

Communication

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Luotonin A. A Naturally Occurring Human DNA Topoisomerase I Poison

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The demonstrated clinical utility of two camptothecin analogues as antitumor agents¹ has prompted intensive efforts to identify additional clinical candidates in this class.² Accordingly, study of the mechanism of action of the camptothecins has been of great interest, as it may afford insights leading to improved therapeutic agents. A key biochemical target for CPT (1) is the covalent binary complex formed between DNA and topoisomerase I during DNA relaxation; stabilization of this complex by CPT is believed to lead to cell death.³



Luotonin A (2) is a pyrroloquinazolinoquinoline alkaloid extracted from the Chinese medicinal plant Peganum nigellastrum.4 Luotonin A is cytotoxic toward the murine leukemia P-388 cell line (IC₅₀ 1.8 μ g/mL), although the mechanism is unknown.^{4a,5} There are obvious structural similarities between CPT (1) and luotonin A, notably in identical rings A-C. The greatest differences are in ring E, which is known to be critical for CPT function as a topoisomerase I inhibitor and antineoplastic agent.⁶ Alterations in the lactone ring or 20-OH group typically render CPT dysfunctional, although a few exceptions have been reported.2g,7 The lactone of CPT has long been known to be quite electrophilic,8 and most structure-activity studies have supported a relationship between lactone electrophilicity and the ability of CPT to stabilize the topoisomerase I-DNA covalent binary complex.6 It has been suggested that this putative relationship may reflect the transient covalent attachment of CPT to the topoisomerase I-DNA complex.^{3b,9} In this context, the lack of functionality in the E-ring of luotonin A argues against its ability to function in the same fashion as CPT.

Presently, we demonstrate that despite the lack of E-ring functionality, luotonin A stabilizes the human DNA topoisomerase I–DNA covalent binary complex and mediates topoisomerase I-dependent cytotoxicity in intact cells. Shown in Figure 1 is the effect of CPT and luotonin A on the stabilization of the topoisomerase I–DNA binary complexes in a ³²P-end labeled 222 bp DNA duplex. In common with CPT, luotonin A¹⁰ effected concentration-dependent stabilization of the enzyme–DNA binary complex. While luotonin A was less potent than CPT, stabilization was observed at the same sites for both; the identical effect, as judged by electrophoretic analysis, argues that the chemistry of cleavage was the same for both. Neither **1** nor **2** had any measurable effect on DNA in the absence of topoisomerase I.

While not thought to contribute to its antitumor activity, CPT also inhibits topoisomerase I-mediated relaxation of supercoiled DNA when present at high concentrations. This is illustrated in Figure 2 at $100-500 \ \mu$ M CPT concentrations in the presence of



1 2 3 4 5 6 7 8 9

substrate was 3'-³²P end labeled on the scissile strand. Human topoisomerase I-mediated cleavage reactions were incubated at 37 °C for 1 h and then digested with proteinase K. Lanes 1 and 2, Maxam–Gilbert sequencing reactions; G, and C + T, respectively. Lane 3, DNA alone; lane 4, 50 μ M CPT; lane 5, 50 μ M luotonin A; lane 6, topoisomerase I and 50 μ M CPT; lane 7, topoisomerase I and 50 μ M luotonin A; lane 8, topoisomerase I and 10 μ M luotonin A; lane 9, topoisomerase I and 2 μ M luotonin A. No significant DNA cleavage was observed in the presence of topoisomerase I + DNA alone. In a parallel experiment, cleavage at the site marked by an arrow was stabilized to the extents of 58%, 14%, and 1% at 50, 10, and 2 μ M luotonin A, respectively, relative to that achieved with 50 μ M CPT.



Figure 2. Effect of luotonin A on human topoisomerase I-mediated DNA relaxation. Supercoiled pSP64 plasmid DNA was incubated at 37 °C for 10 min as indicated. Lane 1, supercoiled pSP64 plasmid DNA alone; lane 2, DNA + 0.1 ng of topoisomerase I; lanes 3–5, DNA + topoisomerase I + 500, 200, and 100 μ M of camptothecin, respectively; lanes 6–8, DNA + topoisomerase I + 500, 200, and 100 μ M of luotonin A, respectively.

0.1 ng of human topoisomerase I. As shown in the Figure, luotonin A (2) had no effect on DNA relaxation under the same conditions, although 2 did weakly inhibit plasmid DNA relaxation by calf thymus DNA topoisomerase I (not shown).

To evaluate the possible cytotoxic effects resulting from stabilization of the enzyme–DNA binary complex, luotonin A was evaluated in a strain of *Saccharomyces cerevisiae* lacking yeast topoisomerase I, but harboring a plasmid having the human topoisomerase I gene under the control of a galactose promoter.¹¹ As shown in Table 1, 1 μ M CPT had no effect when this yeast strain was grown on raffinose. However, 1 μ M CPT caused 74% inhibition after 2 days when the same yeast strain was grown on galactose, resulting in topoisomerase I expression. Luotonin A produced 36% inhibition of growth when employed at 1 μ M Table 1. Human Topoisomerase I-Dependent Cytotoxicity of CPT (1) and Luotonin A (2) toward *S. cerevisiae*^a

compound	concentration (µM)	% inhibition on growth medium	
		raffinose	galactose
CPT (1)	1.0	0	74
luotonin A (2)	1.0	0	36
	0.5	0	23

^{*a*} Inhibition of RS321Nph-TOP1 grown in minimal medium containing 3% raffinose or galactose for 2 days at 30 °C.

concentration in the presence of galactose.¹² In replicate experiments, lutonin A exhibited IC₅₀ values from 5.7 to 12.6 μ M in the presence of galactose. The comparable values for CPT were 0.74–0.86 μ M.

The closely analogous effects of CPT and luotonin A on stabilization of the topoisomerase I–DNA binary complex, and on the production of human topoisomerase I-dependent cytotoxicity in yeast, suggest that the two agents likely function in the same fashion. This conclusion identifies a putative biochemical locus for the cytotoxic action of luotonin A and has important implications both for the mechanism of inhibition of topoisomerase I function by CPT and for the design of new CPT analogues.

At a mechanistic level, it seems clear that no electrophilic E-ring lactone is needed for stabilization of the topoisomerase I–DNA covalent binary complex. Despite the apparent correlation between E-ring lactone electrophilicity and topoisomerase I inhibitory activity,⁶ the present findings add to the weight of evidence that argues against the covalent attachment of CPT to the enzyme–DNA binary complex.

In terms of inhibitor design, recent X-ray crystallographic studies¹³ and computational models¹⁴ suggest a role for the 20(S)-OH group in interaction with DNA topoisomerase I, possibly through hydrogen bonding to the enzyme. This interaction is further supported by the lack of activity of 20(R) CPT,^{15,16} as well as 20deoxyCPT.16,17 The 20-chloro, bromo, and amino derivatives of CPT have been shown to stabilize the topoisomerase I-DNA binary complex and to produce human topoisomerase I-dependent cytotoxicity in yeast, albeit with somewhat reduced potency.¹⁶ While it seems likely that functional groups properly oriented at the 20position of CPT can contribute to the stability of the ternary complex formed with topoisomerase I and DNA, the present results obtained with luotonin A suggest that, even in the absence of any functional group at the 20-position, an aromatic E-ring can confer reasonable stability to the formed ternary complex. This may reflect a stacking interaction first posited by Kohn, Pommier and co-workers14a,18 and now present in many models of the ternary complex.13 It seems reasonable to suggest that the presence of appropriate functional groups on the E-ring might further modulate the interaction of luotonin A with the topoisomerase I-DNA binary complex. In any case, the present results make it clear that an electrophilic E-ring is not essential for stabilization of the topoisomerase I-DNA covalent binary complex.

At a practical level, it may be noted that elaboration of the E-ring of CPT is perhaps the most challenging aspect of the synthesis of this natural product. The several concise syntheses already reported for luotonin A^{5,10,19} suggest that the preparation of luotonin A analogues should be straightforward.

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