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Fundamental Relationships Between Structure, Reactivity, and Biological Activity for the Duocarmycins and CC-1065

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Introduction

The duocarmycins (**1** and **2**)¹ belong to a small family of natural products (Figure 1) that also include yatakemycin (**3**)² and CC-1065 (**4**).³ Their exceptionally potent cytotoxic activity is derived from their ability to bind and alkylate DNA in AT-rich regions of the minor groove.^{4–8} This family of natural products incorporates a remarkable set of molecular features integrated into compact structures capable of simultaneously expressing multiple functions.⁹

Much like their predecessors distamycin and netropsin, the overall curvature and shape of the molecules leads to their preferential binding in the narrower, deeper, AT-rich minor groove where stabilizing van der Waals contacts are maximized (shape-selective recognition).^{8–11} The alkylation subunit vinylogous amide conveys a remarkable stability to a cross-conjugated and otherwise reactive cyclopropane. We have suggested that the disruption of this vinylogous amide conjugation by way of a DNA minor groove binding-induced conformational change brings the cyclopropane into conjugation with the cyclohexadienone activating it for nucleophilic attack, and provides the catalysis for the DNA alkylation reaction (shape-dependent catalysis, Figure 2).^{8,12,13} Significantly, these compounds are relatively unreactive until they reach their biological target, where they are selectively activated for DNA alkylation (target-based activation).¹⁴ Unique to this class, this activation occurs without a chemical change or reaction of the molecule, rather it occurs simply through a binding-induced conformational change in the compound. Studies that define the key relationships between structure and reactivity and quantitate the magnitude of these effects are detailed herein.

Just as important as their unique method of activation is a parabolic relationship between intrinsic reactivity and biological potency that has emerged from studying the natural products and numerous synthetic analogues (Figure 3).¹⁵ Presumably this reflects the requirement for sufficient stability for the compounds to reach their biological target balanced against the need for sufficient reactivity to efficiently alkylate DNA once they do. Establishment of this parabolic relationship defined what this optimal balance of reactivity and stability is for this class of natural products and key studies designed to define this parabolic relationship and exploit its use are also reviewed herein. Combined, these studies highlight remarkable fundamental relationships between structure, reactivity, and biological potency embodied in this class of natural products.

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Importance and Role of the Vinylogous Amide: Structure versus Reactivity and Reaction Regioselectivity

The synthesis, reactivity comparison, and correlation with X-ray structures of a key series of alkylation subunit analogues established the magnitude and role of the vinylogous amide stabilization, revealed the stereoelectronic alignment of the reacting cyclopropane that dictates the regioselectivity of ensuing nucleophilic additions, and defined their dominant role in controlling the chemical reactivity and regioselectivity for this class of natural products. Moreover, their side-by-side comparison provides a depiction of the structural changes and reactivity effects that accompany a DNA binding-induced conformational change in **1-4** and indicates that they are of a magnitude to account for catalysis of the DNA alkylation reaction.

Thus, X-ray crystallographic studies of a structurally related and progressive series of 1,2,9,9a-tetrahydrocyclopropa[*c*]benz[*e*]indol-4-one (CBI)^a analogues afforded information regarding the extent of vinylogous amide conjugation (*c* bond length), the disposition of the attached acyl groups (χ_1 dihedral angle), the orbital alignment of the cyclopropane bonds, and their intimate relationship with both the stability of the molecule and the regioselective addition of nucleophiles to the cyclopropane. This series examined the effects of increasing the C-ring size from a 5-membered ring (CBI) as found in the natural products to a 6- and 7-membered ring system (2,3,10,10a-tetrahydro-1*H*-cyclopropa[*d*]benz[*f*]quinol-5-one (CBQ)¹⁶ and 1,2,3,4,11,11a-hexahydrocyclopropa[*c*]naphtho[2,1-*b*]azepin-6-one (CNA),¹⁷ respectively), Figure 4.

The reactivity increases as measured by solvolysis across the series are remarkable ($>10^4$) despite the subtle structural changes; CBQ¹⁶ and especially CNA¹⁷ exhibit substantial reactivity at pH 7 whereas both CBI and **8** are stable at this pH, Figure 4. Beautifully reflecting this remarkable increase in reactivity, the *c* bond lengths smoothly increase across this series (1.390, 1.415, and 1.428 Å) reflecting the progressive loss in vinylogous amide conjugation (Figure 5). Tracking with this increase in reactivity and loss of vinylogous amide stabilization, the χ_1 dihedral angles progressively increase (21°, 34°, and 86°) across the series as well, mimicking the structural and reactivity changes imposed by a DNA binding-induced conformational change that increases the χ_1 dihedral angle. Just as significant, the composite bond lengths of the cyclopropane increase as one progresses across the series, reflecting their weaker bond strengths, their increasing reactivity, and their increasing conjugation with the activating cyclohexadienone system. Additionally and unique to this series, the cyclopropane alignments with the cyclohexadienone π -system undergo remarkable changes as well, dictating the regioselectivity of nucleophilic attack.

The X-ray crystal structure of *N*-CO₂Me-CBI ($>20:1$ favoring ring-opening attack at C9 vs ring-expansion attack at C9a) reveals a favorable overlap between the orbital of the cyclopropane 8b-9 bond and the developing π -system of the phenol solvolysis product, whereas the orbital of the 8b-9a bond is much less effectively aligned to become a part of the developing aromatic π -system. The lengths of these two bonds reflect this variation in alignment, with that of the cleaved and stereoelectronically aligned 8b-9 bond (1.544 Å) being longer and weaker than the 8b-9a bond (1.521 Å). *N*-Boc-CBQ shows little preference for regioselective addition (3:2 ring-opening attack at C10 vs ring-expansion attack at C10a) and contains a cyclopropane that is nearly ideally conjugated and aligned with the π -system. Both cyclopropane bonds are perfectly bisected by the cyclohexadienone π -system, suggesting that the slight preference for addition to the least-substituted carbon is due to preferential cleavage of the longer and inherently weaker 9b-10 bond (1.543 Å), in addition to a potential steric preference for S_N2 addition to the least-substituted C10 carbon. Analysis of *N*-CO₂Me-CNA reveals a situation similar to, but reversed from that of *N*-CO₂Me-CBI. The bond to the more-substituted carbon (10b-11a) is now better aligned with the developing π -system of the product phenol, and it is

now the longest (1.565 Å) and weakest of the cyclopropane bonds. These observations correlate beautifully with the observed reaction regioselectivity of the CNA alkylation subunit (<1:20 ring-opening attack at C11 vs ring-expansion attack at C11a), which is completely reversed

^aAbbreviations:

DSA	duocarmycin SA
DA	duocarmycin A
epi-DA	<i>epi</i> -duocarmycin A
CBI	1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]benz[<i>e</i>]indol-4-one
CPI	1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]pyrrolo[3,2- <i>e</i>]indol-4-one
MeCPI	7-methyl-1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]pyrrolo[3,2- <i>e</i>]indol-4-one
CI	1,2,7,7a-tetrahydrocyclopropa[1,2- <i>c</i>]indol-4-one
CNA	1,2,3,4,11,11a-hexahydrocyclopropa[<i>c</i>]naphtho[2,1- <i>b</i>]azepin-6-one
CBQ	2,3,10,10a-tetrahydro-1 <i>H</i> -cyclopropa[<i>d</i>]benzo[<i>f</i>]quinol-5-one
iso-CBI	1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]benzo[<i>f</i>]indol-8-one
CMCFI	6-carbomethoxy-1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]furano[3,2- <i>e</i>]indol-4-one
CPyI	methyl 1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]pyrido[3,2- <i>e</i>]indol-4-one-7-carboxylate
MCBI	7-methoxy-1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]benz[<i>e</i>]indol-4-one
CCBI	7-cyano-1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]benz[<i>e</i>]indol-4-one
MCCBI	5-methoxycarbonyl-1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]benz[<i>e</i>]indol-4-one
HMCBI	5-hydroxymethyl-1,2,9,9a-tetrahydrocyclopropa[<i>c</i>]benz[<i>e</i>]indol-4-one
CPzI	7-methyl-1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]pyrazolo[4,3- <i>e</i>]indol-4-one
CBIn	1,2,9,9a-tetrahydro-1 <i>H</i> -cyclopropa[<i>c</i>]benz[<i>e</i>]inden-4-one
iso-DSA	6-methoxycarbonyl-1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]pyrrolo[2,3- <i>g</i>]indol-4-one
MeCTI	7-methyl-1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]thieno[3,2- <i>e</i>]indol-4-one
iso-MeCTI	6-methyl-1,2,8,8a-tetrahydrocyclopropa[<i>c</i>]thieno[2,3- <i>e</i>]indol-4-one
TMI	5,6,7-trimethoxyindole-2-carboxylate

from that observed with CBI. Notably, in all instances and contrary to an early mischaracterization,¹⁸ the ring expansion cleavage of the cyclopropane proceeds with clean inversion of stereochemistry at the reacting center indicative of a S_N2, not S_N1, nucleophilic addition.¹⁹ Thus, in addition to providing insights into their reactivity, the studies highlight the importance of the fused 5-membered C-ring, which serves to enforce a stereoelectronically-controlled nucleophilic addition to the least substituted cyclopropane carbon.

Additional modifications to the structures have provided further insight into the importance of the vinylogous amide and the linking amide (Figure 6).²⁰ Removing the carbonyl of the linking amide in CBI-TMI²¹ (**9**) afforded a compound **10** that failed to alkylate DNA even under forcing conditions (37 °C, 14 days, 10⁻¹ M agent vs 25 °C, 24 h, 10⁻⁶ M for **9**), that was biologically inactive displaying cytotoxic activity that was >10⁵-fold less potent than **9**, and that exhibited little solvolysis requiring exposure to pH 1 to detect a measurable reactivity. In addition to establishing the fundamental importance of the linking amide, the comparisons highlight the truly remarkable stability provided to this unusual ring system by full vinylogous amide conjugation that in turn is reduced (tuned) by N-acylation as found in the natural product structures. Its disruption by a DNA binding-induced conformational change is no longer possible with **10**, accounting for its complete lack of DNA alkylation. The disparate behavior of **9** versus **10** not only revealed that the linking amide plays an important role activating the alkylation subunit for nucleophilic attack, but that it is critical to catalysis of the DNA alkylation reaction.

Equally revealing was examination of 1,2,9,9a-tetrahydro-1*H*-cyclopropa[*c*]benz[*e*]inden-4-one (CBIn, **11**, Figure 7),²² which contains a carbocyclic skeleton lacking the nitrogen crucial for stability of the alkylation subunit. Consistent with intuitive expectations, CBIn proved sufficiently stable for characterization, but remarkably reactive toward nucleophilic addition. CBIn is 3200× less stable than its CBI counterpart at pH 3 and greater than 10³-10⁴× more reactive than CBI at pH 7, where CBI is stable. Solvolysis studies conducted over a pH range of 2-12 revealed no acid concentration dependence between pH 4-12 indicating that the reaction is uncatalyzed (versus acid-catalyzed) in this pH range, and suggesting that there is no basis for invoking an acid-catalyzed activation^{18,23} of these compounds when bound to DNA (pH 7.4, 0.0001-0.00004% protonation of phosphate backbone).

Additional insights into the behavior of this class of natural products was recently reported with the examination of **12**, which lacks the characteristic vinylogous amide stabilization and is incapable of activation by C4-carbonyl protonation (Figure 8). Preparation of **12** revealed that spirocyclization can occur through deprotonation of the indole NH, rather than the conventional phenol, but provides an intrinsically much more reactive compound.²⁴ It also exhibits a rate of solvolysis that is independent of pH. Although more reactive than its natural product counterpart, **12** exhibits an identical DNA alkylation selectivity as duocarmycin SA and a reduced, but significant level of biological activity (L1210 IC₅₀ = 5 nM).

These observations reaffirm several key features shown to contribute to the DNA alkylation selectivity of duocarmycin SA (DSA). First, the DNA alkylation selectivity is derived from their intrinsic non-covalent binding selectivity within the minor groove of DNA,^{10,13,25} and not from the source of alkylation catalysis, which differs between **1** and **12**. Like **12**, derivatives that lack the C4-carbonyl,^{10,25} analogues in which the C4-carbonyl has been relocated,²⁶ and those that utilize an alternative mechanism of catalysis²⁷ exhibit an indistinguishable DNA alkylation selectivity, albeit proceeding at altered rates. Second, the comparable behavior of **1** and **12** indicates that the alkylation selectivity cannot originate from a sequence-dependent DNA backbone phosphate protonation^{18,23,28} of the C4-carbonyl of **1**, as this feature is not present with **12**.

Although this alternative spirocyclization plays no present day role in the behavior of duocarmycin SA, it is possible that it is related to an evolutionary origin of **1** and its related natural products, Figure 9. Since it is reasonable to assume that the *p*-spirocyclization (through displacement of an activated primary alcohol) occurs as the last step of the biosynthesis of duocarmycin SA, it is tempting to suggest that such compounds first emerged without the ability to undergo *p*-spirocyclization, closing to the cyclopropane with participation of the indole NH as in **13** instead (Figure 9). Evolutionary pressures may have improved on this by subsequent aryl hydroxylation to afford the present day phenol precursors where the resulting increase in potency (10 pM vs 5 nM) would have provided the producing bacteria with a beneficial growth advantage.

A provocative indication that there is perhaps merit to this proposal emerges from the examination of the correlation between biological potency (cytotoxic activity, $-\log IC_{50}$) and chemical stability ($-\log k_{solv}$) where **13**, like **12**, would be projected to lie at an early stage of Nature's steps in the optimization of this class of DNA alkylating agents, Figure 10.

Fundamental Parabolic Relationship Between Reactivity and Biological Activity

A wide variety of synthetic duocarmycin SA analogues have been prepared and studied.²⁹ The 1,2,9,9a-tetrahydrocyclopropa[*c*]benz[*e*]indol-4-one (CBI) alkylation subunit has been the most extensively examined, due to the relative ease of preparation and its impressive properties compared to the natural products.³⁰ Initial studies described the effects of C7-substitution, which is para to the C4-carbonyl. The electronic effects of groups at this position were expected to probe their impact on reactivity and to address the question of whether protonation of the carbonyl was important for activation of these agents. Comparison of the parent unsubstituted CBI subunit with two derivatives, 7-cyano-1,2,9,9a-tetrahydrocyclopropa[*c*]benz[*e*]indol-4-one (CCBI, **14**)³¹ and 7-methoxy-1,2,9,9a-tetrahydrocyclopropa[*c*]benz[*e*]indol-4-one (MCBI, **15**)³² (Figure 11) established that the electronic nature of C7-substituents did not have a significant effect on the biological potency or intrinsic reactivity.

Despite the opposing electronic nature of the substituents found in CCBI and MCBI, only a 2-fold difference in solvolysis reactivity between the *N*-Boc derivatives was observed. The cytotoxic activity mirrors this trend, with *N*-Boc-CCBI being only 4-fold more potent than *N*-Boc-MCBI, and the full length agents (TMI, indole₂) were essentially indistinguishable from each other. In fact, a Hammett analysis of the acid-catalyzed solvolysis of MCBI, CBI, and CCBI revealed a very small ρ value (-0.30 at pH 3), indicative of little substituent effect on the reactivity.

More interestingly, both substituents had a beneficial impact on the DNA alkylation properties of the fully elaborated analogues. The DNA alkylation properties (rate, efficiency, and selectivity) of such MCBI and CCBI derivatives were essentially indistinguishable, but both alkylated DNA at rates and efficiencies that exceed the CBI derivatives. This suggested that it is not the electronic nature of the substituent, but rather the simple presence of a C7-substituent that serves to improve the DNA alkylation rate and efficiency. We have suggested that this is the result of the substituent extending the rigid length of the agent and increasing the extent of the DNA binding-induced conformational change when bound in the minor groove.³³ Consistent with this interpretation, incorporation of an electron-withdrawing C5-substituent (CO₂Me, **16**) subtly increased the intrinsic stability, but slowed the rate and efficiency of DNA alkylation, whereas the corresponding C7-derivative (**17**) displayed an analogous subtle decrease in reactivity, but displayed the increased rate and efficiency of DNA alkylation observed with both **14** and **15**.³⁴

In contrast to A-ring substituents, N² substituents have a huge impact on the properties of the molecules by virtue of the direct impact on the vinylogous amide. In addition to those highlighted in Figures 6 and 7, early studies enlisting a series of modified acyl derivatives of CBI indicated that their impact on reactivity was extraordinarily large and linear in a Hammett analysis ($\rho = -3.0$, $r = 0.983$) with the stronger electron-withdrawing substituents imparting the greater stability, and that this increased stability directly correlated with a near linear and equally large increase in cytotoxic potency ($r = 0.979$), Figure 12.³⁵

Until recently, the relationship between chemical stability and biological potency was restricted to the examination of such derivatives that possessed sufficient reactivity to alkylate DNA and was interpreted to reflect the ability of the chemically more stable derivatives to more effectively reach their biological target (DNA).³⁶ However, as the number of such compounds examined increased, the plot of such a correlation more closely resembled a parabolic relationship, but one in which the derivatives lying on the “too stable” side of the relationship were scarce. Consequently, in our efforts to define the full nature of this relationship, we began examining CBI derivatives that might accurately reflect the behavior of less reactive derivatives. A unique series of *N*-aryl CBI derivatives, in which the electronic properties of the aryl *p*-substituent could be systematically varied to alter the reactivity, were found to be remarkably stable relative to the typical *N*-acyl derivatives.¹⁵ This reactivity followed a well-defined correlation with σ_p ($\rho = 0.17$) in which electron-withdrawing substituents enhance and electron-donating substituents decrease the reactivity (solvolysis at pH 2, Figure 13).

X-ray structures of the full series of derivatives revealed that increasingly electron-withdrawing *p*-substituents effectively increase the diagnostic c bond length, decrease the d bond length, decrease the f bond length, and increase the χ_1 dihedral angle, all of which indicate an effect of diminishing the conjugation of the vinylogous amide (Figure 14), leading to increases in reactivity. Conversely, the presence of electron-donating *p*-substituents had the opposite effect, effectively increasing the conjugation of the vinylogous amide and further stabilizing the alkylation subunit to nucleophilic attack. Most significantly, the correlation of reactivity with cytotoxic activity is now reversed with the more reactive derivatives in this unusually stable series exhibiting the more potent activity. Beautifully, this series served to complete the parabolic relationship between intrinsic reactivity (stability) and cytotoxic activity providing the data shown in the shaded box in Figure 14.

In addition to the impact of attached substituents, there are structural features embedded in the skeletons of the natural products that also impact the reactivity of these compounds. It has been suggested that an internal hydrogen bond between the pyrrole NH and the C4-carbonyl of duocarmycin SA serves to activate the agent for nucleophilic attack.³⁷ To probe this feature, *iso*-duocarmycin SA (*iso*-DSA, **27**) was prepared and evaluated. In addition to addressing this question of H-bond activation, examination of this analogue would also serve to probe the effects of incorporating a H-bond donor directed towards the floor of the minor groove, and would determine whether a second vinylogous amide system would serve to further stabilize or destabilize the alkylation subunit.³⁸

Solvolysis studies revealed *N*-Boc-*iso*-DSA (**26**) to be slightly more reactive by a factor of two than its parent *N*-Boc-DSA (**8**). As a result, the *iso*-duocarmycin SA series was predictably less potent than the natural product, Figure 15. *N*-Boc-*iso*-DSA (**26**) was found to be 5-fold less potent than its natural product counterpart, nicely reflecting the 2-fold difference observed in the solvolytic reactivity and the same trend was observed for *iso*-duocarmycin SA (**27**). When **26** was placed on the plot of reactivity versus potency and given the anomalously potent activity of duocarmycin SA, its relative cytotoxic activity even more closely approached that expected of a derivative exhibiting its relative reactivity (Figure 16).

Thus, the enhanced reactivity of **26** versus **8** indicates that the putative internal H-bond of **8**, which has been removed in **26**, does not contribute to the activation of the compound towards nucleophilic addition. In addition, the incorporation of a second vinylogous amide in the molecule does not further stabilize the reactive cyclopropane, which would lead to greater potency and reduced solvolysis reactivity. In fact, the opposite appears to be true, suggesting that the additional cross-conjugated and competing vinylogous amide destabilizes the alkylation subunit.

Using the Parabolic Relationship: Rational Design of an Improved CC-1065

The parabolic relationship between reactivity and potency that emerged from these studies provides the opportunity to rationally design analogues that would sit at or near the pinnacle of the parabola, with a reactivity/potency relationship similar to that of duocarmycin SA. In our efforts, a series of benzothiophene-based analogues were designed and evaluated.³⁹ Intuitively, the replacement of the pyrrole NH of CC-1065 with the larger sulfur atom was expected to release some of the ring strain inherent with the pyrrole, creating a ring system closer in structure to that of CBI, and increasing the stability of the molecule. Additionally, the increased electronegativity of sulfur versus the pyrrole nitrogen was expected to further contribute to an increasingly stable alkylation subunit whose activity was expected to surpass that of its parent compound CC-1065. These intuitive arguments were supported by computational studies (AM1, MNDO) that predicted that the substitution of sulfur for NH in the CPI structure would increase the stability of the alkylation subunit.

The synthesis and examination of 7-methyl-1,2,8,8a-tetrahydrocyclopropa[*c*]thieno[3,2-*e*]indol-4-one (MeCTI, **30**) and 6-methyl-1,2,8,8a-tetrahydrocyclopropa[*c*]thieno[2,3-*e*]indol-4-one (*iso*-MeCTI, **31**) confirmed these expectations.³⁷ Solvolysis of *N*-Boc-MeCTI (**30**) and *N*-Boc-*iso*-MeCTI (**31**) revealed two compounds whose stabilities (pH 3, $t_{1/2}$ = 206 and 209 h, respectively) were substantially increased over that of their parent compound *N*-Boc-MeCPI (**29**, $t_{1/2}$ = 37 h). They also proved slightly more stable than *N*-Boc-DSA (**8**, $t_{1/2}$ = 177 h) and *N*-Boc-CBI (**5**, $t_{1/2}$ = 133 h), making them among the most stable alkylation subunits explored to date. Significantly, these compounds exhibited substantially increased biological potency versus CC-1065 and its derivatives. The natural enantiomers of *N*-Boc-MeCTI (**30**, IC₅₀ = 30 nM) and *N*-Boc-*iso*-MeCTI (**31**, IC₅₀ = 25 nM) were 10-fold more potent than *N*-Boc-MeCPI (**29**, IC₅₀ = 330 nM) and 2-3-fold more potent than *N*-Boc-CBI (**5**, IC₅₀ = 80 nM), beautifully falling in line with expectations based on their relative solvolytic reactivities. This is illustrated on the parabolic curve, where MeCTI and *iso*-MeCTI sit at or near the pinnacle of the relationship, just below the anomalously potent DSA alkylation subunit (Figure 18).

MeCTI-PDE₂ (**28**), the sulfur-based analogue of CC-1065 (**4**), displayed an identical DNA alkylation selectivity relative to the parent compound, displaying the characteristic enantiomeric differences. Significant in these comparisons is the observation that a single atom change in the skeleton of the alkylation subunit structure of CC-1065 (NH→S) predictably increased the stability of the natural product that in turn reliably led to a substantial increase in its biological potency.

Conclusions

The small family of natural products that include the duocarmycins, yatakemycin, and CC-1065 incorporate a remarkable combination of molecular features integrated into their compact structures controlling their DNA binding and ensuing alkylation reaction and their resulting exceptionally potent biological activity. The systematic total synthesis of an extensive series of analogues containing deep-seated structural changes have been utilized to define subtle relationships between structure, reactivity, reaction regioselectivity and biological potency

using fundamental chemical principles. Superimposed on a skeleton that selectively binds DNA minor groove AT-rich sequences (shape-selective recognition), the compounds incorporate a spirocyclopropylcyclohexadienone whose reactivity has been masked by a cross-conjugated and remarkably stabilizing vinylogous amide, taming what would ordinarily be an exceptionally reactive electrophile. We have suggested that disruption of this vinylogous amide conjugation through a DNA binding-induced conformational change activates the cyclopropane for a stereoelectronically-controlled nucleophilic attack serving as the catalysis for the DNA alkylation reaction (shape-dependent catalysis). These studies highlight the remarkable design Nature utilized to mask a reactive electrophile that is subsequently capable of selective activation by a unique mechanism upon reaching its biological target (target-based activation).

In addition to elucidating the mechanism of activation of these natural products, a detailed understanding of the relationship between intrinsic reactivity and biological potency resulted from studying analogues with deep-seated structural modifications. These studies defined a fundamental parabolic relationship between chemical reactivity and biological potency (cytotoxic activity) and identified the optimal balance between reactivity—stability, reflecting a stability required to effectively reach their biological target in a biological milieu while maintaining a sufficient reactivity to effectively alkylate DNA once they do. Remarkably, but perhaps not surprisingly, duocarmycin SA and yatakemycin incorporate an alkylation subunit that lies at the pinnacle of this relationship reflecting Nature's exquisite optimization of their DNA alkylation properties. Unique insights into Nature's potential steps in this evolutionary optimization were revealed with the discovery of an alternative, but less productive indole N⁵H spirocyclization of duocarmycin SA lacking a C4-phenol. Finally, as a result of the increased understanding of these compounds (reactivity vs potency), we described the rational design of a synthetic alkylation subunit (CTI) incorporating a single skeletal atom change that maintains the DNA alkylation properties of its parent natural product (CC-1065), but exhibited a greater chemical stability and resulting increased biological potency. This remarkably stable alkylation subunit analogue lies at the pinnacle of the parabolic relationship and constitutes one of the most stable derivatives to be characterized to date.

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Biography

Dale L. Boger received his B.S. degree in Chemistry from the University of Kansas in 1975, and Ph. D. degree from Harvard University in 1980. Following graduate school, he returned to the University of Kansas as a member of the faculty in the Department of Medicinal Chemistry, moved to Purdue University in 1986, and joined the newly founded Department of Chemistry at The Scripps Research Institute in 1991 as the Richard and Alice Cramer Professor of Chemistry. His research interests span the fields of organic and bioorganic chemistry, medicinal chemistry, the study of DNA-agent and protein-ligand interactions, and antitumor agents.

Karen S. MacMillan received her B.S. degree in Chemistry from the University of California, Davis in 2002 and her Ph.D. degree in Chemistry from The Scripps Research Institute in 2009 under the guidance of Professor Dale L. Boger. Currently, she is working as an Assistant Instructor in the Chemistry Core Facility within the Department of Biochemistry at the University of Texas Southwestern Medical Center.

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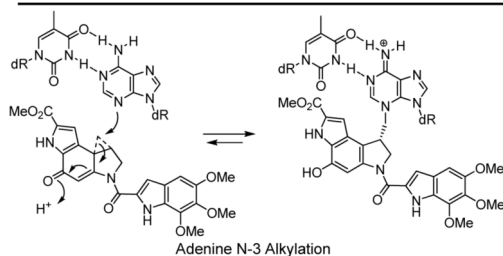
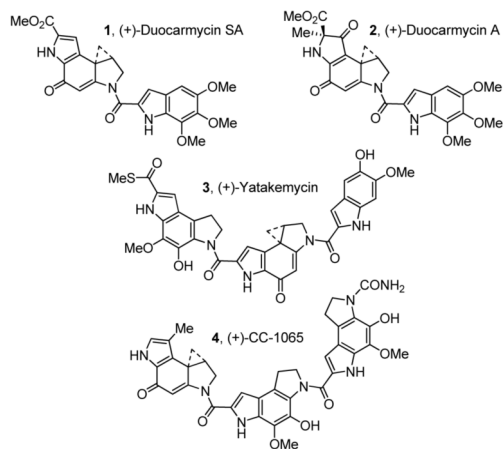
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- | | |
|---|--|
| <ul style="list-style-type: none"> • Minor Groove • Adenine N-3 Alkylation • S_N2 not S_N1 Addition • Stereoelectronic Control • Inherently Reversible Reaction • Catalysis: DNA Binding Induced Conformational Change • DNA Alkylation and Nucleophilic Addition Independent of H⁺ concentration @ pH ≥ 5-6: <i>Not Acid-catalyzed</i> | <ul style="list-style-type: none"> • Direct Relationship of Functional Stability and Potency • Unexpectedly Stable • Vinylogous Amide Stabilization • Both Enantiomers Effective |
|---|--|

DSA: 5'-AAAA vs CC-1065: 5'-AAAAA

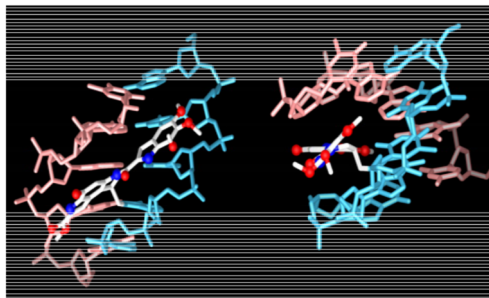


Figure 1.
Natural products and the DNA alkylation reaction.

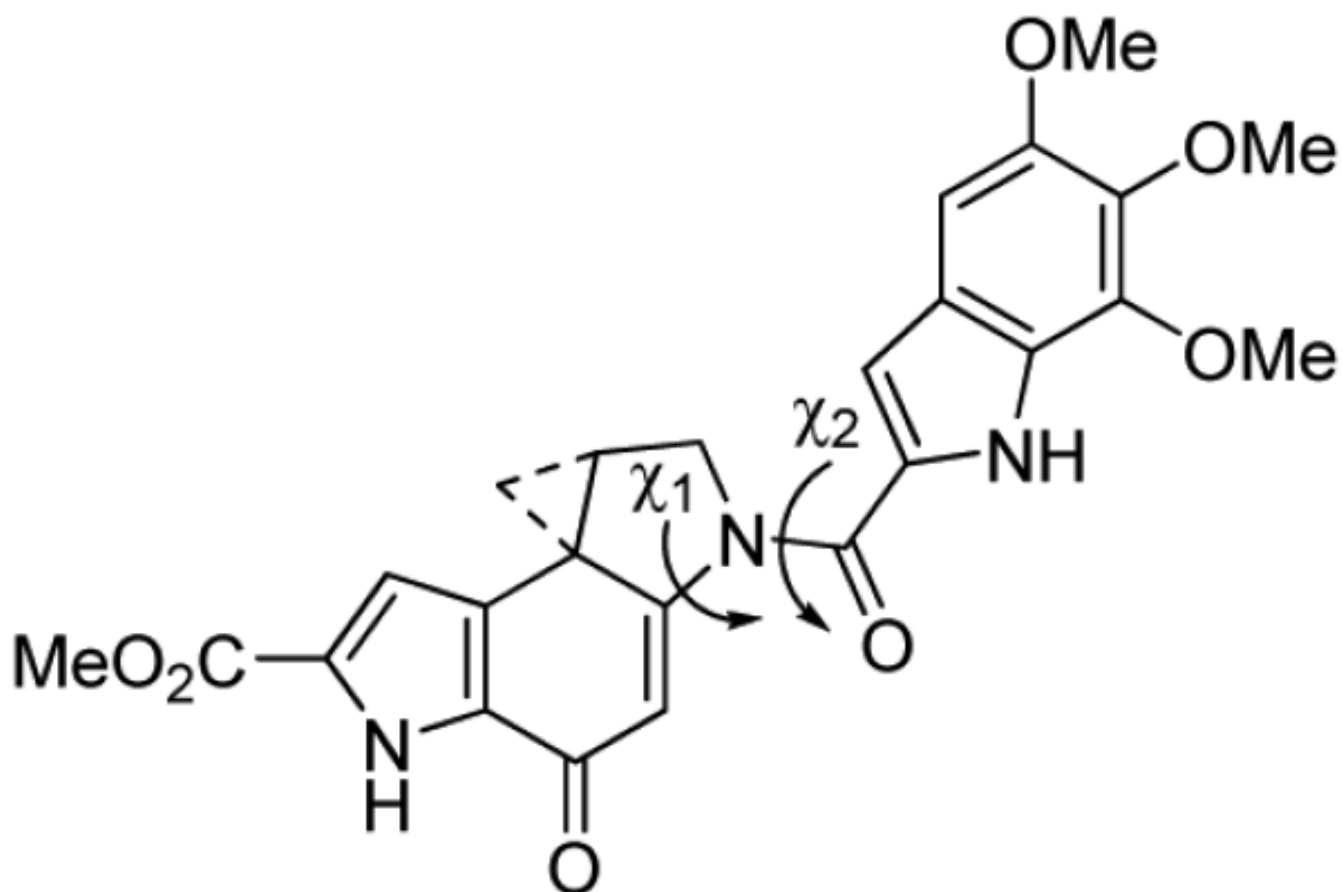


Figure 2.
Shape-dependent catalysis.

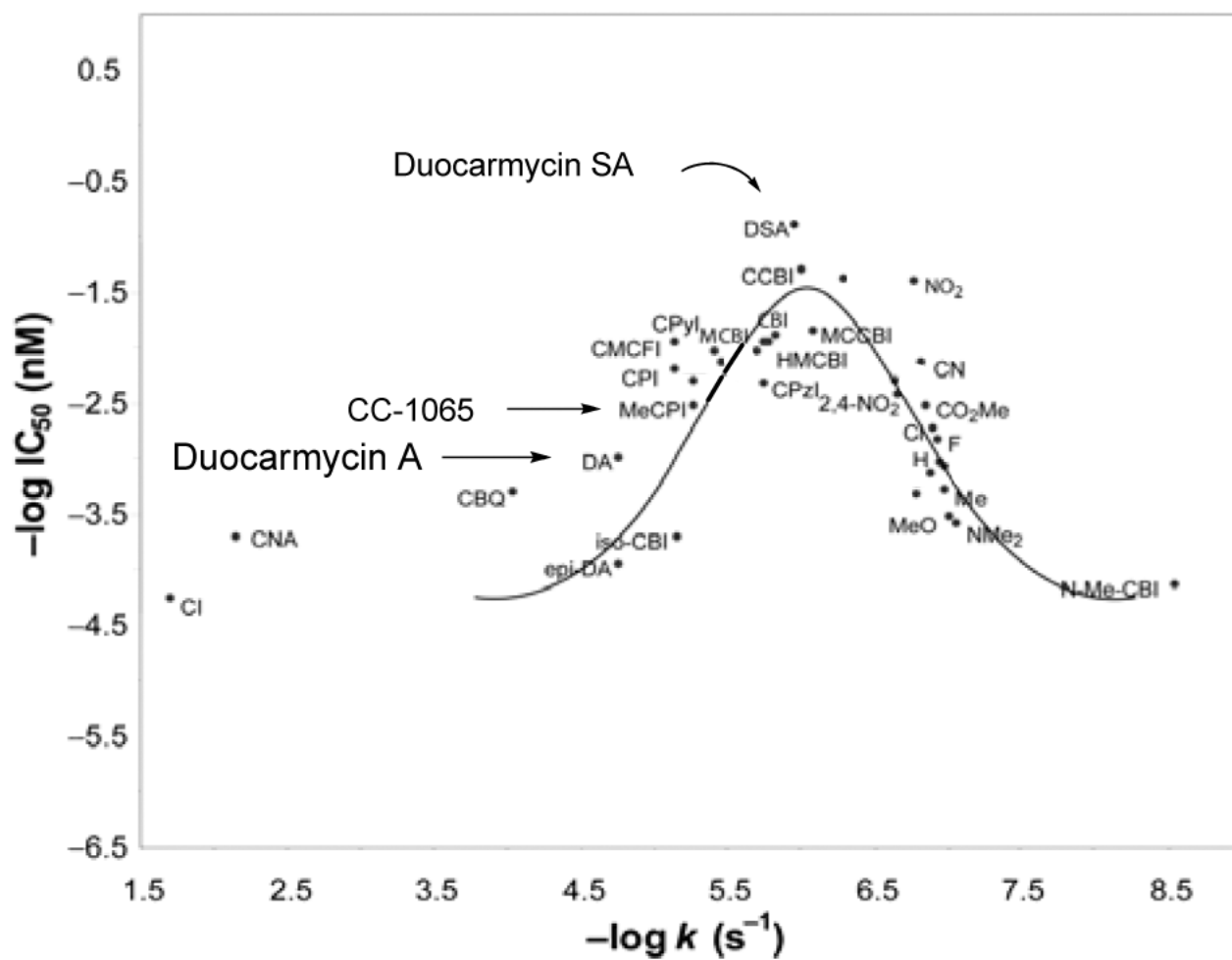


Figure 3.
Parabolic relationship between reactivity and potency.

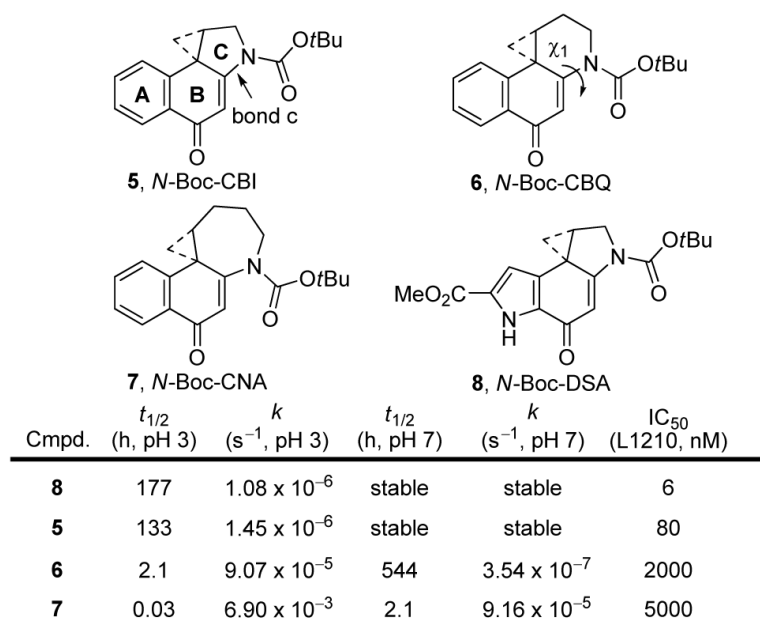


Figure 4.
C-Ring analogues of CBI.

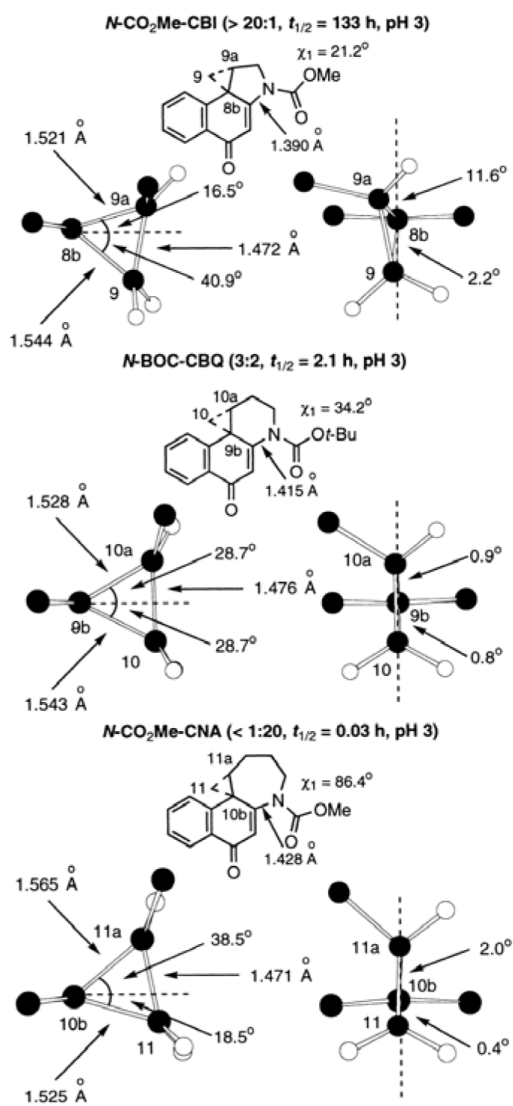


Figure 5.
X-ray crystal structure data.

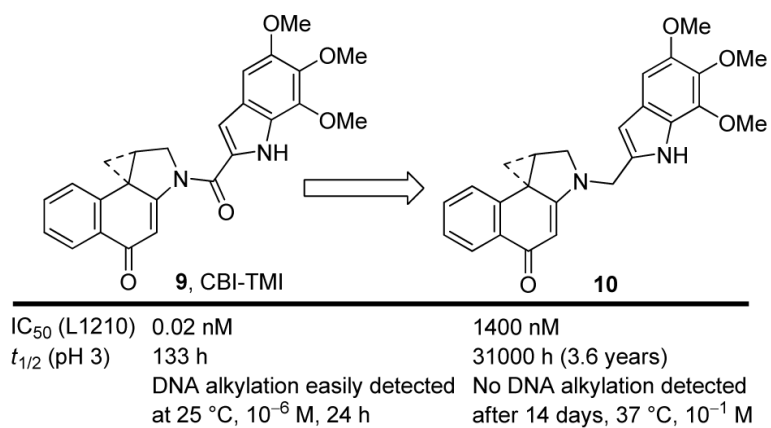


Figure 6.
Importance of the linking amide.

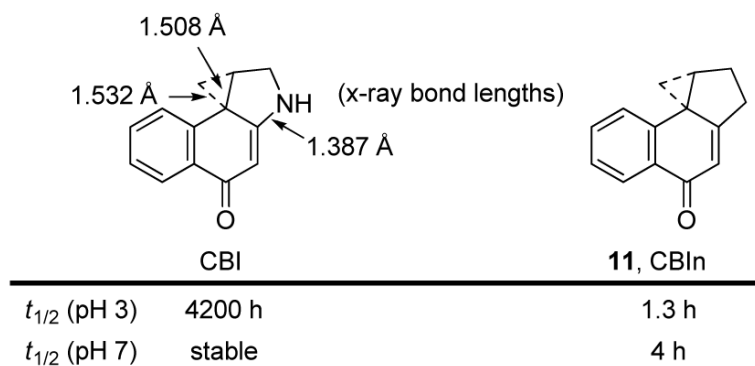
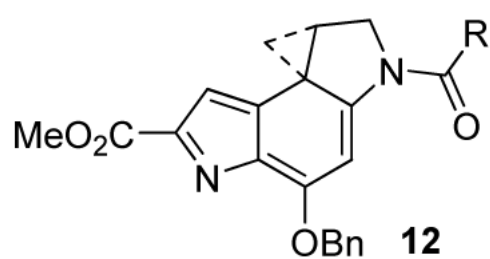
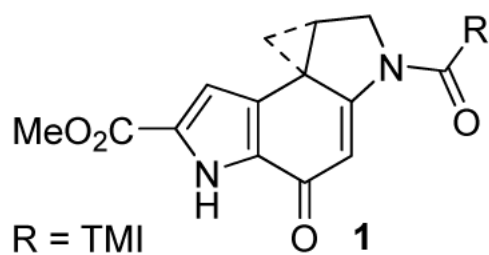


Figure 7.
CBI versus CBIIn.



IC ₅₀ (L1210, pM)	10	5000
<i>t</i> _{1/2} (pH 3)	177 h	25 min
<i>t</i> _{1/2} (pH 7)	stable	43 min
Rel. rate DNA alk.	1	4

Figure 8.
Alternate spirocyclization.

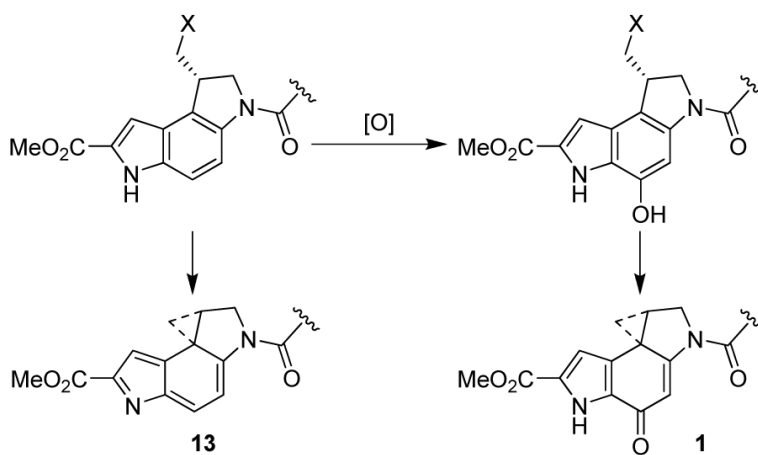


Figure 9.
Potential ancestral role of the alternative spirocyclization.

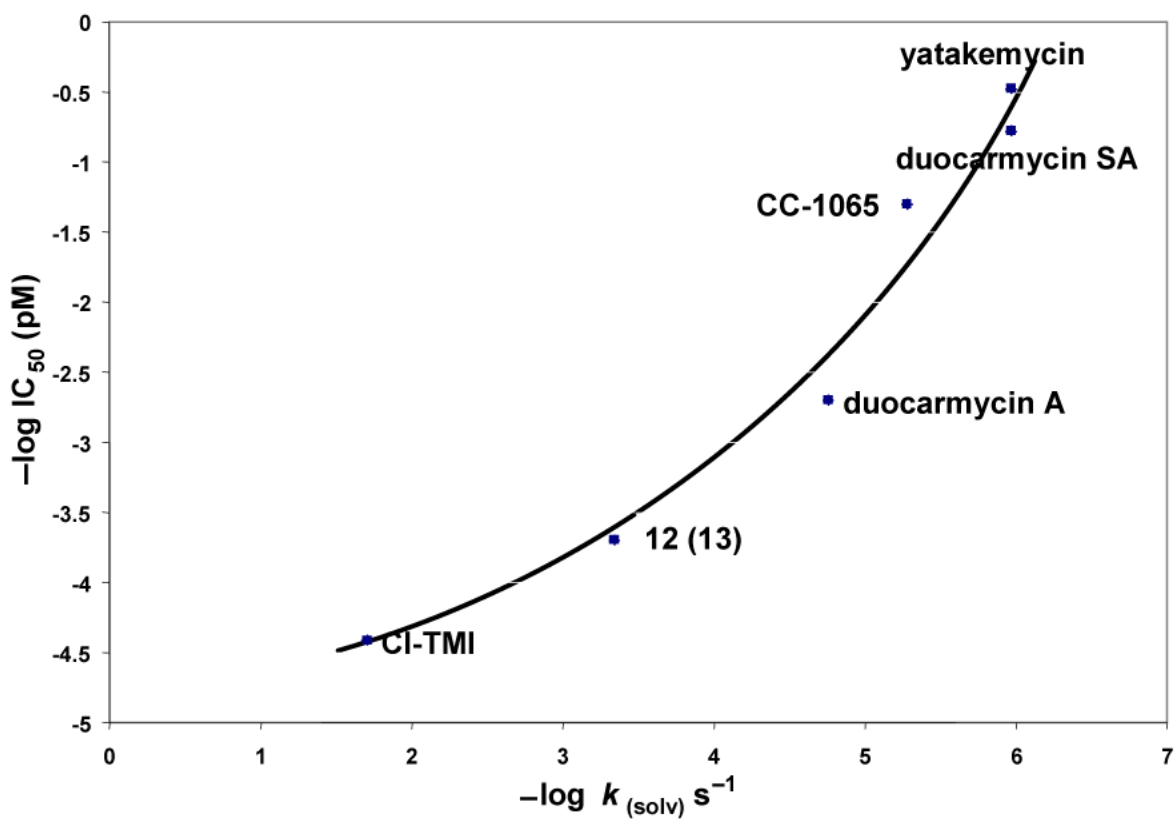


Figure 10.

Plot of $-\log \text{IC}_{50}$ (L1210, cytotoxic activity) versus $-\log k_{\text{solv}}$ (pH 3) that includes the natural products and a representative related compound illustrating the direct correlation between stability and biological potency and highlighting Nature's potential steps in the optimization of this class of DNA alkylating agents.

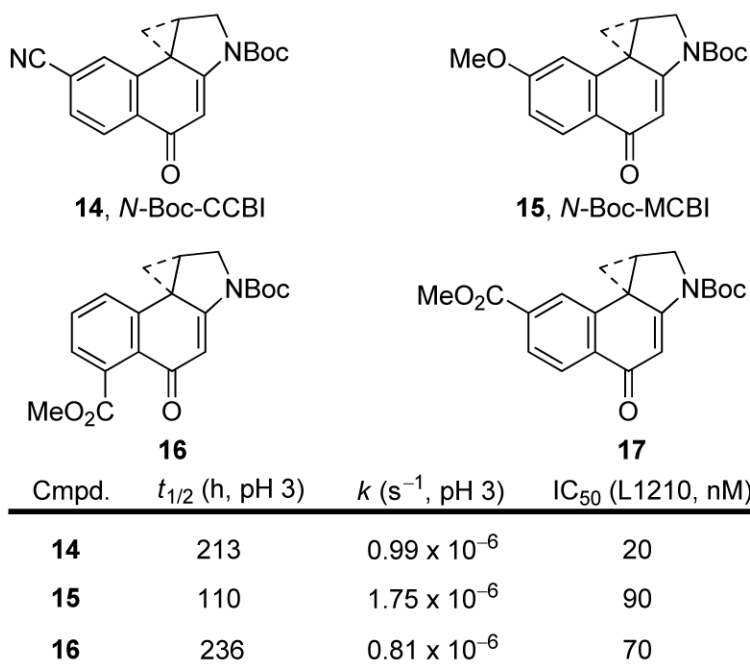
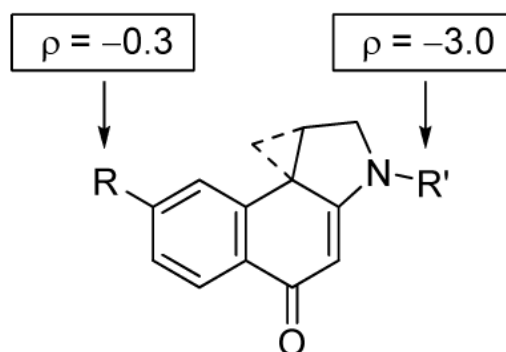


Figure 11.
A-Ring derivatives of CBI.



R' (R = H)	IC ₅₀ (L1210, nM)	t _{1/2} (h, pH 3)	k (pH 3, s ⁻¹)
SO ₂ Et	24	383	0.5 x 10 ⁻⁶
COEt	110	96	2.0 x 10 ⁻⁶
CO ₂ Me	140	57	3.4 x 10 ⁻⁶
CONHMe	200	36	5.4 x 10 ⁻⁶

Figure 12.
N²-Acyl derivatives of CBI.

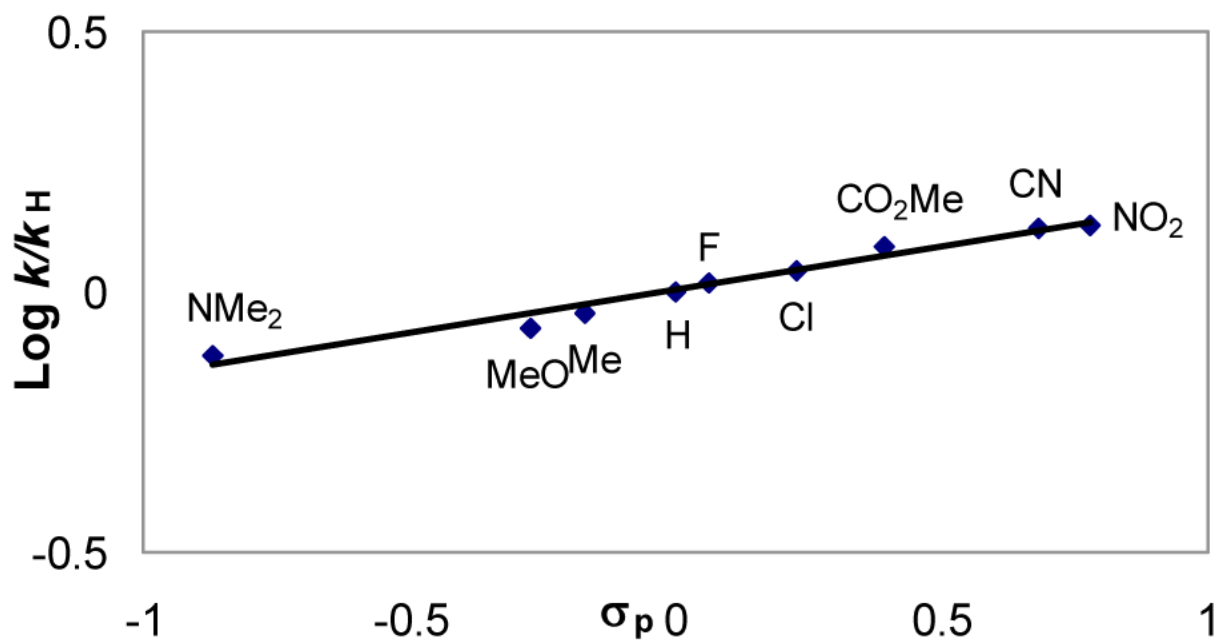


Figure 13.
Hammett plot of para substituted *N*-aryl-CBI derivatives.

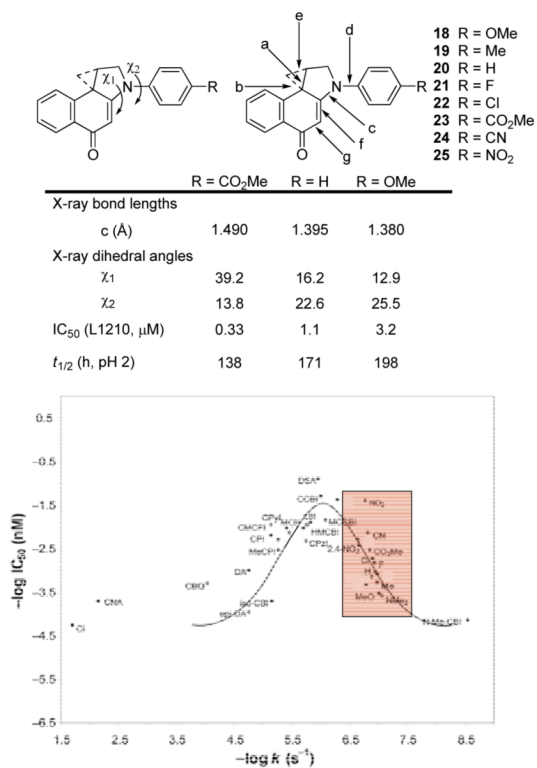
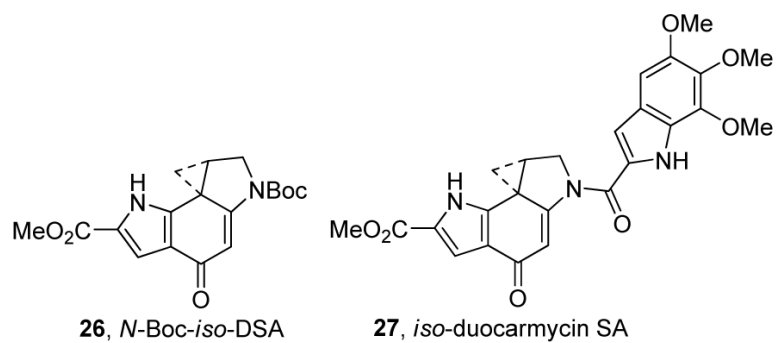


Figure 14. Structural and reactivity effects of para substituted *N*-aryl-CBI derivatives.



Compound	IC ₅₀ (pM)
8 , <i>N</i> -Boc-DSA	6000
26 , <i>N</i> -Boc- <i>iso</i> -DSA	33000
1 , duocarmycin SA	10
27 , <i>iso</i> -duocarmycin SA	50

Figure 15.
N-Boc-*iso*-DSA and *iso*-duocarmycin SA.

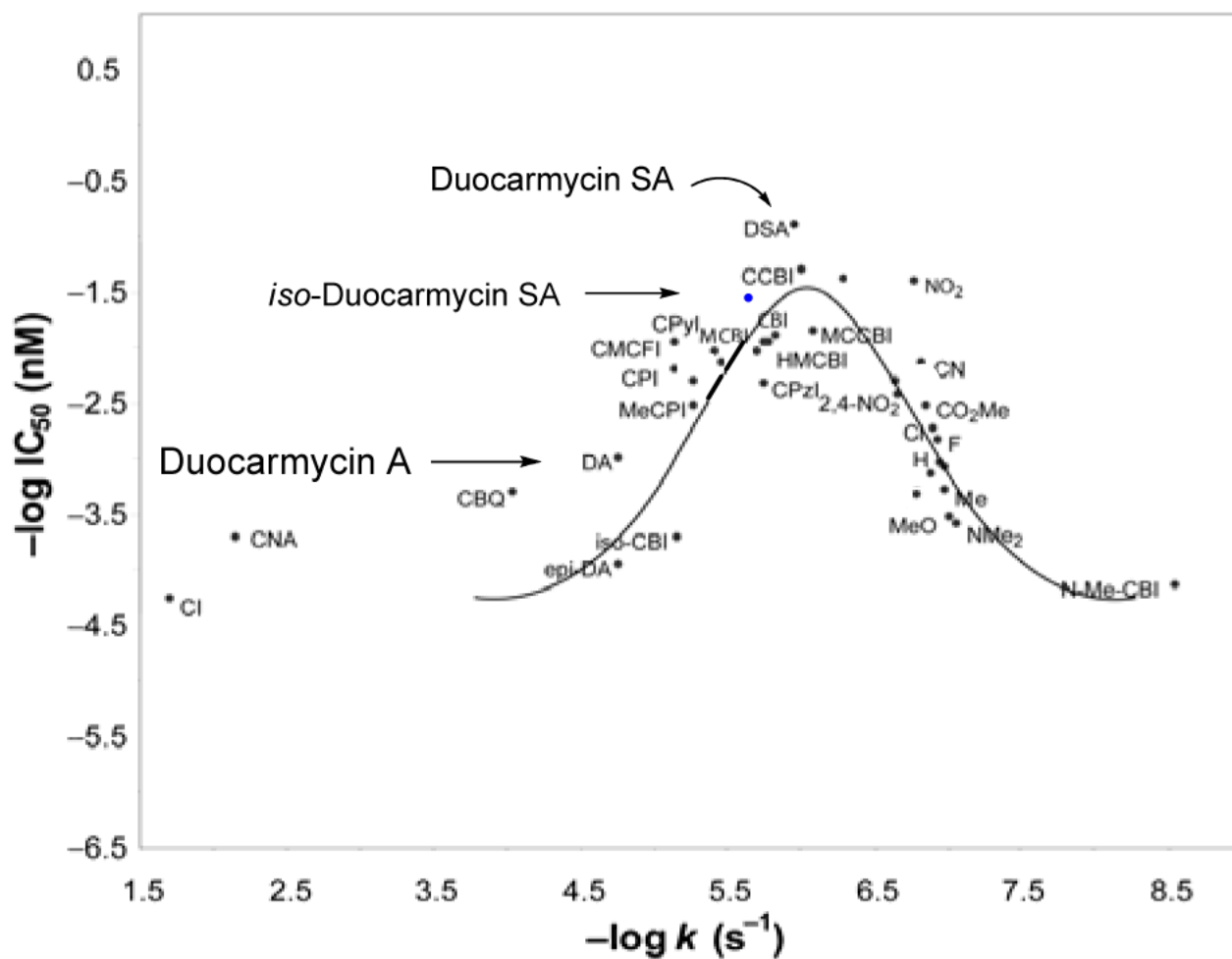


Figure 16.
Placement of *N*-Boc-*iso*-DSA on the parabolic curve.

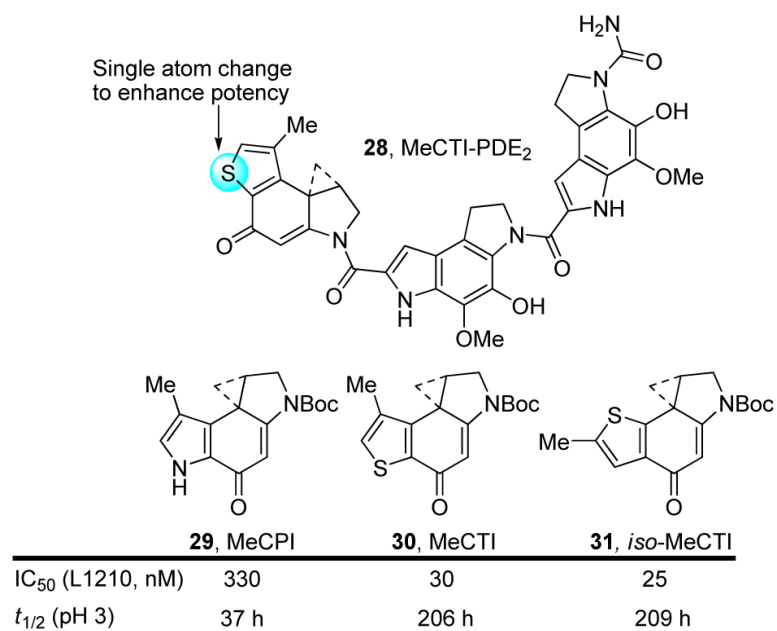


Figure 17.
Rationally designed analogues of CC-1065.

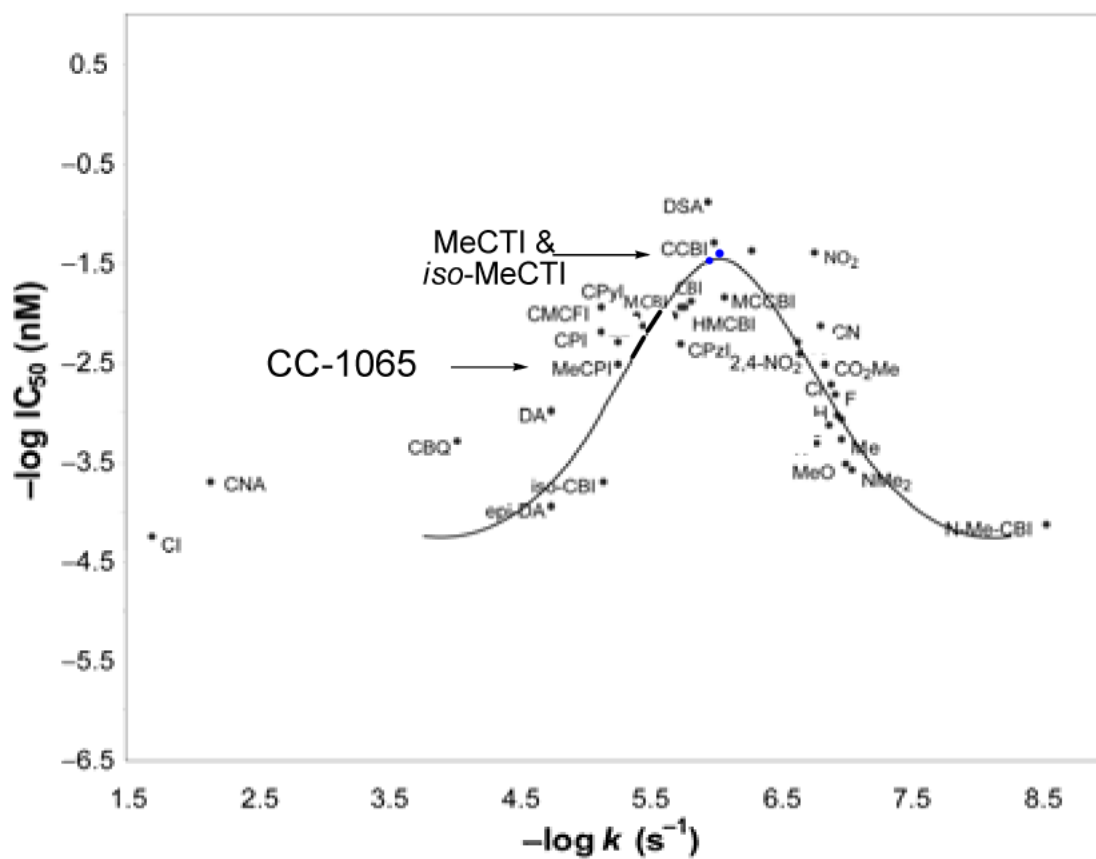


Figure 18.
MeCTI and *iso*-MeCTI on the plot of reactivity versus potency.