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#### 1 A novel tri-unsaturated highly branched isoprenoid (HBI) alkene

#### 2 from the marine diatom Navicula salinicola

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#### 13 ABSTRACT

14 A novel tri-unsaturated  $C_{25}$  highly branched isoprenoid (HBI) alkene has been identified in a laboratory culture of the diatom Navicula salinicola and 15 its structure determined using a combination of NMR spectroscopy and gas 16 17 chromatography-mass spectrometry (GC-MS). This represents the first 18 report of a C<sub>25</sub> HBI in a marine diatom from the *Navicula* genus, although a 19 different tri-unsaturated  $C_{25}$  HBI has been reported previously in the freshwater species N. sclesvicensis and unspecified HBIs have been 20 21 identified in the brackish N. phyllepta. The newly characterised HBI contains a relatively unusual conjugated diene sub-unit, a structural feature 22

only previously reported in some HBIs biosynthesised by a further marine
diatom, *Haslea ostrearia*.

25 Keywords: highly branched isoprenoid; alkene; diatom; Navicula salinicola

26 1. Introduction

27  $C_{25}$ highly branched isoprenoid (HBI) alkenes are common 28 components of marine and lacustrine sediments worldwide and are 29 biosynthesised by certain diatoms mainly belonging to the genera Haslea, Pleurosigma, Berkeleya 30 Navicula, Rhizosolenia, and Pseudosolenia (Volkman et al., 1994; Belt et al., 1996, 2000, 2001; Sinninghe Damsté et al., 31 32 1999; Grossi et al., 2004; Brown et al., 2014; Kaiser et al., 2016). A single tri-unsaturated C<sub>25</sub> HBI (Structure 6; Fig. 1) has been identified in the 33 34 freshwater diatom Navicula sclesvicensis (Belt et al., 2001), and the mainly 35 brackish species N. phyllepta has also been reported as an HBI-producing diatom (Sinninghe Damsté et al., 2004), although no structures were given. 36 37 In contrast, no HBIs have as yet been reported in any marine species within the *Navicula* genus. Here, we identify a novel tri-unsaturated  $C_{25}$  HBI 38 isolated from a laboratory culture of the marine diatom N. salinicola and 39 40 report its structure based on analysis by NMR spectroscopy and gas 41 chromatography–mass spectrometry (GC–MS).

42 2. Experimental

The benthic diatom N. salinicola was collected from a coastal marine
environment (M. Kulikovskiy, June 2016 at Nha Trang, Vietnam,
12°13'14.5"N 109°12'18.3"E) and kept as strain BTD1 in the Laboratory of

Molecular Taxonomy of Aquatic Plants Institute of Plant Physiology (RAS). 46 47 (Further taxonomic information can be found in the Supplementary Information). Initially, N. salinicola was cultured at 15 °C, 150 µmol m<sup>-2</sup> s<sup>-1</sup> 48 continuous light in small flasks (150ml). Samples were harvested by 49 filtration using MF-Millipore<sup>™</sup> membrane filters (25 mm diameter, 0.3 µm 50 51 pore size) during the exponential and stationary growth phases (Fig. 2). 52 Large-scale cultures were then set up in several 2 l conical flasks under the 53 same growth conditions. 80 l of such cultures were harvested by centrifugation (4000 rpm, 10 mins) during the stationary growth phase. For 54 55 the small- and large-scale culture experiments, we used enriched f/2 medium (Guillard and Ryther 1962), along with the following nutrient 56 concentrations: 2646 µM NaNO<sub>3</sub>, 318 µM Na<sub>2</sub>SiO<sub>3</sub>, 108 µM NaH<sub>2</sub>PO<sub>4</sub>. The 57 58 filtered and centrifuged biomass was freeze dried and the resulting dry 59 material extracted via sonication using hexane (3 ml (small-scale cultures); 60 25 ml (large-scale culture)). The total hexane extract (THE) was then 61 concentrated by removing hexane under a stream of nitrogen and partially purified using column chromatography (SiO<sub>2</sub>). The hydrocarbon fraction 62 63 (hexane) was analysed by GC–MS using an Agilent 7890 gas chromatograph 64 equipped with a HP<sub>5MS</sub> fused-silica column (30 m;  $0.25 \mu m$  film thickness; 0.25 mm internal diameter) coupled to an Agilent 5975 series Mass 65 Selective Detector (MSD). NMR data were obtained using a JEOL ECP-400 66 NMR spectrometer with chemical shifts measured relative to those of CDCl<sub>3</sub> 67 68 (<sup>1</sup>H: 7.24 ppm; <sup>13</sup>C: 77.0 ppm). NMR data were collected on the THE

obtained from the large-scale culture. The purity of the newly reported  $C_{25:3}$ HBI (see Section 3) in this THE was estimated to be ca. 90% based on its relative peak area (GC-MS; Supplementary Information) and by the relative integration values of H-23 (Fig. 1) versus the alkenic protons of the co-occurring polyunsaturated linear alkenes ( $\delta$  = ca. 5.3 ppm) in the <sup>1</sup>H NMR spectrum.).

75 3. Results and discussion

76 Following extraction of several small-scale cultures of N. salinicola from the exponential and stationary growth phases, analysis of partially 77 purified THEs by GC-MS revealed the presence of a suite of closely eluting 78 polyunsaturated linear alkenes (e.g. heneicosa-3,6,9,12,15,18-hexaene; n-79 80  $C_{21:6}$ ), trace amounts of a di-unsaturated HBI (2; Fig. 1) and a further 81 compound exhibiting similar mass spectral properties to a range of C<sub>25</sub> HBIs 82 characterised previously. However, although the retention index (RI) of this 83 component ( $RI_{HP5ms} = 2141$ ) did not match that of any previously reported C<sub>25</sub> HBIs, hydrogenation of an aliquot of one THE resulted in the formation 84 of the parent HBI alkane C<sub>25:0</sub>, thus confirming the C<sub>25</sub> carbon skeleton. 85 86 Interestingly, this new HBI was only detected in cultures harvested during 87 the stationary phase and was identified as tri-unsaturated on the basis of its 88 molecular ion ( $M^+$  346; Fig. 3). At this point, it is not clear why this new 89 HBI and the co-occurring diene 2 were not detected during the exponential growth phase, although we note that some variability in cellular HBI 90 91 concentrations and distributions have been reported in a small number of

previous studies (e.g. Wraige et al., 1997,1998,1999; Brown et al., 2020). 92 93 From the large-scale culture of N. salinicola, we obtained ca 0.2 mg of the 94 partially purified HBI triene (3; Fig. 1), which enabled full structural characterisation using <sup>1</sup>H and <sup>13</sup>C NMR spectroscopy. A conjugated diene 95 96 sub-structure (C22–C25; C23–C24; Fig 1) could be readily identified through 97 a particularly characteristic low field resonance in the <sup>1</sup>H NMR spectrum 98 due to H-23 (Wraige et al., 1997; Allard et al., 2001), together with further 99 low field resonances that could be attributed to alkenic methylene protons 100 at C24 and C25 (Table 1). The third double bond could also be identified 101 from its methylene protons at C17. Alternative positions for this third 102 double bond at C1-C2 or C14-C15 can be discounted due to the observation 103 of two isopropyl groups in the <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 1). Further, a 104 double bond at C10–C18 leaves a solitary methyl group at C17 whose <sup>13</sup>C chemical shift would be at ca. 15.5 ppm, by comparison with related 105 106 compounds (e.g. 1; Belt et al., 2012). In contrast, isolated methyl groups at 107 C18 in previously characterised HBIs resonate at ca. 19–20 ppm (e.g. 19.8 108 ppm for HBIs 1 and 2; Fig. 1) (Johns et al., 1999; Belt et al., 2012), 109 consistent with that observed for HBI 3 (i.e. 19.9 ppm; Table 1). The  ${}^{13}C$ 110 NMR spectrum of HBI 3 also contained individual resonances due to the six magnetically inequivalent alkenic carbon nuclei (Table 1) and complete <sup>13</sup>C 111 112 resonance assignments could be proposed by comparison with structurally 113 similar HBIs characterised previously (Fig. 1; Belt et al., 1996, 2012; Wraige 114 et al., 1997; Allard et al., 2001), some of which contain a conjugated diene

115 sub-unit (i.e. 7–8). The GC RI of HBI 3 ( $RI_{HP5ms} = 2141$ ) is substantially 116 higher than those of some other HBI trienes (e.g.  $RI_{HP5ms} = 2114$  (4); 2109 117 (5); 2090 (6)) but is consistent with that reported for the structurally related 118 tetraene 7 ( $RI_{HP-5} = 2159$ ; Allard et al., 2001).

119 The identification of a  $C_{25}$  HBI in N. salinicola represents the first 120 example of HBI production within a marine Navicula species despite the 121 near-ubiquity of this genus within natural diatom populations. Since the 122 Navicula and Haslea genera are quite similar, phylogenetically, with the 123 latter well-known as an HBI-producing genus, the new finding is probably 124 not surprising; however *Navicula* is a far more common genus, potentially making it a more important source of some HBIs in marine sediments. In 125 126 terms of its structure, the conjugated diene sub-structure (C22-C25; C23-127 C24; Fig. 1) is somewhat unusual, although there is some previous 128 precedent for such a feature in other HBIs isolated from a small number of 129 cultures of the marine diatom Haslea ostrearia (Wraige et al., 1997; Allard et al., 2001). In fact, HBI 3 is a close structural analogue of HBI 7 identified 130 131 previously in *H. ostrearia*, albeit as a minor component. Since the retention 132 index of HBI 3 ( $RI_{HP5ms} = 2141$ ) is similar to two HBI tetraenes identified in 133 some cultures of *H. ostrearia* (viz.  $RI_{HP-5} = 2143-2146$ ; Allard et al., 2001), 134 one of which also contained HBI 7, it possible that HBI 3 may also have been present in the corresponding lipid extracts, but not identified due to co-135 elution. We are unware of any geochemical reports of HBI 3 although its 136 137 characterisation described herein may, in the future, lead to its positive

identification in sedimentary archives, an outcome that may potentially add
to the use of HBIs as palaeoenvironmental indicators (c.f. HBIs 1 and 2 for
Arctic and Antarctic sea ice; see Belt, 2018 for a review).

141 4. Conclusions

We report the structural identification of a novel C<sub>25</sub> HBI biomarker 142 143 in the marine diatom N. salinicola, the first example of HBI production within a marine Navicula species, thus expanding the potential number of 144 sources of HBIs in the environment. Further studies into N. salinicola and 145 146 HBIs related species valuable in the of may prove use as 147 palaeoenvironmental proxies.

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- 221 Figure legends

Figure 1. Structures of C<sub>25</sub> HBIs referred to in the text. HBIs are numbered
in order of increasing unsaturation.

Figure 2. Growth curve of the small-scale culture of the marine diatom N.
salinicola. Cell densities were estimated by measuring in vivo fluorescence
between 460 nm to 670 nm using the SpectraMax® iD3 Multi-Mode
Microplate Reader. Cultures were harvested during the exponential (empty
arrow) and/or stationary (solid arrow) growth phases (n=1).

**Figure 3.** Structure and mass spectrum of HBI **3**.





249

250 Table 1. Key NMR data for HBI **3**.

	Carbon shift ( $\delta$ /ppm)	Proton Number	Proton shift (δ/ppm)
1,16*	22.8, 22.7*	1,15,16,19	0.85 (12H, m)
2	28.0	5,7,21	2.05 (5H, m)
3	39.1	17	4.77, 4.72 (2H, 2 x s, br)
4	25.6	18	0.82 (3H, t, J=6.9 Hz)
5	33.0	23	6.33 (1H, dd, J=17.6, 11.0 Hz)
6	152.3	24a	5.01 (1H, d, J=11.0 Hz)
7	47.1	24b	5.16 (1H, d, J=17.6 Hz)
8	29.8	25	4.96 (2H, m, br)
9	34.9		
10	33.0		
11	37.1		
12	24.8		
13	39.4		
14	28.0		
15,19*	22.8, 22.7*		
17	109.0		
18	19.9		
20	29.4		
21	32.0		
22	146.9		
23	139.0		
24	113.1		
25	115.5		

251 \*Assi

\*Assignments may be interchanged