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RESEARCH ARTICLE

How elevated CO_2 affects our nutrition in rice, and how we can deal with it

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Abstract

Increased concentrations of atmospheric CO_2 are predicted to reduce the content of essential elements such as protein, zinc, and iron in C_3 grains and legumes, threatening the nutrition of billions of people in the next 50 years. However, this prediction has mostly been limited to grain crops, and moreover, we have little information about either the underlying mechanism or an effective intervention to mitigate these reductions. Here, we present a broader picture of the reductions in elemental content among crops grown under elevated CO_2 concentration. By using a new approach, flow analysis of elements, we show that lower absorption and/or translocation to grains is a key factor underlying such elemental changes. On the basis of these findings, we propose two effective interventions—namely, growing C_4 instead of C_3 crops, and genetic improvements—to minimize the elemental changes in crops, and thereby avoid an impairment of human nutrition under conditions of elevated CO_2 .

Introduction

On the basis of the results of free-air CO₂ enrichment (FACE) experiments, Myers et al. [1, 2] argued that increasing CO₂ concentration ([CO₂]) over the next 40–60 years would lead to deficiencies of essential elements, including nitrogen (protein), zinc, and iron, in C₃ grains and legumes. In turn, this would threaten human nutrition, especially in low-resource countries where people depend on grain crops for their micronutrient intake. Studies supporting this prediction have been reported since the 1990s, including meta-analyses [3, 4, 5] and FACE experiments [6, 7]. C₃ grains and legumes supply 60% of dietary Zn and/or Fe for roughly 2 billion people [1]. Among these crops, rice is an important staple for half of the world's population and also supplies more than 30% of total protein for 3.4 hundred million individuals [8].

Elemental content varies among organs in a plant. For example, the elemental content differs markedly between brown rice and the "polished" grains that we ordinarily consume. To our knowledge, all previous studies except one [9, 10] have analyzed the mineral content of brown grains. In many countries, the rice plant body ("straw") is supplied as feed to cattle [11]. It is not known whether the response of the elemental content to elevated [CO₂] is the same in brown grains as it is in polished rice or the plant body; if not, predictions of deficiencies might not reflect the full scale of damage to the nutritional qualities of rice caused by elevated [CO₂].

More importantly, the information available on the effect of $[CO_2]$ on grain crops is limited, and little is known about either the underlying mechanism or practicable strategies to avert a future nutritional crisis. Here we explore three points concerning the potential damage to human nutrition caused by elevated $[CO_2]$. First, we present a broader picture of its impact, including data for the plant body of rice and other crops (e.g., feed crops and leaf vegetables); second, we investigate the physiological mechanism underlying the reduction in elemental content; and third, we propose feasible targets to minimize the effect of elevated $[CO_2]$ on the basis of this mechanism.

Materials and methods

Plant materials

Rice (Oryza sativa L.) was grown under FACE conditions in Tsukubamirai, Japan, from 2010 to 2012 [12]. In 2010, nine rice varieties ("Koshihikari"; japonica "Nipponbare"; japonica highyield "Hoshiaoba"; large grain type "Bekoaoba"; indica high-yield "Takanari", "Hokuriku 193", and "IR72"; tropical japonica "Lemont"; and japonica-dominant high-yield "Momiroman") as well as a "Koshihikari" line in which part of chromosome 5 was substituted by "Kasalath" (SLrg5) [13] were cultivated. We also grew "Koshihikari" and a chromosome segment substitution line for GS325 (SL-GS3) in the genetic background of "Koshihikari" in 2012. Seeds of the two chromosome segment substitution lines have been deposited in the Rice Genome Resource Center (http://www.rgrc.dna.affrc.go.jp/index.html). The panicles were harvested at maturity and the brown and/or polished grains were analyzed for elemental content. For the analysis of elemental stocks and flows, three tillers of "Koshihikari" were sampled from three plants for each replication at six stages of growth in 2010. Sampling was conducted at the early vegetative (36 days after transplanting [DAT]), late vegetative (50 DAT), panicle formation (64 DAT), heading (78 DAT), grain filling (92 DAT), and maturity (107 DAT) stages. These samples were divided into six parts (see Fig 1B), and dried portions were powdered and used for elemental measurements. In 2011, the flag leaves of "Koshihikari" were harvested at the heading stage and immediately stored in liquid nitrogen. These leaves were used in microarray analysis and real-time PCR. In addition, we grew three feed crops (Avena sativa, Lolium perenne, and Medicago sativa) and four leaf vegetables (Brassica rapa var. perviridis [2 cultivars], Brassica rapa var. chinensis, Brassica oleracea var. capitata, and Beta vulgaris var. cilia) at two concentrations of CO_2 in semi-closed growth chambers [14]. The CO_2 concentrations were fixed at 370 (ambient) and 700 ppm (elevated). The above-ground parts of these plants were sampled 30 days after sowing, except for Lolium perenne and Medicago sativa, which were sampled at 40 days.

Measurement of elemental content

The content of nitrogen and other elements was analyzed by, respectively, an NC analyzer (SUMIGRAPH NCH-22, Sumika Chemical Analysis Service) and an energy-dispersive X-ray fluorescence spectrometer (EDXRF, element analyzer JSX-3201, JEOL). For the analysis of all elements except nitrogen, three 13-mm-diameter tablets were formed from each replicate, and



Fig 1. Influence of elevated [**CO**₂] **on elemental content in crops.** (a) Comparison of changes in the content of all tested elements under elevated [CO₂] between brown rice (black circles) and polished grains (red diamonds). Data are presented as the mean and one side of 95% confidence intervals (bars) of nine varieties. Red and blue backgrounds denote a significant difference (P < 0.05), where previous work on brown grain has under- or overestimated, respectively, elemental changes in polished grains. For Al, Si, and Br, the values were off the scale of the *y* axis, as indicated by the wavy line. (b) Time-dependent changes in the elemental content of each part of the rice plant under elevated [CO₂]. Data are presented as the mean. (c) Percentage change in the elemental content of polished grains and plant body ("straw") of rice ("*Koshihikari*"). (d) Estimated change in elemental content in feed crops under elevated [CO₂]. (e) Estimated change in elemental content in leaf vegetables under elevated [CO₂]. The raw data are provided in <u>Table 1</u> and <u>S4 Table</u>. In (c)-(e), essential elements are arranged in decreasing order of intake requirement. Elements that decreased significantly (P < 0.05) under elevated [CO₂] are shown in red.

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each tablet was measured three times according to the method of Kashiwagi et al. (2009) [15]. The average of these nine measurements was taken as the result for each replicate. Differences in mineral content between different $[CO_2]$ conditions were tested by Z-test.

Calculation of elemental stocks and flows

The relative stock of each element in the plant was calculated as a product of the relative content and dry weight. The relative rate of elemental translocation between plant parts during different growth stages, which we call "flow," was estimated by using the stock of the element. Flow was calculated as follows: (1) Flow = $(C_2W_2-C1W_1)/(t_2-t_1)$, where C and W are the relative content and dry weight, respectively; C_nW_n is the stock value of the element; $C_2W_2-C_1W_1$ indicates the change in the content of the element between growth stages; and t_2-t_1 is the number of days between the growth stages. A positive flow value indicates that the element has flowed into the plant tissue. Conversely, a negative value means the element has translocated to other tissues or has been discharged by defoliation. The flow of elements among organs was estimated by net flows for each element, as described for the modeling of carbon and nitrogen flows [16, 17]. Hierarchic clustering of the flow pattern was conducted by a group average method based on Euclidean distance. In Fig 2A, we show a representation of the difference in net flows between ambient and elevated [CO₂] conditions.

Estimation of the impact on feed crops and leaf vegetables

Changes in element content were estimated from data for eight feed crops and seven leaf vegetables, including eight new data sets obtained in this study (feed crops: *Avena sativa*, *Lolium perenne*, *Medicago sativa*, and rice straw; leaf vegetables: *Brassica rapa* var. *perviridis* [two cultivars], *Brassica rapa* var. *chinensis*, *Brassica oleracea* var. *capitata*, and *Beta vulgaris* var. *cilia*). Other data were taken from six previous reports (see legends of <u>Table 1</u> and <u>S4 Table</u>). The effects of elevated [CO₂] on elemental content were assessed by Z-test. Elements that were not measured were excluded from the statistical analysis.

Gene expression analyses

Total RNA was extracted from a rice flag leaf at the heading stage. We used a rice genome 44K oligonucleotide microarray (G2519F#15241, Agilent Technologies) containing approximately 42,000 oligonucleotides synthesized on the basis of nucleotide sequences and full-length cDNA data from RAP-DB [18, 19]. Fluorescence probe labeling, hybridization, scanning, and data analysis were performed according to the manufacturer's instructions. We focused only on expression levels that showed a more than 1.5-fold change in expression that was statistically significant (P < 0.05 by Significance Analysis of Microarray). The extracted total RNA was converted into first-strand cDNA, and then quantitative RT-PCR was performed with the same samples used in the microarray. *Actin (Act 1)* was used as a control. The following primers were used: for *Act1*, forward GACTCTGGTGATGGTGTCAGC, reverse GGCTGGAAGAGGA CCTCAGG; for *OsZIP5*, forward CTGGAGCTGGGAGTGGTGGTG, reverse TCAGTCTCATCATGG GAAAAA.

Results

Influence of elevated [CO₂] on elemental content in rice

As compared with ambient conditions, elevated $[CO_2]$ reduced the content of N (protein), S, Mn, and Zn in polished grains by 13.5%, 12.6%, 20.5%, and 5.9%, respectively (Fig 1A). The



Fig 2. Influence of elevated $[CO_2]$ on flow of elements in a rice plant during growth, and genetic improvement to prevent elemental reduction in grains. (a) Change in elemental flow in rice ("*Koshihikari*") under elevated $[CO_2]$ as compared with ambient $[CO_2]$. Arrows indicate the direction of elemental flow: blue shows an increase, red a decrease, and grey no change under elevated $[CO_2]$ versus ambient $[CO_2]$. Purple arrows indicate that the direction of flow was reversed by CO_2 treatment. The extent of the change in flow is indicated by arrow width; the part of the plant showing

the greatest changes for each element is represented by the widest arrow. Abbreviations are defined in Fig 1B. (b) Percentage change in the nitrogen (protein) and sulfur content of grains of wild-type "*Koshihikari*" and a chromosomal segment substitution line containing *rg*518 or *GS*319 grown under elevated [CO₂] relative to ambient [CO₂].

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corresponding decreases in brown rice were, respectively, 2.0%, 3.5%, 7.0%, and 3.0% smaller than those in polished grains. For both types of grain, the decrease in Fe content was not significant; however, polishing reduced Fe content by 10.9%. The influence of elevated [CO₂] varied among plant parts and growth stages (Fig 1B). As compared with the grains, the plant body exhibited a greater decrease in elemental content under elevated [CO₂] (Fig 1C): for example, the content of Mg, S, and Mn in the plant body decreased by 28%, 21%, and 53%, respectively (all P < 0.05), which was approximately 2-, 1.5-, and 4-fold greater than the decreases observed in grains.

Influence of elevated [CO₂] on elemental content in feed crops and leaf vegetables

The analysis of eight feed crops, which combined our new data including rice straw with data from three previous studies (Table 1), showed that elevated [CO₂] was associated with a significant decrease in five major and three minor essential minerals (Fig 1D). Further, the content of Ca and S was reduced by 12.7% and 19.5%, respectively (all P < 0.05; Fig 1D, Table 1). In contrast, as compared with C₃ feed crops, the elemental content of the plant body of C₄ crops was little affected by elevated [CO₂]. The analysis of seven leaf vegetables, again combining previous data with our current results, indicated that elevated [CO₂] was associated with a reduction of about 20% in the concentration of multiple essential minerals, including Mg, P, Ca, Mn, Fe, and Zn (Fig 1E).

Influence of elevated [CO₂] on the expression of transporter genes

Gene expression profiles were assessed in a flag leaf of rice obtained around the heading stage. As compared with ambient conditions, 798 genes were significantly down-regulated by at least one-third, and the expression of 529 genes increased 1.5-fold under FACE conditions. Among the genes showing changes, 14 encoded transporters. Ten of these genes, including K and

 Table 1. Effect of elevated $[CO_2]$ on elemental content in feed crops. ND means no data. *Significant at P<0.05. The values mean the ratio of elemental content under elevated $[CO_2]$ to content under ambient condition (%).

Plant species	Plant type	S	К	Ca	Р	Mg	Zn	Fe	Mn	Cu	Reference
Avena sativa	C3	56.21	82.19	101.90	41.76	95.95	57.17	66.28	54.12	56.80	new data
Lotium perenne	C3	88.17	105.35	97.01	105.82	92.93	87.74	105.56	62.98	64.22	new data
Oryza sativa (straw)	C3	79.49	96.19	86.02	104.11	71.88	75.27	58.53	47.23	77.97	new data
Bromus tectorum	C3	ND	82.35	83.87	ND	ND	ND	ND	84.21	ND	Blank et al. [<u>35</u>]
Festuca vivipara	C3	ND	62.16	54.77	52.04	62.98	ND	ND	ND	ND	Baxter et al. [36]
Medicago sativa	C4	98.31	88.72	98.94	96.53	116.74	89.54	86.58	146.16	96.14	new data
Bouteloua curtipendula	C4	ND	94.03	101.58	88.18	87.57	126.36	108.09	73.98	141.37	Polley et al. [37]
Sorghastrum nutans	C4	ND	109.82	74.44	99.71	101.21	101.25	125.19	77.74	108.38	Polley et al. [37]
Ave.		80.54	90.10	87.32	84.02	89.89	89.55	91.70	78.06	90.81	
P value (ZTEST)		0.0302	0.0623	0.0288	0.1061	0.1380	0.2750	0.4327	0.0765	0.4733	
		*	NS	*	NS	NS	NS	NS	NS	NS	
Ave in C3		74.62	85.65	84.71	75.93	80.93	73.39	76.79	62.14	66.33	
Ave in C4		98.31	97.53	91.65	94.80	101.84	105.71	106.62	99.30	115.30	

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Element	CODE	GENE	Fold-change	P value (SAM)
Fe	Os07g0689600	OsNAS3	2.66	0.0317
Various element	Os09g0333500	PDR-like ABC transporter (PDR3 ABC transporter).	2.14	0.0163
К	Os04g0445000	Potassium channel SKOR (Stelar K(+) outward rectifying channel).	1.64	0.0273
Various element	Os06g0695800	ABC transporter related domain containing protein. (potassium transport)	1.58	0.0306
S	Os01g0719300	Sulfate transporter 3.1 (AST12) (AtST1). (absorption)	0.64	0.0111
Various element	Os09g0472200	ABC transporter.	0.60	0.0225
Mg	Os03g0137700	Mg2+ transporter protein, CorA-like family protein. (absorption)	0.59	0.0034
Zn	Os03g0411800	Zinc transporter 11 precursor (ZRT/IRT-like protein 11)	0.57	0.0021
Heavy metal	Os04g0464100	Heavy metal transport/detoxification protein domain ontaining protein. (absorption/desorption)	0.53	0.0373
Fe, Zn	Os04g0613000	Zinc transporter 1 precursor (ZRT/IRT-like protein 1). OsZIP3	0.53	0.0487
Various element	Os11g0155600	ABC transporter related domain containing protein.	0.51	0.0014
Heavy metal	Os02g0582600	Heavy metal transport/detoxification protein domain containing protein.	0.47	0.0037
Mn, Fe, Cu, Zn	Os05g0472700	Zinc transporter protein OsZIP5.	0.46	0.0003
K	Os08g0466200	K+ potassium transporter family protein. (absorption)	0.35	0.0076

Table 2. List of genes for element transport that is affected by elevated [CO₂].

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heavy metal transporters, were down-regulated, whereas the expression of the other four increased (<u>Table 2</u> and <u>S1 Fig</u>). Among the ten down-regulated genes, five have functions in the import of minerals into cells without the involvement of ABC transporters.

Flow analysis of elements in rice throughout different growth stages

To clarify the physiological factors responsible for the elemental changes, we analyzed the influence of elevated $[CO_2]$ on the elemental flow in rice. Retranslocation of N, S, and Cu from leaf blades was lower under elevated $[CO_2]$ than under ambient $[CO_2]$ conditions during the maturity stage (Fig 2A). Elements were clustered into three broad groups according to their flow pattern under ambient $[CO_2]$ conditions (increased, decreased, or reversed), and notably, Cu, Mo, and Mn moved into a different group under FACE conditions (S2 Fig). The large change in flow pattern indicates a shift in the allocation of elements among tissues.

Discussion

As a first step towards mitigating the damage caused by elevated $[CO_2]$, we require a complete picture of the reduction in the elemental content of crops expected in the near future. As compared with polished grains, data from brown rice have underestimated the decrease in N (protein), S, Mn, and Zn under elevated $[CO_2]$ by 2.0%, 3.5%, 7.0% and 3.0%, respectively (Fig 1A). Our results suggest that increasing $[CO_2]$ levels might have more serious consequences than previous influential predictions that were based on brown rice [1]. In particular, the drop in N in polished rice would negatively affect the nutritional status of the 153 million individuals of Bangladesh, who depend largely on rice for their protein intake and are already estimated to have an individual daily protein intake below the standard of "hungry" as defined by FAO/ WHO (S1 Table).

Although we used the same rice samples as in a previous work [1], we found different changes in several elements. For example, Myers et al. [1] found that Fe content was reduced by 5.2% under elevated [CO₂], whereas we found a slight increase. This may be because the elemental content of rice grains can be affected by environmental factors, such as air temperature [20], during the ripening period. Among the 18 varieties that Myers et al. [1] used, we selected nine varieties with similar heading behavior that were appropriate for predicting damage to actual

production due to FACE, although this narrowed the genetic diversity represented. This difference in aims between Myers et al.'s [1] and our study might also explain our different results.

Our results showed that elevated $[CO_2]$ might also impair human nutrition via leaf vegetables and feed crops that we either eat directly or utilize as animal feed, not only via grain crops. As compared to the grain, elemental content declined markedly in the plant body of rice. For example, Mg and Mn content in the plant body decreased 2- and 4-fold, respectively (Fig 1C). These data, and the difference between polished (endosperm) and brown (endosperm plus bran) grains demonstrate that the elemental reduction due to elevated $[CO_2]$, is specific to each organ. The impact on leaf vegetables and feed crops should also be noted, because these are important sources of micronutrients despite being consumed in smaller amounts than staple grains.

In the analysis of feed crops, which combined our new data including rice straw with data from three previous studies (see Fig 1D, Table 1), elevated $[CO_2]$ decreased the content of five major (S, K, P, Ca, and Mg) and three minor (Zn, Fe, and Mn) essential minerals by 8.3% to 21.9%. It is especially noteworthy that the content of Ca was reduced by 12.7%. At present, much of the world's population is facing a deficiency of Ca [21, 22]. Approximately 25.4 million adults in the United Kingdom and the United States have a high risk of Ca deficiency [21], and 54% of the population of Africa (5.7 hundred million individuals) is at risk of Ca deficiency [22]. Leaf vegetables are an important source of essential minerals; for example, in the United Kingdom and the United States, 10% of Ca and Mg intake comes from leaf vegetables [7]. The main source of Ca is milk, however, with UK adults obtaining half of their Ca intake from milk and dairy products [7]. The elemental content of the livestock's ingested feed correlates with that of the milk that they produce [23, 24]; therefore, the lower elemental content of feed crops grown under elevated $[CO_2]$ is likely to reduce the elemental content of milk and dairy products. Ultimately, elevated $[CO_2]$ might affect human nutrition and health even in high-income countries, where food supplements for micronutrients including Ca are readily available.

The effect of elevated $[CO_2]$ on Mn and S content has not previously been a research focus because we generally obtain an adequate intake of these minerals in daily life [1-5]. Our present data showed that, under elevated $[CO_2]$, the content of these elements decreased markedly across various plant parts across several species (Fig 1). A previous study showed that consumption of a Mn-deficient diet (0.11 mg/day; about 3% of RDA) for 39 days led to a mild form of dermatitis among five of seven men [25]. Sulfur is present primarily as a constituent of S-containing amino acids; a restricted dietary supply of these acids slows the synthesis rate of whole-blood glutathione-SH, an important antioxidant, and diminishes turnover. Glutathione-SH deficiency is a risk factor for chronic liver disease [26]. In the next 40–60 years, therefore, we may face unexpected challenges to our health owing to a reduction in the intake of minerals that we currently obtain in sufficient amounts.

Our new approach, flow analysis of elements, suggested that lower absorption and/or translocation is a key factor underlying the lower elemental content, at least in rice grains, under elevated $[CO_2]$ (Fig 2A). For example, lower retranslocation of N and S from leaf blades during the maturity stage led to a reduction in these elements in grains under elevated $[CO_2]$. During retranslocation, accumulated elements are transported with carbohydrates in phloem sap. These results therefore suggest that elevated $[CO_2]$ might affect elemental retranslocation via carbohydrate translocation. Zn- and Fe-regulated transporter-like protein 5 (OsZIP5) plays a main role in Zn deficiency and in the import of mineral ions such as Mn²⁺, Fe²⁺, and Cu²⁺, and is controlled at the transcriptional level in rice [27]. Under FACE conditions, its expression level was reduced by one-half, which might damage the distribution of these essential minerals (Table 2). In addition, Fe deficiency negatively regulates the expression of rice *nicotianamine synthase gene 3* (*OsNAS3*), which is related to transport [28]. Therefore, high [CO₂] might affect the expression of related transporter genes and hence absorption and/or translocation of minerals.

On the basis of the mechanism underlying the reductions in the content of essential elements, there are two potentially effective interventions to maintain the nutritional levels of food sources grown under elevated $[CO_2]$. In contrast to C_3 crops, the elemental content of the grains in C_4 crops such as maize and sorghum is reported to be little affected by elevated $[CO_2]$ [1]. Accordingly, we found that the elemental content in C₄ plants was rarely reduced by elevated $[CO_2]$ not only in the grain but also in the plant body (<u>Table 1</u>). A similar tendency was reported by Loladze $[\underline{4}]$, who estimated the effect of elevated $[CO_2]$ on the mineral contents in various C₃ and C₄ plants in a large meta-analysis. Thus, to minimize the nutritional degradation caused by increasing $[CO_2]$, the first solution might be to cultivate C_4 instead of C_3 crops. This strategy would be effective for feed crops and for leaf vegetables where we consume the plant body. Because nutrient profiles and suitability for cultivation differ between C₃ and C₄ crops, C₃ crops cannot be replaced in every situation. Under elevated [CO₂], C₄ plants decrease their transpiration rate, similar to C₃ plants, but the parallel increase in both photosynthesis and carbohydrate content is greater in C_3 plants [29]. The carbohydrates that increase under elevated [CO₂] might further dilute elemental content (termed the "dilution effect") [4]. As a result, the decrease in elemental content might be more severe in C_3 plants than in C₄ plants. Because we found that different elements were reduced at different rates, however, control mechanisms other than the dilution effect were probably operating.

The second potential intervention is based on the observation that, across C_3 crops, elevated [CO₂] significantly reduced the content of N and S, elements for which daily nutritional requirements are the highest [26]. The flow pattern of N and S suggests that reduced retranslocation to the panicle from stocks in the plant body at approximately the grain-filling stage is the main mechanism governing the observed reduction in these elements under high $[CO_2]$ (Fig 2A). N and S are retranslated with carbohydrate in sap [30, 31], and a greater sink (grain) size induces higher carbohydrate translocation [32]. We therefore hypothesized that improving carbohydrate translocation or expanding sink size might alleviate the elemental decline in grains grown under elevated [CO₂]. We found that, under FACE conditions, the content of N (protein) and S significantly decreased in grains from a premium Japanese cultivar, "Koshihikari" (control), whereas N and S content was maintained in two "Koshihikari" lines containing a chromosomal segment substitution of rg5, a locus that improves carbohydrate translocation ability to grains [13], or substitution of GS3, a gene controlling grain size [33] (Fig 2B). These results show that improvement in retranslocation is a promising target to alleviate the elemental decline in grains grown under elevated $[CO_2]$. By introducing a locus via molecular markers, it is possible to improve a single trait of a cultivar while maintaining other agronomic traits for several years [27, 34]. Combining such a breeding program with verification using FACE might yield new varieties to minimize the reduction of elements in grains of C₃ crops under elevated [CO₂].

In summary, we have shown that elevated $[CO_2]$ has the potential to cause damage to human nutrition and health via leaf vegetables and feed crops, not only grain crops. The greatest threat comes from the more severe drop in N and Zn in polished rice grains, and the lower content of Ca in feed crops and leaf vegetables. In addition, a reduction in the intake of minor minerals such as Mn and S that we currently obtain in sufficient amounts might cause unanticipated health risks. Flow analysis and transcriptomics suggested that lower absorption and/or translocation of elements is a key factor underlying the lower elemental content in rice grains, and the lower expression of related transporter genes under elevated $[CO_2]$ might also play a role. On the basis of this mechanism underlying the decrease in elemental content, we have proposed two practical strategies—the cultivation of suitable C₄ instead of C₃ crops and the genetic development of C₃ grain crops with molecular markers—to avoid this impending crisis in human nutrition.

Supporting information

S1 Table. Countries whose populations received protein from rice in 2011. Source: United Nations Food and Agriculture Organization food balance sheets and 2011 United Nations estimated population. (http://faostat3.fao.org/browse/FB/*/E) Under elevated [CO2], countries shown on a red background were estimated the protein supplies at the below level of hunger. The standard of hunger (52.5g/day) is calculated by the below; Standard in cal. by WHO/FAO 2,100 kcal per a day (ref. 1) Standard in protein by UNHCR, UNICEF, WFP and WHO <10% of cal. (ref. 2) Cal. per protein 4 kcal/g. (PDF)

S2 Table. The mineral content of brown and polished grain in rice (Value per 100g). Source: USDA National Nutrient Database for Standard Reference (long and medium grain rice) (<u>http://ods.od.nih.gov/</u>). Standard tables of food composition in Japan, Fifth Revised and Enlarged Edition—2005 - (short grain rice). (<u>http://www.mext.go.jp/b_menu/shingi/gijyutu/gijyutu3/toushin/05031802/002/001.pdf</u>) (PDF)

S3 Table. The ratio in mineral content between brown and polished grain in rice. The values indicate the ratio of the mineral content in polished grain to content in brown grain (= 100). (PDF)

S4 Table. Analysis of elevated CO₂ effects on elemental contents in leaf vegetables. ND means no data. ***, **, * Significant at P<0.001, 0.01, 0.05. In plant species, *B*. means *Brassica*. (PDF)

S1 Fig. Expression change of *OsNAS3* (Os07g0689600) and *OsZIP5* (Os05g0472700) under elevated $[CO_2]$ condition relative to ambient $[CO_2]$. *Actin* (*Act1*) was used as the control. Data are presented as mean \pm s.d. (n = 3). Student's t tests were used to calculate P values. (PDF)

S2 Fig. Dendrogram for elemental flows in elevated $[CO_2]$ and ambient $[CO_2]$. The trees were estimated from the elemental flows between plant parts in "Koshihikari". Hierarchic clustering was calculated by group average method based on Euclidean distance. (PDF)

S1 File. Original data of Fig 1A. (XLSX)

S2 File. Original data of Figs <u>1B</u> and <u>2A</u> (Relative contents of 18 elements). (XLSX)

S3 File. Original data of <u>Table 2</u> (Microarray analysis). (XLSX)

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References

- Myers SS, Antonella Z, Itai K, Peter H, Andrew LDB, Arnold BJ et al. Increasing CO₂ threatens human nutrition. Nature 2014; 510: 139–142. <u>https://doi.org/10.1038/nature13179</u> PMID: <u>24805231</u>
- Myers SS, Wesells KR, Kloog I, Zanobetti A, Schwartz J. Effect of increased concentrations of atmospheric carbon dioxide on the global threat of zinc deficiency: a modelling study. The Lancet Global Health 2015; 3: e639–e645. Available from: http://www.thelancet.com/journals/langlo/article/PIIS2214-109X(15)00093-5/ https://doi.org/10.1016/S2214-109X(15)00093-5 PMID: 26189102
- Loladze I. Rising atmospheric CO₂ and human nutrition: toward globally imbalanced plant stoichiometry? Trends Ecol. Evol. 2002; 17: 457–461.
- Loladze I. Hidden shift of the ionome of plants exposed to elevated CO₂ depletes minerals at the base of human nutrition. eLife 2014; 3: e02245. Available from: <u>https://elifesciences.org/articles/02245</u> <u>https://doi.org/10.7554/eLife.02245</u> PMID: <u>24867639</u>
- Taub DR, Miller B, Allen H. Effects of elevated CO₂ on the protein concentration of food crops: a metaanalysis. Global Change Biol. 2008; 14: 565–575.
- Lieffering M, Kim H-Y, Kobayashi K, Okada M. The impact of elevated CO₂ on the elemental concentrations of field-grown rice grains. Field Crops Res. 2004; 88: 279–288.
- Yang L, Wang Y, Dong G, Gu H, Huang J, Zhu J, et al. The impact of free-air CO₂ enrichment (FACE) and nitrogen supply on grain quality of rice. Field Crops Res. 2007; 102: 128–140.
- United Nations Food and Agriculture Organization. Food balance sheets and 2011 United Nations estimated population. <u>http://faostat3.fao.org/browse/FB/*/E</u>.
- Goufo P, Luis MM, Carranca FC, Rosa EA, Trindade H. Effect of elevated carbon dioxide concentration on rice quality: proximate composition, dietary fibers, and free sugars. Cereal Chem. 2014; 91: 293– 299.
- Goufo P, Falco V, Brities C, Wessel DF, Kratz S, Rosa EAS. et al. Effect of elevated carbon dioxide concentration on rice quality: nutritive value, color, milling, cooking, and eating qualities. Cereal Chem. 2014; 91: 513–521.
- 11. Doyle PT, Devendra C, Pearce GR. Rice straw as a feed for ruminants. 1986; ISBN 0-86403-192-9.
- Hasegawa T, Sakai H, Tokida T, Nakamura H, Zhu C, Usui Y et al. Rice cultivar responses to elevated CO₂ at two free-air CO₂ enrichment (FACE) sites in Japan, Funct Plant Biol. 40; 148–159 (2013).
- Ishimaru K, Kashiwagi T, Hirotsu N, Madoka Y. Identification and physiological analyses of a locus for rice yield potential across the genetic background. J Exp Bot. 5; 2745–2753 (2005).
- Sakai H, Yagi K, Kobayashi K, Kawashima S. Rice carbon balance under elevated CO2. New Phytologist. 150; 241–249 (2001).
- Kashiwagi T, Shindoh K, Hirotsu N, Ishimaru K. Evidence for separate translocation pathways in determining cadmium accumulation in grain and aerial plant parts in rice. BMC Plant Biol. 9, 8 (2009). <u>https:// doi.org/10.1186/1471-2229-9-8 PMID: 19154618</u>
- Pate JS, Layzell DB, McNeil DL. Modeling the transport and utilization of carbon and nitrogen in a nodulated legume. Plant Physiol. 63; 730–737 (1979). PMID: <u>16660802</u>

- Jeschke WD, Atkins CA, Pate JS. Ion circulation via phloem and xylem between root and shoot of nodulated white lupin. J. Plant Physiol. 117; 319–330 (1985). <u>https://doi.org/10.1016/S0176-1617(85)</u> 80068-7 PMID: 23195799
- Sakai H, Lee SS, Tanaka T, Numa H, Kim J, Kawahara Y et al. Rice Annotation Project Database (RAP-DB): An integrative and interactive database for rice genomics. Plant Cell Physiol. 54, e6 (2013). https://doi.org/10.1093/pcp/pcs183 PMID: 23299411
- Kawahara Y, Bastide M, Hamilton JP, Kanamori H, McCombie WR, Ouyang S et al. Improvement of the Oryza sativa Nipponbare reference genome using next generation sequence and optical map data. Rice. 6; 4 (2013). https://doi.org/10.1186/1939-8433-6-4 PMID: 24280374
- 20. Weon S, Chamura S. Occurrence of varietal differences in protein, phosphorus, and potassium content in brown rice, and influence of temperature and shading during the ripening period on it. Japan. Jour. Crop Sci. 1980; 49: 199–204.
- Broadley MR, White PJ. Eats roots and leaves. Can edible horticultural crops address dietary calcium, magnesium and potassium deficiencies? Proc. Nutr. Soc. 2010; 69: 601–612. <u>https://doi.org/10.1017/S0029665110001588</u> PMID: 20509990
- Joy EJ, Ander EL, Young SD, Black CR, Watts MJ, Chilimba AD et al. Dietary mineral supplies in Africa. Physiol. Plant. 2014; 151: 208–229. <u>https://doi.org/10.1111/ppl.12144</u> PMID: <u>24524331</u>
- Herwig N, Stephan K, Panne U, Pritzkow W, Vogl J. Multi-element screening in milk and feed by SF-ICP-MS. Food Chem. 2011; 124: 1223–1230.
- Coni E, Bocca A, Lanni D, Caroli S. Preliminary evaluation of the factors influencing the trace element content of milk and dairy products. Food Chem. 1995; 52: 123–130.
- Friedman BJ, Freeland-Graves JH, Bales CW, Behmardi F, Shorey-Kutschke RL, Willis RA et al. Manganese balance and clinical observations in young men fed a manganese-deficient diet. J. Nutr. 1987; 117: 133–143. <u>https://doi.org/10.1093/jn/117.1.133</u> PMID: <u>3819860</u>
- Long SP, Ainsworth EA, Leakey AD, Nösberger J, Ort DR. Food for thought: lower-than-expected crop yield stimulation with rising CO₂ concentrations. Science 2006; 312: 1918–1921. <u>https://doi.org/10. 1126/science.1114722</u> PMID: <u>16809532</u>
- Lee S, Jeong HJ, Kim SA, Lee J, Guerinot ML, An G. OsZIP5 is a plasma membrane zinc transporter in rice. Plant Mole. Biol. 2010; 73; 507–517.
- Inoue H, Higuchi K, Takahashi M, Nakanishi H, Mori S, Nishizawa NK. Three rice nicotianamine synthase genes, OsNAS1, OsNAS2, and OsNAS3 are expressed in cells involved in long-distance transport of iron and differentially regulated by iron. Plant J. 36; 366–381 (2003). PMID: <u>14617093</u>
- McGrath JM, Lobell DB. Reduction of transpiration and altered nutrient allocation contribute to nutrient decline of crops grown in elevated CO₂ concentrations. Plant Cell Environ. 36; 697–705 (2013). <u>https:// doi.org/10.1111/pce.12007</u> PMID: <u>22943419</u>
- Leakey ADB, Ainsworth EA, Bernacchi CJ, Rogers A, Long SP, Ort DR. Elevated CO₂ effects on plant carbon, nitrogen, and water relations: six important lessons from FACE. J. Exp. Bot. 60, 2859–2876 (2009). <u>https://doi.org/10.1093/jxb/erp096</u> PMID: <u>19401412</u>
- Mae T, Ohira K. The remobilization of nitrogen related to leaf growth and senescence in rice plants (*Oryza sativa* L.). Plant Cell Physiol. 22; 1067–1074 (1981).
- Song XJ, Huang W, Shi M, Zhu MZ, Lin HX. (2007). A QTL for rice grain width and weight encodes a previously unknown RING-type E3 ubiquitin ligase. Nature Genet. 39; 623–630 (2007). <u>https://doi.org/ 10.1038/ng2014</u> PMID: <u>17417637</u>
- 33. Fan C, Xing Y, Mao H, Lu T, Han B, Xu C et al. GS3, a major QTL for grain length and weight and minor QTL for grain width and thickness in rice, encodes a putative transmembrane protein. Theor Appl Genet. 112; 1164–1171 (2006). https://doi.org/10.1007/s00122-006-0218-1 PMID: 16453132
- Ishimaru K, Hirotsu N, Madoka Y, Murakami N, Hara N, Onodera H et al. Loss of function of the IAA-glucose hydrolase gene *TGW6* enhances rice grain weight and increases yield. Nature Genet. 45, 707– 711 (2013). <u>https://doi.org/10.1038/ng.2612</u> PMID: 23583977
- **35.** Blank RR, White RH, Ziska LH. Combustion properties of *Bromus tectorum* L.: influence of ecotype and growth under four CO₂ concentrations. Int. J. Wildland Fire 15, 227–236 (2006).
- Baxter R, Gantley M, Ashenden TW, Farrar JF. Effects of elevated carbon dioxide on three grass species from montane. pasture. II. Nutrient uptake, allocation and efficiency of use. J. Exp. Bot. 45, 1267– 1278 (1994).
- Polley HW, Fay PA, Jin VL, Gerald FC. CO₂ enrichment increases element concentrations in grass mixtures by changing species abundances. Plant Ecol. 212, 945–957 (2011).