Atropselective syntheses of (–) and (+) rugulotrosin A utilizing point-to-axial chirality transfer

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Chiral, dimeric natural products containing complex structures and interesting biological properties have inspired chemists and biologists for decades. A seven-step total synthesis of the axially chiral, dimeric tetrahydroxanthone natural product rugulotrosin A is described. The synthesis employs a one-pot Suzuki coupling/dimerization to generate the requisite 2,2'-biaryl linkage. Highly selective point-to-axial chirality transfer was achieved using palladium catalysis with achiral phosphine ligands. Single X-ray crystal diffraction data were obtained to confirm both the atropisomeric configuration and absolute stereochemistry of rugulotrosin A. Computational studies are described to rationalize the atropselectivity observed in the key dimerization step. Comparison of the crude fungal extract with synthetic rugulotrosin A and its atropisomer verified that nature generates a single atropisomer of the natural product.

s a consequence of the hindered rotational barriers of highly substituted biaryls and related compounds, a type of stereochemistry in the form of axial chirality is generated, often termed atropisomerism¹. Atropisomerism exists widely in nature and can lead to biological stereoselectivity for molecular targets^{2–4}. Moreover, in contrast to traditional sp^3 carbon-centred chirality, control of axial chirality and atropisomer selectivity remains highly challenging, especially in the context of complex natural product synthesis^{3–8}.

Tetrahydroxanthones9-secondary fungal metabolites-are an emerging family of natural products. Dimeric tetrahydroxanthones¹⁰ including the secalonic acids^{11,12} (Fig. 1a) display intriguing anticancer and antibacterial properties. During the last decade, tetrahydroxanthones containing biaryl axial chirality including the 2,2'-linked natural product¹³ rugulotrosin A (1) and the 2,4'-linked congener rugulotrosin B (2) (Fig. 1b) have been isolated and characterized. Related hybrid chromone lactone/tetrahydroxanthones including the 2,4'-linked heterodimer gonytolide $E(3)^{14}$ and the 4,4'-linked chromone lactone homodimer gonytolide A (4)¹⁵ have also been reported. Recently, these interesting molecules have attracted significant interest in the synthetic community, as evidenced by reports of syntheses of both monomeric¹⁶⁻²² and dimeric natural products²³. However, syntheses of tetrahydroxanthones bearing axial chirality, such as rugulotrosin A (1) and related compounds, have not been reported.

To establish axial chirality for rugulotrosin A, we envisioned that a point-to-axial chirality transfer strategy could be implemented. In the last few decades, point-to-axial chirality transfer has been widely used in atropselective synthesis. There are several different approaches to point-to-axial chirality transfer: (1) the use of an *ortho*-auxilary^{5,6} or chiral leaving group²⁴ for intermolecular, atropselective coupling; (2) utilization of low-energy conformers derived from existing chirality centres^{25,26} or chiral auxiliaries^{5,6} to execute intramolecular couplings; (3) transfer of chiral sp^3 carbon to axial chirality^{27,28}; and (4) the use of remote and existing chirality centres to influence atropselectiviy using intermolecular oxidative/ metal-mediated couplings²⁹⁻³³. Shaw and colleagues, for example, have reported that the chiral monomer 5 underwent oxidative coupling to generate atropisomer 6 in 76:24 diastereomeric ratio (d.r.), which could be improved to higher selectivity using a chiral catalyst system²⁹ (Fig. 1c). Similarly, Lipshutz and coworkers performed an atropselective cross-coupling to access the korupensamine B precursor 9 using boronic acid 7 and iodotetrahydroisoquinoline partner 8^{31} (Fig. 1c). In the present studies we considered a remote asymmetric induction strategy for point-toaxial chirality transfer for rugulotrosin synthesis in the absence of chiral auxiliaries. In the case of dimerization of tetrahydroxanthone substrate 10, it was of interest to determine whether the chirality of the remote stereocentres could be transferred to the axial chirality of ortho, ortho dimer 11 (Fig. 1d), a product bearing a coupling pattern different from those of previous examples. In this case we planned to make use of transition-metal catalysis to bring the two monomers in close proximity to enhance the potential for chirality transfer.

Results and discussion

Our synthesis was initiated with chromone 12, which was prepared in greater than 50 g batches and purified by recrystallization (Fig. 2). Utilizing diisopropylsilyl ditriflate to activate chromone 12 to a siloxybenzopyrylium species²², vinylogous addition of 2-trimethylsiloxyfuran was achieved, which was followed by hydrogenation to afford chromone lactone 13 on a decagram scale (89% yield, d.r. > 10:1) after purification by trituration (Supplementary pages 4–19). Further treatment of 13 with NaH in THF led to the production of *epi*-blennolide C 14 through Dieckmann cyclization. Due to the propensity of the vinylogous acid moiety of tetrahydroxanthone 14 to both epimerization and tetrahydroxanthone-to-chromone lactone rearrangement, we utilized the neutral methylating

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Me CO₂C OH Metal catalyst



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OR

MeO₂C OH

reagent trimethylsilyldiazomethane to obtain the desired O-methylated product **16** in 61% yield as well as the separable isomer **15** (20% yield). After phenol-directed *ortho*-iodination of **16**, the derived aryl iodide **17** was subjected to kinetic acylative resolution using Birman's catalyst³⁴ on a gram scale (selectivity (s) factor >200)³⁵. Chiral monomer (–)-**17** was isolated in 46% yield (99% enantiomeric excess (e.e.)) and the O-acylated product (+)-**18** was obtained in 46% yield (99% e.e.) after recrystallization.

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After obtaining the two enantiopure tetrahydroxanthone monomers, our initial plan was to make use of our stannane dimerization strategy, which successfully led to syntheses of secalonic acids A and D^{23} . However, after evaluation of various conditions, we observed only trace stannylation of iodides **17** or **18** (Table 1), which may be due to steric hindrance of the aryl iodide substrate. Subsequently, we considered that the less bulky borate may more readily undergo transmetallation. Gratifyingly, iodide (–)-17 was found to be stable under basic conditions and to produce an inseparable mixture of dimeric products (d.r. = 88:12) under one-pot Suzuki coupling conditions using Pd(OAc)₂/SPhos³⁶ and *bis*(pina-colato)diboron. We subsequently attempted to establish the stereochemistry of the major atropisomer by NMR analysis and by acid-mediated deprotection of both atropisomers **19** and **20** to **1** and **23** (Fig. 4). However, both efforts failed, as the axial chirality centre is remote from existing stereocentres, leading to almost identical ¹H NMR spectra for both atropisomers of rugulotrosin A. Fortunately, a single crystal of the major atropisomer (–)-**19** was obtained by recrystallization (CH₂Cl₂/MeOH) from its mixture with **20**. X-ray crystal structure analysis of (–)-**19** (CCDC no.



Figure 2 | Scalable syntheses of enantiopure tetrahydroxanthone monomers. Key steps involve siloxyfuran addition to a benzopyrylium species (12 to (\pm) -13) followed by kinetic acylative resolution of (\pm) -17 using the Birman catalyst.

1022610) was achieved, verifying that the relative stereochemistry was identical to that found in natural rugulotrosin A¹³.

During our evaluation of dimerization conditions we found that water was essential and that the ratio of atropisomers was dependent on the ligands for palladium rather than solvent, temperature or reaction time. Accordingly, we focused our efforts on the influence of ligands on atropisomeric selectivity (Table 1). After extensive ligand screening, we found that monophosphines were crucial for the success of the reaction and that bidentate ligands such as XantPhos or DPEPhos (not shown) failed to provide the desired dimeric products. Besides SPhos (L1), the related ligand RuPhos (L3) provided dimeric products in even higher atropselectivity (93:7). Interestingly, we found when the 2' position of the monophosphine ligand did not contain an oxygen atom (for example, a carbon atom in XPhos L2 or a nitrogen atom in DavePhos, not shown), dimeric products were obtained in trace amounts. After experiments with achiral ligand L4, which provided similar results to SPhos, we anticipated that related chiral ligands may lead to efficient control of atropselectivity³⁷. We began this investigation with (R)- and (S)-BI-DIME ligands $L5^{38,39}$, which have recently shown excellent atropselectivities in biaryl couplings. However, neither ligand was able to perform the one-pot dimerization of (-)-12. Gratifyingly, chiral versions of the SPhos ligands S-Cy-MOP (L6) and R-Cy-MOP (L7)⁴⁰⁻⁴² both successfully produced dimeric products. Ligand R-Cy-MOP provided a satisfactory d.r. (95.5:4.5) and the mismatched ligand S-Cy-MOP afforded a lower d.r. (81:19), both favouring the same major atropisomer. Even with the increased bulk of ligand L8, the d.r. did not improve noticeably. To enhance product yield, we prepared the Buchwald thirdgeneration precatalysts⁴³ with SPhos, R-Cy-MOP and S-Cy-MOP. Indeed, use of the precatalyst increased yields for these cases and maintained atropselectivity. Use of *bis*(neopentyl glycolato) diboron or tetrahydroxydiboron^{44,45} as boron transfer reagents led only to trace amounts of dimeric products.



Figure 3 | One-pot Suzuki dimerization of chiral tetrahydroxanthone monomers (+)-18. Optimal conditions shown using Pd(OAc)₂ and SPhos as achiral ligand.



*Conditions for Pd precatalyst: 0.2 equiv. Pd precatalyst, 0.7 equiv. BPin₂, 3 equiv. K₃PO₄, THF/water 4:1, 70 °C, 5 h; [†]0.6 equiv. BPin₂ was used.

The acylated, enantiopure monomer (+)-18 was also investigated in the Suzuki dimerization, which generated a separable 3:1 mixture of atropisomers in 36% yield (Fig. 3). Fortunately, an X-ray crystal structure of the minor atropisomer (-)-22 (CCDC no. 1022611) was obtained to verify the relative and absolute stereochemistry. Introduction of a bulky ester onto the secondary alcohol of the monomer (for example, OCOPh) did not affect the ratio of atropisomers.

After obtaining and establishing stereochemistry for several protected dimers, we anticipated that we could access all atropisomers and enantiomers of rugulotrosin A. Dimer (-)-19 was treated with 3M HCl/MeOH for 10 min, which afforded rugulotrosin A ((-)-1) in 89% yield (Fig. 4a). In a similar manner, (+)-rugulotrosin A was obtained from acidic hydrolysis of (+)-21 (Fig. 4b). With both enantiomers of rugulotrosin A in hand, natural rugulotrosin A was unambiguously assigned as (+)-1 though comparison of rotation data. Similarly, *atrop*-rugulotrosin A (-)-23 and *ent-atrop*-rugulotrosin A (+)-23 were synthesized using similar protocols from the protected precursors (-)-22 and (+)-22 (Fig. 4c,d). To verify the stability of the axially chiral biaryl moiety, we heated rugulotrosin A (+)-1 in toluene (1 mg ml⁻¹) at 100 °C for 12 h and 150 °C for 3 h. Under both conditions we did not observe the formation of *atrop*-rugulotrosin. This is supported by the high barrier of rotation of rugulotrosin A, calculated as 47.7 kcal mol⁻¹ using the B3LYP/6-31G(d) level of theory (Supplementary pages 52–58).



Figure 4 | Syntheses of (–) and (+)-rugulotrosin A and *atrop-rugulotrosin A.* **a**, Synthesis of *ent*-rugulotrosin A. **b**, Synthesis of rugulotrosin A and comparison rotation data with natural sample. **c**, Synthesis of *atrop*rugulotrosin A. **d**, Synthesis of *ent-atrop*-rugulotrosin A. **e**, Comparison between the natural extract and synthetic rugulotrosins. HPLC-DAD (210 nm) (Zorbax C₁₈ column, gradient elution H₂O/MeCN plus 0.05% HCO₂H) analysis of 21-day *Penicillium* nov. sp. (MST-F8741) cultures extracted with (i) MeCN or (ii) MeOH, compared against (iii) natural (+)-1, (iv) synthetic (+)-1, (v) (–)-23 and (vi) a mixture of (+)-1, (–)-1, (–)-23 and (+)-23. mAU, milli absorption units.

To rationalize the axial chirality selectivity observed in the dimerization, we modelled the geometry of the expected intermediate diaryl Pd(II) complexes⁴⁶ using SPhos (**L1**) as ligand (Fig. 5). Conformational analysis was carried out by systematic six-fold rotation about the C–Pd bonds, followed by optimization of candidate structures at the B3LYP/LanL2DZ level of theory⁴⁷. The five lowest-energy structures are represented in Fig. 5. From this analysis, the three lowest-energy structures **24a–24c** all have dihedral angles (C1, C2, C2', C1') for the monomer fragments that would afford atropisomer (-)-19 (precursor to rugulotrosin) after stereospecific reductive elimination (Fig. 5a). The next two higher energy structures 25a and 25b display a reversed dihedral angle that should lead to atropisomer 20 (Table 1). Relative energies correlate with the steric fit of components. Space filling models of structures 24a-c and 25a,b (Supplementary pages 34-52) show a more crowded arrangement of the latter, in part because of steric interactions between the methyl ester and the cyclohexyl group. The distances between the hydrogen of the methyl ester to the hydrogen on the cyclohexyl of the SPhos ligand are 2.4 and 2.5 Å for conformers 25a and 25b, respectively. Based on literature precedent⁴⁷, we assume that these catalyst-reactant complexes are under thermodynamic equilibration. Additional computational studies will be required to predict barriers to interconversion. It thus appears that product stereochemistry is determined by reactant and catalyst assembly in the Pd(II) complex, and this provides a basis for the prediction of stereochemistry in future atropselective couplings.

After assigning the absolute configuration of rugulotrosin A, we used our library of synthetic rugulotrosins to determine whether Penicillium nov. sp. (MST-F8741) was capable of producing atrop-rugulotrosin A. To this end, separate 21-day fermentations of MST-F8741 were extracted with either MeCN (Fig. 4e(i)) or MeOH (Fig. 4e(ii)) and were subsequently analysed using a high-performance liquid chromatography with photodiode array detection (HPLC-DAD) method optimized for the resolution of rugulotrosin A (Fig. 4e(iii,iv)) and atrop-rugulotrosin A (Fig. 4e (v)). To further enhance sensitivity, these analyses were also carried out using high-performance liquid chromatography with photodiode array detection and electrospray ionization mass spectrometry detection (HPLC-DAD-ESIMS) protocols augmented by single ion extraction (SIE) (Supplementary page 22). As these analytical studies failed to detect any trace of atrop-rugulotrosin in MST-F8741 extracts, we conclude that the rugulotrosin biosynthetic pathway present in Penicillium nov. sp. (MST-F8741) operates with high atropisomer fidelity.

As natural rugulotrosin A had previously been reported to exhibit antibacterial activity¹³, we also tested the synthetic rugulotrosins ((+)-1, (-)-1, (-)-23, (+)-23) against several strains of Gram-positive and Gram-negative bacteria. Consistent with earlier reports, (+)-1 exhibited antibacterial activity against Bacillus subtilis (ATCC 6633) (IC₅₀ = $2.1 \,\mu$ M) and Staphylococcus aureus (ATCC 25923) (IC₅₀ = 6 μ M). Of note, enantiomer (-)-1 was also active against *B. subtilis* (ATCC 6633) ($IC_{50} = 0.7 \mu M$) and S. aureus (ATCC 25923) (IC₅₀ = 18μ M), whereas the antibacterial activity of the atropisomer (-)-23 was weak and limited to B. subtilis (ATCC 6633) (IC₅₀ = 10 μ M) and (+)-23 lacked any appreciable antibacterial activity. None of rugulotrosins (+)-1, (-)-1, (-)-23 or (+)-23 exhibited activity against the Gram-negative bacteria Escherichia coli (ATCC 25922) and Pseudomonas aeruginosa (ATCC 27853), nor did they exhibit cytotoxicity (IC₅₀ > 30 μ M) against human colon (SW620) and lung (NCI-H460) cancer cells.

Conclusion

We have developed a concise, atropselective approach to rugulotrosin A and stereoisomers through point-to-axial chirality transfer, which has facilitated the determination of the absolute configuration of rugulotrosin A. Computational studies modelling the geometry of intermediate diaryl Pd(II) complexes have provided a rationale for the atropselectivity observed in the key Suzuki dimerization. These studies highlight the utility of SPhos and related bulky monophosphine ligands⁴⁸ in palladium couplings both to bring monomers in proximity and enhance remote steric effects leading to atropselectivity. Through HPLC analysis of fungal extracts and

a Conformers leading to (-)-19 (rugulotrosin precursor)



Figure 5 | Computational studies for atropselective Suzuki dimerization. Conformational analysis was optimized at the B3LYP/LanL2DZ level of theory. Dihedral angles were measured by C1, C2, C2' and C1' and are shown in green. **a**, Conformers leading to (–)-19. **b**, Conformers leading to 20.

synthetic samples, we have determined that *Penicillium* nov. sp. (MST-F8741) generates rugulotrosin A in an atropselective manner. Moreover, the atropisomers and enantiomers of rugulotrosin A were found to have different activities against Gram-positive bacteria, illustrating the importance of stereochemistry on target selectivity. Further studies regarding atropselective syntheses of dimeric natural products are currently under investigation and will be reported in due course.

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Author contributions

T.Q. and J.A.P. Jr conceived of the project, designed and carried out the experiments, analysed the data and wrote most of the paper. S.L.S-J. and R.P.J. performed computational studies. Z.G.K. and R.J.C. performed natural extract comparisons and biological studies. All authors discussed the results and commented on the manuscript.

Additional information

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Competing financial interests

The authors declare no competing financial interests.