

# The Biosynthesis of the Fungal Meroterpenoids Boviquinone-3 and -4 follows Two Different Pathways

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## Abstract

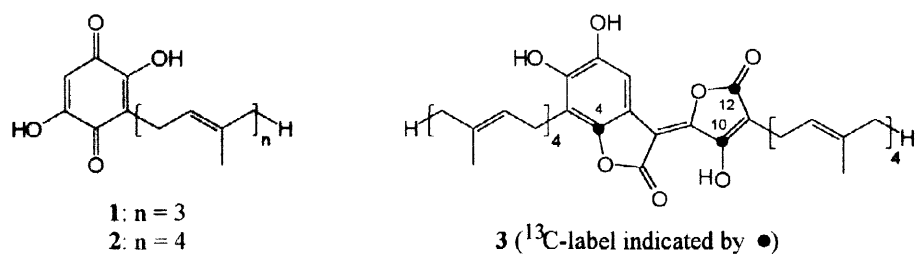
A key step in the biosynthesis of the fungal meroterpenoids boviquinone-3 and -4 is the prenylation of 3,4-dihydroxybenzoic acid. In fruit-bodies of *Suillus bovinus* boviquinone-4 is formed by geranylgeranylation of 3,4-dihydroxybenzoic acid at C-5, whereas in *Chroogomphus rutilus* the farnesyl side chain of boviquinone-3 is introduced at C-2. The biosynthesis of the terpenoid chain of boviquinone-4 follows the mevalonate route. © 1998 Elsevier Science Ltd. All rights reserved.

*Keywords:* biosynthesis; quinones; meroterpenoids; fungal pigments

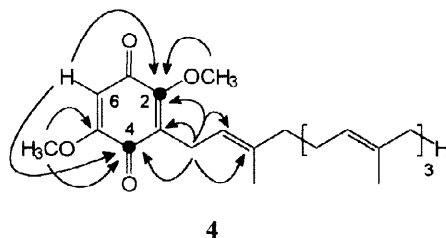
Meroterpenoid quinones like the ubiquinones, plastoquinones and menaquinones are widespread in living organisms and are essential for several life processes. A group of structurally related 2,5-dihydroxy-1,4-benzoquinones with oligoprenyl side chains has been isolated from several basidiomycetes of the order Boletales [1]. Fruit-bodies of *Chroogomphus rutilus* and *C. helveticus* contain the farnesylated boviquinone-3 (helveticone) (1) [2] whereas *Suillus bovinus* produces its geranylgeranyl homologue boviquinone-4 (2) [3,4]. Structurally, the boviquinones resemble the ubiquinones whose quinone ring is known to originate from 4-hydroxybenzoic acid [5,6,7]. In a preliminary feeding experiment we have shown that ring <sup>14</sup>C-labelled 4-hydroxybenzoic is also incorporated into boviquinone-3 (1) by fruit-bodies of *Chroogomphus helveticus* [8]. In this publication we report on a more detailed investigation of boviquinone biosynthesis.

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Feeding of either 4-hydroxy[1- $^{13}\text{C}$ ]benzoic acid (**5**) or 3,4-dihydroxy[1- $^{13}\text{C}$ ]benzoic acid (**6**) [9] at low concentrations (0.04 mmol/fruit-body) to young specimens of *Suillus bovinus* gave rise to specific incorporation of these precursors into boviquinone-4 (**2**) [10]. Because of rapid tautomerization of the 2,5-dihydroxybenzoquinone system the sites of incorporation could only be determined after conversion of boviquinone-4 to its dimethyl ether **4** [4]. In both experiments C-2 and C-4 were equally enriched, and the incorporation was 1.1 and 1.3% ( $^{13}\text{C}$  atom% excess) for the feeding of **5** and **6**, respectively. The labelling pattern of **4** was confirmed by HMBC experiments summarized in Figure 1 in which 6-H as well as the methylene group attached to C-3 exhibit long range couplings to the  $^{13}\text{C}$  enriched carbon atoms C-2 and C-4.



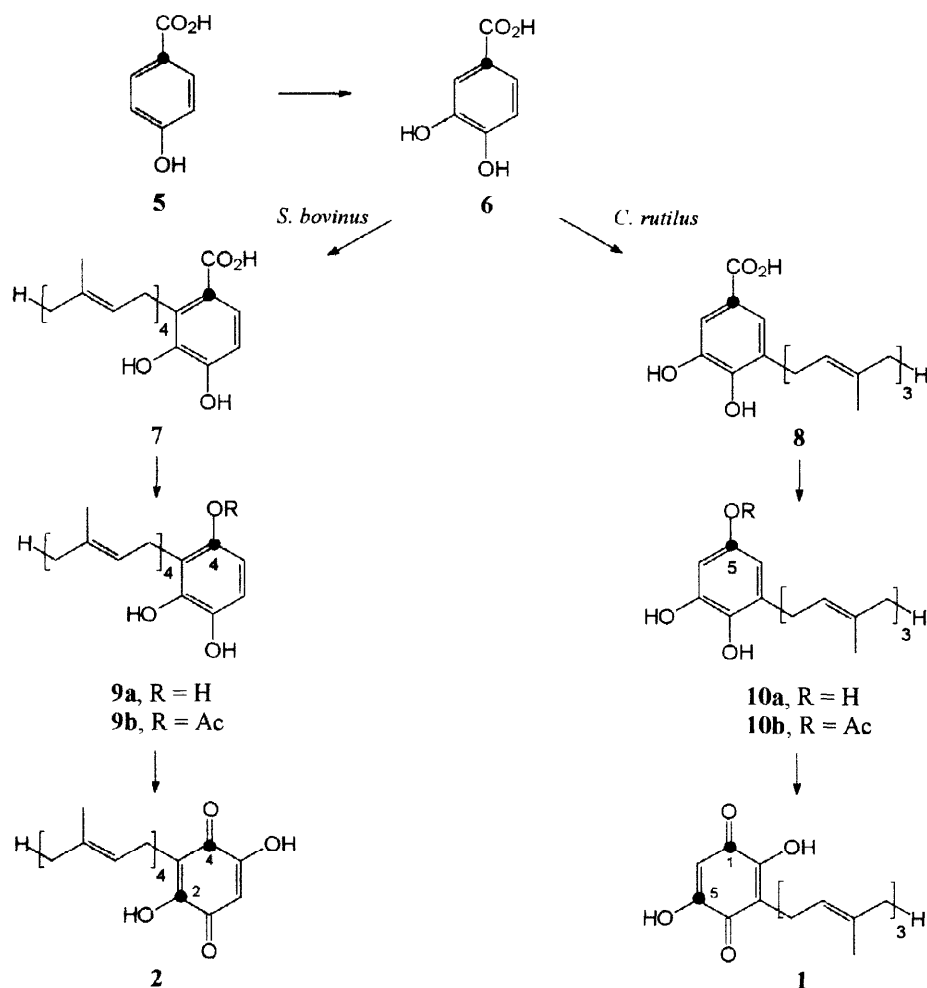
**Figure 1.** HMBC correlations of labelled boviquinone-4 dimethyl ether (**4**) after feeding of 4-hydroxy- or 3,4-dihydroxy[1- $^{13}\text{C}$ ]benzoic acid to fruit-bodies of *Suillus bovinus*.

The feeding experiments indicate that boviquinone-4 (**2**) is formed by prenylation of 3,4-dihydroxybenzoic acid (**6**) at C-2 to yield 2-geranylgeranyl-3,4-dihydroxybenzoic acid (**7**) (Scheme 1). Oxidative decarboxylation of **7** then affords the trihydroxy intermediate **9a** which is further hydroxylated and oxidized to boviquinone-4. 4-Hydroxybenzoic acid (**5**) acts as the precursor of 3,4-dihydroxybenzoic acid.

Interestingly, the application of higher doses of the precursors **5** and **6** (~0.2 mmol/fruit-body) or even other aromatic acids such as benzoic or salicylic acid induces a change of metabolism. Instead of boviquinone-4 (**2**) the fruit-bodies of *S. bovinus* produce large amounts of bovilactone-4,4 (**3**) [11], whose labelling pattern indicated in formula **3** is in agreement with its formation by coupling of trihydroxy compound **9a** with boviquinone-4 (**2**) with concomitant ring cleavage and formation of the lactone rings [2].

The 4-*O*-acetyl derivative of intermediate **9a**, suillin (**9b**), occurs in fruit-bodies of *Suillus variegatus* and several other *Suillus* species [1,12]. Feeding of 4-hydroxy[1- $^{13}\text{C}$ ]benzoic acid and 3,4-dihydroxy[1- $^{13}\text{C}$ ]benzoic acid to young fruit-bodies of *S. variegatus* gave [4- $^{13}\text{C}$ ]-labelled **9b** as expected from Scheme 1 [13].

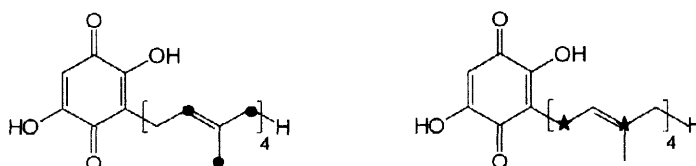
Surprisingly, feeding of 4-hydroxy- or 3,4-dihydroxy[1- $^{13}\text{C}$ ]benzoic acid to *Chroogomphus rutilus* yielded boviquinone-3 (**1**) with equal distribution of the label at C-1 and C-5 (Scheme 1) [14]. This implies that in the biosynthesis of boviquinone-3 prenylation of 3,4-dihydroxybenzoic acid takes place at C-5 to yield 5-farnesyl-3,4-dihydroxybenzoic acid (**8**) as the first intermediate. Subsequent oxidative decarboxylation, hydroxylation at



**Scheme 1.** Proposed pathways for the biosyntheses of boviquinone-4 (2) and boviquinone-3 (1).

C-4 and oxidation affords boviquinone-3 (1). This pathway is supported by the isolation of 5-acetoxy-3-farnesyl-1,2-dihydroxybenzene (10b), the 5-*O*-acetyl derivative of the postulated intermediate 10a [15] from fruit-bodies of *C. rutilus*. The acetate 10b was obtained from both feeding experiments and exhibited  $^{13}\text{C}$  enrichments at C-5 from 5 and 6 of 1.3 and 3.6%, respectively, in agreement with the proposed biosynthesis [14].

Feeding of [1- $^{13}\text{C}$ ]- or [2- $^{13}\text{C}$ ]glucose to fruit-bodies of *Suillus bovinus* resulted in complimentary labelling patterns of the side chain in boviquinone-4 (2) in accord with the formation of isopentenyl pyrophosphate *via* the classical mevalonate route (Figure 2) [16]. This result is supported by the incorporation of  $^{14}\text{C}$ -labelled mevalonate in boviquinone-3 (1) in fruit-bodies of *Chroogomphus helveticus* [8].



**Figure 2.** Labelling patterns of boviquinone-4 (2) after feeding of [1- $^{13}\text{C}$ ]- and [2- $^{13}\text{C}$ ]glucose. Enriched carbon atoms from [1- $^{13}\text{C}$ ]glucose are indicated by • and enriched carbon atoms from [2- $^{13}\text{C}$ ]glucose by \*.

When [ $1\text{-}^{13}\text{C}$ ]tyrosine [10] was added to cultures of *Suillus bovinus*, the bovilactone-4,4 (**3**) isolated after two months disclosed the same  $^{13}\text{C}$  enrichments as in the feeding experiment with 4-hydroxy[ $1\text{-}^{13}\text{C}$ ]benzoic acid (**5**) [17]. This proves the role of tyrosine as precursor of 4-hydroxybenzoic acid in this mushroom [18].

Our results show that the biosynthesis of the boviquinones differs from that of the ubiquinones in the prenylation step, which requires 3,4-dihydroxybenzoic acid instead of the 4-hydroxy derivative. Moreover, the structurally homologous fungal metabolites boviquinone-3 (**1**) and boviquinone-4 (**2**) are produced by two different pathways in closely related mushrooms. Studies on the biosynthesis of tridentoquinone and related meroterpenoids from basidiomycetes will be reported in due course.

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- [10] [ $1\text{-}^{13}\text{C}$ ]-**6** (100 mg) in DMSO (1 ml) was injected in 18 young fruit-bodies. After 3 days in their natural habitat the mushrooms were harvested, worked up, and the resulting boviquinone-4 (50 mg) was converted into its dimethyl ether **4** [4], which was subjected to  $^{13}\text{C}$  NMR spectroscopy. The  $^{13}\text{C}$  enrichment (atom% excess) at C-2 ( $\delta$  155.5) and C-4 ( $\delta$  183.7) was 2.6% for each position. Similarly, feeding of [ $1\text{-}^{13}\text{C}$ ]-**5** yielded **4** with a  $^{13}\text{C}$  enrichment of 1.1%.
- [11] Jägers, E.; Steglich, W. *Angew. Chem.* **1981**, *93*, 1105; *Angew. Chem. Int. Ed. Engl.* **1981**, *20*, 1016. After feeding labelled **5** or **6** for 3 days, **3** showed the following  $^{13}\text{C}$  enrichments: C-4 ( $\delta$  146.9) 50%, C-10 ( $\delta$  161.3) 25% and C-12 ( $\delta$  167.1) 25%.
- [12] Jägers, E.; Pasupathy, V.; Hovenbitzer, A.; Steglich, W. *Z. Naturforsch.* **1986**, *41b*, 645.
- [13] [ $1\text{-}^{13}\text{C}$ ]-**6** (200 mg) in DMSO (300  $\mu\text{l}$ ) was administered *via* syringe to 5 young fruit-bodies. After 3 days the mushrooms were worked up and **9b** (47 mg) was isolated. **9b** showed a  $^{13}\text{C}$  enrichment of 36% at C-4 ( $\delta$  142.1).
- [14] [ $1\text{-}^{13}\text{C}$ ]-**6** (90 mg) in acetone (200  $\mu\text{l}$ ) was applied to 3 young fruit-bodies. After 4 days, **1** (40 mg) and **10b** (12 mg) were isolated and the following  $^{13}\text{C}$  enrichments (atom% excess) were determined: **1** (dimethyl ether derivative) 1.3% at C-1 ( $\delta$  183.73) and C-5 ( $\delta$  158.8); **10b** 3.6% at C-5 ( $\delta$  143.81). Similarly [ $1\text{-}^{13}\text{C}$ ]-**5** (61 mg) yielded **1** (64mg) and **10b** (35 mg) with  $^{13}\text{C}$  enrichments of 0.5 and 1.1%, respectively.
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- [16] [ $1\text{-}^{13}\text{C}$ ]- or [ $2\text{-}^{13}\text{C}$ ]Glucose (200 mg) in  $\text{H}_2\text{O}$  (500  $\mu\text{l}$ ) were injected into 3 young fruit-bodies. After 3 days, **2** was isolated and converted into **4**. In the [ $1\text{-}^{13}\text{C}$ ]glucose experiment,  $^{13}\text{C}$  enrichments of 2-3% were observed for the 4  $\text{C}_\text{H}$  carbons ( $\delta$  119.7-124.4), the 4', 8'- and 12'- $\text{C}_\text{H}_2$  carbons ( $\delta$  ~39.74) and the 5  $\text{C}_\text{H}_3$  carbons ( $\delta$  2 $\times$ 16.0, 16.2, 17.7, 25.7) of the geranylgeranyl chain of **4**. [ $2\text{-}^{13}\text{C}$ ]Glucose experiment:  $^{13}\text{C}$  enrichments of 0.8-1.6% for C-3', C-7', C-11' and C-15' ( $\delta$  131.2-137.3) and the 1'-, 5'-, 9'- and 13'- $\text{C}_\text{H}_2$  carbons ( $\delta$  22.3-26.8). The labelling patterns are in accord with the glycolysis of [ $1\text{-}^{13}\text{C}$ ]- and [ $2\text{-}^{13}\text{C}$ ]glucose to [ $1\text{-}^{13}\text{C}$ ]- and [ $1\text{-}^{13}\text{C}$ ]acetate, respectively, before entering the mevalonate pathway.
- [17] Labelling of **3** after administering of [ $1\text{-}^{13}\text{C}$ ]-tyrosine: C-4 ( $^{13}\text{C}$  enrichment 9%), C-10 and C-12 (each 5%).
- [18] Compare e.g. Strack, D. *Phenolic Metabolism*, Academic Press, San Diego **1997**.