Combinatorial synthesis of novel and potent inhibitors of NADH:ubiquinone oxidoreductase

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Background: NADH:ubiquinone oxidoreductase (complex I) is the first of three large enzyme complexes located in the cell's inner mitochondrial membrane which form the electron transport chain that carries electrons from NADH to molecular oxygen during oxidative phosphorylation. There is significant interest in developing small molecule inhibitors of this enzyme for use as biological probes, insecticides and potential chemopreventive/chemotherapeutic agents. Herein we describe the application of novel natural product-like libraries to the discovery of a family of potent benzopyran-based inhibitors.

Results: Initially a combinatorial library of benzopyrans, modeled after natural products, was synthesized using a solid phase cycloloading strategy. Screening of this diversity oriented library for inhibitory potency against NADH:ubiquinone oxidoreductase activity in vitro using bovine heart electron transport particles provided several lead compounds which were further refined through a series of focused libraries.

Conclusions: Using this combinatorial library approach, a family of potent 2,2dimethylbenzopyran-based inhibitors was developed with IC_{50} values in the range of 18–55 nM. Cell-based assays revealed that these inhibitors were rather noncytotoxic in the MCF-7 cell line; however, they were quite cytostatic in a panel of cancer cell lines suggesting their potential as chemotherapeutic/chemopreventive candidates.

Introduction

NADH:ubiquinone oxidoreductase (complex I) is the first of three large enzyme complexes located in the inner mitochondrial membrane which form the electron transport chain that carries electrons from NADH to molecular oxy¹Department of Chemistry and The Skaggs Institute for Chemical Biology, The Scripps Research Institute, 10550 North Torrey Pines Road, La Jolla, CA 92037, USA

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gen during oxidative phosphorylation (Figure 1) [1]. Complex I is the most intricate enzyme known, consisting of over 40 individual protein sub-units with one non-covalently bound flavin mononucleotide and at least five iron-sulfur clusters. This enzyme serves an essential role



Figure 1. Role of NADH:ubiquinone oxidoreductase (complex I) on oxidative phosphorylation.

in cellular physiology such that structural or functional deficiencies in it have been implicated in the pathogenesis of diseases such as Parkinson's, focal dystonia and Leber's hereditary optic neuropathy [2].

A variety of natural and synthetic inhibitors of complex I (see representative examples in Figure 2) have found multiple applications [3]. First, inhibitors have been used to elucidate the role of this enzyme in normal cell physiology and also to mimic complex I deficiencies in order to study mitochondrial diseases [4]. Second, the structure-activity relationships (SAR) developed using inhibitors have provided important insights into the functional architecture of this complicated enzyme system; furthermore, affinity probes, constructed based upon these inhibitors, have proven valuable in identifying and characterizing the components of complex I [5-7]. Third, NADH:ubiquinone oxidoreductase continues to be a preferred target for the development of commercial insecticides and acaricides [8]. Fourth, and more recently, it has been demonstrated that inhibition of NADH:ubiquinone oxidoreductase causes a corresponding reduction in the activity of ornithine decarboxylase (ODC), possibly through interruption of signal transduction pathways [9]. ODC is responsible for the biosynthesis of polyamine growth factors required for normal cellular proliferation [10]. Since the overexpression of ODC activity in tumor cells contributes to aberrant proliferation, the ability of complex I inhibitors to reduce ODC activity makes them potential candidates for development as chemotherapeutic and/or chemopreventive



Figure 2. Representative natural and synthetic inhibitors of NADH:ubiquinone oxidoreductase.



Figure 3. Selected natural product inhibitors (7–18) of NADH:ubiquinone oxidoreductase isolated from Cubé resin and combinatorial library prototype (19).

agents [11–13]. Moreover, there is increasing evidence that modulation of complex I activity, through small molecules, can induce apoptosis thus further supporting the potential of these inhibitors as anticancer agents [14].

The goal of this study was to identify novel lead compounds through the screening of natural product-like combinatorial libraries. The motivation stemmed from the previous characterization of a family of naturally occurring inhibitors isolated from Cubé resin, an extract of the roots of *Lonchocarpus utilis* and *Lonchocarpus urucu*, which has been used as a botanical insecticide for decades [15,16].



Scheme 1. General strategy for the solid phase combinatorial synthesis of natural product-like small molecules as potential inhibitors of NADH:ubiquinone oxidoreductase. *o*-Prenylated phenols are cycloloaded onto a polystyrene-based selenenyl bromide resin to provide immobilized benzopyran scaffolds which are functionalized and then coupled to a second aromatic unit, thereby affording bridged structures **23**. Library members are released from the resin by oxidation of the selenium moiety to the corresponding selenoxide which undergoes spontaneous *syn*-elimination to afford the desired benzopyran systems (**19**).

Separation of the constituents of this resin provided the major components rotenone (1, Figure 2) and deguelin (7, Figure 3) along with minor constituents, including dehydrodeguelin (8), oxadehydrodeguelin (9), lonchocarpusone (10), 4-hydroxylonchocarpene (11), 4-hydroxy-3methoxylonchocarpene (12), stilbenes 13–15 and isoflavones 16–18 (Figure 3) among others. Inspection of natural products 7–18 revealed an interesting homology as all possessed a 2,2-dimethylbenzopyran motif linked to a terminal aromatic ring through a variety of molecular bridges. De-



Scheme 2. Representative procedures for the solid phase synthesis of amides, esters, and thioesters. Reagents and conditions: (a) 0.5 equivalent (equiv) of selenenyl bromide resin (1.1 mmol/g), CH_2CI_2 , 25°C, 20 min; (b) 10.0 equiv of LiOH, THF:H_2O (20:1), 50°C, 48 h; (c) 10.0 equiv of (COCI)_2, cat. DMF, CH_2CI_2 , 40°C, 2 h; (d) 10.0 equiv of piperazine, 1.0 equiv of 4-DMAP, CH_2CI_2 , 25°C, 12 h; (e) 5.0 equiv of 3,4-(OMe)_2C_6H_3CH_2XH (X=O, NH, S), 15.0 equiv of Et_3N, 1.0 equiv of 4-DMAP, CH_2CI_2 , 25°C, 12 h; (f) 5.0 equiv of 3,4-(OMe)_2C_6H_3XH (X=O, NH, S), 15.0 equiv of 4-DMAP, CH_2CI_2 , 25°C, 12 h; (f) 5.0 equiv of 3,4-(OMe)_2C_6H_3XH (X=O, NH, S), 15.0 equiv of 4-DMAP, CH_2CI_2 , 25°C, 12 h; (g) 5.0 equiv of 3,4-(OMe)_2C_6H_3CH_2XH (X=O, NH, S), 10.0 equiv of 4-DMAP, CH_2CI_2 , 25°C, 20 min. DCC = 1,3-dicyclohexylcarbodiimide, 4-DMAP = 4-(dimethylamino)pyridine, DMF = *N*,*N*-dimethylformamide.



Figure 4. Structures and IC₅₀ values of screening library. The synthesis of individual members is described in the schemes and references indicated.

spite the diversity in these bridging units, all of the compounds (7-18) inhibited complex I activity in in vitro assays [15,16]. Not surprisingly however, the potency of inhibition was strongly influenced by the orientation through which the 'bridge' unit positioned the two ring systems with respect to each other. For example, deguelin $(IC_{50} = 6.9 \text{ nM})$ is almost 500-fold more active than lonchocarpusone (IC₅₀ = 3300 nM). Given this effect, it was envisioned that screening a benzopyran-based combinatorial library wherein this 'bridge' unit was varied (i.e. structure 19, Figure 3) might lead to the identification of interesting SAR, and potentially to new lead compounds in this series. Notably, it was anticipated that these lead compounds, synthesized through combinatorial chemistry, would be more amenable (as compared to the parent natural products) to further synthetic modifications as required for optimization of physical and pharmacological properties.

Results and discussion

First generation discovery library: the bridge region III (compounds 25–76)

Having recently completed the synthesis of several large benzopyran-based combinatorial libraries, we were well positioned to assemble a suitable, first generation library for evaluating this hypothesis [17-19]. Thus, a 52-membered library was selected for preliminary screening as shown in Figure 4. Members were selected so as to simultaneously evaluate both the nature of the 'bridge' unit as well as the importance of substituents on the terminal aromatic ring system. Most members of this discovery library had been previously synthesized via a seleniumbased, solid phase strategy as outlined in Scheme 1. In this approach, a series of ortho-prenylated phenols (20) were cycloloaded (through a six-endo-trig electrophilic cyclization reaction) onto a polystyrene-based selenenyl bromide resin to afford resin-bound benzopyran scaffolds (21) [20,21]. These scaffolds were then functionalized and coupled with a second aromatic unit so as to create diverse bridge types (23). Upon completion, structures of type 23 could be further derivatized if necessary, and then released from the solid support by oxidation of the selenoether to the corresponding selenoxide which could undergo facile syn-elimination. The synthesis of compounds 25–28, 33, 35-41, 46-48, 50-54, 59-62, 64-66 and 73 (Figure 4) via this solid phase route has been described previously [17-21]. Representative procedures for the solid phase synthesis of the remaining members of the described library are illustrated in Scheme 2-4. All library members for biological assay were chromatographically and spectroscopically (¹H-nuclear magnetic resonance) homogeneous.

As shown in Scheme 2, the first type of library members to be constructed was a series of acyl derivatives. Initially, *ortho*-prenylated phenol **77** was cycloloaded onto a selenenyl bromide resin [22] to afford benzopyran scaffold **78**. The methyl ester of **78** was then hydrolyzed (LiOH, THF:H₂O, 40°C) to the corresponding carboxylic acid which was converted to acid chloride 80 by treatment with oxalyl chloride in the presence of catalytic amounts of N,N-dimethylformamide (DMF). This acid chloride then participated in three parallel reaction pathways. In the first sequence, treatment of acid chloride 80 with piperazine and 4-(dimethylamino)pyridine (4-DMAP) provided amide 81. The secondary amine of structure 81 was then coupled to 3,4-dimethoxybenzoic acid in the presence of 1,3-dicyclohexylcarbodiimide (DCC) and 4-DMAP to provide diamide 84 which was released from the solid support by treatment with H₂O₂, giving diamide 45. In the second pathway, acid chloride 80 was treated (in parallel) with a series of aryl nucleophiles including 3,4dimethoxyphenol, 3,4-dimethoxyaniline and 3,4-dimethoxythiophenol to provide structures of type 83, which upon oxidative cleavage afforded ester 30, amide 56 and thioester 71, respectively. In the last pathway, acid chloride 80 was reacted with a series of benzylic nucleophiles including 3,4-dimethoxybenzyl alcohol, 3,4-dimethoxybenzyl amine and 4-methoxy- α -toluene thiol to provide structures of type 82 which were oxidatively cleaved to afford ester 42, amide 31 and thioester 69, respectively. In addition to the representative compounds shown here, this procedure



Scheme 3. Representative procedures for the solid phase synthesis of ether (75), ester (29), and sulfonate (70) compounds. Reagents and conditions: (a) 0.5 equiv of selenenyl bromide resin (1.1 mmol/g), CH_2Cl_2 , 25°C, 20 min; (b) 5.0 equiv of 3,4-dimethoxybenzyl chloride, 5.0 equiv of K₂CO₃, DMF, 60°C, 12 h; (c) 5.0 equiv of 3,4-dimethoxybenzoic acid, 5.0 equiv of DCC, 1.0 equiv of 4-DMAP, CH_2Cl_2 , 25°C, 24 h; (d) 5.0 equiv of 3,4-dimethoxybenzenesulfonyl chloride, 10.0 equiv of Et₃N, 1.0 equiv of 4-DMAP, CH_2Cl_2 , 25°C, 12 h; (e) 6.0 equiv of H₂O₂, THF, 25°C, 20 min.

was repeated in an analogous and parallel manner to prepare compounds 32, 43–44, 49, 55, 57–58 and 68–69 (Figure 4).

As shown in Scheme 3, additional library members with other types of bridging units were constructed from a phenolic benzopyran system. Initially, resin-bound phenol 86 was prepared through the cycloloading of prenylated hydroquinone 85. Library members containing an ether-type bridge were then constructed by alkylation of 86 with 3,4dimethoxylbenzyl chloride in the presence of K₂CO₃ followed by oxidative cleavage to afford ether 75. Ester derivatives were constructed by acylation of phenol 86 with 3,4-dimethoxybenzoic acid, DCC and 4-DMAP to afford, after oxidative cleavage, ester 29. Lastly, sulfonate derivatives were constructed by treatment of phenol 86 with 3,4dimethoxybenzenesulfonyl chloride, Et₃N, and 4-DMAP followed by oxidative cleavage to afford sulfonate 70. In addition to the representative compounds shown here, this procedure was also repeated in an analogous and parallel manner to prepare compounds 63, 67, 74 and 76 (Figure 4).

As shown in Scheme 4, several sulfur-containing library members were synthesized through a halogen-metal exchange reaction. Initially, prenylated bromophenol 88 was cycloloaded onto the selenenyl bromide resin to give ben-

Table 1

Summary of biological data from evaluation of first generation library and proposed lead structures.



Bridge	IC ₅₀ (nM)	Bridge	IC ₅₀ (nM)
Chalcone	850–(>3000)	Alkyne	2300
Stilbene	2200–(>3000)	Ester	55–(>3000)
Coumarin	1900–(>3000)	Amide	2300–(>3000)
Heterocycle	1100–(>3000)	Ether	160–800

First generation lead compounds



Strategic regions for SAR optimization





Scheme 4. Solid phase synthesis of thioamide (72) and thiazole (34) systems. Reagents and conditions: (a) 0.5 equiv of selenenyl bromide resin (1.1 mmol/g), CH_2Cl_2 , 25°C, 20 min; (b) 10.0 equiv of *n*-BuLi (1.6 M in hexanes), THF, $-78 \rightarrow 0^{\circ}C$, 2 h; then 10.0 equiv of 3,4-dimethoxyphenylisothiocyanate, THF, $-78 \rightarrow 25^{\circ}C$, 1 h; (c) 6.0 equiv of H₂O₂, THF, 25°C, 20 min.

zopyran 89. Aryl bromide 89 was subjected to a halogenmetal exchange reaction by treatment with *n*-BuLi at -78° C and then warming to 0°C over 2 h at which time 3,4-dimethoxyphenylisothiocyanate was added. Cleavage of the supposed thioamide 90 by treatment with H₂O₂ unexpectedly afforded a separable mixture of thioamide 72 and thiazole 34 the latter being presumably formed as a result of excess base present in the halogen-metal exchange reaction.

This primary library was screened for inhibitory potency against NADH:ubiquinone oxidoreductase activity as described below. This screening revealed several structure-activity trends (summarized in Table 1 and discussed in detail in the SAR section) which ultimately provided esters 42 and 55 as lead compounds. As described below, a series of focused libraries were then synthesized using a combination of solid and solution phase chemistry in an attempt to evaluate particular structural sub-regions (denoted I–IV in Table 1) of the lead structure, thus allowing SAR to be developed.

First focused library, aryl substituents, regions II and IV (compounds 91–127)

The first of these follow-up libraries focused on regions **II** and **IV** of the lead structure. Of particular interest was how



Scheme 5. Solution phase synthesis of 3,4,5-trimethoxybenzyl ester (55) and modified pyran analogs and their biological activities. Reagents and conditions: (a) 2.0 equiv of 3-chloro-3methyl-1-butyne, 2.0 equiv of K2CO2, 1.7 equiv of KI, 0.02 equiv of Cul, DMF, 65°C, 3 h, 88%; (b) Et₂NPh, 195°C, 1 h, 100%; (c) 2.0 equiv of LiOH, THF:H₂O (10:1), 40°C, 12 h, 95%; (d) 1.1 equiv of 3,4,5-trimethoxybenzyl alcohol, 1.2 equiv of DCC, 0.1 equiv of 4-DMAP, CH2Cl2, 25°C, 12 h, 89%; (e) 0.1 equiv of OsO₄, 1.1 equiv of NMO, *t*-BuOH:THF:H₂O (10:3:1), 25°C, 2 h, 71%; (f) 3.0 equiv of AcCl, 5.0 equiv of pyridine, CH₂Cl₂, $0 \rightarrow 25^{\circ}$ C, 1 h, 97%; (g) 5.0 equiv of triphosgene, 10.0 equiv of pyridine, CH₂Cl₂, 0°C, 1 h, 80%; (h) 1.1 equiv of NBS DMSO:H₂O (10:1), 0°C, 1 h, 80%; (i) 1.0 equiv of m-CPBA, 3.0 equiv of Na2CO3, CH2Cl2, 25°C, 24 h, 69%; (j) 0.2 equiv of 10% Pd-C, H₂, MeOH:hexane (1:1), 25°C, 6 h, 96%. DMSO = methyl sulfoxide, NMO = 4-methylmorpholine N-oxide, NBS = Nbromosuccinimide, m-CPBA = m-chloroperoxybenzoic acid. Compounds 133-139 were synthesized and assayed as racimates with the relative stereochemistry as shown.



Scheme 6. Synthesis of long chain 3,4-dimethoxyphenyl esters. Reagents and conditions: (a) 1.5 equiv of 3,4-dihydropyran, 0.1 equiv of PPTS, CH_2CI_2 , 25°C, 2 h; (b) 2.0 equiv of 3,4-dimethoxyphenol, 2.0 equiv of K_2CO_3 , acetone, 40°C, 12 h; (c) 1.2 equiv of TsOH·H₂O, THF:MeOH (9:1), 25°C, 1 h; (d) 0.5 equiv of 80, 10.0 equiv of Et₃N, 1.0 equiv of 4-DMAP, CH_2CI_2 , 25°C, 12 h; (e) 6.0 equiv of H_2O_2 , THF, 25°C, 20 min.

substituent patterns on these two aromatic systems would effect inhibitory activity. Hence, a series of esters (91–127, Figure 5) were synthesized in parallel on solid support using an identical protocol to that described in Scheme 2. The aromatic building blocks employed contained a variety of substituents including alkyl groups, halogens, alkoxy groups, hydroxyl groups, alcohols and nitro groups in order to assess both steric and electronic effects.

Second focused library, dimethylpyran modifications, region I (compounds 133–140)

The synthesis of the second of the follow-up libraries relied upon solution phase chemistry and was designed so as to evaluate how modifications in the pyran ring system might effect biological activity. Thus, as shown in Scheme 5, lead ester 55 was first resynthesized on a large scale from 4-hydroxymethyl benzoate 128 by initial O-alkylation with 3-chloro-3-methyl-1-butyne to give ether 129 [23]. Heating of alkyne 129 in N,N-diethylaniline at 195°C for 1 h induced an aromatic Claisen rearrangement to provide benzopyran 130 in quantitative yield [23]. The methyl ester of 130 was hydrolyzed (LiOH, THF:H₂O, 40°C, 12 h) to the free acid 131 which was coupled to 3,4,5-trimethoxybenzyl alcohol in the presence of DCC and 4-DMAP to furnish ester 55. With the ester 55 in hand, several derivatization reactions were employed to provide the series of pyranmodified analogs 133-140. First, the olefin of ester 55 dihydroxylated [24] (cat OsO4, NMO, twas BuOH:THF:H₂O, 25°C, 6 h) to afford diol 133 which was subsequently acetylated to diacetate 134 and reacted with triphosgene to provide carbonate 135. In a second event, ester 55 was treated [25] with NBS in the presence



of H_2O to provide a mixture of bromohydrins 136 and 137 as a result of partial bromination of the terminal aromatic ring (i.e. structure 136). Bromohydrin 137 was subsequently acetylated (AcCl, pyridine, CH_2Cl_2 , 25°C, 1 h) to afford acetate 138. Additionally, ester 55 was converted to the corresponding epoxide 139 by treatment with *m*-CPBA [26]. Finally, the olefin of ester 55 was saturated by hydrogenation over 10% Pd–C to provide the corresponding saturated pyran system 140. Scheme 7. Solution phase synthesis of 3,4,5trimethoxyphenylketone (158) and analogs thereof. Reagents and conditions: (a) 1.2 equiv of 2-methyl-3-butyn-2-ol, 1.2 equiv of trifluoroacetic anhydride, 1.5 equiv of DBU, MeCN, 0°C, 6 h, 100%; (b) Et₂NPh, 195°C, 1 h, 100%; (c) 3.0 equiv of BH₃·THF, THF, $0 \rightarrow 25^{\circ}$ C, 2 h, 100%; (d) 1.1 equiv of Dess-Martin periodinane, CH₂Cl₂, 25°C, 1 h, 90%; (e) 1.1 equiv of n-BuLi (1.6 M in hexanes), THF, -78°C, 15 min; then 2.0 equiv of 155, THF, -78°C, 15 min, 45%; (f) 5.0 equiv of Mel, 2.0 equiv of NaH, THF, $0 \rightarrow 25^{\circ}$ C, 12 h, 85%; (g) 4.0 equiv of RONH₂ (R=H, Me, Bn), 5.0 equiv of K₂CO₃, EtOH, 75°C, 2 h, R=H: 52%, R=Me: 78%, R=Bn: 81%; (h) 3.0 equiv of Lawesson's reagent, toluene, 100°C, 3 h, 42%; (i) 1.2 equiv of Tebbe reagent, THF, $-40 \rightarrow 0^{\circ}$ C, 30 min, 88%; (j) 10.0 equiv of H₄N₂·H₂O, 10.0 equiv of KOH, di(ethylene glycol), 245°C, 6 h, 21%; (k) 0.2 equiv of 10% Pd-C, H₂, MeOH:hexane (1:1), 25°C, 6 h; (I) 2.0 equiv of 10% Pd-C, H₂, MeOH, 25°C, 12 h.

Third focused library, the bridge region III (compounds 146–179)

A final library in this series sought to further investigate the nature of the bridge unit. Of particular interest was the effect of length, structure and polarity of this bridge as illustrated in Scheme 6–9. In order to access the relationship between bridge length and biological activity, a set of longer chain analogs 146–148 was synthesized as illustrated in Scheme 6 where alcohols of type 141 were prepared by alkylation of the corresponding bromoalcohols and then coupled to resin-bound acid chloride 80 as previously described. Subsequently, we sought to determine whether or not an ester functionality was optimal as a bridge unit. This was accomplished by the solution phase synthesis of vari-



Scheme 8. Solution phase synthesis of difluoro- and dichlorostyrene analogs 168 and 169. Reagents and conditions: (a) 2.0 equiv of Ph₃P, CCl₄, 60°C, 15 h, 60%; (b) 5.0 equiv of (EtO)₂P(O)CHF₂, 4.0 equiv of *t*-BuLi, DME, -78° C, 1 h; then 85°C, 15 h, 20%.



Scheme 9. Synthesis of 3,4,5-trimethoxyphenylthioether **178** and 3,4,5-trimethoxyphenylsulfone **179**. Reagents and conditions: (a) 1.0 equiv of 3,4,5-trimethoxybenzyl chloride, 3.0 equiv of K₂CO₃, 0.2 equiv of TBAI, DMF, 25°C, 2 h, 74%; (b) 2.0 equiv of 3-chloro-3-methyl-1-butyne, 2.0 equiv of K₂CO₃, 1.7 equiv of KI, 0.02 equiv of Cul, DMF, 65°C, 3 h, 40%; (c) Et₂NPh, 195°C, 1 h, 100%; (d) 2.5 equiv of OXONE, 2.5 equiv of NaHCO₃, THF:H₂O (2:1), 25%. TBAI=tetrabutylammonium iodide.

ous other linking units, including ketones, oximes, thioketones, alkyl chains, thioethers and sulfones as shown in Schemes 7–9. As illustrated in Scheme 7, various keto analogs and their derivatives were constructed starting from 4-bromophenol **149**. Following the procedure of Ding, 4-bromophenol (149) was alkylated with the in situ generated triflate of 2-methyl-3-butyn-2-ol to afford 150, which was heated to 195°C in N,N-dimethylaniline inducing an aromatic Claisen rearrangement and providing benzopyran 151 in quantitative yield [27]. Subsequent treatment of 151 with n-BuLi afforded aryl lithium 152, which was immediately quenched by addition of aldehyde 155 (synthesized from phenylacetic acid 153, as indicated in Scheme 7) to provide benzyl alcohol 156. Dess-Martin periodinane oxidation of alcohol 156 afforded ketone 158. Incidentally, the methyl ether analog of alcohol 156, compound 157, was also synthesized from alcohol 156 via alkylation with MeI in the presence of NaH. With ketone 158 available, a series of modified keto-analogs were then constructed. Thus, oximes 159-161 were prepared by condensation of ketone 158 with the appropriate hydroxy or alkoxy amines in the presence of K₂CO₃. Thioketone 162 was prepared by reaction of ketone 158 with Lawesson's reagent in toluene at 100°C for 3 h, whereas the corresponding olefin analog 163 was constructed by treatment of ketone 158 with Tebbe reagent. Wolf-Kishner reduction of 158 (H₄N₂, KOH, \cdot) afforded the alkyl variant 164, while hydrogenation over 10% Pd-C provided the saturated version (i.e. 165) of ketone 158 as well as of analogs 163 and 164 (i.e. 167 and 166, respectively). Additional variations of ketone 158 were also constructed to address the importance of tether length and bridge substitution. These analogs were synthesized via an analogous protocol to that shown in Scheme 7 and their structures (170-174) are illustrated in Figure 5.

Several halogenated derivatives of styrene 163 were also prepared as illustrated in Scheme 8. Hence, ketone 165 was treated with Ph₃P in CCl₄ at 60°C to afford dichloro-analog 168 [28]. Additionally, ketone 165 was added to a solution of diethyl(difluoromethyl)phosphonate and *t*-BuLi in DME at -78° C and then warmed to 25°C,



Figure 6. Lead compounds (and their IC₅₀ values) selected for evaluation in cell-based assays.

and finally to 85°C to ultimately afford difluoro-analog **169** [29].

A final set of compounds for addressing the nature of the bridge unit was prepared as illustrated in Scheme 9 in an attempt to optimize the activity observed for ether 75 during screening of the first generation library. The thioether variant of this analog was prepared from 4-mercaptophenol (175) by selective *S*-alkylation with 3,4,5-trimethoxybenzyl chloride and K_2CO_3 to provide thioether 176 [30]. Subsequent *O*-alkylation was effected with 3chloro-3-methyl-1-butyne to give alkynyl ether 177. Heating of alkyne 177 in *N*,*N*-diethylaniline at 195°C for 1 h induced an aromatic Claisen rearrangement furnishing benzopyran 178 in quantitative yield. A portion of thioether 178 was then oxidized with OXONE[®] to afford the corresponding sulfone 179.

SAR

SAR optimization was carried out on each of the four regions (I–IV) of interest guided by the inhibitory potency for NADH:ubiquinone oxidoreductase activity in vitro using bovine heart electron transport particles (see Singer [31] for details).

The original library emphasizing bridge region III gave the IC₅₀ values for 25-76 indicated in Figure 4 and the SAR summarized in Table 1. Compounds containing a conjugated and/or rigid bridging unit (i.e. chalcones, stilbenes, coumarins, heterocycles or alkynes) exhibited low activity $(IC_{50} > 1000 \text{ nM})$ similar to those for most of the benzopyran-containing natural products (11-15, Figure 3). More interestingly, methylenes in the bridge unit, for greater conformational flexibility, resulted in higher activity dependent on the substituent pattern on the terminal aromatic ring, i.e. 220 nM IC₅₀ for 3,4-dimethoxybenzyl esters 42 and 55 nM IC₅₀ for 3,4,5-trimethoxybenzyl ester 55, the most potent inhibitor in the screening library. The corresponding amides (31 and 44, Figure 4), presumably more polar than their ester counterparts, were significantly less active (40- and 10-fold, respectively), suggesting the possible importance of lipophilicity at region III.

The first of the focused libraries (91-127, Figure 5) exam-

Table 2

Selected data for g	rowth inhibition (GI ₅₀) of com	pounds 55, 158,	, 165 and 178 in	n NCI cancer cell lines ^a
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Cancer cell type	GI ₅₀ (nM)					
	165	158	178	55	Average ^b	
Leukemia						
HL-60	340	14	840	1810	750	
K-562	730	1 540	5 090	2900	2800	
MOLT-4	96	340	1 230	6800	2560	
RPMI-8226	940	3040	2660	4760	2850	
NSC lung						
EKVX	66	680	3 930	8620	3 3 2 0	
HOP-62	830	280	6 6 9 0	31 300	9780	
NCI-H226	330	1 090	5 560	15 500	5620	
NCI-H322M	180	560	2210	49 600	13100	
Colon						
HCT-116	740	1 370	4 850	7 930	3720	
HCT-15	440	12600	3970	8940	6490	
SW-620	660	23 300	6 400	21 300	12900	
CNS						
SF-295	690	23 100	4 580	23 300	12900	
SNB-19	660	17 100	6110	11700	8890	
U-251	210	430	2 580	14 300	4 380	
Melanoma						
LOX IMVI	180	19700	3 0 5 0	9100	8010	
MALME-3M	320	13400	2 370	7 700	5950	
SK-MEL-5	490	1910	5 160	3200	2690	
UACC-257	210	780	3010	14 500	4630	
Breast						
MCF-7	3 080	880	6570	9100	4910	
NCI/ADR-RES	670	2 500	2 350	6800	3080	
HS 578T	600	860	7 400	25 000	8470	
MDA-MB-435	2 7 9 0	37 200	7 110	14 800	15 500	
Average ^c	690	7 380	4 300	13600		

^aAssays were performed by the Developmental Therapeutics Program of the National Cancer Institute, USA. The compounds were provided as DMSO solutions and evaluated for their in vitro cytostatic properties against 60 human cell lines using the NCI standard protocol. ^bAverage GI₅₀ value for compounds **55**, **158**, **165** and **178** in a particular cell line.

^cAverage GI₅₀ value for a single compound in all cell lines.



Figure 5. Summary of structure activity trends observed for benzopyran-based inhibitors of NADH: ubiquinone oxidoreductase.

ined the substituent patterns for aromatic regions II and IV. Somewhat surprisingly, only esters 110 ($IC_{50} = 44 \text{ nM}$) and 120 ($IC_{50} = 49 \text{ nM}$) exhibited marginally improved activity over the original 3,4,5-trimethoxybenzyl ester 55

(IC₅₀ = 55 nM). Varying substituents on the aromatic benzopyran ring only moderately influenced activity, e.g. introducing a halogen at R¹ (42 \rightarrow 104; 55 \rightarrow 112) or a hydroxyl group at R² (55 \rightarrow 110). A bulky substituent of the



Figure 7. Molecular modeling of deguelin (7), ester 55, and ketone 158.

oxygen at R^4 (42 \rightarrow 121; 55 \rightarrow 122) resulted in only slightly diminished inhibitory activity implying the greater importance of electronic rather than steric factors at this position. More generally, the 3,4,5-trimethoxyphenyl substituent is at or near the optimum for region IV.

The second focused library considered modifications of the pyran ring system (Figure 5). Introduction of a polar functionality on the pyran ring (i.e. 133 and 134) significantly compromised activity, and less polar substituents (i.e. 135–139) afforded no improvements. However, conversion to its saturated counterpart ($55 \rightarrow 140$) resulted in a compound which retained potent biological activity (IC₅₀ = 53 nM) yet was presumably metabolically more stable owing to the removal of the oxidatively sensitive pyran olefin.

A final focused library reexamined the bridging unit (region III) with keto-analogs revealing several interesting trends (Figure 5). First, ketone 158 was more active than the original ester lead with an IC₅₀ value of 39 nM. Alcohol 156, ether 157, oximes 159–161, thioketone 162, and compound 164 were significantly less active than ketone 158. Removal of the pyran olefin of ketone 158 via hydrogenation provided compound 165 which was almost twice as active as the parent olefin with an IC₅₀ value of 24 nM. Intriguingly, conversion of the ketone to the corresponding olefin also increased activity as demonstrated for compound 163 (IC₅₀ = 19 nM) and its saturated pyran counterpart 167 (IC₅₀ = 18 nM). Substitution of the olefin protons with halogens (i.e. structures **168** and **169**) resulted in dichloro- and difluoro-analogs with reduced activity (IC₅₀ values 2700 and 48 nM, respectively). On a separate note, thioether **178** was quite active with an IC₅₀ of 43 nM, whereas the more polar sulfone **179** was inactive.

The overall results summarized in Figure 5 prompted selection of **55**, **158**, **163**, **165**, **167** and **178** (Figure 6) for further biological testing. Despite their structural simplicity, the inhibitory activity of these compounds approached that for the most potent natural product in this series, namely deguelin (7, Figure 3) with an IC₅₀ value of 6.9 nM. Moreover, unlike the structurally complex deguelin, these lead compounds were simple to construct and thus could be readily modified to improve pharmacological properties, solubility or biostability.

The identified lead compounds were evaluated in several cell-based assays. The first involved determination of the cytotoxic concentrations (LC₅₀ values) in MCF-7 human epithelial breast cancer cells using the MTT assay [16]. The six compounds tested showed low cytotoxicity in the MCF-7 cells (LC₅₀ > 30000 nM). Secondly, compounds **55**, **158**, **165** and **178** were also evaluated in the NCI 60-cell cancer panel for the concentration leading to 50% growth inhibition (GI₅₀; see representative examples in Table 2) [32]. Moderate to good activity was observed in growth inhibition in various cell lines. The leukemia cells are generally the most sensitive (GI₅₀ from 750 to 2850

nM) and the CNS cells the least sensitive (GI₅₀ from 4380 to 12900 nM) to these compounds and the overall potency order is 165 > 178 > 158 > 55. The combination of rather potent growth inhibition and possibly low cytotoxicity makes these compounds interesting lead structures as potential chemopreventive/chemotherapeutic agents.

Rationalization of SAR by molecular modeling

In an effort to better understand the molecular basis for the potent inhibitory activity exhibited by several of these compounds, molecular modeling studies were undertaken. Specifically, we hoped to compare the three-dimensional orientation of these leads to that of the structurally more complex natural product deguelin [7] (7, Figure 3) which remains the most potent compound in the benzopyran series. As illustrated in Figure 7, the semi-rigid structure of deguelin (7) was first minimized with molecular mechanics calculations (Insight II, CFF93 force field [33]) which provided its lowest energy conformation as the skewed structure denoted as B in Figure 7. Subsequently, ester 55 (Figure 6) and ketone 158 (Figure 6) were also minimized with the two lowest energy levels of each presented in Figure 7 (i.e. structures A and D for ester 55 and structures C and F for ketone 158). Comparison of these minimized structures revealed that the second lowest energy conformation of ester 55 (structure **D**) and the lowest energy conformation of ketone 158 (structure C) resembled quite closely the bent configuration of deguelin 7 (structure B). In fact, the overlay of the lowest energy conformations of deguelin 7 (B) and ketone 158 (C), presented in diagram E (Figure 7), illustrate a strong homology between the two structures. The fact that the lowest energy configuration of ketone 158 best matched deguelin, whereas the less populated second lowest energy configuration of ester 55 matched deguelin was consistent with the fact that ketone 158 with an IC_{50} value of 39 nM was more potent than ester 55 which possessed an IC₅₀ value of 55 nM.

Significance

Herein we have described how synthetic combinatorial libraries, modeled after natural products, can be utilized to discover novel and structurally simple lead compounds as inhibitors of the enzyme NADH:ubiquinone oxidoreductase (the current structure activity relationships complement previous work in the rotenone series; for a discussion see [34,35]). After discovery, several of these lead inhibitors were optimized through the synthesis of focused libraries which refined specific sub-units of the lead compounds. Ultimately a collection of benzopyran-based inhibitors with IC₅₀ values 18-55 nM were identified. Several of these inhibitors were then evaluated in cell-based assays to evaluate their cytotoxic and cytostatic properties. Interestingly, while most of the compounds were relatively non-cytotoxic, several exhibit potent cytostatic activities across a variety of cancer types and cell lines. The ability of these small molecules to inhibit cancer cell growth, perhaps mediated by the ability to interrupt ODC activity, makes them potential candidates for further development as chemopreventive/chemotherapeutic agents. Further evaluation of these compounds is in progress along with the application of similar natural product-like combinatorial libraries to the discovery of lead compounds in other biological systems.

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