

## Biomimetic Synthesis of Myrtucommulones D–E

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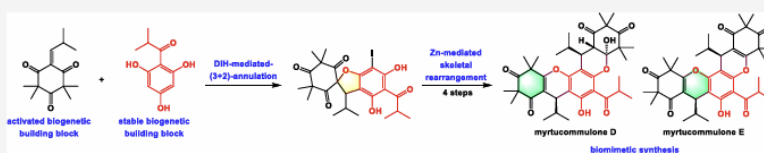
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**ABSTRACT:** A concise and efficient biomimetic synthesis of myrtucommulones D–E has been achieved, proceeding in just 6–7 linear steps from readily available biogenetic building blocks. The key feature of the synthesis was the Zn-mediated skeletal rearrangement reaction, without the need for rare metal photocatalysts and visible light. Based on this biomimetic synthesis, four compounds demonstrated moderate to excellent cytotoxic activities against osteosarcoma cells (U2OS and 143B).

### INTRODUCTION

Xanthenes, a prominent class of heterocyclic compounds, are widely distributed in natural products and exhibit diverse pharmacological activities.<sup>1</sup> Among them, myrtucommulones D–E (1–2; Figure 1) isolated from *Myrtus communis* L.<sup>2</sup> represent structurally unique angular 6/6/6/6-fused pentacyclic xanthenes, characterized by two to four stereocenters and a benzopyrano[2,3-*a*]xanthene core. These compounds display potent antibacterial and  $\alpha$ -glucosidase inhibitory activities,<sup>2,3</sup> yet their therapeutic potential remains underexplored due to limited natural abundance. Notably, the angular pentacyclic motif is conserved in related natural products (e.g., 3–5, highlighted in red, Figure 1),<sup>4</sup> underscoring the demand for versatile synthetic platforms to enable biological studies.

Over recent decades, xanthenes have attracted sustained interest from the synthetic chemistry community owing to their promising pharmacological properties and structural complexity, particularly as exemplified by the demanding 6/6/6/6-fused pentacyclic derivatives.<sup>5</sup> While linear skeletons, such as that of tomentosone C (6)<sup>6</sup> (Figure 1), can typically be constructed,<sup>5a</sup> the synthesis of the angular 6/6/6/6-fused pentacyclic natural xanthenes has presented persistent challenges, with only sporadic reports addressing their preparation.<sup>7</sup> This regioselectivity bias is hypothesized to arise from a critical hydrogen-bond interaction between the C7 ketone and C4 hydroxyl groups, which enhances the electron density on the phenolic oxygen, thereby favoring the formation of the linear product (Scheme 1A).<sup>8</sup>

Our group previously reported the asymmetric synthesis of myrtucommulones D–E using isomyrtucommulone B as a precursor.<sup>9</sup> However, racemization issues emerged during the *retro*-Friedel–Crafts reaction of myrtucommulone B, necessitating chiral resolution to obtain optically pure isomyrtucommulone B (Scheme 1B). To address this limitation, we recently developed a visible-light photocatalyzed skeletal rearrangement strategy employing a rare metal iridium photocatalyst and iron, which enabled the synthesis of *ent*-myrtucommulones E–F (Scheme 1C).<sup>10</sup> Notably, the photocatalyst and Zn can also facilitate the rearrangement reaction; however, excessive reducing capacity of Zn led to predominant side-reactions and diminished yields.<sup>10</sup> This suggests that Zn may intrinsically drive skeletal rearrangement without photocatalyst mediation, enabling direct skeletal rearrangement. Based on this concept and our continued efforts, we herein report the concise biomimetic synthesis of ( $\pm$ )-myrtucommulones D–E, utilizing biogenetic building blocks<sup>11</sup> and a Zn-mediated skeletal rearrangement reaction as a key step. Subsequent bioactivity studies revealed that four compounds showed moderate to excellent efficacy against osteosarcoma growth *in vitro*.

Given the concurrent isolation of flavesone (7a),<sup>12</sup> myrtucommulone B' (9),<sup>13</sup> myrtuspirone A (11),<sup>14</sup> isomyrtucommulone B (12),<sup>15</sup> 6-methylisomyrtucommulone B (13),<sup>16</sup> myrtucommulone B (14),<sup>17</sup> and syncarpic acid (16)<sup>18</sup> from Myrtaceae plants, the potential biosynthetic pathways of myrtucommulones D–E (1–2) are outlined in Scheme 2. The stable biogenetic building block 7a may be activated through

### RESULTS AND DISCUSSION

Given the concurrent isolation of flavesone (7a),<sup>12</sup> myrtucommulone B' (9),<sup>13</sup> myrtuspirone A (11),<sup>14</sup> isomyrtucommulone B (12),<sup>15</sup> 6-methylisomyrtucommulone B (13),<sup>16</sup> myrtucommulone B (14),<sup>17</sup> and syncarpic acid (16)<sup>18</sup> from Myrtaceae plants, the potential biosynthetic pathways of myrtucommulones D–E (1–2) are outlined in Scheme 2. The stable biogenetic building block 7a may be activated through

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## Notes

The authors declare no competing financial interest.

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