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Asymmetric Synthesis of Griffipavixanthone Employing a Chiral Phosphoric Acid-Catalyzed Cycloaddition

Michael J. Smith, Kyle D. Reichl, Randolph A. Escobar, Thomas J. Heavey, David F. Coker, Scott E. Schaus*, and Jr. John A. Porco*

Department of Chemistry and Center for Molecular Discovery (BU-CMD), Boston University, 590 Commonwealth Avenue, Boston, Massachusetts 02215, United States

Abstract

Asymmetric synthesis of the biologically active xanthone dimer griffipavixanthone is reported along with its absolute stereochemistry determination. Synthesis of the natural product is accomplished via dimerization of a *p*-quinone methide using a chiral phosphoric acid catalyst to afford a protected precursor in excellent diastereo- and enantioselectivity. Mechanistic studies, including an unbiased computational investigation of chiral ion-pairs using parallel tempering, were performed in order to probe the mode of asymmetric induction.

The dimeric natural product (+)-griffipavixanthone (**1**)¹ has become a desirable synthetic target due to its interesting framework and anticancer properties. Compound (+)-(**1**) has been shown to increase intracellular reactive oxygen species (ROS) and cleave caspase-3, thereby inducing apoptosis in lung cancer cells.² Recent studies have shown that (+)-(**1**) is a B-Raf and C-Raf inhibitor in esophageal cancer cell lines with comparable effects to FDA-approved drugs.³

We have previously reported the biomimetic synthesis of (±)-**1** wherein treatment of *p*-quinone methide (*p*-QM) **2** with Lewis or Brønsted acids triggered an isomerization–cycloaddition–cyclization cascade affording the polycyclic core (Scheme 1). The reaction sequence began with acid-mediated isomerization of the vinyl *p*-QM **2** to 1,3-diene **3**. Cationic, stepwise [4+2] cycloaddition of **2** and **3** afforded the *anti*-cycloadduct **4** and *syn*-cycloadduct **5**, the latter which underwent intramolecular arylation to afford griffipavixanthone tetramethyl ether **6**.⁴

The complexity and unique nature of this cationic cascade prompted our interest to undertake an asymmetric synthesis. In this regard, we considered that enantioselectivity could be introduced via chiral ion-pairing catalysis.^{5–7} Asymmetric [4+2] cycloadditions are

*Corresponding Authors: porco@bu.edu, seschaus@bu.edu.

ASSOCIATED CONTENT

Supporting Information

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Video clip from the parallel tempering (PT) simulation of **TS-1** and CPA **F** (AVI)

Experimental procedures, computational data, and characterization data for new compounds (PDF)

The authors declare no competing financial interest.

mediated by both Brønsted and Lewis acids.^{8,9} Scheme 2 depicts select literature reports of chiral Brønsted acid-mediated asymmetric reactions involving *p*-QMs. Sun and co-workers have reported CPA-mediated, enantioselective 1,6-arylations of *p*-QMs (Scheme 2a).^{10,11} In addition, Li and co-workers have demonstrated asymmetric α -alkylations to *p*-QMs generating two contiguous stereocenters (Scheme 2b).¹² There is a single report of a diastereoselective, CPA-mediated intramolecular [4+2] cycloaddition.¹³ These studies further prompted our interest to develop an intermolecular, asymmetric [4+2] cycloaddition of a *p*-QM (Scheme 2c). Herein, we (1) describe a CPA-catalyzed asymmetric synthesis of (+)-**1**, (2) demonstrate catalyst-dependent diastereoselectivity, (3) assign absolute stereochemical configuration to (+)-**1**, and (4) provide computational insight for the asymmetric induction observed using parallel-tempering (PT) simulation methods.

Yamamoto and co-workers have demonstrated that chiral Brønsted acid-mediated cycloadditions require specific pK_a 's as dictated by the substrate.¹⁴ With this in mind, our investigation began by surveying chiral Brønsted acids with varying pK_a values including CPAs, imidodiphosphoric acids (IDPs), and *N*-triflyl phosphorimides.^{14–17} While the latter two catalyst classes chiefly led to isomerization of *p*-QM **2** to diene **3**, use of CPAs led to production of cycloadducts **4–6**. A study of electronically variable CPAs was performed (Table 1). We found that treatment of **2** with CPA **A**¹⁸ (entry 1) generated dimer **6** in high enantioselectivity along with products **3–5**. Electron rich CPAs **B** and **C**¹⁹ lowered the reactivity for cycloadducts (entries 2 and 3). With the most promising result observed using —CF₃ catalyst (**A**), we expanded our assessment of electron-deficient, aryl-substituted CPAs and found that the highly electron-deficient Yamamoto catalyst (CPA **D**)²⁰ favored the *anti*-cycloadduct **4** (entry 4). However, the moderately electron deficient CPA **E**²¹ afforded several products, while still maintaining high enantioselectivity for dimer **6** (entry 5). Interestingly, use of CPA **F**¹⁹ bearing the strongly electron-withdrawing nitro (NO₂) substituent provided high diastereo- and enantioselectivity (entry 6). At elevated temperatures (entry 7), the reaction was driven to completion affording (+)-**6** in 72% yield and 99% ee.

To determine the absolute stereochemistry of **1** prepared using CPA **F**, global demethylation of (+)-**6** using our previously reported conditions⁴ was performed. By measuring the optical rotation of **1**, we found that (*R*)-CPA **F** provided the natural (+)-enantiomer of **6** (Figure 1a). To elucidate the absolute stereochemistry of (+)-**1**, an electronic circular dichroism (ECD) spectrum was obtained. Comparison with the theoretical ECD spectrum (CAM-B3LYP/def2-TZVP/PCM (MeOH)//CAM-B3LYP/def2-SVP/PCM (MeOH) indicates that (+)-**1** has 8*R*,8*aR*,12*S* stereochemistry (Figure 1b).²²

At this juncture, we were also interested to determine the enantioselectivity of the *anti*-cycloadduct, the only dimeric product formed upon treatment of **2** with CPA **D**. Upon isolation of the dimeric *p*-QM cycloadduct **4**, heterogeneous reduction with H₂/Pd/C afforded cycloadduct **7** in 33% yield (2 steps) and 71% ee (Scheme 3). The absolute stereochemistry of (+)-**7** was determined using ECD analysis.²²

We also used *in situ* ¹H NMR analysis to monitor reactions comparing both TFA and CPA **F** as catalysts. Using TFA as a catalyst, production of the *anti*-cycloadduct **4** appears to be

more favorable than the *syn*-cycloadduct **5** (Figure 2a). Remarkably, CPA **F** shows reversal of the TFA-catalyzed preference for *anti*-cycloadduct **4**, instead showing initial formation of the *syn*-cycloadduct **5** (Figure 2b). Presumably, CPA catalyst **F** reduces the energetic barrier to the *syn*-cycloadduct **5** (*vide infra*). ¹H NMR data (Figure 2) also indicated formation of diene **3** *in situ*. To probe the potential for decomposition or dimerization, diene **3** was subjected to the standard conditions; in the event, **6** was isolated in 69% yield and 96% ee.²²

As we identified catalyst diastereoselectivity, we were also interested to probe the point at which enantioselectivity was introduced. Treatment of (±)-**5** with CPA **F** resulted in the formation of (+)-**6** in 78% ee (Scheme 4a). In a control experiment, (±)-**4** was subjected to the same conditions, affording (+)-**6** in 99% ee. These results indicate that reorganization to the *syn*-cycloadduct **5** is the enantiodetermining step. To further understand enantioinduction, the reaction was stopped after 3 min and the *syn*-cycloadduct **5** was purified and isolated in 53% yield and 99% ee (Scheme 4b). This reaffirmed that the formation of the *syn*-cyclohexene moiety is the enantiodetermining step (cf. **5**, Scheme 1). In addition, we performed kinetics studies using React IR analysis. The data obtained is consistent with a first-order reaction with respect to catalyst.²²

Previous literature has shown that CPAs promote reactivity through the Brønsted acidic site, the Lewis basic site, or both. This interaction generates transient diastereomeric ion-pairs, one of which is more favorable.²³ Notably, density functional theory (DFT) calculations have been used to elucidate the primary phosphate interaction and the secondary stereo-electronic interactions imparting enantioselectivity.^{24–27}

To investigate the mode of enantioinduction, we used a computational methodology that has not been employed as a tool in organic synthesis. As we were unsure of the mode of asymmetric induction leading to **5**,⁴ we used an enhanced-sampling²⁸ computational method to explore configurations with CPA **F** in an unbiased manner. Parallel tempering (PT) simulations²⁹ were conducted using a molecular-mechanics force field.³⁰ In these calculations, multiple independent simulations are run in parallel at a range of temperatures; this enables frequent crossing of energetic barriers, thereby fully sampling the thermal equilibrium distribution. Applications of PT methodology primarily include biomolecular dynamics to further understand protein structure.^{31–35} A representative video from a simulation involving the preferred enantiomeric TS and CPA **F** is provided.³⁶

The ensembles of configurations of the transition-states (**TS-1** = major, *ent*-**TS-1** = minor) and CPA **F** generated by these PT simulations contain many structures exhibiting simultaneous double hydrogen-bonds (DHBs), where the catalyst acts as a Brønsted acid and a Lewis base. Representative structures of **TS-1**-CPA **F** and *ent*-**TS-1**-CPA **F** complexes sampled during these simulations are presented in Figure 3a,b, respectively. Hydrogen bonding interactions between two phenols on both xanthone fragments³⁷ with the phosphate moiety of the CPA provide a template for long-range chirality transfer.

Consequently, histograms of the probability distributions were used to compute free energy surfaces of hydrogen bonding distances for both **TS-1** and *ent*-**TS-1** (Figure 3c,d, respectively). As expected, the lowest free energy structures are observed when CPA **F** is

doubly hydrogen bound. The simulation of the **TS-1**–CPA **F** reveals evidence of significantly more complexes in which **TS-1** hydrogen bonds to a single phosphate oxygen²² (Figure 3c) compared to *ent*-**TS-1** (Figure 3d). The increased frequency of these intermediate SHB structures results from the enhanced flexibility of **TS-1**–CPA **F** compared to the *ent*-**TS-1**–CPA **F**. The intermolecular dihedral distribution in Figure 4a shows that the conformational space for **TS-1** relative to CPA **F** permits free rotation of the *bis*-xanthone transition state (cf. Figure 4c for atoms involved). In Figure 4b, insurmountable barriers hinder relative rotation of *ent*-**TS-1**–CPA **F**, dramatically reducing conformational space (cf. Figure 3b) in the presence of CPA **F**. Figure 4c depicts the rotational freedom of **TS-1** around the catalyst while maintaining long-range chirality transfer. These findings suggest that enantioinduction is derived, in part entropically, by reducing the configuration space available to *ent*-**TS-1**. Thus, **TS-1**–CPA **F** can interact by a SHB or DHB while the minor can only bind to the transition state in a DHB configuration.

In addition, π -stacking interactions with the catalyst were also found to be prominent.²² Previous studies have observed electrostatic donor–acceptor aromatic stacking effects wherein nitrophenyl substituents are among the strongest acceptors.³⁸ These strong π -interactions play an important role in stabilizing the **TS-1**–CPA **F** complex, driving enantioselectivity.

The CPA **F** catalyst also promotes enthalpic stabilization of the major **TS-1** enantiomer. A random selection of complexes from PT simulations were DFT-optimized (B3LYP/6–31G(d)/IEF-PCM [CH₂Cl₂] to obtain energies for the catalyst-transition state complexes in the presence of a polarizable continuum model (PCM) solvent. On average, the major **TS-1**–CPA **F** complex is energetically more favorable than its minor enantiomeric counterpart by nearly 7 kcal/mol.²²

On the basis of this model, we were interested in developing a catalyst that may perturb enantioinduction by steric interactions (Scheme 5). Use of the sterically modified CPA **G** permitted the formation of dimer **6**, albeit in lower yield and ee (23% and 42%, respectively). Erosion of enantioselectivity may be explained by steric interference between the substrate and the methyl groups of CPA **G**. To rationalize this outcome, PT simulations were also conducted. Interestingly, the major enantiomeric transition state (**TS-1**–CPA **G**) deviated from that of CPA **F**. In the **TS-1**–CPA **F** system, the nitro aryl groups are stacking with the xanthone moieties (Figure 5a). However, the added sterics of CPA **G** prevent π -stacking, resulting in the nitro aryl groups being forced out of plane with the xanthone (Figure 5b). DFT assessment revealed only a 1.5 kcal/mol energy difference between transition states, providing computational support for the erosion of enantioselectivity. The calculated free energy surfaces and energy distributions for the **TS-1**–CPA **G** system are provided in the Supporting Information.²²

In summary, we have reported asymmetric syntheses and absolute stereochemistry assignment of (+)- and (–)-griffipavixanthone, a biologically active xanthone dimer. The key cascade reaction utilizes a chiral phosphoric acid (CPA)- catalyzed cycloaddition of a *para*-quinone methide (*p*-QM) monomer in which a specific aryl nitro-substituted CPA catalyst provides high diastereo- and enantioselectivity. In addition, we have interrogated the

mechanism and mode of enantioinduction through catalyst electronic effects, *in situ* ^1H NMR reaction profiles, kinetic studies, and computational studies using parallel tempering (PT) simulations to broadly sample chiral ion pair complexes. Further studies on the synthesis of xanthone dimers as well as applications of parallel tempering (PT) to address problems in organic synthesis are currently in progress and will be reported in due course.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1.

Absolute stereochemical determination of (+)-1. (a) Synthesis of (+)-1. (b) Comparison of experimental and computational ECD spectra for (+)-1.

Figure 2.
In situ ^1H NMR analysis over time upon treatment of **2** with TFA (a) and CPA **F** (b).

Figure 3. Representative configurations of **TS-1** (*pink*) doubly hydrogen bonding with CPA **F** (green). DFT-optimized (a) **TS-1** – CPA **F** (0 kcal/mol) and (b) *ent*-**TS-1**–CPA **F** (7 kcal/mol). Free energy surface cuts at $T = 298$ K for (c) **TS-1**–CPA **F** simulation shows free energy minima with single hydrogen bonds (SHB) and DHB. (d) *ent*-**TS-1**–CPA **F** simulation reveals a single DHB minimum only.

Figure 4.

(a and b) Free energy surfaces as a function of intermolecular dihedral angle, φ_D , for **TS-1** and **ent-TS-1** with CPA **F**, respectively. Areas with a barrier of >20 kcal/mol were not sampled. Greater rotation and flexibility is observed in **TS-1** (c) as assessed by the free energy obtained from the intermolecular dihedral angle than in **ent-TS-1**, where free rotation relative to CPA **F** is inaccessible at the reaction temperature.

Figure 5. Structural comparison of DFT-optimized (a) **TS-1-CPA F** and (b) **TS-1-CPA G** highlighting the altered complexation resulting from the sterically modified catalyst.

Scheme 1.
Brønsted Acid-Catalyzed Dimerization Cascade

a. Sun and Coworkers: **b. Li and Coworkers:** **c. This work:** **Scheme 2.**Representative CPA-Catalyzed Asymmetric Reactions of *p*-Quinone Methides

Scheme 3. Reductive Trapping of *anti*-Cycloadduct 4 To Determine Enantioselectivity

Scheme 4. Enantioenrichment of (±)-*p*-Quinone Methide Dimers

Scheme 5.
Disruption of Enantioselectivity Using a Modified CPA

Table 1.

Assessment of BINOL-Derived CPAs

Reaction scheme showing the conversion of compound **2** to products **3**, **4**, **5**, and **6** using a catalyst (5 mol%) in CH₂Cl₂ (50 mM) at room temperature for 20 hours.

Chemical structures of the catalysts (CPAs) A through F are shown, along with their substituents:

- A: R₁=CF₃, R₂=R₃=H
- B: R₁=OMe, R₂=R₃=H
- C: R₁=R₂=R₃=H
- D: R₁=R₃=Me, R₂=NO₂
- E: R₁=Br, R₂=R₃=H
- F: R₁=NO₂, R₂=R₃=H

The products are shown as follows:

- 3**: A substituted coumarin derivative.
- 4** and **5**: Diastereomeric dimers (anti and syn).
- 6**: A dimeric product.

entry	CPA	3 (%)	4 (%)	5 (%)	6 ^a (%) (% ee)
1	A	7	30	9	34 (96)
2	B	51	18	8	—
3	C	49	11	3	—
4	D	10	48	—	—
5	E	8	36	10	36 (95)
6	F	—	23	5	61 (99)
7 ^b	F	—	—	—	72 ^c (99)

^a ¹H NMR yields determined using 1,2,4,5-tetramethylbenzene as internal standard.

^b Reaction temperature is 40 °C.

^c Isolated yield.