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Synthesis and evaluation of a ring-constrained Hsp90 C-terminal inhibitor that exhibits neuroprotective activity

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ABSTRACT

KU-596 is a second-generation C-terminal heat shock protein 90 KDa (Hsp90) modulator based on the natural product, novobiocin. KU-596 has been shown to induce Hsp70 levels and manifest neuroprotective activity through induction of the heat shock response. A ringconstrained analog of KU-596 was designed and synthesized to probe its binding orientation and ability to induce Hsp70 levels. Compound 2 was found to exhibit comparable or increased activity compared to KU-596, which is under clinical investigation for the treatment of neuropathy.

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The 90 KDa heat shock protein (Hsp90) is a molecular chaperone that is responsible for the folding and maturation of more than 300 client proteins and consequently, represents a promising target for the development of both cancer and neurodegenerative agents.¹⁻⁷ Six candidates that target the Hsp90 N-terminus are currently under clinical investigation to assess their efficacy for the treatment of cancer.8,9 Despite these efforts, Hsp90 N-terminal inhibitors induce an undesired, pro-survival heat shock response (HSR), which has impeded their clinical potential for cancer, as increased Hsp90 levels result from the administration of such inhibitors.¹⁰ This same HSR is also reponsible for the upregulation of other heat shock proteins, including Hsp70, which exhibit pro-survival activity and can refold protein aggregates that accumulate during the pathology of several neurodegenerative diseases.11,12

In contrast to N-terminal inhibitors, Hsp90 C-terminal inhibitors can segregate the HSR from the degredation of client proteins, allowing the opportunity to develop small molecules that manifest selective activity towards cancer or neurodegeneration.¹³⁻²¹ KU-32 is derived from novobiocin, the first Hsp90 C-terminal inhibitor identified (Figure 1), and was shown to exhibit cytoprotective activity at concentrations that induce the HSR.^{22,23} In fact, KU-32 can induce the HSR at \geq 500-fold lower concentrations than that required to promote client protein degradation. Consequently, KU-32 represents a

novel probe to investigate the potential neuroprotective activity of such compounds.

New analogs of KU-32 have since been developed and structure-activity relationships of these inhibitors have clearly defined attributes required for neuroprotective versus anticancer activity.^{24,25} For example, the acetamide moiety of KU-32 is necessary to induce the HSR, but replacing this group with a benzamide results in client protein degradation without induction of the HSR. Although no co-crystal structure has been solved for inhibitors bound to the Hsp90 C-terminus,²⁶⁻²⁸ models have been developed that support the binding of KU-32 at this location. One model suggests an unexplored pocket to reside in close proximity to KU-32, which provides an opportunity to design compounds that exhibit increased activity. Accordingly, a second generation of novobiocin analogs was developed to contain a biaryl core, which led to small molecules that manifest enhanced neuroprotective activity.24 Among the novologues prepared, KU-596 was found to be most active in both luciferase reporter and mitochondria bioenergetic assays (Figure 1).24, 25 When KU-596 was docked into the Hsp90 C-terminal model, the noviose sugar was found to project into a subpocket that contains amino acid side chains that contribute hydrogen bonding interactions. In addition, the fluorine atom on KU-596 appeared to form a hydrogen bond with Lys560 (Figure 3A), which is supported by prior structure-activity relationship studies.25

Since it is well-known that conformationally constrained molecules often exhibit lower entropic penalties as well as

improved affinity upon binding, an analog of KU-596 was pursued. The inclusion of a lactam to constrain the biaryl ring system also represented an opportunity to introduce hydrogen bonding interactions with the peptide backbone while rigidifying the biaryl ring system. Prior studies with KU-32 analogs supported substitution at this position, as replacement of the 8-methyl group with other functionalities led to improved activity (Figure 2).^{20, 29} Such data supports the existence of unoccupied space in the binding site and correlates well with the proposed model.



Figure 2. Design of ring-constrained novologue

The phenanthridone containing compound, 2, was designed and then docked into the Hsp90 C-terminal binding site to further investigate the mode of binding (Figure 3B). Interestingly, the fluorine atom on 2 pointed towards Gln488 rather than the aforementioned Lys560 (Figure 3B), which can still form hydrogen bonding interactions via the amide NH.





В



(A) KU-596 docked into the Hsp90 C-terminal binding site.
(B) 2 docked into the Hsp90 C-terminal binding site.
The dash line indicates the distance between the fluorine atom and hydrogen from the amino residue

Figure 3. Molecular modeling for KU596 and compound 2

In addition, the lactam carbonyl participated in a hydrogen bonding network with Gln488. As expected, the sugar moiety maintained a similar orientaion inside the pocket as was predicted for KU-596 (Figure 3A). Similarly, the 3-amido side chain projected into the same hydrophobic region as observed with KU-596.

Compound 2 was pursued via the synthetic route described in Scheme 1. Commercially available 3-nitro-4hydroxy benzaldehyde was converted to benzyl ether 3 via benzyl bromide, followed by iron-mediated reduction of the nitro substituent to form the corresponding aniline, 4. Subsequent amide coupling was performed via conversion of 2-bromo-6-fluoro benzoic acid to the acid chloride, followed by reaction with aniline 4 to ultimately yield 5. The aldehyde of 5 was then converted to acetal 6 to mask its reactive nature during subsequent steps. The intramolecular Heck coupling was known to be catalyzed by palladium in the presence of silver carbonate.³⁰ However, attempts to perform an intramolecular Heck coupling without protection of amide 6 was unsuccessful. Therefore, the amide was methylated to give 7 before cyclization was successfully achieved to produce 8 under modified conditions.³¹ Compound 8 was then converted to 9 via Henry olefination. Compound 11 was then synthesized from 9 via a two-step reduction before acetylation with acetyl chloride. Finally, the activated noviose sugar 13 was prepared³² and appended to phenol **12** upon removal of benzyl group to yield the final product, 2.

To evaluate the ability of **2** to promote Hsp70 induction compared to **1**, western blot analyses were performed in immortalized dorsal root ganglia sensory neuronal 50B11 cells. As depicted in Figure 4, compound **2** is more effective than **1** at the induction of Hsp70 levels. The slight decrease of Hsp70 at the highest concentration of **2** is a common attribute ascribed to this class of modulators, which typically give a U-shape curve upon dosing with Hsp90 C-terminal inhibitors.²⁴

Scheme 1: Synthesis of 2



Reaction conditions: (a) BnBr, K_2CO_3 , Acetonitrile, reflux, overnight, 94%; (b) Fe, NH₄Cl, THF/EtOH/H₂O, reflux, 2h, 75%; (c) 2-bromo-6-fluoro benzoyl chloride, Et₃N, CH₂Cl₂, rt, overnight, 83%; (d) trimethyl orthoformate, TsOH, MeOH, 50 °C, 2h, 87%; (e) NaH, MeI, THF, 0 °C to rt, overnight; (f) Pd(PPh₃)₄, Ag₂CO₃, dioxane, 90 °C, overnight; (g) TsOH monohydrate, acetone, rt, 2h, 76% over two steps; (h) CH₃NO₂, NH₄OAc, 80 °C, overnight, 86%; (i) NaBH₄, Dioxane/MeOH, 0 °C to rt, 2h, 83%; (j) Zn, 1N HCl, EtOH/CH₂Cl₂, rt, 4h; (k) Acetyl chloride, Et₃N, CH₂Cl₂, rt, 3h, 60% over two steps; (l) Pd/C, H₂, EtOAc, r.t., overnight, 75%; (m) **13**, BF₃·OEt₂, CH₂Cl₂, rt, 3h; (n) Et₃N, MeOH, rt, overnight, 30% over two steps.



Figure 4. Western Blot analysis of Hsp70 induction for 2. 50B11 cells were seeded at 2 x 10^5 cells per well and cultured for 24hr at 37 °C prior to treatment with DMSO or the indicated concentrations of KU-596 or 3 for an additional 24 hr. The cells were harvested to determine Hsp70 protein expression level by immunoblotting with and Hsp70/Hsp72 biotin-conjugated monoclonal antibody.

Excessive oxidative stress can also cause mitochondria dysfunction in diabetic sensory neurons, which is a key contributor to the pathogenesis of diabetic peripheral neuropathy (DPN).³³ Thus, oxidative stress can be induced by treating 50B11 cells with 0.5mM hydrogen peroxide to decrease mitochondrial bioenergetics (mtBE) as measured by the Seahorse assay and as indicated by a decreased response to the protonophore, FCCP. Both 1 (Figure 5A) and 2 (Figure 5B) can effectively improve mtBE in a dose-dependent manner. However, drug 2 may be more effective than 1, since

it can significantly increase cellular maximal respiratory capacity (MRC), starting at concentrations of ~ 100 nM (Figure 5C). The improvement of mitochondrial bioenergetics in sensory neurons may be one of the key mechanisms that contributes to the neuroprotective efficacy of both 1 and 2 as well as related molecules.





Figure 5. Improvement in mitochondrial bioenergetics by 2. The 50B11 cells were seeded into a 96 well plate and after 12 hrs the cells were changed into high glucose DMEM medium with 1% FBS, containing the indicated concentrations of 1 (A) or 2 (B) for 24 hr at 37°C. The cells were stressed with 0.5mM H₂O₂ for 2 hr, the medium changed to serum free unbuffered DMEM with 5.5 mM glucose and 1mM pyruvate and the cells incubated at 37°C for 1 hr prior to assessing oxygen consumption rate (OCR) with an XF96 Extracellular Flux Analyzer. (C) Maximal respiratory capacity (MRC) was calculated for each treatment. Results are from one experiment using 5 wells of cells per treatment. *, p < 0.05 vs H₂O₂ alone.

In conclusion, a ring-constrained novologue was synthesized and then evaluated by western blot analyses and mitochondrial bioenergetic assays. Compound 2 exhibited increased activity for mitochondrial bioenergetics against oxidative stress as compared to 1 at all concentrations except a slightly better activity was observed at 100 nM. This result is consistent with the proposed docking studies and supports a similar binding pose for both 1 and 2.

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Supplementary Material

Procedure for the docking study, synthesis of 3-12, western blot analysis, Seahorse assay, characterization and NMR spectra of **3-12**.

Accepter