



# Biosynthetic origin of benzoquinones in the explosive discharge of the bombardier beetle *Brachinus elongatulus*

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## Abstract

Bombardier beetles are well-known for their remarkable defensive mechanism. Their defensive apparatus consists of two compartments known as the reservoir and the reaction chamber. When challenged, muscles surrounding the reservoir contract sending chemical precursors into the reaction chamber where they mix with enzymes resulting in an explosive discharge of a hot noxious chemical spray containing two major quinones: 1,4-benzoquinone and 2-methyl-1,4-benzoquinone (toluquinone). Previously, it has been speculated that the biosynthesis of all benzoquinones originates from one core precursor, 1,4-hydroquinone. Careful ligation of the base of the reservoir chamber enabled us to prevent the explosive reaction and sample untransformed reservoir fluid, which showed that it accumulates significant quantities of 1,4-hydroquinone and 2-methyl-1,4-hydroquinone. We investigated the biosynthetic mechanisms leading to quinone formation by injecting or feeding *Brachinus elongatulus* beetles with stable-isotope-labeled precursors. Chemical analysis of defensive secretion samples obtained from 1,4-hydroquinone-*d*<sub>6</sub>-administered beetles demonstrated that it underwent conversion specifically to 1,4-benzoquinone. Analogously, results from *m*-cresol-*d*<sub>3</sub> injected or fed beetles confirmed that *m*-cresol is metabolized to 2-methyl-1,4-hydroquinone, which is then oxidized to 2-methyl-1,4-benzoquinone in the hot spray. Our results refute the previous claim that 1,4-hydroquinone is the precursor of all substituted benzoquinones in bombardier beetles and reveal that they are biosynthetic products of two independent pathways. Most likely, the aforementioned biosynthetic channel of hydroxylation of appropriate phenolic precursors and subsequent oxidation is not restricted to bombardier beetles; it could well be a general pathway that leads to the formation of all congeners of benzoquinones, one of the most widely distributed groups of defensive compounds in arthropods.

**Keywords** Benzoquinone · Biosynthesis · Bombardier beetles · *Brachinus elongatulus* · Defensive secretions

## Introduction

Carabid beetles (Carabidae) are well known for using chemical weapons against their enemies (Eisner 1958; Dazzini-Valcurone and Pavan 1980; Dettner 1985; Will et al. 2000). Among the defensive behaviors exhibited by carabids, the most fascinating is the explosive chemical spraying of beetles belonging to the subfamilies Brachininae and Paussinae (Schildknecht 1957; Schildknecht and Koob 1969; Schildknecht et al. 1970). Commonly known as “bombardier beetles,” these beetles have the remarkable ability to explosively discharge with a popping sound a hot noxious chemical spray containing benzoquinones from glandular openings near their abdominal tip (Arndt et al. 2015). In fact, this ejection system has been considered to show fundamental similarity to the pulse jet propulsion mechanism of the German V-1 “buzz” bomb of World War II (Eisner and Aneshansley 1982; Dean et al. 1990).

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Although the defensive secretions of many species of the genus *Brachinus* have been investigated, that of *Brachinus elongatulus* has not been analyzed previously. However, the morphology of the defensive glands of *Brachinus elongatulus* Chaudoir and other bombardier beetles have been described in a series of recent studies (Arndt et al. 2015; Di Giulio et al. 2015; McManus et al. 2018). Similar to all members of the beetle suborder Adephaga (Ober and Maddison 2008; Arndt et al. 2017), bombardiers possess a pair of homologous pygidial glands composed of secretory cells, a reservoir and efferent duct (Balestrazzi et al. 1985). Over the course of evolution of this system (Moore and Wallbank 1968; Maddison et al. 2009), the bombardiers acquired a highly sclerotized chamber known as the reaction chamber, which is separated from the relatively larger reservoir compartment by a one-way valve (Arndt et al. 2015). Bombardiers do not store the final components used in the spray. Instead, they produce the explosive mixture on demand, at the moment of ejection. It has been presumed that the reservoir contains hydroquinones and hydrogen peroxide that function as the precursors for benzoquinone synthesis (Schnepf et al. 1969). When bombardier beetles are provoked, the muscles surrounding the reservoir contract, enabling the reservoir contents to pass through the one-way valve and enter the reaction chamber (Arndt et al. 2015; Di Giulio et al. 2015). The instant hydroquinones and hydrogen peroxide encounter the enzymes in the reaction chamber (Schildknecht and Holoubek 1961), an explosive exothermic reaction takes place producing a mixture of benzoquinones, molecular oxygen, and water, heated up to 100 °C (Schildknecht and Holoubek 1961; Schildknecht et al. 1968, 1970; Aneshansley et al. 1969).

Benzoquinones are not unique to bombardier beetles. In fact, benzoquinones are widely distributed in nature (Abraham et al. 2011; Thomson 1997). They have been reported from bacteria, plants, and many other arthropods (Blum 1987; Happ 1968; Chadha et al. 1961; Dettner 2010) including millipedes (Deml and Huth 2000; Wu et al. 2007; Sekulić et al. 2014; Shear 2015; Bodner et al. 2016) and harvestmen (Fieser and Ardao 1955; Eisner et al. 2004; Rasputnig et al. 2015; Rocha et al. 2013; Gnaspini and Cavalheiro 1998). Within insects, benzoquinones are widespread in cockroaches (Roth and Stay 1958; Nojima et al. 2005), earwigs (Eisner et al. 2000; Gasch et al. 2013), termites (Olagbemiro et al. 1988), and beetles of many families (Blum 1981; Dettner 1991). Apart from their defensive activity, benzoquinones may even act as sex pheromones in scarabaeoid beetles (Vuts et al. 2014).

Although many investigations have been carried out to identify the quinones in beetles, little attention has been given to the biosynthetic origin of these compounds (Happ 1968; Bradshaw 1985; Blum 1987; Morgan 2010). Many investigators have found biosynthetic studies of benzoquinones in arthropods to be challenging and less rewarding than one might

hope. For example, in a very early study on the biosynthetic studies of simple benzoquinones in Jamaican millipedes, Dufey (1974) could not shed much light on the reaction pathways after 2 years of labor. On the other hand, many hypothetical pathways have been proposed with little or no experimental evidence to support these assumptions. For example, Deml and Huth (2000) speculated that both 1,4-benzoquinone (Fig. 1a) and 2-methyl-1,4-benzoquinone (toluquinone; Fig. 1b) originate from the same precursor 1,4-hydroquinone (Fig. 1c) (Shear 2015). However, in a very early study, Meinwald et al. (1966) indicated that in the beetle *Eleodes longicollis*, 1,4-benzoquinone is primarily made from the preformed rings of the aromatic amino acids such as tyrosine or phenylalanine, whereas substituted quinones originate from acetate by the polyketide pathway (Blum 1987). More recently, the biosynthesis of benzoquinones in harvestmen (Opiliones) has been demonstrated to follow a multistep pathway that involves condensation of acetate and propionate units (Rasputnig et al. 2015; Rocha et al. 2013). Moreover, Rocha et al. (2013) indicated that 2-ethyl-1,4-benzoquinone is biosynthesized by the harvestman *Magnispina neptunus* via 3-ethylphenol.

Thus, we predicted that benzoquinones in bombardier beetles might originate in an analogous manner, and the putative biosynthetic precursor of 2-methyl-1,4-benzoquinone (Fig. 1b) could be *m*-cresol. To test this hypothesis, we injected a solution of *m*-cresol-*d*<sub>8</sub> (Fig. 1e) in saline to the hemocoel of several individuals of *B. elongatulus* and subsequently analyzed the defensive spray. Moreover, we tested the long-held hypothesis that the immediate precursors of benzoquinones accumulate in the reservoir. To test this hypothesis, we ligated the base of the reservoir, isolating it from reaction chamber, to verify whether the reservoir contains an accumulation of hydroquinones, which are usually found only as trace components in the sprayed defensive secretion.

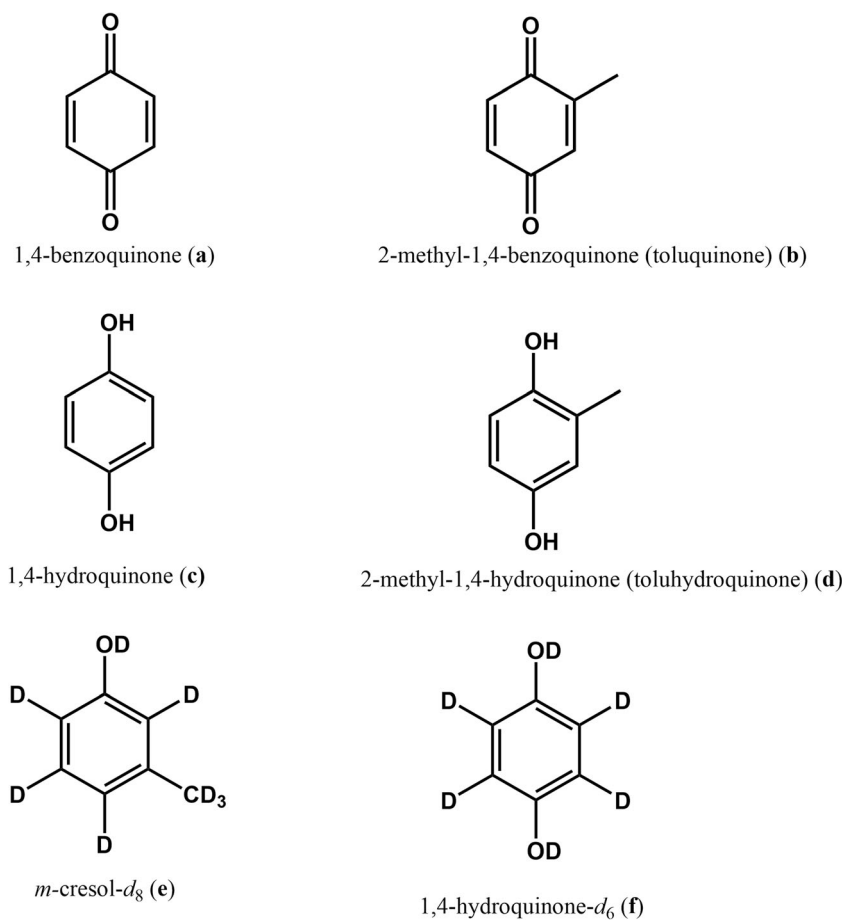
## Materials and methods

### Chemicals

*m*-Cresol-*d*<sub>8</sub> (Fig. 1e) and 1,4-hydroquinone-*d*<sub>6</sub> (Fig. 1f) were purchased from Cambridge Isotopes (Tewksbury, MA), or Sigma-Aldrich Chemical Co (St. Louis, MO). The stock solutions of the isotope-labeling compounds were made in saline (2 mg/100 µL) by dissolving 20 mg of each compound in 1.0 mL of saline solution.

### Collection and handling of beetles and sample preparation

*Brachinus elongatulus* beetles were collected in Pima Co., AZ, and transferred to laboratory facilities at UC Berkeley, CA. Beetles were maintained on moist sphagnum, potting

**Fig. 1** Structures of chemicals used in this study

soil, and sand mix in an environmental chamber at a day/night regime of 13 h light/11 h dark, and day/night temperatures of 23 °C/16 °C. When not receiving any specific treatment, they were held in aggregation and fed dry, high-protein dog food (Blue Buffalo Co, Wilton, CT), banana, and chopped meal worms approximately once per week.

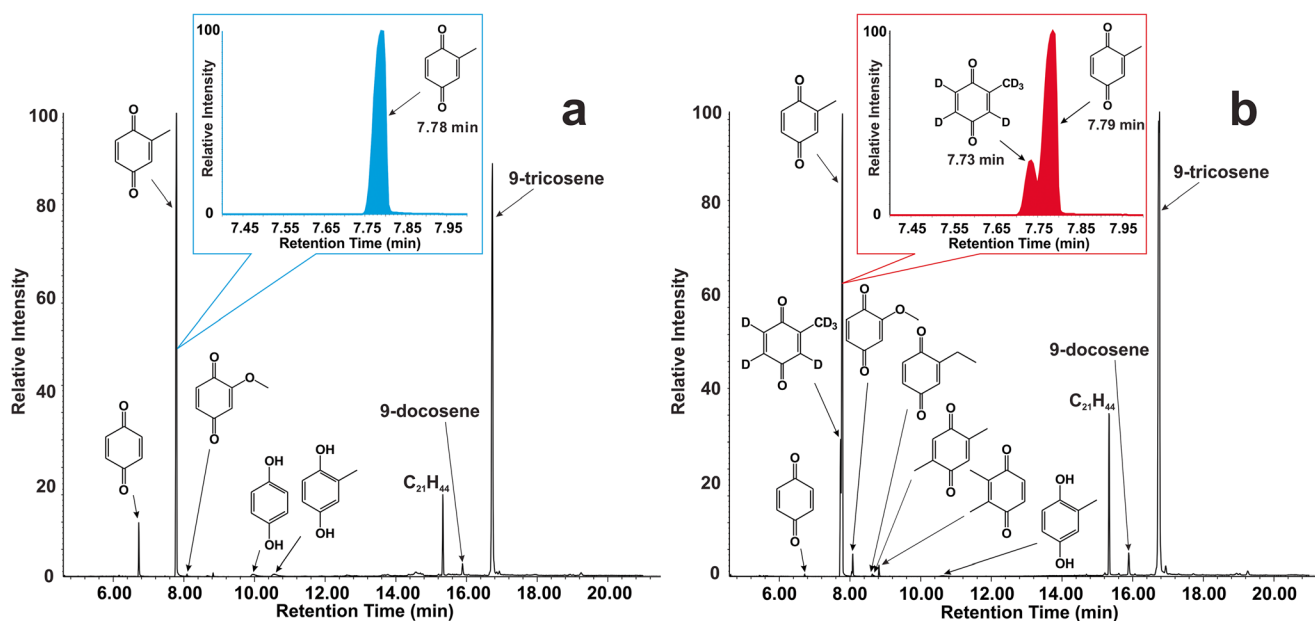
For feeding controls and treatments, the dry-dog food was ground and sifted. Particles larger than 0.5–1.0 mm were retained and used for feeding. Approximately 12.0 mg of the sifted dog food was placed in caps cut from 0.2-mL PCR tubes. For each feeding session, 1.5 µL of saline or *m*-cresol-*d*<sub>8</sub> in saline solution (20 mg/mL) was applied to the food placed in the caps immediately prior to presenting it to the beetle. In some cases, beetles were observed to feed immediately, but typically they feed at night and the food dish was left in the container with the beetle for two nights. Uneaten food and dishes were removed prior to the subsequent feedings. Starting on the day individual beetles were segregated, they were presented food on days 1, 3, 5, 8, 14, 21, and 24, up to the day of dissection or sampling. Altogether, 18 beetles were used for the feeding experiments (2 females, 1 male, and 15 sex-undetermined specimens).

Solutions were directly injected using a 5.0-µL borosilicate glass syringe with a 33-gauge, 12° bevel point needle. A

single syringe and set of needles were dedicated to each treatment to prevent cross-contamination. The needle was inserted into the abdomen of the beetles in the membranous area between the 3rd and 4th ventrites. After injections, needles and syringes were cleaned with a series of washes using water, methanol, and hexane. By this procedure, *m*-cresol-*d*<sub>8</sub> in saline was injected into 23 beetles (1 female, 3.0 µL; 1 female, 3.5 µL; 2 females, 4.0 µL; 1 male, 2.0 µL; 18 sex-undetermined specimens, 5.0 µL each).

Samples of glandular products of beetles were obtained either by dissecting out the glands or by catching ejected spray on filter paper. Sprays were collected by placing a small strip of clean filter paper behind the beetle and stimulating it to spray by slightly pinching a leg with forceps. Filter paper samples were immediately placed in glass vials with solvent, according to the method of Eisner et al. (1977). Beetles were either chilled in a freezer as described below and held with a pair of forceps as they returned to room temperature, at which point they spray or were tethered as done by Aneshansley et al. (1969) and induced to spray.

Live beetles were carefully handled to prevent any discharges and placed in a 0 °C freezer for approximately 3 min to anesthetize and immobilize them. For dissections, an immobilized beetle was then placed on a pre-chilled watch



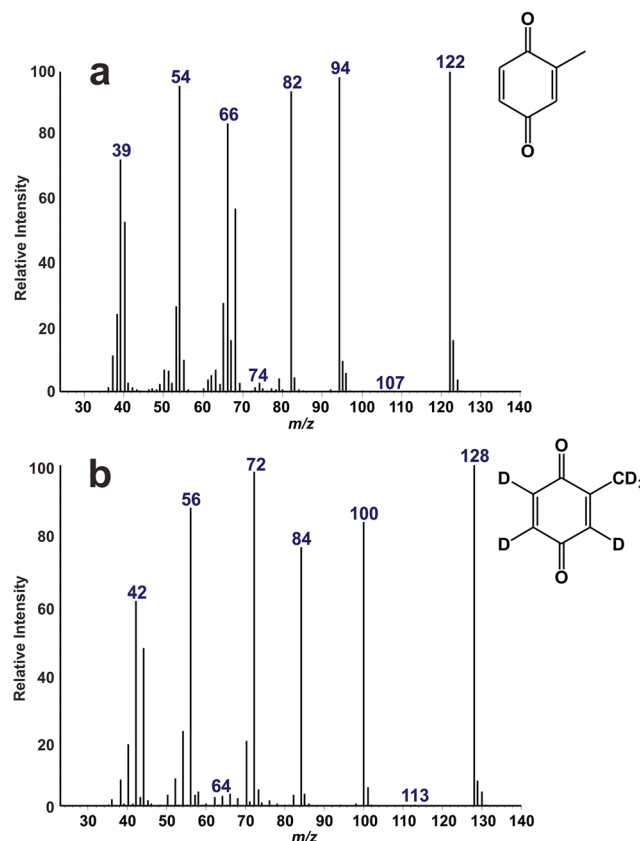
**Fig. 2** Gas chromatograms obtained from GC-MS analysis of a hexane extract of defensive glands of *B. elongatulus* prepared 7 days after the injection of 5  $\mu$ L of saline (a) or a solution of *m*-cresol- $d_8$  in saline (b).

Insets show expansions of the 7.4–8.0 min region of the chromatograms. Analyses were performed on a fused-silica capillary column coated with DB-1

glass, and the abdomen was separated from the rest of the body. The abdomen was dissected further on a glass plate held at room temperature. The tergum was cut along the epipleura and lifted off to reveal the internal organs. A very small amount of distilled water was used to prevent the tissue from sticking to the forceps, and a drop of water was placed on the open abdomen to prevent it from drying during dissection. Any residual water on the dissected structures was wicked away with a small strip of clean filter paper prior to placing it into the solvent.

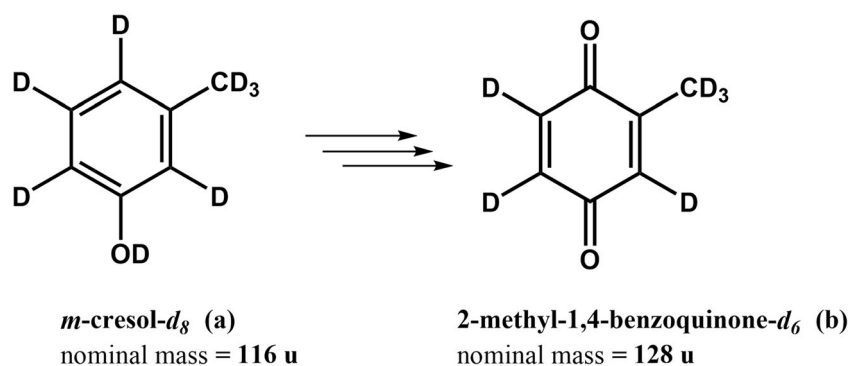
Either one or both of the left and right reservoirs and reaction chambers were excised as a whole gland complex or individually ligated and separated. Whole glands were removed by pulling the reaction chamber away from the tergite and lifting the entire reservoir and reaction chamber out of the abdomen and placing the whole system into a glass vial containing hexane (25–40  $\mu$ L). The relative fullness of the reservoirs was evaluated, and if very full each whole gland was removed and placed in a vial. If a single beetle appeared to have relatively little fluid in both of the left and right reservoirs, both were pooled into a vial as one sample.

In order to sample the reservoir fluid without the inclusion of the reaction chamber contents, a separate reservoir-ligation dissection was performed. To ligate the reservoir, an overhand knot was tied in a single strand of human hair, and the loop of the knot was slipped over the reservoir while it was still in the abdomen. The knot was carefully cinched down around the base of the reservoir just above the reaction chamber. By gently pulling the hair, the reservoir was separated from the abdomen, and the reservoir was broken at its junction with the



**Fig. 3** Electron-ionization (70 eV) mass spectra corresponding to the 2-methyl-1,4-benzoquinone peak in Fig. 2b (a), and that for the deuteriated compound that eluted immediately before the 2-methyl-1,4-benzoquinone peak (b)

**Fig. 4** Conversion of *m*-cresol-*d*<sub>8</sub> (a) to 2-methyl-1,4-benzoquinone-*d*<sub>6</sub> (b)



reaction chamber. In this way, the reservoir contents were held intact, and there were no apparent signs of the benzoquinone-forming reaction occurring (no color change). The strand of hair was trimmed near the knot, and the ligated reservoir was placed in a glass vial containing hexane (25–40  $\mu$ L). Seven beetles used as controls received only the saline solution. Beetles fed with saline-treated food were then subjected to dissection of the whole gland (one specimen, sex was not determined), dissection of the ligated reservoir (one female and one male), or spray on filter paper (one female and one male). Beetles injected with 5.0  $\mu$ L saline were subject to similar treatment.

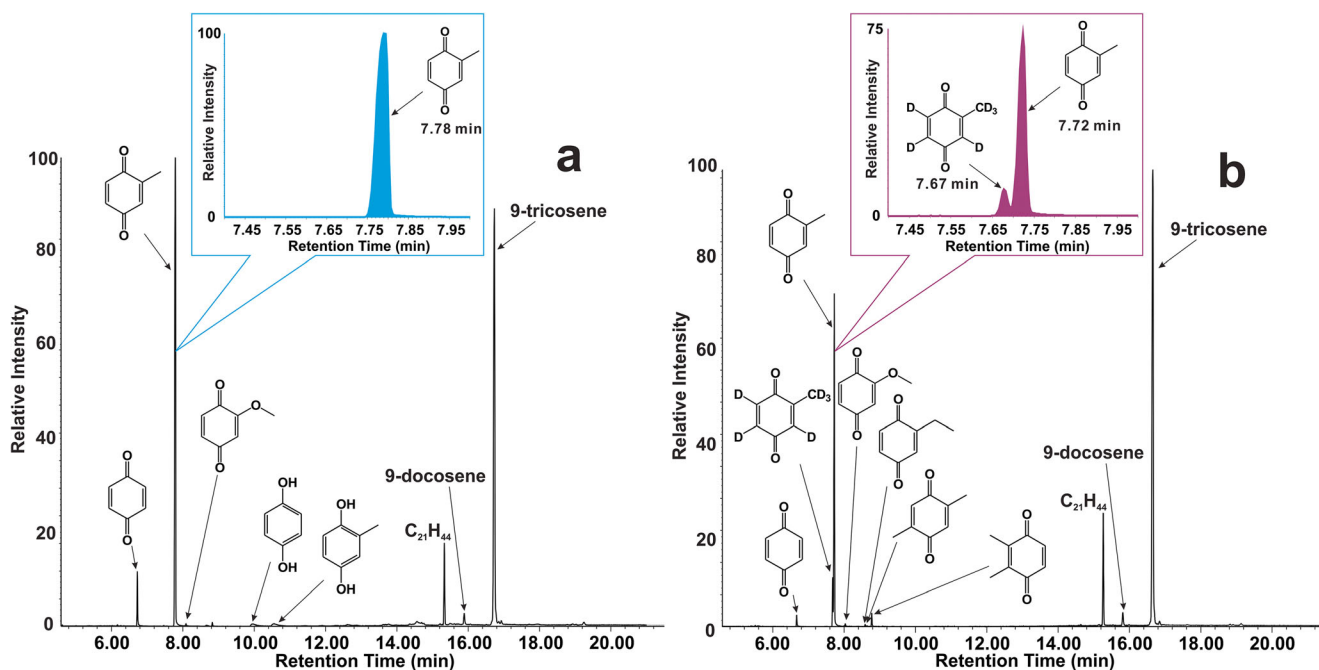
Eleven beetles were treated with 1,4-hydroquinone-*d*<sub>6</sub> injections: three females (two injected with 5.0  $\mu$ L, one injected with 3.5  $\mu$ L), two males (injected with 5.0  $\mu$ L), three beetles of undetermined sex (injected with 5.0  $\mu$ L), and three females (injected with 5.0  $\mu$ L and subsequently fed with dog food

mixed with 5  $\mu$ L of 1,4-hydroquinone-*d*<sub>6</sub>). These beetles were then maintained individually and fed with untreated dog food on days 1, 3, and 5, and then dissected on days 7–9 and the glandular apparatus of each specimen was removed and extracted as above.

Samples, either whole gland dissections or filter papers containing sprayed secretions, were placed in hexane (25–40  $\mu$ L) and shipped to Hoboken, NJ, in an insulated box chilled with Koolit-gel packs.

### Chemical analysis

Samples were analyzed by GC-MS on an Agilent HP 6890 Gas Chromatograph (GC) fitted with a fused-silica capillary column coated with DB-1 (dimethylpolysiloxane, 30 m  $\times$  0.25 mm ID, film thickness 0.25  $\mu$ m) linked to Agilent HP 5973 Mass Selective Detector (MSD). Samples were



**Fig. 5** Gas chromatograms obtained from GC-MS analysis of a hexane extract of defensive glands of *B. elongatulus* after injecting 5  $\mu$ L of saline (a) or after feeding of laced food admixed with *m*-cresol-*d*<sub>8</sub> for about

3 weeks (b). Insets show expansions of the 7.4–8.0 min region of the chromatograms. Analysis were performed on a fused-silica capillary column coated with DB-1

introduced by splitless injection at 250 °C. The GC oven was kept initially at 40 °C for 2.00 min and raised at a rate of 20 °C/min to a final temperature of 280 °C and held for 7.5 min. Alternatively, some samples were analyzed on a Shimadzu QP5050 mass spectrometer linked to a Shimadzu GC-17A gas chromatograph (GC) installed with a DB-624 (6% cyanopropyl-phenyl, 94% dimethylpolysiloxane; film thickness 1.12 µm) coated 25 m × 0.20 mm ID capillary column. The oven temperature program was 100 °C initially and increased at a rate of 10 °C/min to a final temperature of 250 °C.

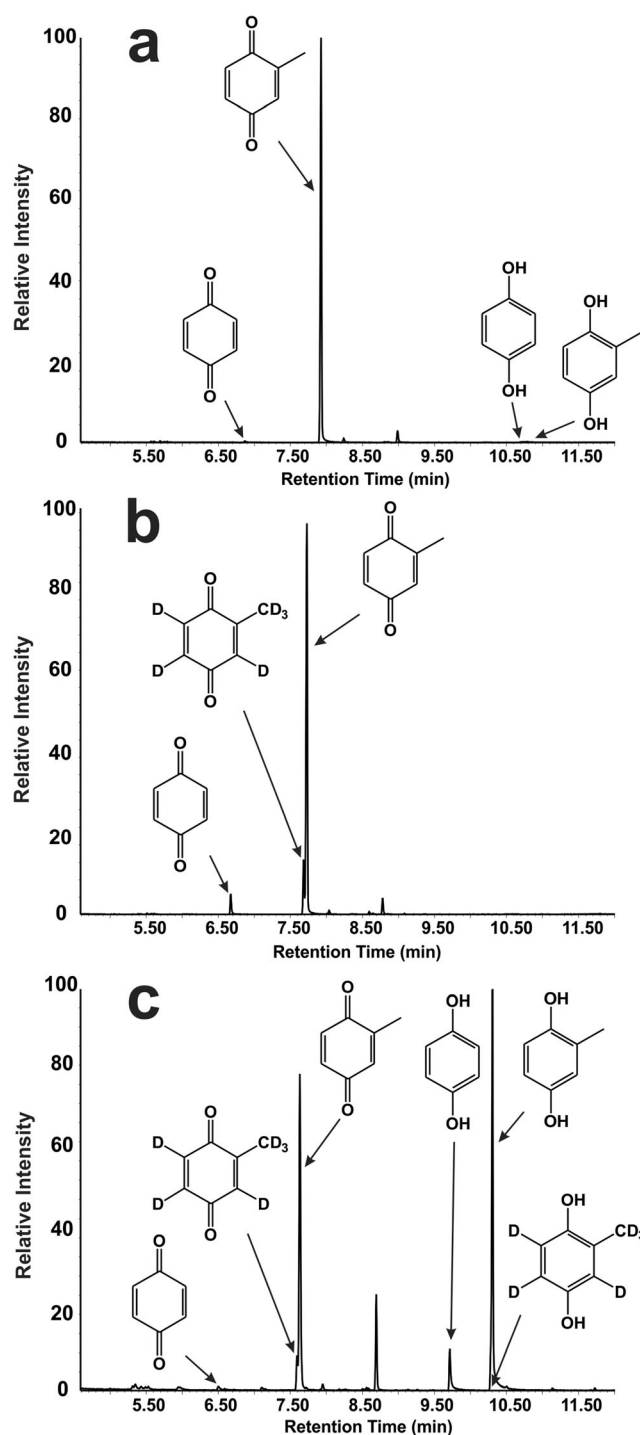
### Derivatization with dimethyl disulfide

An aliquot of secretion extract (5 µL) was mixed with dimethyl disulfide (2 µL) in a reaction vial (100 µL). A solution of iodine in ether (5%, 10 µL) was added, and the mixture was kept overnight at room temperature. Afterward, the reaction mixture was decolorized with a minimum volume of aqueous sodium thiosulfate (5%). Products were extracted into hexane and examined by GC-MS on a DB-1 coated capillary column (Francis and Veland 1987; Buser et al. 1983). The initial oven temperature was 150 °C, which was immediately raised to 295 °C at a rate of 15 °C/min and held constant until all derivatives eluted.

### Results

The analysis of an extract made from the defensive apparatus of *B. elongatus* by GC-MS showed that the major component in the secretion is 2-methyl-1,4-benzoquinone (Fig. 2a).

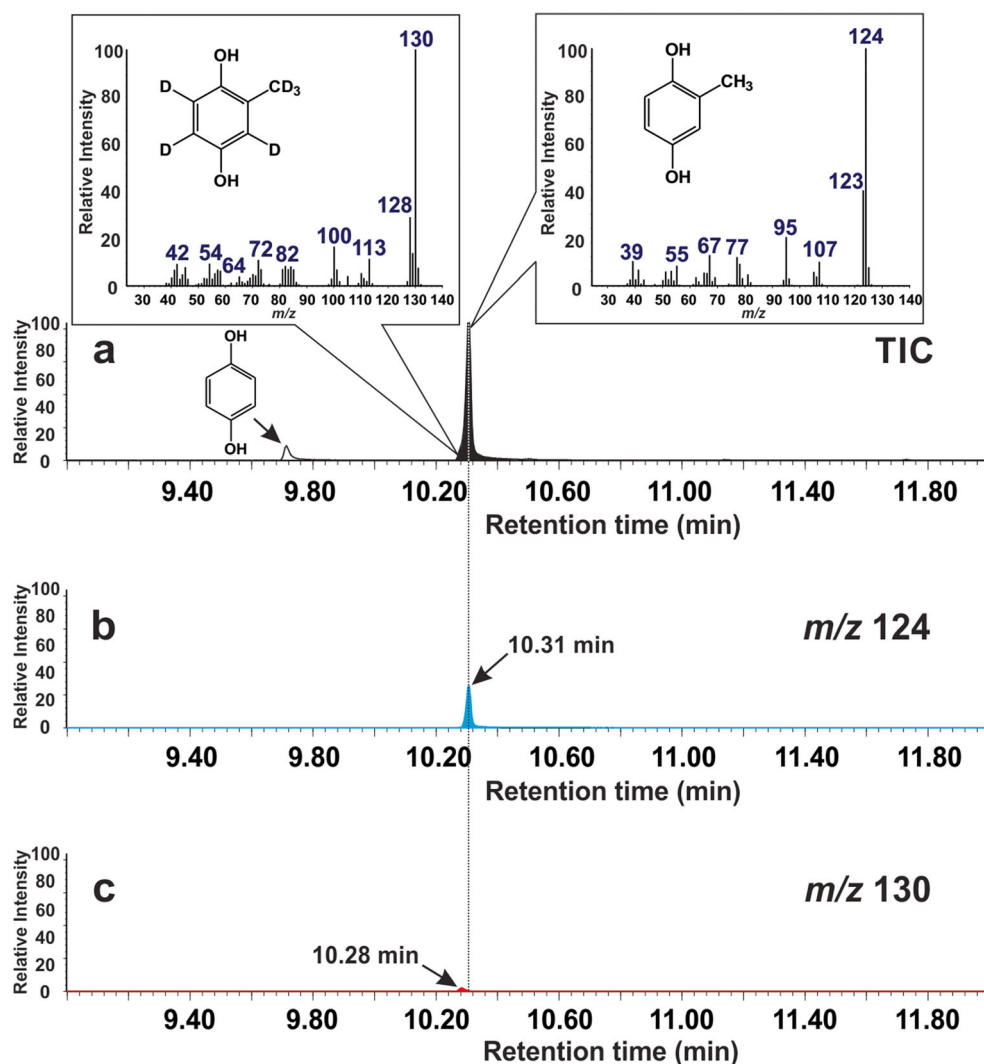
A typical gas chromatogram recorded from control specimens showed two major peaks for 1,4-benzoquinone (Fig. 1a) and 2-methyl-1,4-benzoquinone (Fig. 1b), in addition to a few minor signals for methoxy-1,4-benzoquinone, ethyl-1,4-benzoquinone, 2,5-dimethyl-1,4-benzoquinone, 2,3-dimethyl-1,4-benzoquinone, and traces of 1,4-hydroquinone (Fig. 1c) and 2-methyl-1,4-hydroquinone (Fig. 1d) (Fig. 2a). All compounds were identified by comparing retention times and mass spectra with those of synthetic standards (see Supplementary Table S1 for analytical details). The hydrocarbon peaks were identified as heneicosane, 9-docosene, and 9-tricosene (Supplementary Figures S1 and S2). Compared with the gas chromatographic profiles generated from hexane extracts prepared from control specimens, those recorded from beetles injected with *m*-cresol-*d*<sub>8</sub> showed an additional chromatographic peak of significant intensity, which eluted immediately before the 2-methyl-1,4-benzoquinone peak. Typical profiles recorded from two beetles kept for 7 days before dissection are depicted in Fig. 2b (analytical details of all other samples and results are summarized in Supplementary Table S2).



**Fig. 6** Chromatograms obtained from GC-MS analysis of a hexane extract of defensive glands of a control *B. elongatus* sample (a) and that of a *m*-cresol-*d*<sub>8</sub>-fed specimen (b). c Chromatogram recorded from an extract of the reservoir fluid of a *m*-cresol-*d*<sub>8</sub>-fed specimen that was prevented from mixing with the reaction chamber contents by ligation. Analysis were performed on a fused-silica capillary column coated with DB-1

The electron-ionization mass spectrum corresponding to the additional chromatographic peak that eluted immediately before the 2-methyl-1,4-benzoquinone peak showed

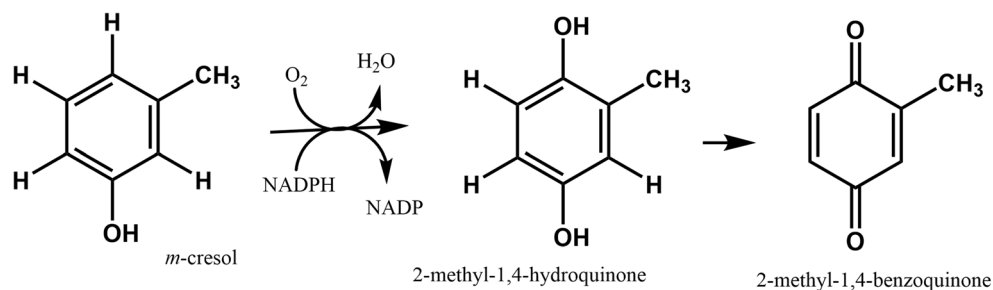
**Fig. 7** An expansion of the 9.0- to 12.0-min region of the total-ion chromatogram depicted in Fig. 6c (a) and ion chromatograms extracted from acquired data for  $m/z$  124 (b) and  $m/z$  130 (c). Insets show background-subtracted mass spectra corresponding to two specific points of the chromatogram. Analysis were performed on a fused-silica capillary column coated with DB-1

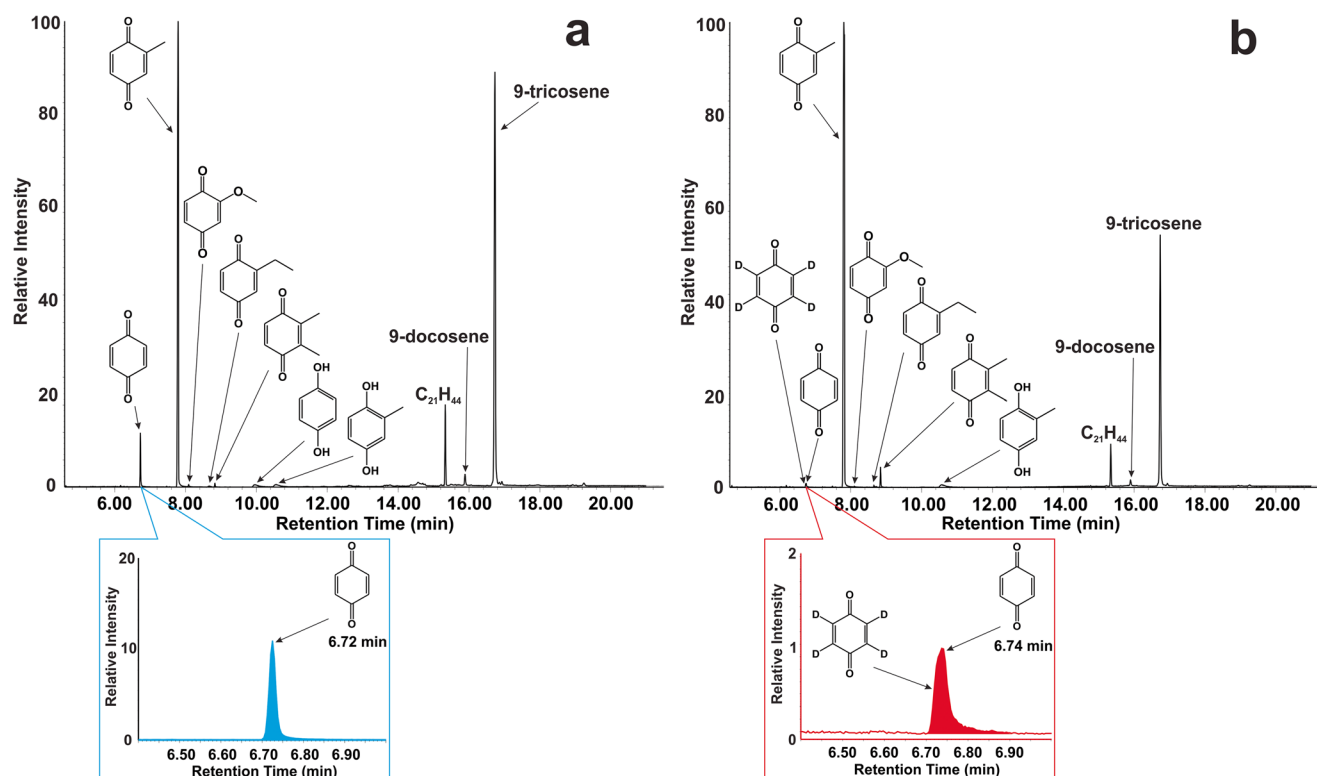


a signal at  $m/z$  128 for its molecular ion (Fig. 3b). On the basis of mass spectral interpretations provided by Beynon and Williams (1960), it was evident that of the eight deuterium atoms initially present in the *m*-cresol- $d_8$  (Fig. 4a) precursor compound, only six of them were incorporated into the product (Fig. 4b). This result conclusively showed that *B. elongatulus* beetles are able to convert *m*-cresol- $d_8$  (Fig. 4a (nominal mass 116 u), injected into the hemocel), into 2-methyl-1,4-benzoquinone- $d_6$  (Fig. 4b; nominal mass 128 u) (Fig. 4).

Remarkably, *B. elongatulus* beetles were capable of converting *m*-cresol- $d_8$  (Fig. 4a; nominal mass 116 u) into 2-methyl-1,4-benzoquinone- $d_6$  (Fig. 4b; nominal mass 128 u) even when the precursor compound was only added to their food. In fact, incorporations were detected even after 24 h of feeding. Figure 5 b shows data recorded from a specimen that was continuously fed food mixed with the deuterium-labeled compound for about 3 weeks. Moreover, the deuterium labeling in *m*-cresol- $d_8$  (Fig. 1e) in both experiments (injection and feeding) was

**Fig. 8** Conversion of *m*-cresol to 2-methyl-1,4-benzoquinone via 2-methyl-1,4-hydroquinone





**Fig. 9** Gas chromatograms obtained from GC-MS analysis of a hexane extract of defensive glands of *B. elongatulus* after injection of 5  $\mu$ L of saline (a) or solution of 1,4-hydroquinone- $d_6$  in saline (b). Analysis were performed on a fused-silica capillary column coated with DB-1

incorporated specifically only to 2-methyl-1,4-benzoquinone (Fig. 1b) and not into 1,4-benzoquinone (Fig. 1a) or any other quinone present in the secretion in trace amounts.

Chromatograms recorded from extracts of the reservoir fluid from ligated reservoirs showed pronounced peaks for the two hydroquinones (1,4-hydroquinone (Fig. 1c) and 2-methyl-1,4-hydroquinone (Fig. 1d)), which are usually found only as trace components in the sprayed defensive secretion (Supplementary Figure S3). For example, in the chromatograms recorded from extracts of the whole glandular complex, the hydroquinone peaks were negligibly small (Figs. 2a and 5a). Evidently, our ligation technique reduces the formation of benzoquinones significantly; however, the technique does not provide a hermetic seal and prevent their formation completely (Fig. 6c).

We hypothesized that these hydroquinones are the immediate precursors of the benzoquinones. In order to test our hypothesis, we extracted ligated reservoirs of *m*-cresol- $d_8$ -fed beetles. The chromatogram recorded from these samples showed a more pronounced peak for 2-methyl-1,4-hydroquinone (Fig. 6c) and also a small peak for the presence of 2-methyl-1,4-hydroquinone- $d_6$  (Fig. 1f). The formation of 2-methyl-1,4-hydroquinone- $d_6$  was confirmed by extracting an ion-intensity profile for the  $m/z$  130 ion (Fig. 7c). The retention time of the 2-methyl-1,4-

hydroquinone- $d_6$  peak was slightly shorter than that of the 2-methyl-1,4-hydroquinone peaks as that expected from a deuteriated analog. Moreover, the background-subtracted mass spectrum corresponding to the  $m/z$ -130-ion peak confirmed that of the eight deuterium atoms present in the precursor *m*-cresol- $d_8$ , six of them are incorporated to 2-methyl-1,4-hydroquinone- $d_6$  (Fig. 7a insets).

Evidently, *m*-cresol undergoes hydroxylation to 2-methyl-1,4-hydroquinone first (presumably in the secretory cells, by a NADPH and oxygen mediated biosynthetic process; Artigot et al. 2009), which then undergoes oxidation to 2-methyl-1,4-benzoquinone in the reaction chamber (Fig. 8). In an analogous manner, 1,4-hydroquinone is oxidized to 1,4-benzoquinone in the reaction chamber. When 1,4-hydroquinone- $d_6$  (Fig. 1f) was administered to *B. elongatulus* and the defensive gland extract was obtained after a few days and analyzed by GC-MS, the chromatographic peak for 1,4-benzoquinone was accompanied by another peak for 1,4-benzoquinone- $d_4$  (Fig. 9b). The presence of 1,4-benzoquinone- $d_4$  was confirmed by extracting a mass chromatogram for the  $m/z$  112 ion (Supplementary Figure S4c). Moreover, the background-subtracted mass spectrum corresponding to the  $m/z$  112 ion confirmed that of the six deuterium atoms present in the precursor, four of them have been incorporated to 1,4-benzoquinone- $d_4$  (Supplementary Figure S4a).

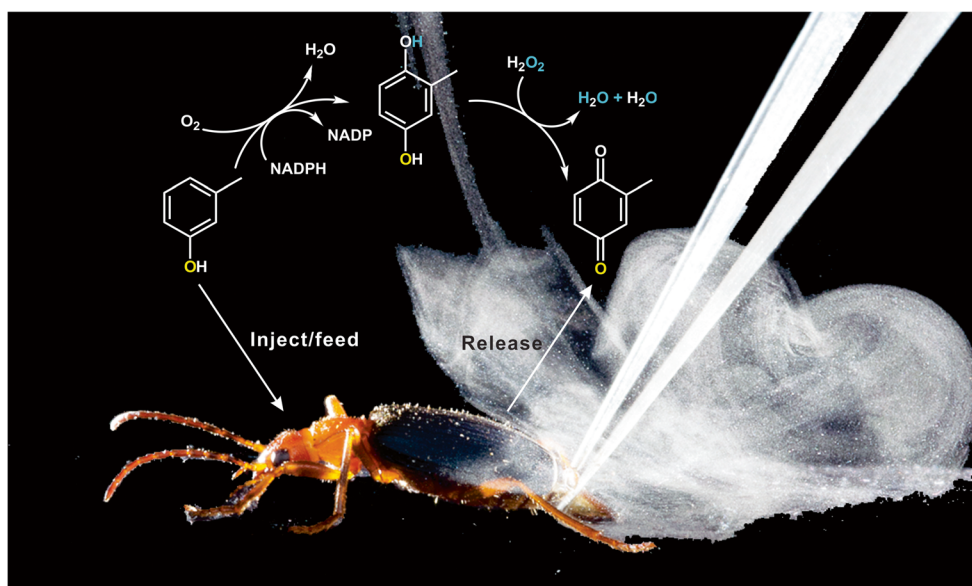


Fig. 10 Biosynthesis of 2-methyl-1,4-benzoquinone

## Discussion

In 2000, Deml and Huth (2000) speculated that the biosynthetic processes of all benzoquinones originate from one core precursor. According to their proposal, 1,4-hydroquinone is the central molecule that gets metabolized and converted to all benzoquinones (Shear 2015). However, we predicted that 2-methyl-1,4-benzoquinone in bombardier beetles might not originate 1,4-benzoquinone. To test our hypothesis, we injected a solution of *m*-cresol-*d*<sub>8</sub> to several beetles. Subsequent chemical analysis showed that six deuterium atoms from *m*-cresol-*d*<sub>8</sub> were incorporated into 2-methyl-1,4-benzoquinone. Furthermore, our mass chromatographic analysis of data acquired from the defensive secretion of 1,4-hydroquinone-*d*<sub>6</sub>-injected bombardier beetles revealed the presence of only one deuterium-labeled compound. In other words, the deuterium atoms from 1,4-hydroquinone-*d*<sub>6</sub> were specifically incorporated only into 1,4-benzoquinone and not into 2-methyl-1,4-benzoquinone or any other quinone present in the secretion. Since no other benzoquinone acquired the deuterium-labeling, we can conclude that each benzoquinone originates only from a specific hydroquinone precursor and not by an alkylating mechanism using 1,4-hydroquinone as the core precursor as suggested by Deml and Huth (2000). In addition, we confirm that bombardier beetles have the ability to convert *m*-cresol to 2-methyl-1,4-hydroquinone, which is then oxidized to 2-methyl-1,4-benzoquinone in their defensive spray (Fig. 10).

It is known that *m*-cresol is a cytotoxic compound that interacts with membrane lipids at the cellular level and causes membrane leakage (Paiva et al. 2016). However, *B. elongatulus* appears to be resistant to the ill effects of *m*-cresol. Whether the labeled *m*-cresol was provided in their food or via direct injection into the hemocoel, the beetles were

able to convert it into hydroquinones stored in the reservoir. It is likely that the *m*-cresol added to their food passed from the digestive system through the midgut wall into the hemocoel and was transported through the hemolymph to the secretory cells of the reservoir. McManus et al. (2018) showed that the secretory cells and Malpighian tubules are always intertwined in *B. elongatulus* (see Fig. 2 in McManus et al. (2018)). This physical connection further suggests that the digestive system may also be somehow involved in the production of *Brachinus* defensive chemicals. Whatever the phenol transport pathway may be, the speed at which the beetles can incorporate *m*-cresol into benzoquinone—in as little as 24 h—is remarkable and attests to the importance of defensive chemistry for the survival of these beetles.

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## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflict of interest.

**Statement on the welfare of animals** All applicable international and national guidelines for the care and use of animals were followed.

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