

Identification of a Covalent Importin-5 Inhibitor, Goyzensolide, from a Collective Synthesis of Furanoheliangolides

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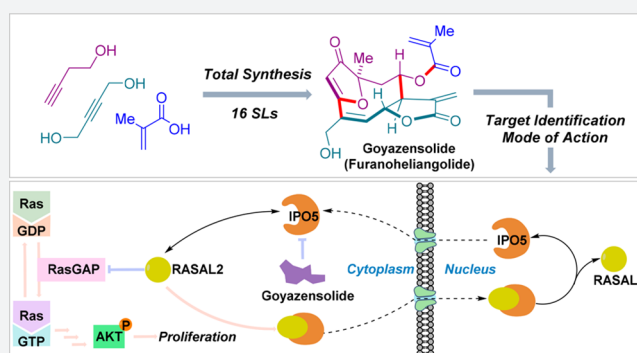


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ABSTRACT: Sesquiterpenes are a rich source of covalent inhibitors with a long history in traditional medicine and include several important therapeutics and tool compounds. Herein, we report the total synthesis of 16 sesquiterpene lactones via a build/couple/pair strategy, including goyzensolide. Using an alkyl-tagged cellular probe and proteomics analysis, we discovered that goyzensolide selectively targets the oncoprotein importin-5 (IPOS) for covalent engagement. We further demonstrate that goyzensolide inhibits the translocation of RASAL-2, a cargo of IPOS, into the nucleus and perturbs the binding between IPOS and two specific viral nuclear localization sequences.



INTRODUCTION

Goyzensolide (1, Figure 1), a member of the furanoheliangolide sesquiterpene family, has been isolated from several plants used in traditional medicine and was first identified in a screen for schistosomicidal agents.¹ It has also been shown to be effective in an animal model bearing a colon tumor (HT-29 xenograft)² and as a potential treatment for neurofibromatosis type 2 (NF2),³ for which there is currently no FDA-approved intervention. There is clear evidence that goyzensolide inhibits the expression of genes under the control of *c-myc*, a proto-oncogenic transcription factor;⁴ however, its exact mode of action remains undefined. 15-Deoxygoyzensolide (3, Figure 1) was found to be slightly more cytotoxic than goyzensolide,⁵ suggesting that small modifications can enhance activity. 15-Deoxygoyzensolide was also found to be active in a number of quantitative-cell-based reporter assays.⁶

The activity of sesquiterpenes often hinges on the presence of Michael acceptors that can engage a target covalently.^{7,8} A well-studied example is helenalin, used therapeutically as an anti-inflammatory, which inhibits the NF- κ B pathway by covalent interaction with cysteines in p65, thereby precluding the interaction of p65 with DNA.⁹ Another example is ainsliadimer A, which was shown to selectively inhibit IKK kinase by reaction with Cys46 present in an allosteric pocket.¹⁰ Our own studies on deoxyelephantopin led to the discovery that it reacts with PPAR- γ and antagonizes the function of this nuclear receptor by engaging a cysteine located in the zinc-finger motif.¹¹

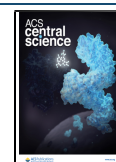
While some sesquiterpenes are readily available from natural sources, understanding the covalent interactome requires the synthesis of probes that cannot always be achieved by direct functionalization of the natural product and requires a total synthesis.

RESULT AND DISCUSSION

Collective Synthesis of Furanoheliangolides. Furanoheliangolides are secondary metabolites produced from farnesyl pyrophosphate by cyclization, involving the germacrenyl cation intermediate and subsequent oxidative reactions (Figure 1a).¹² The promiscuity of this biosynthetic pathway yields a range of closely related analogues akin to combinatorial library synthesis.¹³ The collective synthesis of furanoheliangolides and related natural products (1–17, Figures 1 and S1), including goyzensolide (1), was envisioned via a gold-catalyzed transannulation of key intermediates 18 to generate the strained bicyclic framework (Figure 1b).¹⁴ Disconnection of the C7–C8 and C3–C4 bonds (Barbier-type allylation and Sonogashira coupling, respectively) would reduce the complexity to readily available synthons (20–23) amenable to a build/couple/pair strategy.^{15,16}

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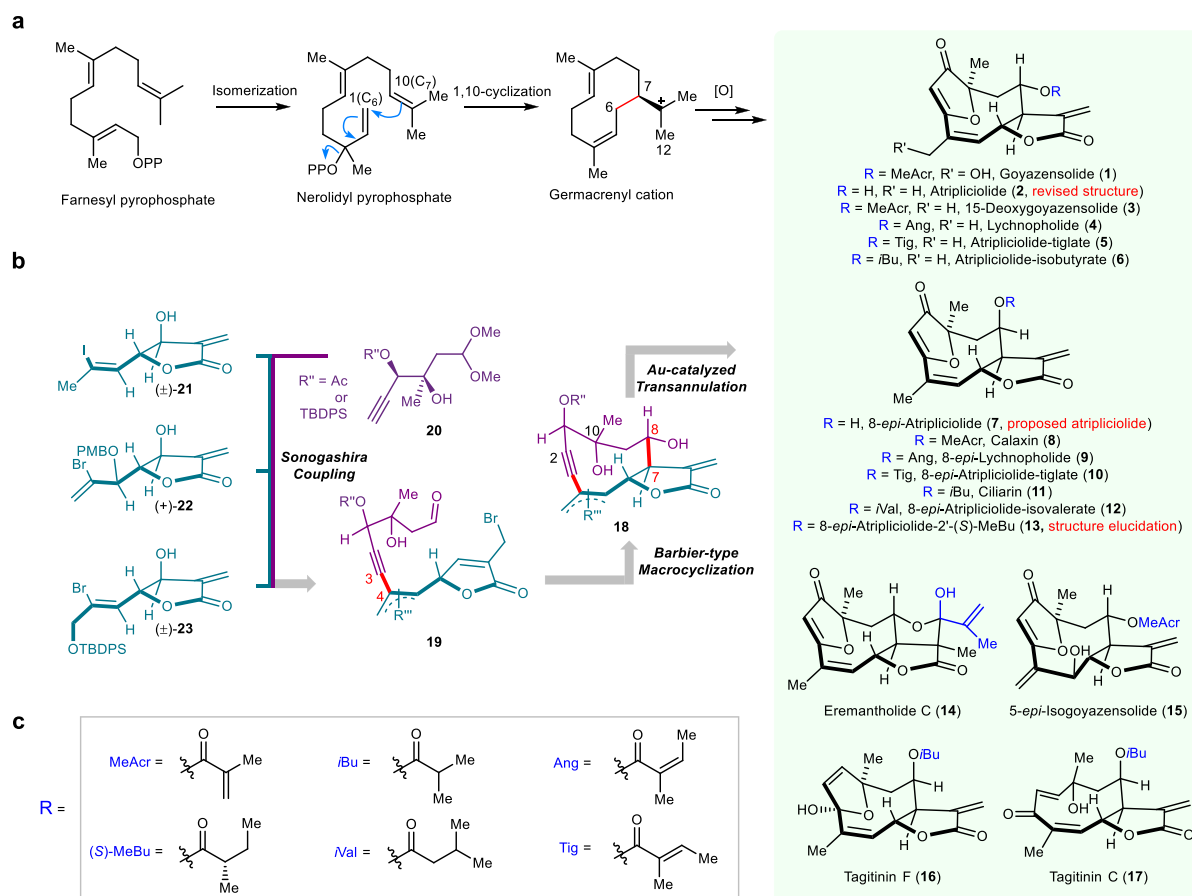


Figure 1. Biosynthesis and retrosynthetic analysis for the collective synthesis of furanoheliangolides through a build/couple/pair strategy. (a) Schematic biosynthesis of furanoheliangolides and related sesquiterpene (1–17) from farnesyl pyrophosphate. (b) Retrosynthetic analysis for the collective synthesis of 1–17. (c) Structure of the different side chains (R).

In the synthetic direction, key intermediate **20a** (Scheme 1) was obtained in five steps from vinyl alcohol **24** via a sequence involving a one-pot Dess–Martin periodinane (DMP) oxidation, acetal protection followed by a Sonogashira coupling with trimethylsilylacetylene (**24** → **25**), Sharpless asymmetric dihydroxylation (SAD) of the enyne,¹⁷ selective secondary alcohol protection (**25** → **26**), and desilylation of the acetylene.¹⁸ The sequence delivers the intermediate **20a** with 39% overall yield and 91% *ee* on a decagram scale. The second intermediate **21** was prepared in two steps from aldehyde **27** and allyl bromide **28** by Barbier allylation/lactonization (**27** + **28** → **29**) followed by SeO₂ allylic oxidation (**29** → **21**). While asymmetric synthesis of this synthon is possible, we used a racemic product with an interest for both diastereomers resulting in the pairing of alkyne **20a** and vinyl iodide **21**.

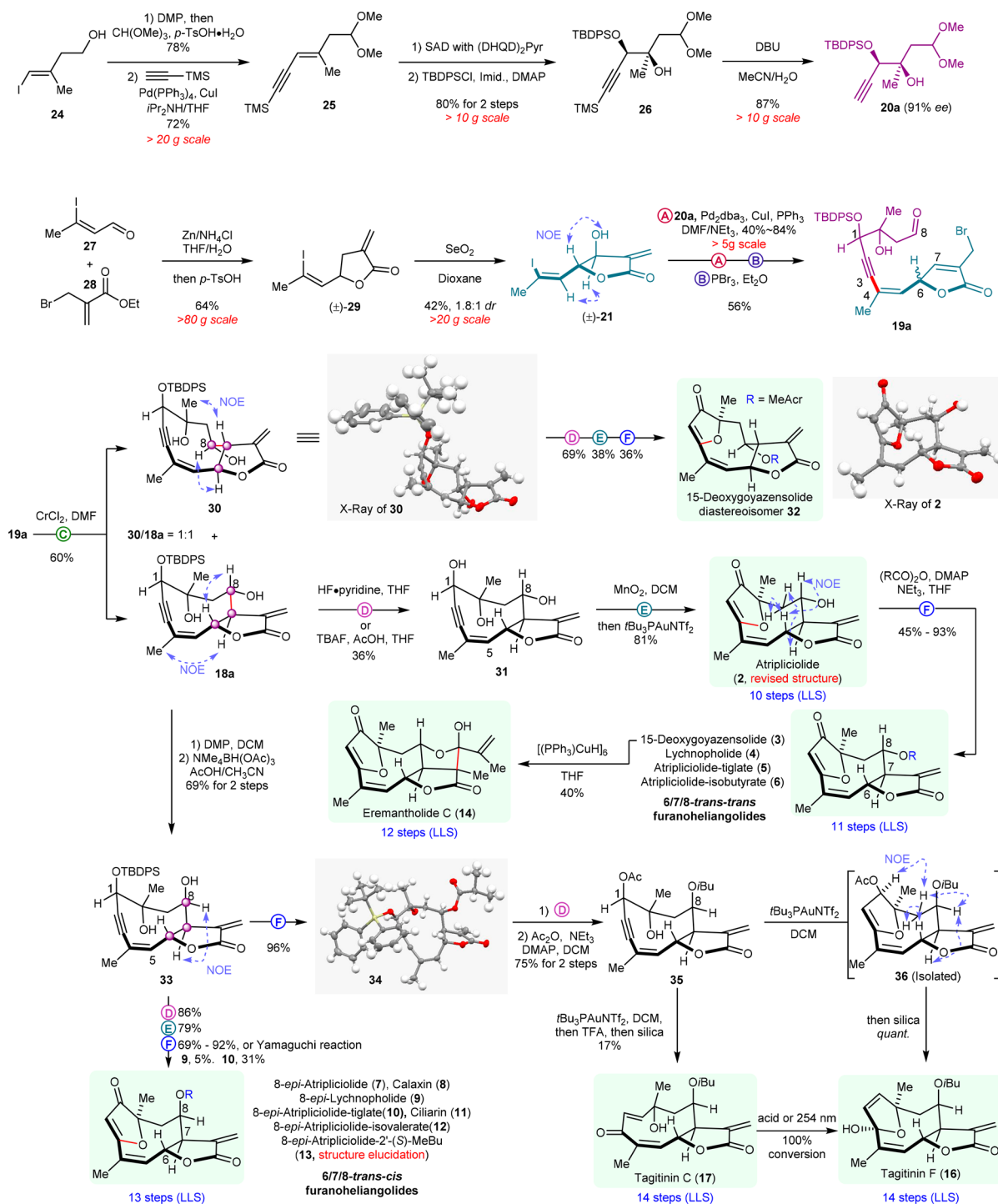
Sonogashira coupling (**20a** + **21**, reaction A) followed by PBr₃ (reaction B) to convert the allylic alcohol in a bromide (S_N2') with concomitant deprotection of the dimethyl acetal afforded the cyclization precursor **19a**. Treatment of **19a** with CrCl₂ (reaction C) in DMF afforded a clean transformation to the desired germacrene framework as a separable mixture of the two diastereomers (**18a** and **30**) arising from the undefined stereocenter at C6. While nuclear Overhauser effect (NOE) patterns clearly discerned the two diastereoisomers, the structure of **30** was confirmed by X-ray diffraction (CCDC2049428, Supplementary Section m). The strained 10-member ring of the germacrene framework is challenging to

synthesize, and to the best of our knowledge, this is the first example merging the α -methylene- γ -butyrolactone formation with the closure of a 10-membered germacrene framework. Further diversification to the furanoheliangolide proved to be highly efficient following the removal of the silyl protecting group (**18a** → **31**, reaction D). Selective oxidation of the propargyl alcohol (MnO₂) followed by treatment with *t*Bu₃PAuNTf₂ (reaction E) afforded the furanoheliangolide **2**.

It is noteworthy that the spectral data of **2** matched the reported data of atriplicioidide.¹⁹ The stereochemical assignment was further validated by X-ray analysis (CCDC2049427, Supplementary Section m). Analysis of other possible diastereoisomers on route to **32** or **7** did not match the NMR data, thus allowing a revision of the structure of atriplicioidide as **2** (Supplementary Section c). Acylation of the unique hydroxyl group of atriplicioidide with the different possible side chains (reaction F) afforded 15-deoxygoyazensolide (**3**) and related natural products **4–6** (Supplementary Section e). 15-Deoxygoyazensolide (**3**) was further converted to eremantholide C (**14**) using Stryker's reagent.²⁰ The reaction sequence (D–F) to convert the germacrene framework to furanoheliangolide was also used to convert **30** into a hitherto unreported 15-deoxygoyazensolide diastereoisomer **32**.

Based on a number of furanoheliangolides having an *R*-stereochemistry at C8, we sought to invert the stereochemistry of the hydroxyl at this position in intermediate **18a** (Scheme

Scheme 1. Diversification of Germacrene Framework and Synthesis of Furanoheliangolides



1). The most efficient protocol proved to be a DMP oxidation followed by an Evans–Saksena reduction²¹ leveraged on the chelation of the C10 hydroxyl, to afford 33 as a single diastereoisomer. The stereochemical assignment was verified by X-ray diffraction of the acylated intermediate 34 (CCDC2049429, Supplementary Section m). Applying the reaction sequence (D–F) to convert this germacrene framework (33) to furanoheliangolides afforded six new natural products (8–13, Supplementary Section e). These syntheses also allowed the stereochemistry to be established in the side chain of 8-*epi*-atripliciolide-2'-(*S*)-MeBu,²² which remained

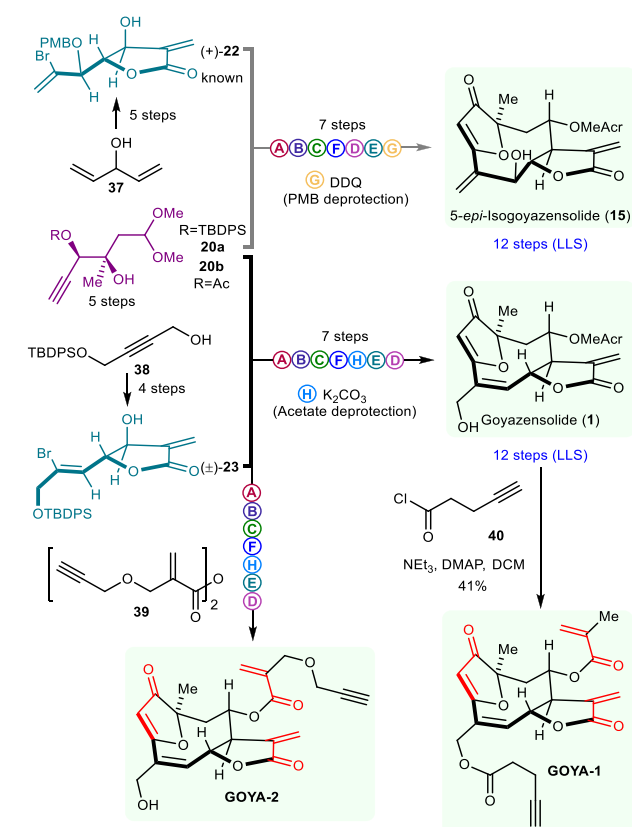
undefined hitherto. Comparison of other diastereoisomers provided an unequivocal match (Supplementary Section d).

Germacrene 34 was used to access yet another set of natural products employing a slight modification in the reaction sequence (Scheme 1): acetylation of the C1 hydroxyl (35) rather than oxidation to a propargyl ketone as in the previous sequence. While we anticipated that 35 could be converted to tagitinin C (17) using the gold catalysis ($t\text{Bu}_3\text{PAuNTf}_2$) directly via a Meyer–Schuster rearrangement,²³ we obtained tagitinin F (16) instead. However, using the same catalyst in DCM followed by a TFA treatment did yield tagitinin C (17),

albeit in low yield due to its instability ($17 \rightarrow 16$).²⁴ The synthetic **16** and **17** were identical to authentic samples (Supplementary Section e). This insight came from analysis of the reaction pathway by NMR (Figure S7). Based on the observed reaction intermediates, we propose that the mechanism (Figure S8) proceeds by a gold activation of the alkyne to yield intermediate **36**. This intermediate then undergoes a [3,3]-sigmatropic rearrangement followed by a hydrolysis to afford tagitinin F (**16**). Given sufficiently acidic conditions, the acetal can open to facilitate the *Z*- to *E*-olefin isomerization, thus yielding tagitinin C (**17**).

Exploiting the same synthetic sequence with a different pairing strategy afforded *5*-*epi*-isogoyazensolide (**15**) and goyazensolide (**1**) as shown in Scheme 2. *5*-*epi*-Isogoyazensolide

Scheme 2. Further Pairing and Stereoselective Synthesis of *5*-*epi*-Isogoyazensolide (**15**), Goyazensolide (**1**), and Probes GOYA-1 and GOYA-2



lide (**15**) comes from the pairing of **22** with **20a** (propargylic hydroxyl protected with *tert*-butyldiphenylsilyl), whereas goyazensolide comes from the pairing of **23** with **20b** (propargylic hydroxyl protected with an acetate). Synthon **22** was prepared from 1,4-pentadien-3-ol (**37**) in five steps by a known protocol²⁵ (asymmetric center introduced by Sharpless asymmetric dihydroxylation) followed by a diastereoselective allylic oxidation (SeO_2). Synthon **23** was prepared following the same sequence as for **21** from known alcohol **38**.

It is important to note that the pairing of **22** and **20a** yields a single diastereoisomer affording **15** enantioselectively. In both cases, the spectroscopic data fully matched that of the natural isolate (Supplementary Section e). These syntheses also enabled the preparation of goyazensolide probes tagged with

an alkyne group (GOYA-1 and GOYA-2) for subsequent labeling studies.

Thus, this simple build/couple/pair sequence afforded germacrene frameworks (**18a**, **30**, **33**, etc.), which can be further diversified into 16 natural products (**1–6**, **8–17**) within 10–14 steps, none of which had been previously synthesized.

Goyazensolide Interacts Covalently with Importin-5 (IPO5). The presence of multiple Michael acceptors in goyazensolide (**1**) and their similarity to those of other covalent inhibitors in the sesquiterpene lactone family led us to speculate that goyazensolide interaction with its target could be also covalent. Indeed, the α -*exo*-methylene- γ -butyrolactone moiety of goyazensolide is present in many sesquiterpenes (>1000), including helenalin (**H**, Figure 2e), and the disposition of this functionality with respect to the second Michael acceptor (methacrylate side chain) is precisely the same as in deoxyelephantopin (**D**, Figure 2e). The furanone endows goyazensolide with a third Michael acceptor. Having access to the natural product, we decided to synthesize a goyazensolide probe with the intention of preparing an activity-based probe for a proteome-wide target screen (Figure 2a).^{26,27} With this in mind, we first acylated the primary allylic alcohol with 4-pentynoyl chloride (**40** \rightarrow GOYA-1, Scheme 2), as had been previously done successfully for helenalin derivatives.²⁸ To our surprise, alkyne-tagged GOYA-1 did not afford meaningful protein labeling in a crude extract (30 min of labeling, CuAAC coupling with Cy3- N_3 and SDS-PAGE analysis, Figure 2b, left). Increasing the probe concentration and reaction time did not improve the results.

We thus explored another position to introduce the alkyne and *de novo* synthesized GOYA-2 (Scheme 2), with the alkyne extending from the methyl acrylate side chain. Treatment of crude lysate from PC3 cells with GOYA-2 followed by reaction with Cy3- N_3 and SDS-PAGE revealed a strong band at around 120 kDa that was specifically competed when the lysate had been previously treated with the unmodified goyazensolide (Figure 2b, right). Lysates from five other cancer cell lines (HeLa, K562, U2OS, HT-29, SW620) showed a similar profile, suggesting this target was not cell-line-specific (Figure S15). Competition with other sesquiterpenes (Figure 2e) containing the same α -*exo*-methylene- γ -butyrolactone (helenalin, lactucin), the similar unsaturated side chain (thapsigargin), or the combination of both (deoxyelephantopin, cnicin, cynaropicrin) showed that none of them competed for the same target as goyazensolide (Figure 2f).

In order to enrich the targeted proteins, we performed a pull-down using streptavidin beads of the cell lysate treated with GOYA-2 after reaction with desthiobiotin-Cy3- N_3 (Figure S13) and visualized the result on an SDS-PAGE. A triple in-gel digestion of the Cy3 band with trypsin/GluC/chemotrypsin and analysis of the sample by MS/MS afforded a list of peptides corresponding to proteins, which were filtered by molecular weight (between 105–145 kDa) and coverage (>15 unique peptides detected for each protein), resulting in four candidates (Figure 2c). The protein with highest coverage was IPO5 (63 unique peptides, 41% coverage) followed by pyruvate carboxylase (52 unique peptides, 33% coverage), UBA1 (19 unique peptides, 18% coverage), and ACLY (18 unique peptides, 13% coverage). The identity of the target was confirmed by a Western blot with an antibody specific for IPO5. The same experiment with anti-UBA1 antibody or anti-ACLY antibody did not yield signal above background (Figure

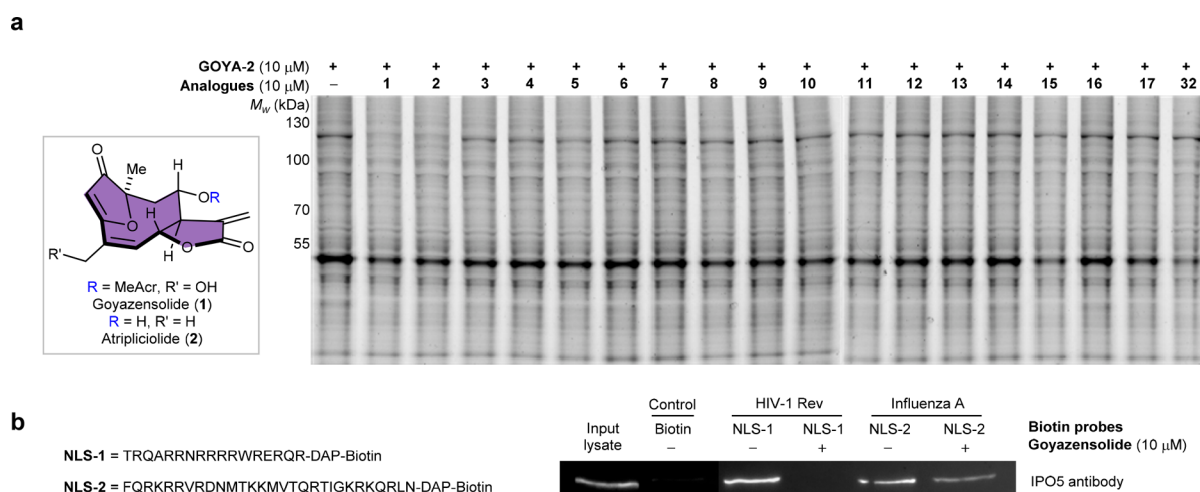


Figure 3. Competition experiments with the other natural products synthesized, and goyzensolide inhibits the interaction of IPO5 with specific NLSs. (a) SW620 lysate was incubated for 30 min with DMSO or one of the natural products (10 μM , 30 min) and then labeled with 10 μM GOYA-2 followed by a CuAAC reaction with Cy3-N₃ for 1 h at RT in the dark and analyzed by SDS-PAGE using fluorescence as a readout. The labeling/competition experiment shows selective binding of goyzensolide (1) and atriplicolide (2) to IPO5. (b) Left, sequence of two viral biotinylated NLSs, NLS-1 (HIV Rev) and NLS-2 (PB1 influenza A). Right, immunoblot of competitive interaction between NLS and IPO5. Magnetic streptavidin beads were saturated with corresponding NLSs and then incubated for 2 h with HT29 lysate treated with DMSO or goyzensolide (10 μM , 30 min).

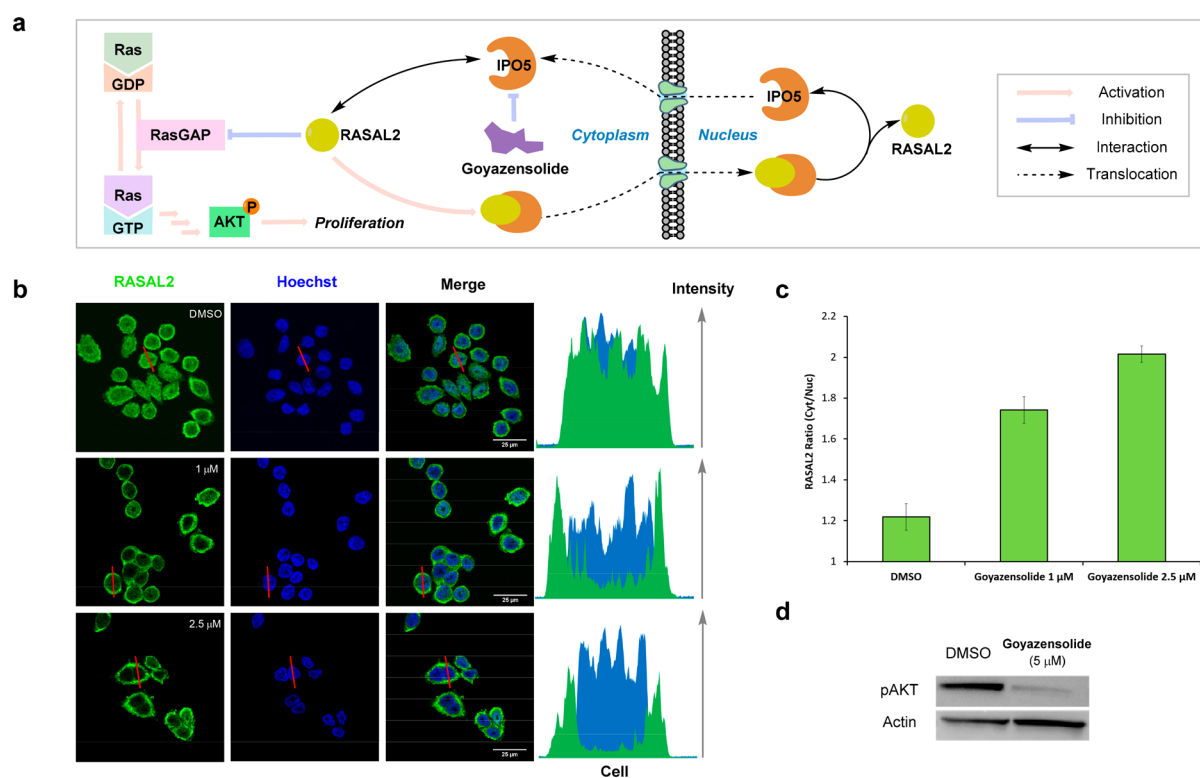


Figure 4. Covalent engagement of IPO5 with goyzensolide inhibits its transport activity in cells. (a) Illustration of IPO5 involvement in the Ras pathway. (b) Confocal microscopy images of SW620 cells treated with DMSO or 1 or 2.5 μM goyzensolide for 3 h, imaging of localization of RASAL2 using a RASAL2 primary antibody followed by an Alexa fluor488 secondary antibody (green), nuclear stain with Hoechst (blue), and fluorescence quantification of the images. Fluorescence intensity profile across the red line is shown on the right. (c) Ratio of fluorescence in cytosol versus nucleus (number of cells > 15). (d) Immunoblot of p-AKT (p-AKT Ser473 antibody) expression upon treatment of SW620 cells with DMSO or goyzensolide (5 μM , 4 h).

essential cargo to the nucleus, and disruption of the interaction between the NLS of the influenza A protein PB1 and IPO5 inhibits viral replication.²⁹ IPO5 is also important for the transport of viral proteins of HCV and HIV.³⁰ To test whether goyzensolide was effective at inhibiting the interaction

between IPO5 and the NLSs of viral proteins, the sequences NLS-1 (NLS of Rev; HIV) and NLS-2 (NLS of PB1; influenza A) (Figures 3b and S20) were coupled to biotin tags and loaded on magnetic beads for pull-down experiments with IPO5. In the absence of an inhibitor, both NLS sequences

effectively pulled-down IPO5, as observed in a Western blot of the elution fraction (Figure 3b). Addition of goyazensolide (10 μ M) to the cell lysates prior to the addition of the beads inhibited the pull-down of IPO5. Thus, the covalent interaction between goyazensolide and IPO5 inhibits its interaction with these two NLSs.

Goyazensolide Inhibits the Cellular Function of IPO5.

IPO5 has been shown to be involved in the transport of RAS protein activator-like 2 (RASAL2) to the nucleus, a negative regulator of RasGAP. Thus, nuclear translocation of RASAL2 frees RasGAP to convert inactive Ras-GDP to active Ras-GTP as shown in Figure 4a.³¹ The ratio of cytosolic RASAL2 and nuclear RASAL2 was clearly shown to depend on IPO5 using siRNA silencing of IPO5 (increased cytosolic concentration of RASAL2) and IPO5 overexpression (decreased cytosolic concentration of RASAL2).³¹ In order to understand if goyazensolide could phenocopy the silencing activity of siRNA targeting IPO5, an SW620 colon cancer cell line was used to measure the RASAL2 ratio in the cytosol vs the nucleus in the presence and the absence of goyazensolide (Figure 4b). In the absence of treatment, the cytosolic/nuclear ratio of RASAL2 is 1.2. Upon treatment with 1.0 or 2.5 μ M goyazensolide for 3 h, the ratio increased to 1.8 and 2.0, respectively (Figure 4c). Thus, goyazensolide treatment led to an increased cytosolic /nuclear ratio of RASAL2 in a dose dependent fashion.

Since the inhibition of RASAL2 translocation to the nucleus should have a downstream effect on the level of pAKT, we also monitored the level of pAKT (Ser473) in the presence or absence of drug. As shown in Figure 4d, treatment with 5 μ M goyazensolide led to a dramatic reduction of pAKT. Collectively, these experiments demonstrate that IPO5 is the target of goyazensolide and that covalent engagement of IPO5 inhibits its transport activity *in cellulo*.

Sesquiterpenes are a rich source of bioactive compounds, and their syntheses continue to challenge synthetic chemists (for recent examples, see refs 32–35). From a synthetic chemistry perspective, goyazensolide stands out as a highly oxidized member of the germacranolides with protean C3 and C10 bridged furan ring featuring an anti-Bredt double bond. The present work demonstrates that despite its complexity, such natural products can now be accessed synthetically in less than 13 steps. The strategy employed is compatible with a build/couple/pair strategy facilitating the collective synthesis of multiple natural products, including goyazensolide. This required a cyclization to form a 10-membered ring, which is notoriously difficult. The use of Barbier allylation to simultaneously install the α -*exo*-methylene- γ -butyrolactone and achieve this cyclization is unprecedented in this context, and the work reported indicates that it is a robust strategy to access the strained framework. The use of an alkyne in the ring facilitates the pairing chemistry by mild and efficient Sonogashira coupling, and the alkyne offers divergent functionalization pathways to furanoheliangolide or related heliangolides. The chemistry described enabled the collective synthesis of 16 complex sesquiterpene lactones, many with important biological activity (See Figure S1 for a complete set of reference to biological activity).

The use of some of these compounds in traditional medicine gives them the benefit of a preliminary toxicology profile in humans. However, the lack of information on the mode of action often precludes a more insightful and targeted use of such compounds. In the case of goyazensolide, there was clear

evidence of anticancer activity from mouse xenografts and cellular assays, but the mode of action remained unknown. The finding that goyazensolide selectively engages IPO5 covalently and inhibits its interaction with cargo rationalizes its anticancer activity and points to potential antiviral applications. IPO5 is highly expressed in colorectal cancer³¹ and esophageal cancer.³⁶ As an antiviral target, there is clear evidence that IPO5 is recruited by viruses and that its trafficking activity is essential in the viral replication cycle.^{29,30}

The fact that a small molecule can outcompete a high-affinity interaction with larger molecular entities such as the interaction of transportin with an NLS sequence of a protein underpins the unique features of covalent inhibitors. Transport of cargo between the cytoplasm and nucleus is a highly regulated process with the nuclear pore complex acting as a molecular sieve restricting diffusion. Translocation thus requires active transport leveraged on the action of a superfamily of karyopherin (importin and exportin).³⁷ The ability to modulate this traffic with small molecules is empowering,^{38,39} but there are only few examples thus far. The first example of a small molecule regulating a karyopherin is leptomycin B, a polyketide bearing a Michael acceptor, which selectively inhibits exportin-1 (XPO-1, also known as chromosome region maintenance 1 (CRM1))⁴⁰ by reacting with its target.⁴¹ This discovery led to significant medicinal chemistry efforts resulting in a molecule (selinexor) that is FDA-approved for multiple myeloma.⁴²

The second example, ivermectin, was identified from a high-throughput screen for inhibitors of protein nuclear import. Characterization of the nuclear transport inhibitory properties of ivermectin demonstrated that it is a broad-spectrum inhibitor of importin α/β nuclear import with potent antiviral activity toward both HIV-1 and dengue virus, both of which are strongly reliant on importins.⁴³ Notably, this FDA-approved drug also inhibits SARS-CoV-2 replication. Studies on SARS-CoV proteins have revealed a potential role for importin $\alpha/\beta 1$ heteroduplex in nucleocytoplasmic shuttling of the SARS-CoV's nucleocapsid protein.⁴⁴ IPO5 has clearly been shown to be implicated in the transport of influenza A's RNA polymerase,⁴⁵ and mutations affecting binding to IPO5 severely attenuate viral growth,²⁹ suggesting this activity could be phenocopied with small molecule inhibitors of IPO5. The crystal structure of IPO5 coupled to docking studies of the NLS revealed critical interactions of influenza's PA–PB1 subcomplex with IPO5.⁴⁶

CONCLUSION

In summary, we report the first total synthesis of goyazensolide and 15 related natural products. Our studies revise the stereochemical assignment of two natural products. Significantly, this synthesis enabled the study of the covalent interactome of goyazensolide, revealing IPO5 as its molecular target. The covalent engagement inhibits the interaction of IPO5 with cargos, provides a rationale for its cytotoxic activity in several cancer cell lines, and points to potential antiviral activity.

The resurging interest in covalent inhibitors brings renewed interest to sesquiterpenes, particularly those that have already been used in traditional medicine, affording a preliminary assessment of toxicity. Covalent inhibition is particularly attractive if target binding is in competition with strong endogenous interactions such as in protein–protein interactions or transcription factors interacting with DNA.

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acscentsci.1c00056>.

Additional figures and tables, experimental details for the synthesis of all new compounds including experimental procedures, characterization and spectral data, detailed information for biological materials, cell culture and preparation of lysates, pull-down experiments, in-gel tryptic digestion, LC-MS/MS analysis, mass spectrometry analysis, preparation of NLS-1 and NLS-2, pull-down experiment with viruses' NLS, microscopy and data analysis, and pAKT assay (PDF)

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Notes

The authors declare no competing financial interest.

The raw data has been deposited on Zenodo (<https://zenodo.org/record/4314704>).

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