 3,8a-disubstituted indolizinone†

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C-myc promoter G-quadruplex is a very important target for developing anti-cancer drugs. However, highly selective recognition of DNA c-myc G-quadruplex with parallel configuration is also a very challenging problem. Here, we showed a new type of N-containing alkaloid, 3,8a-disubstituted indolizinone, which adopted a distorted configuration. 6-Methyl-3-(naphthalen-2-yl)-8a-(4 methylpyridin-2-yl)-indolizinone could selectively recognize DNA c-myc G-quadruplex leading to an obvious fluorescence enhancement by 11-fold, and meanwhile the G-quadruplex can be stabilized up to 90 °C. Further, it could down-regulate the transcription of oncogene c-myc in both human non-small cell line (A549) and human laryngeal epithelial cell line (Hep-2). PAPER

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promoter is a parallel stranded structure with two G3NG3 single-nt strand-reversal loops, and parallel-stranded structures are found to be more common in the promoter G-quadruplexes.¹ Some known G-quadruplex-interactive compounds have become prospective anticancer agents that display relatively low cytotoxicity. In particular, Quarfloxin, a first-in-class G-quadruplex-interactive drug reached phase 2 clinical trials, which was considered to be a c-myc G-quadruplex ligand at first.⁶ Quarfloxin is a fluoroquinolone derivative with antineoplastic activity, and another known c-myc G-quadruplex ligand quindoline is a derivative of the natural product cryptolepine,⁴ and both of them are N-containing alkaloids. Here we reported a new type of N-containing alkaloid, indolizinone (Scheme 1), which can selectively recognize DNA c-myc G-quadruplex (DNA MycG4) and down-regulate oncogene c-myc.

Results

Indolizinones are N-heterocycles that have only recently appeared in the chemical literature,⁷ and to the best of our

Scheme 1 Structure of 6-methyl-3-(naphthalen-2-yl)-8a-(4-methylpyridin-2-yl)-indolizinone (indolizinone).

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knowledge, any application of which wasn't found to be reported. Recently, we firstly reported a copper-catalyzed domino reaction to construct functionalized 3,8a-disubstituted indolizinones, and this protocol was simple and highly efficient.⁸ In the solid state, 3,8a-disubstituted indolizinones adopted distorted configuration, and thus they provided no obviously strong fluorescence spectra in solution due to the lack of a larger conjugated π system. Most of DNA MycG4 ligands that have been previously reported are N-heterocycles with planar π systems, which can't distinguish DNA MycG4 from other configurations of G-quadruplexes. In this report, we found 6-methyl-3-(naphthalen-2-yl)-8a-(4-methylpyridin-2-yl) indolizinone (indolizinone) here could selectively recognize DNA MycG4, leading to its greatly enhanced fluorescence intensity. We firstly compared the fluorescence intensity of indolizinone with five oligonucleotides of known G4 structures (Table S1†): intermolecular parallel configuration H7, intramolecular hybrid-type configuration M24, intramolecular antiparallel configuration A24, intramolecular parallel configuration Kit and Myc, relative to the signal of indolizinone alone. As shown in Fig. 1, indolizinone itself at 4 μ M gives a very weak fluorescence signal. While addition of 8 μ M DNA MycG4, and then standing for 24 hours, the fluorescence signal of indolizinone lead to an obvious enhancement by 11-fold. Addition of other types of G-quadruplexes, such as intermolecular parallel type H7 led to 0.5-fold enhancement, and for intramolecular types, parallel DNA c-Kit 4.5-fold, hybrid-type M24 1.6-fold and anti-parallel type A24 2.5-fold, respectively. These results indicated that indolizinone preferred to recognize DNA MycG4. RSC Advances

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Furthermore, fluorescence titrations were carried out to evaluate the binding constants of indolizinone with these five nucleic acid sequences (Fig. S1–S3†). A 1 : 1 binding model for indolizinone with DNA G-quadruplex was confirmed by the job's plot analysis. The typical titration curves and fitting results according to the 1 : 1 binding model are presented. The binding constants for indolizinone with Myc and Kit are (9.91 \pm 0.65) \times 10^5 and $(5.77 \pm 1.60) \times 10^5$, respectively. The binding strength to Myc was obviously more strong than to Kit. In comparison with other intramolecular DNAG4s, the binding strength to A24 or M24 was much weaker, with binding constants (2.43 ± 0.37) \times 10⁴ and (1.88 \pm 0.65) \times 10⁴, respectively. Of note, the binding

Fig. 1 Fluorescence intensity of indolizinone with five varied DNA Gquadruplexes in a 20 mM Tris HCl (40 mM K⁺, pH 7.0) solution at room temperature, $E_x = 423$ nm.

Then we subsequently performed this assay on a larger set of 25 nucleic acid sequences. Structures of these oligonucleotides have been characterized, and these oligonucleotides have been used in the G4-Fluorescence Intercalator Displacement (FID) assay.⁹ All sequences are listed in Table 2, and results are presented in Fig. 2. Fifteen of the 25 sequences are likely to form G4s. The fluorescence enhancements of indolizinone interacted with the parallel configuration G4 in the promoter region (c-myc, c-kit1, c-kit2, bcl-2, AKT1, VAV1, 35B1, 27Kras) were 11 fold, 4.0-fold, 3.7-fold, 2.5-fold, 4.0-fold, 5.2-fold, 4.3-fold, and 3.4-fold, respectively. And the human telomeric DNA Gquadruplex 22Ag and 45Ag, which adopt anti-parallel configuration in the presence of K^+ , led to the fluorescence enhancements of indolizinone with the value F/F_0 1.5-fold and 2.8-fold, respectively. And meanwhile, in the presence of another five Gquadruplex forming sequences (21CTA, TBA, Ceb25, CT4, CGG12), and other non-G-quadruplex DNA templates including single-stranded (G-tripl, dT30), double-stranded (ds26, 19AT, dx12, PS1c), stem-loop (GCdx), triplex, and trinucleotide (CTG12, CAG12), all these oligonucleotide sequences led to a $F/F₀ < 2$. Indolizinone showed higher response towards intramolecular DNA G-quadruplex with parallel configuration, especially most to DNA MycG4, comparing to other types of Gquadruplex and non-G-quadruplex templates. While it presented very weak response to intermolecular parallel Gquadruplex H7, just leading to 0.5-fold enhancement, probably attributing to its lack of loop-binding sites.

To evaluate the effect of indolizinone on the specific recognition of DNA MycG4, we applied circular dichroism to monitor the melting temperature (T_m) of DNA MycG4 in the presence of 1 equiv. molar ratio of indolizinone. The change of T_m in the folded and unfolded quadruplex structure upon interacting with ligand provides evidence of thermal stabilization of DNA structure.¹⁰ DNA MycG4 was incubated with indolizinone for 24 h, the T_m values for DNA MycG4 were calculated as shown in Fig. 3. Indolizinone shows the preference to stabilize DNA MycG4 by raising the temperature up to 90 $^{\circ}$ C monitored at 265 nm. In the contrast, indolizinone shows a weak ability obviously to stabilize other types of G-quadruplexes (see Fig. S4†).

Based on the emission enhancement and the absorption spectral results, fluorescence intensity decay measurements

Table 1 F/F_{0s} and binding constants for the indolizinone with different DNAG4s determined from the fitted curves, $E_m = 494$ nm

Oligonucleotides	F/F_0^a	Binding constants
Myc	11	$(9.91 \pm 0.65) \times 10^5$
Kit	4.5	$(5.77 \pm 1.60) \times 10^5$
A24	2.5	$(2.43 \pm 0.37) \times 10^4$
M24	1.6	$(1.88 \pm 0.65) \times 10^4$
H7	0.5	< 10

^a The concentrations of indolizinone and DNAG4 are 4 μ M and 8 μ M, respectively.

Table 2 Oligonucleotides used in the study

were conducted to demonstrate the strong binding of indolizinone with DNA MycG4. The fluorescence decay of indolizinone was fast (1 ns) in solution due to the highly feasible torsional

Fig. 3 CD melting profile of 4 μ M DNA MycG4 alone or in the presence of 1 equiv. molar ratio of indolizinone in a 20 mM Tris HCl (40 mM K+, pH 7.0) solution.

relation channel in the excited state. However, the lifetimes increased to the nanosecond level in the presence of DNA MycG4. As shown in Fig. 4, the fluorescence decay was much lower for indolizinone in the presence of 1 equiv. or 2 equiv. molar ratio of DNA MycG4, with the τ value of 3.54 ns and 3.71 ns, respectively. This result indicated that the interaction between indolizinone and DNA MycG4 hinders the rotation of indolizinone and increases its lifetime.

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Fig. 4 Fluorescence decay traces of indolizinone (4 μ M) in the absence and presence of DNA MycG4 in a 20 mM Tris HCl (40 mM K⁺, pH 7.0) solution.

Oncogene c-myc encodes an important transcriptional regulator c-myc protein, which is involved in cell proliferation, senescence, and apoptosis.¹¹ Because c-myc promoter region contains G-quadruplex forming G-rich sequence, its binding with indolizinone could increase its stability resulting in downregulation of oncogene c-myc transcription, which would be identified using real-time RT-PCR.¹² To test the above hypothesis and understand the effect of indolizinone on oncogene cmyc, two cancer cell lines with different translocation break points within the oncogene c-myc were tested. Human nonsmall cell line (A549) and human laryngeal epithelial cell line (Hep-2) were obtained from the National Infrastructure of Cell Line Resource (Beijing, China). As shown in Fig. 5, indolizinone

Fig. 5 qRT-PCR to determine the transcription of oncogene c-myc in the A549 and Hep-2 cells treated with indolizinone.

could down-regulate the transcription of oncogene c-myc in both A549 and Hep-2 cells, after treated with indolizinone at 0.5, 1.0, 2.0, 4.0, and 8.0 μM for 48 h. The total RNA was extracted and reverse transcripted to cDNA. This cDNA was then used as a template for specific RCR amplification of the c-myc sequence and controlled by β -actin.

Conclusions

Indolizinones are N-heterocycles that have only recently appeared in the chemical literature, which can be prepared from the simple and achievable starting materials efficiently. Here, we firstly reported the indolizinone has the potential ability to recognize DNA MycG4 by fluorescence spectra and varied temperature CD spectra. When indolizinone complexed with DNA MycG4 in the ratio of $1:2$, and thus it can give an obvious fluorescence enhancement by 11-fold, and meanwhile the DNA MycG4 can be stabilized up to 90 \degree C. Further, we found indolizinone could down-regulate the transcription of oncogene c-myc in both of human non-small cell line (A549) and human laryngeal epithelial cell line (Hep-2) by real-time RT-PCR experiment. We believed that indolizinone will have a potential application in the exploration of anticancer drugs. For Advances
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Conflicts of interest

There are no conflicts to declare.

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Notes and references

- 1 Selected recent reviews, see: (a) Q. Cao, Y. Li, E. Freisinger, P. Z. Qin, R. K. O. Sigel and Z. W. Mao, Inorg. Chem. Front., 2017, 4, 10–32; (b) Y. W. Chen and D. Z. Yang, Curr. Protoc. Nucleic Acid Chem., 2012, 17.5.1–17.5.17; (c) K. M. Miller and R. Rodriguez, Expert Rev. Clin. Pharmacol., 2011, 4, 139–142; (d) J. T. Davis, Angew. Chem., Int. Ed., 2004, 43, 668–698; (e) L. H. Hurley, R. T. Wheelhouse, D. Sun, S. M. Kerwin, M. Salazar, O. Y. Fedoroff, F. X. Han, H. Y. Han, E. Izbicka and D. Von Hoff, Pharmacol. Ther., 2000, 85, 141–158; (f) T. M. Ou, Y. J. Lu, J. H. Tan, Z. S. Huang, K. Y. Wong and L. Q. Gu, ChemMedChem, 2008, 3, 690–713.
- 2 Q. Li, J. F. Xiang, Q. F. Yang, H. X. Sun, A. J. Guan and Y. L. Tang, Nucleic Acids Res., 2013, 41, D1115–D1123.
- 3 Selected organic probes: (a) J. Mohanty, N. Barooah, V. Dhamodharan, S. Harikrishna, P. I. Pradeepkumar and A. C. Bhasikuttan, J. Am. Chem. Soc., 2013, 135, 367–376; (b) J. Dai, M. Carver, L. H. Hurley and D. Z. Yang, J. Am. Chem. Soc., 2011, 133, 17673–17680; (c) D. Koirala,

S. Dhakal, B. Ashbridge, Y. Sannohe, R. Rodriguez, H. Sugiyama, S. Balasubramanian and H. Mao, Nat. Chem., 2011, 3, 782–787; (d) S. J. Xu, Q. Li, J. F. Xiang, Q. F. Yang, H. X. Sun, A. J. Guan, L. X. Wang, Y. Liu, L. J. Yu, Y. H. Shi, H. B. Chen and Y. L. Tang, Nucleic Acids Res., 2015, 43, 9575–9586; (e) R. Haudecoeur, L. Stefan, F. Denat and D. Monchaud, J. Am. Chem. Soc., 2013, 135, 550–553; (f) M. Wang, Z. Mao, T. S. Kang, C. Y. Wong, J. L. Mergny, C. H. Leung and D. L. Ma, Chem. Sci., 2016, 7, 2516.

- 4 (a) R. V. Brown, T. Wang, V. R. Chappeta, G. H. Wu, B. Onel, R. Chawla, H. Quijada, S. M. Camp, E. T. Chiang, Q. R. Lassiter, C. Lee, S. Phanse, M. A. Turnidge, P. Zhao, J. G. Garcia, V. Gokhale, D. Z. Yang and L. H. Hurley, J. Am. Chem. Soc., 2017, 139, 7456–7475; (b) T. M. Ou, Y. J. Lu, C. Zhang, Z. S. Huang, X. D. Wang, J. H. Tan, Y. Chen, D. L. Ma, K. Y. Wong and J. C. Tang, J. Med. Chem., 2007, 50, 1465–1474; (c) K. H. Leung, H. Z. He, B. He, H. J. Zhong, S. Lin, Y. T. Wang, D. L. Ma and C. H. Leung, Chem. Sci., 2015, 6, 2166. Paper Worden College, Y. Samondr, R. Redigion, J. K. C. Ham, D. Von Hoff, K. Andres and W. G. Ricence. Published on 2017. This article is licensed under a Creative Commons and T. Man, No Creative Commons Articles. 2013. At
	- 5 L. M. Resar, E. Emison, S. Kim, Q. Li, J. E. Prescott, D. Wonsey and K. Zeller, Exp. Cell Res., 1999, 253, 63–77.
	- 6 D. Drygin, A. Siddiqui-Jain, S. O'Brien, M. Schwaebe, A. Lin, J. Bliesath, C. B. Ho, C. Proffitt, K. Trent, J. P. Whitten,
- J. K. C. Lim, D. Von Hoff, K. Anderes and W. G. Rice, Cancer Res., 2009, 69, 7653–7661.
- 7 A. R. H. Narayan and R. Sarpong, Org. Biomol. Chem., 2012, 10, 70, and references cited therein.
- 8 B. Q. Hu, L. X. Wang, G. Shen, Y. L. Tang and L. Yang, RSC Adv., 2015, 5, 100097.
- 9 (a) V. Gabelica, R. Maeda, T. Fujimoto, H. Yaku, T. Murashima, N. Sugimoto and D. Miyoshi, Biochemistry, 2013, 52, 5620-5628; (b) A. R. Faverie, A. Guédin, A. Bedrat, L. A. Yatsunyk and J. L. Mergny, Nucleic Acids Res., 2014, 1– 8; (c) S. J. Xu, Q. Li, J. F. Xiang, Q. F. Yang, H. X. Sun, A. J. Guan, L. X. Wang, Y. Liu, L. J. Yu, Y. H. Shi, H. B. Chen and Y. L. Tang, Sci. Rep., 2016, 6, 24793–24801.
- 10 L. X. Wang, Q. Shang, Q. Li, J. F. Xiang, Y. Liu, A. J. Guan, H. X. Sun, L. J. Yu and Y. L. Tang, Tetrahedron, 2015, 71, 4982.
- 11 T. Simonsson and M. Henriksson, Biochem. Biophys. Res. Commun., 2002, 290, 11–15.
- 12 T. M. Ou, J. Lin, Y. J. Lu, J. Q. Hou, J. H. Tan, S. H. Chen, Z. Li, Y. P. Li, D. Li, L. Q. Gu and Z. S. Huang, J. Med. Chem., 2011, 54, 5671–5679.