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Ocean acidification increases iodine accumulation in kelp-based coastal food webs

Running head: Ocean acidification increases iodine in kelp

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ABSTRACT

Kelp are main iodine accumulators in the ocean, and their growth and photosynthesis are likely to benefit from elevated seawater CO_2 levels due to ocean acidification. However, there are currently no data on the effects of ocean acidification on iodine metabolism in kelp. As key primary producers in coastal ecosystems worldwide, any change in their iodine metabolism caused by climate change will potentially have important consequences for global geochemical cycles of iodine, including iodine levels of coastal food webs that underpin the nutrition of billions of humans around the world. Here, we found that elevated pCO$_2$ enhanced growth and increased iodine accumulation not only in the model kelp *Saccharina japonica* using both short-term laboratory experiment and long-term *in situ* mesocosms, but also in several other edible and ecologically significant seaweeds using long-term *in situ* mesocosms. Transcriptomic and proteomic analysis of *S. japonica* revealed that most vanadium-dependent haloperoxidase genes involved in iodine efflux during oxidative stress are down-regulated under increasing pCO$_2$, suggesting that ocean acidification alleviates oxidative stress in kelp, which might contribute to their enhanced growth. When consumed by abalone (*Haliotis discus),* elevated iodine concentrations in *S. japonica* caused increased iodine accumulation in abalone, accompanied by reduced synthesis of thyroid hormones. Thus, our results suggest that kelp will benefit from ocean acidification by a reduction in environmental stress however, iodine levels in kelp-based coastal food webs will increase, with potential impacts on biogeochemical cycles of iodine in coastal ecosystems.

INTRODUCTION

Anthropogenic emissions of CO_2, associated ocean acidification (OA) and global warming, are increasing at rates unprecedented in the geological record (Sunday et al., 2017; Thomsen et al., 2017). These changes are expected to directly affect primary producers by changing growth rates, photosynthesis and metabolism, and indirectly by reducing biodiversity and altering ecosystem structure (Enochs et al., 2015; Martínez-Botí et al., 2015; Myers et al., 2017; Ullah, Nagelkerken, Goldenberg, & Fordham, 2018). Furthermore, OA is predicted to cause an increase in the accumulation of toxic phenolic compounds across multiple trophic levels, from phytoplankton to zooplankton (Jin et al., 2015). Higher temperatures are also expected to increase the concentrations of nutrients such as nitrogen (N), potassium (K), and magnesium (Mg) stored in living biomass (Zhang et al., 2018) and impact the global biotic metabolic rates (Dillon, Wang, & Huey, 2010). Metabolic rate changes and nutrient variation under climate change will affect human health through consumption of these organisms (Jin et al., 2015; McKibben et al., 2017; Zhu et al., 2018) and potentially impact food security (Bloom, Burger, Asensio, & Cousins, 2010; Loladze, 2002; Myers et al., 2017; Phalkey, Aranda-Jan, Marx, Höfle, & Sauerborn, 2015). However, climate impacts on the nutrient composition of primary producers (such as, seaweeds) in coastal ecosystem has, to date, been little explored.

Seaweeds, including, brown, red, and green algae are a rich source of iodine (Küpper, 2015; Küpper et al., 2008; Nitschke & Stengel, 2015; Ye et al., 2015) and are widely harvested for food and exploited for commercial alginate and iodine. Iodine is an essential nutrient required for the synthesis of thyroid hormones (THs), triiodothyronine (T3) and thyroxine (T4) (Berg

et al., 2017). Iodine deficiency in humans can result in unexpected health problems such as

hypothyroidism and goiter, whereas high iodine intake from seaweeds can lead to

hyperthyroidism, a reversible condition which can cause symptoms such as nontoxic or

diffuse nodular goiter, latent Graves' disease and long standing iodine deficiency (Leung &

Braverman, 2014). Natural consumers of seaweeds such as fish and shellfish are also a rich

dietary source of iodine for humans (Nitschke & Stengel, 2015). It is therefore essential to

understand how the iodine content of seafood will change under global climate change. This

information can for instance be used by the World Health Organization (WHO) to provide

recommendations on appropriate levels of seaweeds consumption to maintain a sufficient

daily iodine intake (Fig. 1).

With respect to iodine, kelp (order Laminariales) are particularly important as they are not

only being the greatest iodine accumulators among living organisms, but also playing an

important role in the global biogeochemical cycle of iodine (Küpper, 2015; Küpper et al.,

2008). In *Laminaria* tissue, iodine is mostly stored as iodide on the thallus surface and in the

apoplast. Iodide efflux occurs under oxidative stress and iodide in the peripheral tissues acts

as an inorganic antioxidant to detoxify both aqueous oxidants and ozone, stimulating the

release of molecular iodine and volatile iodinated compounds to the atmosphere (Cosse et al.,

2009; Küpper & Kroneck, 2014). Oxidative burst and associated iodine metabolism also play

a direct defensive role in both controlling the growth of potentially pathogenic bacteria living

at the thallus surface and scavenging a variety of reactive oxygen species (ROS) (Küpper et

al., 2008; Küpper, Müller, Peters, Kloareg, & Potin, 2002; Strittmatter et al., 2016). During

oxidative burst, algal cells rapidly release large amounts of activated oxygen species (AOS), such as superoxide (O_2^-), hydrogen peroxide (H_2O_2) or hydroxyl radicals (OH^-). This release is elicited by exposure to oligomeric degradation products of alginate and possibly other molecular signals. It has been suggested that the mechanism of iodine antioxidation is linked to the production of vanadium-dependent haloperoxidases (vHPOs), which comprises seventeen vanadium-dependent bromoperoxidases (vBPOs) and fifty-nine iodoperoxidases (vIPOs) in the *S. japonica* genome (Butler & Carter-Franklin, 2004; Cosse et al., 2009; Ye et al., 2015) (Fig. 1).

Despite widespread interest in the biological response of kelp to climate change, there is currently no information on the *in situ* molecular response associated with iodine metabolism. Here, three ecologically and socio-economically important kelp species (*S. japonica*, *U. pinnatifida*, and *M. pyrifera*), as well as another four coastal seaweeds (*Ulva pertusa* and *Ulva intestinalis* in Chlorophyta; *Gracilaria lemaneiformis* and *Gracilaria chouae* in Rhodophyta) were used to study the effect of increasing pCO$_2$ on iodine accumulation of seaweeds using both short-term laboratory experiments and one long-term, *in situ* experiment. In a feeding experiment, the transfer of iodine between *S. japonica* and its consumer *Haliotis discus* was investigated. We used these experiments to explore: (i) the effect of elevated pCO$_2$ on iodine accumulation in kelp-based coastal food webs; (ii) whether short-term laboratory experiments can be compared with *in situ* ocean mesocosms; (iii) the molecular mechanism involved in iodine metabolism at the transcriptional and protein level in response to elevated pCO$_2$. Together, the research presented here will be used to assess the global biogeochemical cycle of iodine under future climate change and provide information

for making recommendations on appropriate levels of seaweeds consumption to reach adequate daily iodine intake.

MATERIALS AND METHODS

Algal material and culture conditions

For the laboratory experiments, young sporophytes of *S. japonica* were collected from semi-enclosed Sungo Bay, located on the northwestern coast of the Yellow Sea, China (37°01′–37°09′ N, 122°24′–122°35′ E) in December 2016, when the mean seawater temperature was 10.3°C. Similar sized algal samples (average length is 10 cm) were transported back to the laboratory within three hours, in a tank of cold seawater. In the laboratory, the intact samples were washed with sterile seawater until they were free from visible epiphytes and then pre-cultured in aquaria supplemented with f/2 medium (Guillard, 1975) at 10 ± 1°C with vigorous air bubbling for 2 days before the start of the experiment. The lighting conditions were set at 100 µmol photons m^{-2} s^{-1} supplied by white fluorescent lamps, with a photoperiod of 12 h light and 12 h darkness.

Effect of increasing $p\text{CO}_2$ and temperature on iodine accumulation

To study the individual effect of temperature on iodine accumulation in *S. japonica* sporophytes, a gradient of five temperatures (5°C, 10°C, 15°C, 20°C, and 23°C) was established, mimicking the annual variation of temperature during the growth period. To

examine the combined effect of increasing pCO_2 and temperature, three temperatures (10°C, 15°C, 20°C) and five pCO_2 levels (400 μatm, 700 μatm, 1,000 μatm, 1,500 μatm, and 2,000 μatm), were selected (Table S1). These pCO_2 levels were chosen as they reflect current and future pCO_2 levels up to the year 2,300 under IPCC (Intergovernmental Panel on Climate Change) scenario RCP 8.5. The experiment was conducted in flasks using three biological replicates per treatment. Before experiments, algal samples were pre-cultured in seawater under various pCO_2 and temperature conditions for 96 hours. Pre-cultured algae were then inoculated into 500-mL Erlenmeyer flasks containing 400 mL of adjusted f/2 seawater medium and supplemented with 20 μmol L^{-1} KI (half saturation concentration derived from iodine uptake kinetics, in Fig. S1) for 24 h. At the end of the experiment, algal samples in each treatment were collected and rinsed for iodine determination within 24 h. Individual flasks were cultured inside a CO_2 chamber (HP1000G-D, China), programmed to supply 400 μatm, 700 μatm, 1,000 μatm, 1,500 μatm, or 2,000 μatm pCO_2 by bubbling at each designated temperature for 24 h. The pH and temperature were measured at the beginning and end of the experiment with a pH meter (Orion ROSS, Fisher Scientific Instruments). To determine total alkalinity (TA) at each treatment, 20 ml of culture medium was filtered with GF/F membrane and measured using an 848 Titrino plus automatic titrator (Metrohm, Riverview, FL, USA). Chemical carbonate system parameters were calculated using the CO2SYS Package in MS Excel (Pierrot, Lewis, & Wallace, 2006) based on pH, temperature, salinity, and TA (Table S1).

Potential antioxidant property of iodide in *S. japonica*

To determine the potential antioxidant properties of iodide in *S. japonica*, oligoguluronate elicitor (GG, Shanghai Zzbio Co, Ltd, China) was used as exogenous defense elicitors to induce oxidative stress and iodine efflux from algal tissue into the culture medium (referred to previous study of Küpper, Kloareg, Guern, & Potin, 2001). Algal samples were inoculated into 150-mL Erlenmeyer flasks containing 100 mL sterile seawater without (set as control) or with exogenous oligoguluronate elicitors at a final concentration of 100 μg ml^{-1} (GG, set as treatment). Three elicitation experiments were conducted at 10°C and 100 μmol photons m^{-2} s^{-1}. To understand the effect of GG on iodine efflux over time, three seawater samples were randomly taken from the untreated control and GG treatments at four time intervals (0 h, 1 h, 3 h, and 5 h). To understand the effect of elevated pCO$_2$ in the presence of GG, iodine efflux was measured after 3 hours under five pCO$_2$ conditions (400 μatm, 700 μatm, 1,000 μatm, 1,500 μatm, or 2,000 μatm, bubbled at 10°C for 24 h). To investigate the prolonged effect of ocean acidification on iodine efflux, the short-term vs long-term response of iodine efflux in *S. japonica* under ocean acidification conditions was compared. Under short-term-ambient-carbon (SAC) and short-term-elevated-carbon (SEC) treatments, algae were pre-cultured at CO$_2$ concentrations of 400 μatm and 1,000 μatm respectively, for 7 days. Under long-term-ambient-carbon (LAC) or long-term-elevated-carbon (LEC) treatments, algae were pre-cultured under the same pCO$_2$ levels for 30 days. Following pre-culturing, iodine efflux under ambient (400 μatm) or elevated (1,000 μatm) pCO$_2$ was measured after 3 hours.

***In situ* mesocosm experiments**

Three kelp species, *S. japonica*, *U. pinnatifida*, and *M. pyrifera*, as well as other coastal

seaweeds (*Ulva pertusa* and *Ulva intestinalis* in Chlorophyta; *Gracilaria lemaneiformis* and

Gracilaria chouae in Rhodophyta) were collected off the coastline of Sungo Bay for the

mesocosm experiment (Fig. S2). With the exception of *M. pyrifera*, these algae are widely

consumed by humans. *M. pyrifera* was selected as it is a preferred food source of marine

invertebrates, such as sea urchins and abalone, which are harvested by the local fishing

industry. The mesocosms were designed following (Xu et al., 2017). Six net cages (three per

treatment) were used, applying two pCO_2 levels, ambient pCO_2 (400 μatm, bubbled with air)

and elevated pCO_2 (1,000 μatm, bubbled with air/CO_2 premixed gas using a CO_2 Enrichlor,

CE-100B; Wuhan Ruihua Instrument & 25 Equipment Ltd) (Fig. S3). One hundred and

twenty individuals of each algal species of similar size were grown in each cage for

approximately five months, from December 2016 to May 2017. During this period, six

individuals of each algal species from each net cage (n=6) were randomly collected for iodine

determination, and 10 individuals (n=10) from each net cage were selected and weighed to

monitor growth. To determine the effect of kelp size on iodine accumulation during the

culture period, kelp which were pre-acclimated under ambient or elevated pCO_2 for 48 hours

in the sea were sampled at different time points: 0 day, 30 days, 60 days, and 90 days for *S.

japonica*, 0 day, 30 days, and 60 days for *U. pinnatifida*, and 0 day, 20 days, and 40 days for

M. pyrifera. Seawater carbonate chemistry in the ambient (400 μatm) or elevated (1,000

μatm) pCO_2 culture environments were monitored every two days and the carbonate

chemistry was calculated using the CO2SYS Package in MS Excel based on pH, temperature,

salinity, and TA (Fig. S4) (for further details of the mesocosm design, see methods in supporting information).

To assess the effects of consumption of *S. japonica* by abalone, sporophytes were grown at either ambient (400 μatm) or elevated (1,000 μatm) pCO_2. Five feeding experiments were conducted *in situ*: 1) abalones were fed with ambient pCO_2-cultured *S. japonica* and were also cultured in an ambient pCO_2-cage; 2) abalones were fed with ambient pCO_2-cultured *S. japonica* and were cultured in an elevated pCO_2-cage; 3) abalones were fed with elevated pCO_2-cultured *S. japonica* and were also cultured in an elevated pCO_2-cage; 4) abalones were fed with elevated pCO_2-cultured *S. japonica* and were cultured in an ambient pCO_2-cage; 5) abalones were fed with ambient pCO_2-cultured *S. japonica* and were cultured in the sea with no cage; 6) abalones were fed with elevated pCO_2-cultured *S. japonica* and were cultured in the sea with no cage. The experiments, with six replicates (n = 6), were conducted independently for 30 days, starting on April 10th 2017. The experimental abalones were fed with fresh algal tissue at intervals of 3 days and collected for iodine determination at intervals of 10 days. At the end of the experiments, the abalones were cleaned with sterile seawater and the tissue was removed quickly and entirely from the shell, weighed and frozen in liquid nitrogen, and stored at -80 °C for total iodine and thyroid hormones (THs) determination (see methods in supporting information).

Transcriptomic and proteomic analysis of iodine metabolism in *S. japonica*

To explore the effect of ocean acidification on the iodine metabolism on the *S. japonica* transcriptome, on day 90 of the mesocosm experiment, *S. japonica* sporophytes were randomly selected, cleaned with sterile seawater, frozen in liquid nitrogen and stored at -80 °C for subsequent transcriptomic (with three biological replicates for each pCO$_2$) and proteomic (with two biological replicates for each pCO$_2$) analysis (see methods in supporting information). The transcriptome data have been deposited in the Sequence Read Archive (SRA) at the National Center for Biotechnology Information (NCBI) and are accessible through accession number (SRP127301); the mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium via the PRIDE (Vizcaíno et al., 2016) partner repository with the dataset identifier PXD011219.

Statistical analysis

Statistical models were carried out with R software (R Core Development Team, 2014) and model selection was based by Akaike Information Criterion (AIC). The effect of temperature and the combined effect of increasing pCO$_2$ and temperature on iodine accumulation were analyzed using a mixed effects model using the lmer function within the package lme4 and lmerTest. When modelling the effect of temperature alone, temperature was used as a fixed factor, when modelling the combined effect of temperature and pCO$_2$, both temperature and pCO$_2$ were used as fixed factors (the interaction between temperature and pCO$_2$ was dropped

from the model based on AIC). The effect of measurement type, either iodide or total iodine accumulation, were used as random factors in both models.

Using the lm function within R, three liner models were used to compare changes in iodine efflux in *S. japonica* treated with oligoguluronates (GG) or untreated (control). To test for the effect of time (hours) on iodine efflux, time, treatment (GG or control), and the interaction between time and treatment were included in a linear model. The effect of increasing pCO_2 (400 to 2000 µatm), treatment (GG or control), and the interaction between pCO_2 and treatment were included in a second linear model. The effect of ambient and elevated pCO_2, treatment (GG or control), time (days), the interaction between pCO_2 and time, and the interaction between treatment and pCO_2 were included in a third linear model.

In order to compare the responses of the three species of seaweeds used in the mesocosm experiment (*S. japonica*, *M. pyrifera* and *U. pinnatifida*), the relative change in iodine accumulation and weight (g) over time were calculated, relative to the responses at time point zero, under control conditions (400 µatm). To test for the effect of pCO_2 (ambient or elevated), over time (days), on relative changes in iodine accumulation, the effect of pCO_2, time, species and the interactions between time and species were included in a linear model. To test for the effect of pCO_2, over time, on the relative weight of the seaweeds, the effect of pCO_2, time, species and the interaction between pCO_2, time and species were included in a linear model.

Three linear models were used to test the effect of diet and $p\text{CO}_2$ on 1) iodine enrichment 2) T3 concentration and 3) T4 concentration in the abalone *(H. discus)*. For iodine enrichment, the effect of diet, time, $p\text{CO}_2$, the interaction between diet, time and $p\text{CO}_2$, and environment (caged or non-caged) were included in the model. For both T3 and T4 hormones, the effect diet, $p\text{CO}_2$, environment (caged or non-caged) and the interaction between environment and diet were included in the model. Data collected for iodine enrichment and T3 concentration best fit a lognormal distribution and for this reason a generalized linear model which takes into consideration a lognormal distribution was carried out, using the glm function in the stats package.

RESULTS

Effect of increasing $p\text{CO}_2$ and temperature on iodine accumulation of *S. japonica* in laboratory

Elevated $p\text{CO}_2$ caused changes in iodine accumulation in *S. japonica* when both the $p\text{CO}_2$ and temperature were altered at the same time (Fig. 2a; $F_{1,86} = 14.515$, $P = 0.0003$), with no significant effect of temperature on iodine accumulation (Fig. 2a; $F_{1,86} = 1.507$, $P = 0.223$). As $p\text{CO}_2$ increased from 400 µatm to 1,000 µatm at the control temperature (10 °C), both iodide and total iodine increased and reached the highest observed iodine accumulation at 1,000 µatm. As $p\text{CO}_2$ increased to 1,500 µatm, iodide and total iodine accumulation decreased and leveled off as $p\text{CO}_2$ increased to 2,000 µatm. When temperature was altered in isolation, there was a significant effect on iodide and total iodine accumulation in *S. japonica*

(Fig. 2b; $F_{1, 27} = 19.47$, $P = 0.0002$). As temperature increased, from 5 °C to 15 °C under the control pCO$_2$ concentration (400 µatm), iodide and total iodine accumulation increased by 64% and 58%, respectively (Fig. 2b). At 20 °C and 23 °C, iodine accumulation decreased slightly but it was not significantly different from the maximum iodine accumulation observed at 15 °C. However, there were no significant effects of increasing pCO$_2$ or temperature (isolation or combination) on iodine solubility and availability in seawater medium (Fig. S5).

Potential antioxidant properties of iodide in *S. japonica* in laboratory

Under oxidative stress, induced by the addition of oligoguluronates (GG), iodine efflux in *S. japonica* was 61% greater than the untreated control group, after 5 hours (Fig. 3a; $F_{1,225} = 22.428$, $P = 0.0001$). Iodine efflux in *S. japonica* treated with GG increased over three hours and then decreased slightly after five hours, with little change in the untreated control group over time (Fig. 3a; *effect of time on the iodine efflux,* $F_{1,47} = 4.685$, $P = 0.0427$) and is supported by a significant interaction between the treatment (GG and control) and time (Fig. 3a; $F_{1,20} = 4.562$, $P = 0.0452$).

Iodine efflux in *S. japonica* increased with increasing pCO$_2$ until 1500 µatm and decreased again at 2000 µatm (Fig. 3b; $F_{1,10} = 2.95$, $P = 0.098$). Iodine efflux was 51% - 72% higher in *S. japonica* treated with GG than the untreated group, depending on the pCO$_2$ levels (Fig. 3b;

$F_{1,475} = 148.375$, P < 0.0001), and this is supported by a significant interaction between the

pCO_2 conditions and treatment (GG and untreated) (Fig. 3b; $F_{1,17} = 5.173$, $P = 0.031$).

Long-term exposure to elevated pCO_2 (LEC) intensified iodine efflux from *S. japonica.*

Iodine efflux from *S. japonica* treated with GG was 29% higher after 30 days exposure to

elevated pCO_2, compared with 7 days of exposure to elevated pCO_2 (SEC) and 44% greater

than both short- and long-term exposure to ambient pCO_2 conditions (7 days and 30 days;

SAC and LAC) (Fig. 3c; $F_{1,48} = 21.119$, $P = 0.0002$). Very little change in iodine efflux was

observed in the untreated control group (Fig. 3c; $F_{1,18} = 107.3027$, P < 0.0001). This is

supported by significant interactions between pCO_2 and time (Fig. 3c; $F_{1,12.08} = 5.812$, $P =$

0.0268) and pCO_2 and treatment (GG and untreated) (Fig. 3c; $F_{1,28.99} = 5.812$, $P = 0.0268$).

**Effect of increasing pCO_2 on iodine accumulation of seaweeds using *in situ* mesocosm
experiments**

In the mesocosm experiment, elevated pCO_2 increased tissue growth (relative changes in

fresh weight) of all three kelp species compared to ambient pCO_2 (Fig. 4 a, b and c, *top

panels*; $F_{1, 828} = 332.087$, $P < 0.0001$). This effect was consistent over time, which is

supported by a significant interaction between time and pCO_2 conditions (Fig. 4 a, b and c,

top panels; $F_{1, 828} = 175.387$, $P < 0.0001$). Changes in relative weight were significantly

different between the three species (Fig. 4 a, b and c, *top panels*; $F_{2, 828} = 504.273$, $P <$

0.0001), including significant interactions between the species of seaweeds and time (Fig. 4

a, b and c, *top panels*; $F_{2,828} = 38.434$, $P < 0.0001$), and significant interactions between pCO$_2$ conditions, time and species of seaweeds (Fig. 4 a, b and c, *top panels*; $F_{2,828} = 28.354$, $P < 0.0001$). *M. pyrifera* showed the largest increase in weight under elevated pCO$_2$ at day 40, compared with *S. japonica* at day 90 and *U. pinnatifida* at day 60. *S. japonica* and *U. pinnatifida* showed similar changes in weight under ambient and elevated pCO$_2$ until day 30, thereafter *U. pinnatifida* showed larger increases in weight than *S. japonica* by day 60 (Fig. 4a).

Iodine accumulation increased in the three seaweeds under elevated pCO$_2$ compared to ambient pCO$_2$ (Fig. 4 a, b and c, *bottom panels*; $F_{1,353} = 36.362$, $P < 0.0001$). The highest iodine accumulation was observed at time zero, thereafter relative iodine accumulation declined with time (Fig. 4 a, b and c, *bottom panels*; $F_{1,353} = 791.782$, $P < 0.0001$) however, iodine accumulation declined more quickly under ambient pCO$_2$ conditions compared to elevated pCO$_2$. The three seaweeds species responded with different intensities to changes in pCO$_2$ (Fig. 4 a, b and c, *bottom panels*; $F_{2,353} = 16.322$, $P < 0.0001$). *S. japonica* showed the highest iodine accumulation at time zero, both under ambient pCO$_2$ (7.91 mg g^{-1} dw $\pm$ 0.13 SE) and elevated pCO$_2$ (7.87 mg g^{-1} dw $\pm$ 0.19 SE), but it also had the greatest decline in iodine accumulation over 90 days (total decline of 85% and 73% for ambient and elevated pCO$_2$ respectively). Compared to *S. japonica, U. pinnatifida* and *M. pyrifera* showed modest iodine accumulation at time zero. Iodine accumulation declined under both ambient and elevated pCO$_2$ by 58% and 26% respectively, over 60 days in *U. pinnatifida* (Fig. 4b) and 63% and 56%, over 40 days in *M. pyrifera* (Fig. 4c). This variation in iodine accumulation

over time is supported by a significant interaction between species of seaweeds and time (Fig. 4 a, b and c, *bottom panels*; $F_{2,353} = 16.553$, $P < 0.0001$). In addition to kelp species (Fig. S6), we also found similar responses to elevated pCO_2 in other seaweeds species, where both algal biomass and iodine accumulation increased with the elevated pCO_2 (Fig. S7).

Iodine transfer between *S. japonica* and its consumer *H. discus* in a mesocosm experiment

In the feeding experiment, abalone fed with *S. japonica*, cultured under elevated pCO_2 (1000 µatm) had a higher iodine accumulation than those fed on seaweeds cultured under ambient pCO_2 (400 µatm) (Fig. 5a, $F_{1,106} = 155.24$, $P < 0.0001$). In addition, iodine accumulation in the abalone was stimulated when the feeding experiments were conducted under elevated pCO_2 (1000 µatm) (Fig. 5a, $F_{1,106} = 29.53$, $P < 0.0001$). Iodine accumulation in all experimental abalone increased over the 30 days of cultivation (Fig. 5a, $F_{1,105} = 1003.48$, $P < 0.0001$), and abalone fed with seaweeds grown under high pCO_2 and under elevated pCO_2 conditions showed the highest amount of iodine accumulation (Fig. 5a; *interaction between diet, time and pCO_2*, $F_{1,102} = 26.29$, $P < 0.0001$). There was no difference in iodine accumulation between feeding experiments conducted in a cage or in the field with no cage (Fig. 5a, $F_{1,103} = 1.31$, $P = 0.255$).

It was found that both T3 (Fig. 5b; $F_{1,33} = 831.98$, $P < 0.0001$) and T4 (Fig. 5b; $F_{1,33} = 482.77$, $P < 0.0001$) hormone concentrations declined when the abalone were fed with *S. japonica* cultured under elevated pCO_2, regardless of the pCO_2 conditions of the feeding experiment. Whilst there was a decline in thyroid concentration under elevated pCO_2, for both T3 (Fig. 5b; $F_{1,34} = 143.29$, $P < 0.0001$) and T4 concentrations ($F_{1,34} = 183.36$, $P < 0.0001$), the largest drop in thyroid concentration occurred when abalone were fed with algae cultured under high pCO_2 and under elevated pCO_2 conditions. Linear regression analysis further demonstrated that dietary inclusion of *S. jap*onica with higher iodine content caused an increase in iodine intake into the abalone (Fig. S8a) and resulted in lower concentration of T3 (Fig. S8b) and T4 (Fig. S8c) hormones. Moreover, results also revealed that there were significant negative relationships between iodine concentration and T3 (Fig. S9a) and T4 content (Fig. S9b). Overall, THs concentrations were higher in the field with no cage for both T3 (Fig. 5b; $F_{1,32} = 324.30$, $P < 0.0001$) and T4 (Fig. 5b; $F_{1,32} = 284.49$, $P < 0.0001$) hormones. However, THs concentration drop in field conditions when abalone were fed with algae cultured under elevated pCO_2, and this is supported by a significant interaction between the pCO_2 level that dietary seaweeds were cultured under and the conditions of the feeding experiment (i.e. caged or not caged), for both T3 concentrations (Fig. 5b; $F_{1,31} = 23.93$, $P < 0.0001$) and T4 concentrations (Fig. 5b; $F_{1,31} = 47.16$, $P < 0.0001$).

In situ transcriptional or proteomic response of _S. japonica_ to elevated pCO$_2$

Transcriptome data showed that 11 _vIPO_ (vanadium-dependent iodoperoxidases) and 12 _vBPO_ (vanadium-dependent bromoperoxidases) UniGenes were significantly differentially expressed between ambient and elevated pCO$_2$ of culture environments (Fig. 6a). Among these genes, 8 _vIPOs_ and 8 _vBPOs_ were down-regulated, whereas 3 _vIPOs_ and 4 _vBPOs_ were up-regulated under elevated pCO$_2$ in the mesocosms (Table S2, S3). The variation trend was verified for 8 genes by RT-PCR (Fig. S10). In the proteomic data, it was found that four vHPOs proteins were down-regulated at elevated pCO$_2$ (Fig. 6b). Therein, the trend of relative expression of two vBPOs (SJ19768 and SJ19850) was consistent with those in the transcriptome. In contrast, vIPOs (SJ00628) and vBPOs (SJ10798) did not significantly change at the transcriptomic level.

DISCUSSION

Brown algae are predicted to benefit from elevated pCO$_2$ under global change (Enochs et al., 2015; Johnson, Russell, Fabricius, Brownlee, & Hall-Spencer, 2012; Linares et al., 2015; Porzio, Buia, & Hall-Spencer, 2011; Xu et al., 2017) and therefore will thrive in natural pCO$_2$ rich environments (Enochs et al., 2015; Johnson, Russell, Fabricius, Brownlee, & Hall-Spencer, 2012; Linares et al., 2015; Porzio, Buia, & Hall-Spencer, 2011). It is supported by data that tissue growth of _S. japonica_ (Fig. 2 and Fig. 4a), _U. pinnatifida_ (Fig. 4b), and _M. pyrifera_ (Fig. 4c), and photosynthesis of _S. japonica_ (Fig. S11) all increased under elevated CO$_2$ conditions. However, it was unknown how enhanced growth of kelp, under conditions of

ocean acidification, affect the accumulation and transfer of iodine across coastal marine food webs. Increasing pCO$_2$ will not change the solubility and availability of inorganic iodine in seawater (Fig. S5). However, the data presented here reveal for the first time that increasing pCO$_2$ causes iodine accumulation in coastal seaweeds and leads to the accumulation of iodine in consumers of kelp, which potentially will result in elevated iodine intake of consumers of kelp if pCO$_2$ continues to rise (1,000 µatm in 2100 under RCP 8.5) as predicted by climate models (Fig. 1).

In laboratory experiment, we found that at pCO$_2$ greater than 400 µatm (the current pCO$_2$ value), changes in temperature had no effect on iodine accumulation and predictions of iodine accumulation in *S. japonica* could be made based on changes in pCO$_2$ alone (Fig. 2). It is suggested that this is because pCO$_2$ is the dominant environmental driver when pCO$_2$ and temperature change simultaneously. This finding is supported by Brennan and Collins (2015) who demonstrated that changes in the growth rate of the microalgae *Chlamydomonas reinhardtii* could be predicted using the dominant environmental driver in test environments with up to eight environmental drivers (Brennan & Collins, 2015). Global change will involve many simultaneous environmental changes, such as changes in CO$_2$, pH, temperature, and nutrient (Boyd, Lennartz, Glover, & Doney, 2015; Gruber, 2011; Hutchins & Fu, 2017). Results presented here highlight the importance of measuring the response to multiple environmental drivers in order to understand how they interact. Predictions on the combined effects of environmental drivers using traditional additive or multiplicative models

could potentially overestimate the effects of ocean warming and ocean acidification on iodine accumulation (Fig. S12).

Elevated iodine accumulation in *S. japonica* under ocean acidification was associated with down-expression regulation of vHPOs (vIPOs and vBPOs), which has never been observed before in any alga (Fig. 6). While vHPOs specific physiological role is still unclear, these genes have previously been studied for their role in the oxidative stress response in brown algae including, *S. japonica* (Ye et al., 2015) and *Laminaria digitata* (Cosse et al., 2009). In addition, the physiological antioxidant role of iodine in algae is suggested to be linked to the presence of particular *vHPO*s (Colin et al., 2003, 2005). Under oxidative stress, induced by elicitor-triggered oxidative burst, iodine efflux increased in *S. japonica*, even under high pCO_2 (Fig. 3), supporting the physiological role of iodine in algae. In some cases, the iodine efflux was up to twice as high as in the control group (Fig. 3a). Thus, iodine efflux is probably an efficient strategy to cope with oxidative stress (Küpper et al., 2008). Furthermore, the oligoalginate-triggered oxidative burst and associated iodide release is also known to be an efficient strategy against infection, suggesting that under ocean acidification kelp may be able to cope better with infections as they have more stored iodine (Küpper et al., 2008; Küpper, Müller, Peters, Kloareg, & Potin, 2002).

Elevated iodine content in kelp has the potential to be transferred to their consumers such as abalone. *H. discus* fed with algae cultured under elevated CO_2 showed higher levels of iodine accumulations and reduced levels of T3 and T4 hormones compared to animals fed with algae cultured under ambient CO_2 (Fig. 5). Iodine is essential for the synthesis of thyroid hormones (THs), which play important roles in development, metamorphosis and metabolism of vertebrates and invertebrates, including humans (Bath et al., 2017; Huang et al., 2015). Previous studies have demonstrated that sea urchin larvae could potentially receive TH precursors or the active hormones from some of the microalgal species that they consumed (Heyland & Moroz, 2006). Whilst, few data were available on the effect of iodine intake on the THs of invertebrates, some studies on humans demonstrate that high-dose kelp supplementation significantly decreased the total triiodothyronine levels of subjects (Eliason, 1998). The increased iodine transfer and reduced thyroid hormones could potentially impact nutrient composition of costal primary producers as well as in kelp-based coastal food webs.

Our results suggest that projected pCO_2 (1,000 μatm in 2100 under RCP 8.5) will increase the growth of coastal seaweeds and their iodine accumulation, with the potential for higher levels of iodine to be transferred from seaweeds to consumers. Thus, there is a potential risk of iodine overconsumption in consumers of kelps if pCO_2 increases as projected for the coming decades. Furthermore, biogeochemical cycling of iodine in coastal ecosystems might change as a consequence of enhanced accumulation in coastal marine food webs. This may even have consequences for coastal non-marine habitats as volatile iodinated compounds can be released to the atmosphere (Cosse, Potin, & Leblanc, 2009; Küpper & Kroneck, 2014).

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REFERENCES

Bath, S. C., Hill, S., Infante, H. G., Elghul, S., Nezianya, C. J., & Rayman, M. P. (2017).
Iodine concentration of milk-alternative drinks available in the UK in comparison with
cows' milk. *British Journal of Nutrition*, *118*, 525–532.

Berg, V., Nøst, T. H., Skeie, G., Thomassen, Y., Berlinger, B., Veyhe, A. S., … Hansen, S.
(2017). Thyroid homeostasis in mother–child pairs in relation to maternal iodine status:
the MISA study. *European Journal of Clinical Nutrition*, *71*, 1002–1007.

Bloom, A. J., Burger, M., Asensio, J. S. R., & Cousins, A. B. (2010). Carbon dioxide
enrichment inhibits nitrate assimilation in wheat and *Arabidopsis*. *Science*, *328*,
899–903.

Boyd, P. W., Lennartz, S. T., Glover, D. M., & Doney, S. C. (2015). Biological ramifications
of climate-change-mediated oceanic multi-stressors. *Nature Climate Change*, *5*, 71.

Brennan, G., & Collins, S. (2015). Growth responses of a green alga to multiple
environmental drivers. *Nature Climate Change*, *5*, 892.

Butler, A., & Carter-Franklin, J. N. (2004). The role of vanadium bromoperoxidase in the
biosynthesis of halogenated marine natural products. *Natural Product Reports*, *21*,
180–188.

Colin, C., Leblanc, C., Michel, G., Wagner, E., Leize-Wagner, E., Van Dorsselaer, A., &
Potin, P. (2005). Vanadium-dependent iodoperoxidases in *Laminaria digitata*, a novel

biochemical function diverging from brown algal bromoperoxidases. *JBIC Journal of Biological Inorganic Chemistry*, *10*, 156–166.

Colin, C., Leblanc, C., Wagner, E., Delage, L., Leize-Wagner, E., Van Dorsselaer, A., ... & Potin, P. (2003). The brown algal kelp *Laminaria digitata* features distinct bromoperoxidase and iodoperoxidase activities. *Journal of Biological Chemistry*, *278*, 23545.

Cosse, A., Potin, P., & Leblanc, C. (2009). Patterns of gene expression induced by oligoguluronates reveal conserved and environment-specific molecular defense responses in the brown alga *Laminaria digitata*. *New Phytologist*, *182*, 239–250.

Dillon, M. E., Wang, G., & Huey, R. B. (2010). Global metabolic impacts of recent climate warming. *Nature*, *467*, 704.

Eliason, B. C. (1998). Transient hyperthyroidism in a patient taking dietary supplements containing kelp. *The Journal of the American Board of Family Practice*, *11*, 478–480.

Enochs, I. C., Manzello, D. P., Donham, E. M., Kolodziej, G., Okano, R., Johnston, L., ... & Valentino, L. (2015). Shift from coral to macroalgae dominance on a volcanically acidified reef. *Nature Climate Change*, *5*, 1083.

Gruber, N. (2011). Warming up, turning sour, losing breath: ocean biogeochemistry under global change. *Philosophical Transactions of the Royal Society of London A: Mathematical, Physical and Engineering Sciences*, *369*, 1980–1996.

Guillard, R. R. L. (1975). Culture of phytoplankton for feeding marine invertebrates. In W. L.

Smith & M. H. Chanley (Eds.), *Culture of marine invertebrate animals* (pp. 26–60).

New York, NY: Plenum Press.

Heyland, A., & Moroz, L. L. (2006). Signaling mechanisms underlying metamorphic

transitions in animals. *Integrative and Comparative Biology*, *46*, 743–759.

Huang, W., Xu, F., Qu, T., Zhang, R., Li, L., Que, H., & Zhang, G. (2015). Identification of

thyroid hormones and functional characterization of thyroid hormone receptor in the

pacific oyster *Crassostrea gigas* provide insight into evolution of the thyroid hormone

system. *PloS One*, *10*, e0144991.

Hutchins, D. A., & Fu, F. (2017). Microorganisms and ocean global change. *Nature

Microbiology*, *2*, 17058.

Jin, P., Wang, T., Liu, N., Dupont, S., Beardall, J., Boyd, P. W., ... & Gao, K. (2015). Ocean

acidification increases the accumulation of toxic phenolic compounds across trophic

levels. *Nature Communications*, *6*, 8714.

Johnson, V. R., Russell, B. D., Fabricius, K. E., Brownlee, C., & Hall-Spencer, J. M. (2012).

Temperate and tropical brown macroalgae thrive, despite decalcification, along natural

CO_2 gradients. *Global Change Biology*, *18*, 2792–2803.

Küpper, F. C. (2015). Iodine in Seaweeds–Two Centuries of Research. In Springer Handbook

of Marine Biotechnology (pp. 591-596). Springer, Berlin, Heidelberg.

Küpper, F. C., & Kroneck, P. M. (2014). Iodine bioinorganic chemistry: physiology, structures, and mechanisms. *Iodine Chemistry and Applications*, 555–589.

Küpper, F. C., Carpenter, L. J., McFiggans, G. B., Palmer, C. J., Waite, T. J., Boneberg, E. M., ... & Potin, P. (2008). Iodide accumulation provides kelp with an inorganic antioxidant impacting atmospheric chemistry. *Proceedings of the National Academy of Sciences*, *105*, 6954–6958.

Küpper, F. C., Kloareg, B., Guern, J., & Potin, P. (2001). Oligoguluronates elicit an oxidative burst in the brown algal kelp *Laminaria digitata*. *Plant Physiology*, 125, 278–291.

Küpper, F. C., Müller, D. G., Peters, A. F., Kloareg, B., & Potin, P. (2002). Oligoalginate recognition and oxidative burst play a key role in natural and induced resistance of sporophytes of Laminariales. *Journal of Chemical Ecology*, *28*, 2057–2081.

Leung, A. M., & Braverman, L. E. (2014). Consequences of excess iodine. *Nature Reviews Endocrinology*, *10*, 136.

Linares, C., Vidal, M., Canals, M., Kersting, D. K., Amblas, D., Aspillaga, E., ... & Hereu, B. (2015). Persistent natural acidification drives major distribution shifts in marine benthic ecosystems. *Proc. R. Soc. B*, *282*, 20150587.

Loladze, I. (2002). Rising atmospheric CO_2 and human nutrition: toward globally imbalanced plant stoichiometry?. *Trends in Ecology & Evolution*, *17*, 457–461.

Martínez-Botí, M. A., Foster, G. L., Chalk, T. B., Rohling, E. J., Sexton, P. F., Lunt, D. J., ...
& Schmidt, D. N. (2015). Plio-Pleistocene climate sensitivity evaluated using
high-resolution CO_2 records. *Nature*, *518*, 49.

McKibben, S. M., Peterson, W., Wood, A. M., Trainer, V. L., Hunter, M., & White, A. E.
(2017). Climatic regulation of the neurotoxin domoic acid. *Proceedings of the National
Academy of Sciences*, *114*, 239–244.

Myers, S. S., Smith, M. R., Guth, S., Golden, C. D., Vaitla, B., Mueller, N. D., ... & Huybers,
P. (2017). Climate change and global food systems: potential impacts on food security
and undernutrition. *Annual Review of Public Health*, *38*, 259–277.

Nitschke, U., & Stengel, D. B. (2015). A new HPLC method for the detection of iodine
applied to natural samples of edible seaweeds and commercial seaweed food products.
Food Chemistry, *172*, 326–334.

Phalkey, R. K., Aranda-Jan, C., Marx, S., Höfle, B., & Sauerborn, R. (2015). Systematic
review of current efforts to quantify the impacts of climate change on undernutrition.
Proceedings of the National Academy of Sciences, *112*, 4522–4529.

Pierrot, D. E., Lewis, E., & Wallace, D. W. R. (2006). MS Excel program developed for CO_2
system calculations. Oak Ridge, TN: Carbon Dioxide Information Analysis Center, Oak
Ridge National Laboratory, U.S. Department of Energy. doi:
10.3334/CDIAC/otg.CO2SYS_XLS_CDIAC105a

Porzio, L., Buia, M. C., & Hall-Spencer, J. M. (2011). Effects of ocean acidification on macroalgal communities. *Journal of Experimental Marine Biology and Ecology*, *400*, 278–287.

R Core Development Team. (2014). R: A language and environment for statistical computing: R Foundation for Statistical Computing. http://www.R-project.org/.

Strittmatter, M., Grenville-Briggs, L. J., Breithut, L., Van West, P., Gachon, C. M., & Küpper, F. C. (2016). Infection of the brown alga *Ectocarpus siliculosus* by the oomycete *Eurychasma dicksonii* induces oxidative stress and halogen metabolism. *Plant Cell & Environment*, *39*, 259–271.

Sunday, J. M., Fabricius, K. E., Kroeker, K. J., Anderson, K. M., Brown, N. E., Barry, J. P., ... & Klinger, T. (2017). Ocean acidification can mediate biodiversity shifts by changing biogenic habitat. *Nature Climate Change*, *7*, 81.

Thomsen, J., Stapp, L. S., Haynert, K., Schade, H., Danelli, M., Lannig, G., ... & Melzner, F. (2017). Naturally acidified habitat selects for ocean acidification–tolerant mussels. *Science Advances*, *3*, e1602411.

Ullah, H., Nagelkerken, I., Goldenberg, S. U., & Fordham, D. A. (2018). Climate change could drive marine food web collapse through altered trophic flows and cyanobacterial proliferation. *PLoS Biology*, *16*, e2003446.

Vizcaíno, J. A., Csordas, A., del-Toro, N., Dianes, J. A., Griss, J., Lavidas, I., … & Hermjakob, H. (2016). 2016 update of the PRIDE database and related tools. *Nucleic Acids Research*, *44*, D447–D456.

Xu, D., Schaum, C. E., Lin, F., Sun, K., Munroe, J. R., Zhang, X. W., ... & Ye, N. (2017). Acclimation of bloom-forming and perennial seaweeds to elevated pCO$_2$ conserved across levels of environmental complexity. *Global Change Biology*, *23*, 4828–4839.

Ye, N., Zhang, X., Miao, M., Fan, X., Zheng, Y., Xu, D., ... & Wang, Y. (2015). *Saccharina* genomes provide novel insight into kelp biology. *Nature Communications*, *6*, 6986.

Zhang, K., Song, C., Zhang, Y., Dang, H., Cheng, X., & Zhang, Q. (2018). Global-scale patterns of nutrient density and partitioning in forests in relation to climate. *Global Change Biology*, *24*, 536–551.

Zhu, C., Kobayashi, K., Loladze, I., Zhu, J., Jiang, Q., Xu, X., ... & Fukagawa, N. K. (2018). Carbon dioxide (CO$_2$) levels this century will alter the protein, micronutrients, and vitamin content of rice grains with potential health consequences for the poorest rice-dependent countries. *Science Advances*, *4*, eaaq1012.

SUPPORTING INFORMATION

Additional Supporting Information may be found in the online version of this article:

Table S1. Carbonate system data (mean ± standard errors) in the laboratory set up using the CO2SYS Package.

Table S2. The genes used for real-time quantitative PCR (RT-qPCR) of *S. japonica* cultured in field experiment under ambient (bubbled with air) and elevated $p\mathrm{CO_2}$ (bubbled with mix air of 1,000 μatm CO_2).

Table S3. The primers used for real-time quantitative PCR (RT-qPCR) of *S. japonica* cultured in field experiment under ambient (bubbled with air) and elevated $p\mathrm{CO_2}$ (bubbled with mix air of 1,000 μatm CO_2).

Fig. S1. Kinetics of iodine influx by *S. japonica* sporophyte under of various concentrations of KI. (a) Iodine influx rate (IIR) of *S. japonica* sporophyte. (b) The Lineweaver-Burk plots between KI substrate and iodine influx rate which was used for Lineweaver-Burk K_m and IUR_{max} determination. In a laboratory set up, kinetics of iodine influx of *S. japonica* under a series of iodide concentration in surrounding medium were determined over a duration of 24 hours. With increasing concentration of iodide (1-30 μmol L^{-1} KI), the influx rate increased significantly until the external KI concentration exceeded 10 μmol L^{-1} (One-Way ANOVA, df=5, P<0.001). Furthermore, iodine influx followed a Michaelis-Menten equation and Lineweaver-Burk plot was used for $K_{1/2}$ (19.94 μmol L^{-1}) and IIR_{max} (15.04 μmol g^{-1} dry weight h^{-1}) determination (R^2=0.9547, df=1, F=84.229, P<0.05).

Fig. S2. Experimental design of the field experiment. pCO$_2$ treatments were carried out in triplicate tanks (n=3 per pCO$_2$ level), under ambient pCO$_2$ (400 µatm, bubbled with air) and elevated pCO$_2$ (bubbled with air/CO$_2$ premixed gas using a CO$_2$ Enrichlor). In the field experiment, we held all seaweed species simultaneously as logistics did not allow for each species to be grown on its own at each pCO$_2$ level at sufficient replication. The letters in the net cages represent experimental algal species used in the mesocosms including *S. japonica* (a), *U. pinnatifida* (b), *M. pyrifera* (c), *G. lemaneifor* (d), *G. chouae* (e), *U. pertusa* (f), and *U. intestinalis* (g).

Fig. S3. Example for coastal mesocosms on the sea. (a) photograph of net cages and CO$_2$ Enrichlor used for ocean acidification experiment; (b) Variation of nutrient (mean ± standard errors, n=6); (c) Variation in temperature; (d) Variation of irradiance. During the culture period (160 days), the seawater temperature initially decreased from 10.36 ± 0.44 °C to 2.08 ± 0.17 °C after 80 days and then increased from 2.37 ± 0.48 °C to 12.73 ± 0.22 °C. The average irradiance was 98 µmol photons m^{-2} s^{-1} and the light condition would not limit algal growth.

Fig. S4. Variation of seawater carbonate chemistry (mean ± standard error) under ambient (400 µatm bubbled with air) or elevated pCO$_2$ (1,000 µatm bubbled with air/CO$_2$ premixed gas using a CO$_2$ Enrichlor) conditions for more than 5 months in the coastal field experiment. (a) pH; (b) Total alkalinity (TA); (c) DIC; (d)HCO$_3^-$; (e) CO$_3^{2-}$; (f) CO$_2$. The shaded areas indicate the standard deviation of three replicates.

Fig. S5. The individual and combined effect of increasing pCO$_2$ and temperature on iodine solubility and availability. Colored columns show the observed variation of iodide (a) or iodate (b) concentration of three biological replicates ($\pm$ SE) under increasing pCO$_2$ at different temperatures of 5°C, 10°C, 15°C, 20°C and 25°C.

Fig. S6. Algal fresh weight (top panels) and iodine accumulation (bottom panels) under ambient pCO$_2$ (400 µatm, bubbled with air, blue) and elevated pCO$_2$ (bubbled with mix air of 1,000 µatm CO$_2$, red) for (a) *S. japonica*, (b) *U. pinnatifida*, and (c) *M. pyrifera* in mesocosm experiments. Colored circles show the average of biological replicates at each time point ($\pm$ SE), with n=30 for fresh weight and n=18 for iodine accumulation.

Fig. S7. Algal fresh weight (top panels) and iodine accumulation (bottom panels) under ambient pCO$_2$ (400 µatm, bubbled with air, blue) and elevated pCO$_2$ (bubbled with mix air of 1,000 µatm CO$_2$, red) for (a) *U. pertusa*, (b) *U. intestinalis*, (c) *G. lemaneiformis*, and (d) *G. chouae* in mesocosm experiment. Colored circles show the average of biological replicates at each time ($\pm$ SE), with n=30 for fresh weight and n=18 for iodine accumulation.

Fig. S8. Linear regression analysis between the iodine concentration of *S. japonica* and iodine concentration or THs synthesis in the consumer *H. discus*. (a) Linear regression analysis between the iodine concentration in *S. japonica* and iodine concentration in *H. discus*; (b) Linear regression analysis between the iodine concentration in *S. japonica* and T3 concentration in *H. discus*; (c) Linear regression analysis between the iodine concentration in *S. japonica* and T4 concentration in *H. discus*.

Fig. S9. Linear regression analysis between the iodine concentration and THs synthesis in *H. discus*. (a) Linear regression analysis between the iodine concentration and T3 concentration; (b) Linear regression analysis between the iodine concentration and T4 concentration.

Fig. S10. Relative expression level of several *vHPO* genes by real-time quantitative PCR (RT-qPCR).

Fig. S11. The individual and combined effects of temperature and pCO$_2$ on F$_v$/F$_m$ of *S. japonica* in laboratory set up. (a) The individual effect of increasing temperature on F$_v$/F$_m$ where culture pCO$_2$ was set at 400 µatm and bubbled with air; (b) The individual effect of increasing pCO$_2$ on F$_v$/F$_m$ where the temperature was maintained at 10 °C; (c) The combined effects of high temperature of 15 °C and increasing pCO$_2$ on F$_v$/F$_m$; (d) The combined effects of high temperature of 20 °C and increasing pCO$_2$ on F$_v$/F$_m$.

Fig. S12. Comparing the observed effect and the predicted effect of increasing pCO$_2$ and temperature on the kelp *S. japonica*. Filled circles show the observed response in total iodine accumulation of three biological replicates (± SE) under increasing pCO$_2$ at different temperatures; 10°C (grey), 15°C (yellow) and 20°C (blue). Colored solid lines show the predicted response of iodine accumulation under increasing pCO$_2$ at different temperatures; 15°C (yellow) and 20°C (blue) using (a) the additive model and (b) the multiplicative model. The dashed grey line shows the observed response of increasing pCO$_2$ under control temperature conditions (10°C). This is equivalent to predictions based on the dominant environmental driver (in this example the dominant driver is pCO$_2$) and demonstrates that

predictions on iodine accumulation in *S. japonica* are most accurate when based on the

response to $p\text{CO}_2$ alone. Note that the y-axis differ between panels a and b.

Fig. 1. Conceptual diagram showing altered iodine metabolic pathway and global iodine geochemical cycle under ocean acidification. The yellow up arrows indicate the up-regulated pathway; the yellow down arrows indicate the down-regulated pathway; the black arrow with dashed lines indicates the possible outcome under further ocean acidification. The abbreviations ROS represents the reactive oxygen species; vHPOs represents the vanadium-dependent haloperoxidases; THs represents the thyroid hormones.

Fig. 2. The combined effect of increasing $p\text{CO}_2$ and temperature and the effect of increasing temperature alone on iodine accumulation in the kelp *S. japonica*. Filled circles show the average iodide (top panels) and total iodine (bottom panels) accumulation in three biological replicates ($\pm$ SE). (a) Colored circles show the combined effects of increasing $p\text{CO}_2$ at different temperatures; 10°C (red), 15°C (blue) and 20°C (green). Horizontal dashed lines indicate the iodine accumulation under control temperature (10°C) and $p\text{CO}_2$ (400 µatm) conditions. (b) Colored circles (orange) show that iodide and total iodine accumulation increases with increasing temperature under control $p\text{CO}_2$ levels (400 µatm).

Fig. 3. The effect of increasing $p\text{CO}_2$ on iodine efflux of *S. japonica* upon oligoguluronate-triggered oxidative burst in laboratory experiment. Colored circles show the average iodine efflux of three biological replicates ($\pm$ SE) under oligoguluronate elicitor (GG, blue) and Control (red) conditions. (a) The change in iodine efflux of *S. japonica* under GG and control conditions over 5 hours at 10°C ; (b) The effect of increasing $p\text{CO}_2$ on iodine

efflux of *S. japonica* after 3 hours of GG elicitation at 10°C; (c) The short-term elevated

pCO$_2$ vs long-term elevated pCO$_2$ effect on iodine efflux 10°C. Treatments shown in the

x-axis denote the culture conditions of *S. japonica*; SAC = short-term (7 days) under ambient

carbon (400 µatm), SEC = short-term (7 days) under elevated-carbon (1,000 µatm), LAC =

long-term (30 days) under ambient-carbon (400 µatm), and LEC = long-term (30 days) under

elevated-carbon (1,000 µatm).

Fig. 4. Relative changes in weight (top panels) and relative changes in iodine

accumulation (bottom panels) under ambient pCO$_2$ (400 µatm, bubbled with air, blue)

and elevated pCO$_2$ (bubbled with mix air of 1,000 µatm CO$_2$, red), in a mesocosm

experiment. Three kelp species (a) *S. japonica*, (b) *U. pinnatifida*, (c) *M. pyrifera* were

cultured in mesocosm experiment. Colored circles show the average of biological replicates

at each time (± SE), with n=30 for the relative changes in weight and n=18 for the relative

changes in iodine.

Fig. 5. Changes in iodine accumulation and thyroid hormones (THs) synthesis in *H.*

***discus* fed with seaweeds cultured under either ambient and elevated pCO$_2$.** Colored

symbols show the average iodine accumulation of six biological replicates (± SE) fed with

either *S. japonica* cultured under 400 µatm (circles) or 1,000 µatm (triangles) pCO$_2$. Feeding

trials were conducted inside a cage under either 1,000 µatm (red symbols) or 400 µatm (blue

symbols) pCO$_2$ or in the field with no cage under 400 µatm (gray symbols) pCO$_2$. (a)

Changes in iodine accumulation in *H. discus* were measured every 10 days for 30 days. (b)

Changes in the concentrations of triiodothyronine (T3) and thyroxine (T4) at day 30, relative

to data collected from feeding experiments with algae cultured under 400 µatm, in the field.

Fig. 6. The effect of increasing $p\text{CO}_2$ on the relative expression of vHPO at transcriptomic or proteomic level in *S. japonica* cultured under ambient (400 μatm) or elevated (1,000 μatm) $p\text{CO}_2$ scenarios, using an *in situ* mesocosm experiment. (a) Variation of *vHPO* gene expression; (b) Variation of vHPO protein expression. * (p-value <0.05) and ** (p-value <0.001) represent significant differences between treatments of ambient (400 μatm) or high (1,000 μatm) $p\text{CO}_2$.

(a)

(b)