

G OPEN ACCESS

Citation: Zhang Z, Sun Y, Jiang X, Wang W, Wang Z-Y (2021) Sugar inhibits brassinosteroid signaling by enhancing BIN2 phosphorylation of BZR1. PLoS Genet 17(5): e1009540. https://doi.org/10.1371/journal.pgen.1009540

Editor: Li-Jia Qu, Peking University, CHINA

Received: September 21, 2020

Accepted: April 6, 2021

Published: May 14, 2021

Peer Review History: PLOS recognizes the benefits of transparency in the peer review process; therefore, we enable the publication of all of the content of peer review and author responses alongside final, published articles. The editorial history of this article is available here: https://doi.org/10.1371/journal.pgen.1009540

Copyright: © 2021 Zhang et al. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the manuscript and its <u>Supporting</u> Information files.

Funding: This work was supported by grants from National Institute of Health (NIH, R01GM066258 to Z-Y.W., https://www.nigms.nih.gov), the National

RESEARCH ARTICLE

Sugar inhibits brassinosteroid signaling by enhancing BIN2 phosphorylation of BZR1

Zhenzhen Zhang^{1,2}, Ying Sun³, Xue Jiang¹, Wenfei Wang^{1*}, Zhi-Yong Wang^{2*}

1 College of Life Sciences, Fujian Agriculture and Forestry University (FAFU), Fuzhou, China, 2 Department of Plant Biology, Carnegie Institution for Science, Stanford, California, United States of America, 3 Hebei Key Laboratory of Molecular and Cellular Biology, Key Laboratory of Molecular and Cellular Biology of Ministry of Education, College of Life Science, Hebei Normal University, Hebei Collaboration Innovation Center for Cell Signaling, Shijiazhuang, China

* wenfeiwang@fafu.edu.cn (WW); zywang24@stanford.edu (ZW)

Abstract

Sugar, light, and hormones are major signals regulating plant growth and development, however, the interactions among these signals are not fully understood at the molecular level. Recent studies showed that sugar promotes hypocotyl elongation by activating the brassinosteroid (BR) signaling pathway after shifting Arabidopsis seedlings from light to extended darkness. Here, we show that sugar inhibits BR signaling in Arabidopsis seedlings grown under light. BR induction of hypocotyl elongation in seedlings grown under light is inhibited by increasing concentration of sucrose. The sugar inhibition of BR response is correlated with decreased effect of BR on the dephosphorylation of BZR1, the master transcription factor of the BR signaling pathway. This sugar effect is independent of the sugar sensors Hexokinase 1 (HXK1) and Target of Rapamycin (TOR), but requires the GSK3-like kinase Brassinosteroid-Insensitive 2 (BIN2), which is stabilized by sugar. Our study uncovers an inhibitory effect of sugar on BR signaling in plants grown under light, in contrast to its promotive effect in the dark. Such light-dependent sugar-BR crosstalk apparently contributes to optimal growth responses to photosynthate availability according to light-dark conditions.

Author summary

Genetic studies of the brassinosteroid (BR) deficient mutants revealed its essential role in seedling development in the dark, but subsequent studies showed no significant difference in BR level between seedlings grown under light and darkness. We recently observed that light does affect BR levels in Arabidopsis, but in a sugar dependent manner. In the dark, sugar increases BR level as well as BR sensitivity by stabilizing the steroid response factor BZR1 through the Target of Rapamycin (TOR) signaling pathway. However, the BR level is decreased by sugar under light and by darkness on sugar-free medium. These observations raised the question of how the combinations of light and sugar modulate BR signaling. We addressed this question using genetic physiological analyses and found interestingly that sugar inhibits brassinosteroid response in light-grown plants by

Natural Science Foundation of China grant (NO. 31700254, http://www.nsfc.gov.cn)and FAFU-International Collaborative Program (KXb16005A, https://www.fafu.edu.cn) to W.W., the Postdoctoral Innovative Talent Support Program (BX201700052, http://www.chinapostdoctor.org. cn/index.html) and the China Postdoctoral Science Foundation (2018M642551, http://jj. chinapostdoctor.org.cn/website/index.html) to Z.Z. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing interests: The authors have declared that no competing interests exist.

stabilizing the glycogen synthase kinase 3 homolog BIN2 and attenuating the dephosphorylation of BZR1, but independently of TOR. Our results indicate that sugar acts through distinct pathways to promote and inhibit BR signaling in dark and light conditions. Our work illustrates an intricate three-way crosstalk whereby the combination of light and sugar signals modulate the brassinosteroid signaling pathway to optimize growth according to both environmental and metabolic conditions.

Introduction

Plant growth is highly sensitive to environmental light conditions, the levels of endogenous hormones, and the availability of photosynthates (sugars). Sugar not only provides essential material and energy for growth, but also functions as signaling molecules. The sugar signaling pathways mediate plant responses to starvation (low sugar) or excess of sugar, mostly through modulating hormonal pathways [1-5]. How light modulates the sugar-hormone interactions to optimize growth is not well understood.

Brassinosteroids (BR) are a major class of growth-promoting hormones that regulate a wide range of developmental and physiological processes, including photomorphogenesis. BR plays an essential role in plant developmental responses to darkness, so called skotomorphogenesis, as the BR-deficient mutants show strong de-etiolation or constitutive photomorphogenesis phenotypes in the dark [6]. While it was widely speculated that light would reduce BR levels to promote photomorphogenesis, experimental measurement showed surprisingly no significant difference in BR level between seedlings grown under light and those grown in the dark [7]. Further studies uncovered light-BR crosstalk through interactions between downstream components of the signaling pathways [8,9]. However, recent studies suggest that BR level and sensitivity are modulated by the combination of light and sugar conditions [2,10].

BR-responsive gene expression is mediated by the Brassinazole-Resistant 1 (BZR1) family transcription factors [11]. Both nuclear localization and DNA-binding activity of BZR1 are inhibited due to phosphorylation by the GSK3-like kinase Brassinosteroid-Insensitive 2 (BIN2) [12–16]. BR signaling through the BRI1 receptor kinase leads to inactivation and degradation of BIN2 [11,17,18], and dephosphorylation of BZR1 by protein phosphatase 2A (PP2A) [19]. Unphosphorylated BZR1 accumulates in the nucleus, where it recruits the TOP-LESS family repressors to inhibit gene expression [20,21] and interacts with transcription factors of other hormonal and light signaling pathways to promote shoot cell elongation [8,22–24].

BZR1 protein level is regulated through several mechanisms. Sugar signaling through Target of Rapamycin (TOR) stabilizes BZR1. When seedlings are shifted from light to darkness and undergo starvation, BZR1 is degraded due to TOR inactivation [2,25]. The degradation of BZR1 and its homolog under starvation and stress conditions involves the autophagy pathway [2,26]. Phosphorylated BZR1 is degraded by the proteasome following PUB40-mediated ubiquitination in roots of Arabidopsis [27]. The dephosphorylated BES1, a homolog of BZR1, can be ubiquitinated and degraded by the SINAT E3 ligase in a light dependent manner [28]. In addition, BZR1 is also modified and stabilized by the small ubiquitin-like modifier (SUMO), and salt stress induces deSUMOylation of BZR1 to inhibit growth [29].

We recently observed that sugar decreases the BR level in Arabidopsis plants grown under light. However, after shifting light-grown Arabidopsis seedlings into darkness, the BR levels increased in seedlings grown on media containing sugar but decreased in those grown on sugar-free media, suggesting light-dependent effects of sugar on the BR pathway [2].

Therefore, we further tested how sugar affects BR responses under light conditions. We found that, in contrast to the positive effects of sugar on growth and BZR1 accumulation in the dark, high levels of sugar attenuated the BR promotion of hypocotyl elongation and the BR-induced BZR1 dephosphorylation in Arabidopsis grown under constant light. Further, sugar increased the level of BIN2. The inhibitory effects of sugar on hypocotyl elongation and BZR1 dephosphorylation are independent of the sugar sensors Hexokinase 1 (HXK1) and TOR. The results suggest that sugars act through distinct pathways to promote and inhibit BR signaling under different light-dark conditions. Such an intricate three-way crosstalk is likely important for optimizing growth according to both environmental condition and endogenous metabolic status.

Results

Sucrose inhibits BR-induced hypocotyl elongation in light

To test the effects of sugar on BR responses under light-grown conditions, we grew Arabidopsis seedlings on media containing various concentrations of sucrose with or without BR for five days under constant light. BR increased the hypocotyl lengths of seedlings grown on a sugar-free medium but had little effect on the seedlings grown on high concentrations (90 and 150 mM) of sucrose (Figs <u>1A</u> and <u>S1</u>). These sucrose-dependent phenotypes were not caused by osmotic effects since the same concentrations of mannitol did not reduce BR's promotion of hypocotyl elongation (Fig <u>1B</u>). Compared to plants expressing wild type BZR1-CFP, transgenic plants expressing bzr1-1D-CFP, a hypermorphic mutant form that is more effectively dephosphorylated by PP2A [12,19], had slightly shorter hypocotyls without BR treatment and longer hypocotyls after growth on BR-containing medium (Fig <u>1A</u> and <u>1C</u>), consistent with previous observations [<u>30</u>]. Sucrose had much weaker inhibitory effects on BR responsive



Fig 1. Sucrose inhibits the BR-induced hypocotyl elongation and dephosphorylation of BZR1 in light. (A-B) Hypocotyl length of transgenic Arabidopsis seedlings expressing BZR1-CFP (A and B) or bzr1-1D-CFP (C) grown on medium containing 100 nM epi-brassinolide (eBL) and various concentrations of sucrose (A and C) or mannitol (B). **** P<0.0001, *** P<0.001, * P<0.05 (Student's t test). Error bars indicate the standard error of the mean (SEM, three replicates). (D-F) Immunoblot analysis of BZR1-CFP protein in the same batches of seedlings as in penal A to C. Histone H3 was probed as a loading control. P-BZR1 indicates phosphorylated BZR1; BZR1 indicates dephosphorylated BZR1. S, sucrose; M, mannitol.

https://doi.org/10.1371/journal.pgen.1009540.g001

hypocotyl elongation of the *bzr1-1D-CFP* than *BZR1-CFP* seedlings (Fig 1A and 1C). Taken together, the results suggest that sugar inhibits BR-induced hypocotyl elongation by inactivating BZR1.

Sucrose attenuates BR-induced dephosphorylation of BZR1 in light

As BR promotes hypocotyl elongation through dephosphorylation of the BZR1 family transcription factors, we further examined whether sugar affects the phosphorylation status of BZR1. As shown in Fig 1D, exogenous BR treatment promoted dephosphorylation of BZR1 (as shown by the band shift in gel) in seedlings grown on mannitol or low concentrations of sugar (0 and 30 mM), but the effect on BZR1 dephosphorylation is decreased by high concentrations of sucrose (90 mM and 150 mM) (Figs 1D and 1E and S2). Furthermore, sucrose significantly decreased the BR effects on the expression of BZR1 target genes *CPD*, *DWF4* and *SAUR-AC* (S3 Fig), consistent with the effects on BZR1 phosphorylation. The bzr1-1D protein was more dephosphorylated than BZR1 under all conditions, although high concentrations of sugar also increased the phosphorylation of bzr1-1D (Fig 1F). These results indicate that sucrose inhibits BR-induced BZR1 dephosphorylation and activation in light-grown seedlings.

To further test whether sucrose decreases the sensitivity to BR, we treated seedlings grown on media containing 30 mM and 90 mM sucrose with BR for different times. BR caused more rapid dephosphorylation of BZR1 protein in seedlings grown under the low sucrose condition (30 mM) than the high sucrose (90 mM) condition (Fig 2). The total BZR1 protein level is lower on 30 mM sucrose media than on 90 mM sucrose (1:1.8) before BL treatment but



Fig 2. High level of sucrose inhibits BR-induced dephosphorylation of BZR1. (A) Immunoblot analysis of BZR1-CFP in seedlings grown in 30 mM or 90 mM sucrose medium after treatment with 1 μ M brassinolide (BL) for the indicated time. Numbers below the image show the relative level of total BZR1 protein. (B) Quantification of the ratio between dephosphorylated and phosphorylated BZR1 (P-BZR1) using results shown in (A) and two additional biological repeats. (C) Immunoblot of BZR1-CFP in seedlings grown for five days on media containing 30 mM or 90 mM sucrose and different concentrations of BL after two-day germination in 30 mM sucrose medium. The relative levels of total BZR1 protein are shown below the image. (D) Quantification of the BZR1/P-BZR1 ratio using results shown in (C) and two additional biological replicate samples. Error bars indicate the SEM.

https://doi.org/10.1371/journal.pgen.1009540.g002

reached similar levels after 60 min BL treatments (Fig 2A), suggesting that the difference in phosphorylation is not due to a difference in BZR1 protein level. Base on three biological replicate results, the 15-minute BR treatments of seedlings grown in the low-sugar condition caused similar BZR1 dephosphorylation (ratio of dephosphorylated BZR1 to phospho-BZR1, BZR1/P-BZR1) to that caused by 60-minute BR treatments of seedlings grown in the high sugar condition (Fig 2B).

We further analyzed BZR1 in seedlings grown on media containing low (30 mM) or high (90 mM) concentrations of sucrose and various concentrations of brassinolide (BL). The results show that a higher BL concentration is required in high sugar condition than in low sugar condition to induce a similar increase of the BZR1/P-BZR1 ratio (Fig 2C and 2D). In the presence of BL (10 nM or higher), BZR1 was more dephosphorylated under low-sugar conditions than high-sugar conditions while the total BZR1 protein level remained similar. Collectively, these results demonstrate that high concentrations of sucrose inhibit BR signaling upstream of BZR1, specifically by inhibiting BR-induced BZR1 dephosphorylation, in light-grown seedlings.

Sucrose suppresses BR signaling upstream of BIN2

Phosphorylation of BZR1 at Ser173 causes its binding and inhibition by the 14-3-3 proteins [16]. It has been reported that sugar increases the amount of many proteins that bind to the 14-3-3 proteins [31,32]. It was proposed that sugar may promote the binding of the 14-3-3 proteins to their target proteins and protect them from degradation by proteolysis [32]. Therefore, sugar may inhibit dephosphorylation or degradation of phospho-BZR1 by enhancing its binding to the 14-3-3 proteins. To test this hypothesis, we analyzed sugar effects on hypocotyl elongation and BZR1 dephosphorylation in transgenic plants that express the *BZR1*^{S173A}–*CFP* protein, which contains the S173A mutation that abolishes 14-3-3 binding [16]. The hypocotyl elongation induced by BR was inhibited by sucrose in the *BZR1*^{S173A}–*CFP* plants, and the BR-induced dephosphorylation of *BZR1*^{S173A} was also inhibited by sucrose, similar to wild-type BZR1 (S4 Fig). Thus, these results indicate that sucrose inhibits BR-induced dephosphorylation and hypocotyl elongation independent of the 14-3-3 proteins.

BR induces BZR1 dephosphorylation by inactivating BIN2. We thus tested whether BIN2 mediates the phosphorylation of BZR1 on high sugar medium. It is reported that BIN2 interacts directly with BZR1 through a 12-amino acid BIN2-docking motif (DM) near the BZR1 C terminus. Deletion of this motif (bzr1- Δ DM) abolishes the interaction of BZR1 with BIN2 and prevents BIN2 phosphorylation of BZR1 *in vivo* [33]. Sugar was unable to cause phosphorylation of the bzr1- Δ DM protein (Fig 3A and 3B), suggesting that BIN2 mediates the sugar-promoted phosphorylation of BZR1. Further, the *bin2-triple* mutant with loss-of-function of BIN2 and its close homologs, *bin2bil1bil2*, showed reduced sensitivity to sugar (Fig 3C and 3D). In the presence of sugar, BR induced more dramatic hypocotyl elongation and BZR1 dephosphorylation in the *bin2bil1bil2* seedlings than wild type (Figs 3C and 3D) and S5). These results suggest that sugar inhibition of BR responsiveness is dependent on BIN2 family proteins. Interestingly, we found that sugar increased the BIN2 level in the absence and presence of exogenous BR (Fig 3E). Together these results support a scenario that sugar causes accumulation of BIN2 protein to increase BZR1 phosphorylation and reduce BR responsive growth.

Sucrose restrains BR signaling in light through unknown sugar signaling pathway

To determine which sugar-signaling pathway mediates the suppression on BR signaling in light, we carried out genetic tests of known sugar-signaling mutants. TOR and HXK1 are



Fig 3. Sucrose inhibition of BR signaling depends on the regulation of BIN2. (A) Schematic presentation of BZR1-CFP and bzr1- Δ DM-GFP. (B) Immunoblot analysis of BZR1 protein in *BZR1-CFP*/Col-0 and *bzr1-\DeltaDM-GFP*/Col-0 seedlings grown on medium containing 100 nM eBL and different concentrations of sucrose (mM S). Histone H3 was probed as a loading control in the Immunoblot analysis. (C-D) Phenotypes and hypocotyl length of wild type (WS) and *bin2bil1bil2 (bin2-triple)* mutant seedlings grown on media containing no sucrose (-S) or 90 mM sucrose (+S) and 0 (-BL) or 10 nM BL (+BL). Bar = 5 mm. **** *P*<0.0001, *** *P*<0.001, *** *P*<0.01 (Student's t test). Error bars indicate the SEM (three replicates). (E) Anti-BIN2 immunoblot analysis of BIN2 proteins in seedlings grown on media containing 0 (-S) or 90 mM (+S) sucrose and 0 (-) or 10 nM (+) BL as indicated. The immunoblot was probed with an anti-Actin antibody as a loading control.

https://doi.org/10.1371/journal.pgen.1009540.g003

known to mediate plant growth response to moderate and high concentrations of sugars, respectively [25,34]. We thus performed sugar and BR treatments in the HXK1-deficient mutant *gin2-1* [25] and the estradiol-inducible TOR silencing line *tor-es* [34]. As HXK1 is the sensor of glucose, we grew seedlings with glucose and sucrose separately to distinguish their effect on BR signaling. The *gin2-1* mutant and wild type plants showed similar sugar inhibition of BR-promoted hypocotyl elongation (Fig 4A and 4B), unlike that *bzr1-1D* mutant, which is less inhibited by sugar compared to wild type (Fig 4A and 4B). In addition, the effects of BR and sugar on the BZR1 phosphorylation level were similar in the *gin2-1* background compared to wild type, with significant amounts of phosphorylated BZR1 remaining in the presence of both BR and high sugar (Fig 4C). The results suggest that HXK1 is not required for sugar inhibition of BR signaling.

Consistent with TOR promoting plant growth, when TOR is inactivated by estradiol-inducible RNAi suppression in the *tor-es* seedlings, seedlings are smaller compared to *tor-es* untreated with estradiol (Fig 4D). However, these *tor-es* plants showed similar sugar inhibition of BR signaling as wild type, based on their hypocotyl elongation and BZR1 dephosphorylation status in response to sugar and BR treatments (Fig 4D–4F). These results suggest that TOR is not involved in sugar suppression of BR signaling in light-grown plants.

Discussion

Plant growth is highly sensitive to environmental signals and endogenous nutrient availability. Hormones, as internal growth regulators, are highly modulated by both environmental conditions and sugar availability to optimize growth and survival. Low sugar levels tend to be limiting for growth when plants are shaded or left in extended darkness, whereas surplus of photosynthate can also be inhibitory to growth. How plants deal with different sugar statuses under different environmental conditions is a key question relevant to crop yield.

We previously showed that sugar depletion in the dark causes BZR1 degradation and growth arrest; under such conditions, exogenous supply of sugar promotes BZR1 accumulation, thus enhancing BR promotion of shoot organ elongation [2,10]. Such sugar promotion of BR signaling in the dark is consistent with the need for maximum shoot elongation under shaded conditions while sugar is available, but arrest of such elongation when the sugar level is low. By contrast, we show in this study that for plants grown under full light, hypocotyl/shoot elongation is not a priority and a surplus of sugar is inhibitory to BR promotion of hypocotyl elongation. Such opposite effects of sugar on BR-dependent hypocotyl elongation are mediated by distinct mechanisms. In the dark, sugar increases BZR1 accumulation through TOR signaling. Under light, sugar increases BZR1 phosphorylation by increasing the level of BIN2 (Fig 5). It's worth noting that sugar also increases BR hormone accumulation in the dark but decreases BR level under light [2]. Thus, sugar has consistent effects on BR level and BR sensitivity, but these sugar effects are switched under dark and light conditions.

Genetic evidence suggests that distinct sugar signaling pathways are involved in the regulation of BZR1 in dark and light. While TOR mediates sugar-dependent stabilization of BZR1 in the dark [2] and BIN2 has been reported to be direct target of TOR-S6K signaling [3], inducible silencing of TOR made no obvious difference in the sugar effect on BZR1 phosphorylation or hypocotyl elongation responses to BR. TOR is inactivated when the sugar level is low, to trigger starvation response and growth arrest [34]. HXK1 is activated by high levels of glucose and mediate glucose inhibition of seedling growth and cotyledon greening [25]. There is also evidence showing that HXK1 mediates glucose positive regulation of BR signaling in promoting lateral root development [4]. It's somewhat surprising that HXK1 is also not required for



Fig 4. Sucrose inhibits BR signaling independent of HXK1 and TOR. (A) Hypocotyl lengths of seedlings grown on medium containing 0 mM or 90 mM of mannitol (90M), glucose (90G), or sucrose (90S) and no BL (-BL) or 100 nM BL (+BL). (B) Quantitation of BR sensitivity. Relative hypocotyl elongation is calculated as the ratio between the increase of length caused by BR and the length without BR treatment. Error bars indicate the SEM (five independent experiments, $n \ge 20$). (C) Immunoblot analysis of BZR1 protein in *BZR1-CFP/Col-0* and *BZR1-CFP/gin2-1*, grown under conditions as shown in (A). (D) Hypocotyl lengths of *tor-es* seedlings grown on media containing 0 mM or 90 mM (90S) sucrose, 0 (-BL) or 100 nM BL (+BL), and 0 or 1 μ M estradiol (+Estradiol). (E) Quantitation of BR sensitivity based on data in (D) using the method described for (B). Error bars indicate the SEM (three independent experiments, $n \ge 20$). (F) Images of seedlings and immunoblot analysis of BZR1 protein in seedlings described in (D).

https://doi.org/10.1371/journal.pgen.1009540.g004

sugar inhibition of BR-induced hypocotyl elongation and BZR1 dephosphorylation (Fig 4A–4C).

Alternative sugar signaling pathways may mediate inhibition of BR signaling. The SnRK1 pathway is inhibited by sugar but activated by sugar/energy deficiency, to deal with nutrient deficiency stress conditions [35]. SnRK1 is not known to be involved in responses to high sugar levels.

Recent studies show that glucose signaling is also mediated by Regulator of G-protein Signaling 1 (RGS1), a seven transmembrane guanosine-triphosphatase-activating protein that



Fig 5. A model of sugar suppressing BR signaling to modify plant growth in light. In light, BR signaling inhibits BIN2 activity, which induces accumulation of active BZR1 to promote plant growth, whereas sugar induces the phosphorylation of BZR1 to suppress BR signaling probably through increasing BIN2 activity by unknown sugar signaling pathway. The black bars and arrows indicate previously reported mechanisms; the blue arrow indicates findings made in this study.

https://doi.org/10.1371/journal.pgen.1009540.g005

keeps GPA1 (G α of G protein) in its inactive state [36–38]. Glucose-induced endocytosis of RGS1 releases its inhibition of G-protein self-activation which is triggered by phosphorylation at its C-terminal region by either the WNK kinases [38,39] or several LRR-RLKs [40–42]. Brassinosteroid insensitive 1 Like 3 (BRL3) and BRI1-associated Kinase 1 (BAK1) phosphorylates RGS1 promoting its endocytosis [41,42]. However, early seedling development and root growth of *rgs* mutants showed similar response to BL as wild type [41]. Recently, it was reported that glucose at low concentration increases the interaction between BRI1 and BAK1 in a manner dependent on BR biosynthesis [43], consistent with sugar increasing BR level in dark-treated seedlings [2]. Interestingly, high concentrations of glucose caused endocytosis of BRI1 and BAK1. Whether these cell surface receptors contribute to the sugar inhibition of BR signaling remains to be tested [43].

O-GlcNAc and O-fucose modifications have recently emerged as major sugar sensing mechanisms that impact on several hormonal pathways in plants [44,45]. Little is known about the interactions between these O-glycosylation pathways and BR signaling.

Our study indicates that light modulates sugar-BR crosstalk, however, the underlying mechanism remains unclear. Recently, several reports showed that light signaling inhibits BR signaling through photoreceptors regulating the activity of BZR1/BZR2 (BES1). Upon light activation, Phytochrome B, cryptochromes, and UVR8 bind with the dephosphorylated BZR1/BES1 and inhibit their DNA-binding activity [9,46–49]. CRY1 also interacts with BIN2 and enhances the interaction of BIN2 with BZR1 [47]. It would be very interesting to test in future

studies whether such direct interactions with phototransduction components rewire the crosstalk between sugar and BR pathways. The interactions between light and sugar pathways in modulating BR responses are important aspects of plant growth regulation that require further molecular investigation.

Materials and methods

Plant materials and growth conditions

Arabidopsis thaliana ecotype Columbia-0 (Col-0), *bzr1-1D* [12], *bzr1-1D-CFP*/Col-0, *BZR1-CFP*/Col-0 [30], *BZR1^{S173A}-CFP*/Col-0, Ler, *gin2-1* [25], *bzr1-ΔDM-GFP* [33], *BZR1-CFP/tor-es* [2] and *tor-es* [50] were all grown in a greenhouse with a 16-hr light/8-hr dark cycle at 22–24°C for general growth and seed harvesting. All the plants were in Col-0 ecotype background except that *gin2-1* is in Landsberg erecta ecotype and *bin2,bil1,bil2* is in Wassilewskija (Ws).

Sugar and BR Treatments and hypocotyl elongation assays

Seeds sterilized by 75% (v/v) ethanol were grown on solid 1/2 MS medium (pH 5.7) with 0.4% phytagel (Sigma) and 1% sucrose. After three days of incubation at 4°C and two days of germination in light at 22–24°C, seeds were transferred into 1/2 MS medium containing different concentrations of sugar and BR and were grown in continuous light for another 5 or 6 days as indicated. TOR silencing was induced in *tor-es* by adding 1 μ M β -estradiol (Sigma, E8875) in the medium. Pictures were taken and the hypocotyl lengths were measured with the Image J. Raw data is shown in S1 Data. Seedlings were harvested for Western Blot.

Primer design and Real-time quantitative PCR

Total RNA of above different sugar and BR treated seedlings was isolated by Spectrum Plant Total RNA Kit (Sigma-Aldrich, Shanghai, China). The primer sequences of BR responsive genes were listed in <u>S1 Table</u>. The qRT-PCR reactions were performed on QuantStudio 6 Flex Real-Time PCR System with TaKaRa Real-time qPCR Master Mix Kit. For each condition, the qRT-PCR experiments were performed with biological triplicates. Raw data is shown in <u>S1</u> Data.

Protein extraction and immunoblot analysis

For protein extraction, plants were frozen in liquid nitrogen, ground, weighed, and added into corresponding 2× SDS buffer (0.125 mM Tris-HCl [pH 6.8], 4% SDS, 20% Glycerol and 2% β -mercapto-ethanol). Samples were heated for 10 min at 95°C, centrifuged at 10000g for 10 min, separated on a 7.5% (detecting BZR1 protein) or 15% (detecting Histone 3H protein) acrylamide gel and then blotted on PVDF membranes (Millipore, IPVH0010) in 192 mM glycine and 25 mM Tris-HCl with a Trans-blot Turbo blotting system (Bio-Rad) for 13 min. Membranes were blocked for 1 hour at room temperature in a blotting buffer (140 mM NaCl, 10 mM KCl, 8 mM Na2HPO4, 2 mM KH2PO4, 0.5% skim milk, and 0.1% Tween20, pH 7.4). The gel blots were incubated with the primary antibodies (anti-GFP, Transgene, HT801, at 1:1000; anti-BZR1 and anti-BIN2 (home-made) at 1 µg/ml; anti-Actin (Sigma, A2228) at 1:5000 dilutions; anti-Histone H3 (Sangon BBI antibody, AB51007) at 1:2000 dilutions). The secondary antibodies were used at 1:5000 dilutions for 1 hour.

Supporting information

S1 Fig. Sucrose inhibits the BR-induced hypocotyl elongation. (A) Phenotypes of BZR1-CFP and *bzr1-1D*-CFP grown on media containing no sugar (-S), 30 to 150 mM sucrose (S) or mannitol (M), as in Fig 1. Bar = 5 mm. (B) Phenotypes of Col-0 grown on medium containing 0 (-BL) or 10 nM brassinolide (+BL) and 0 (-S) or 90 mM sucrose (+S) for six days. (C) Hypocotyl length of seedlings shown in panel B. Error bars indicate the standard error of the mean (SEM, three independent experiments, $n \ge 20$). **** *P*<0.0001 (Student's t test). (TIF)

S2 Fig. Sucrose inhibits the dephosphorylation of BZR1 protein induced by BR. Immunoblot analysis of BZR1 protein with BZR1 antibody in Col-0 grown on medium containing 0 (-BL) or 10 nM brassinolide (+BL) and 0 (-S) or 90 mM sucrose (+S) for six days. (TIF)

S3 Fig. Sucrose inhibits BR responses of BZR1 target genes. (A-C) Relative expression of *CPD*, *DWF4* or *SAUR-AC* analyzed by qRT-PCR in Col-0 seedlings grown on medium containing 0 (-BL) or 10 nM brassinolide (+BL) and 0 (-S) or 90 mM (+S) sucrose for six days. Error bars indicate the SEM (three independent experiments). ** P < 0.01, * P < 0.05 (Student's t test).

(TIF)

S4 Fig. Sucrose inhibits BR signaling independent of 14-3-3 protein. (A) Hypocotyl length of *BZR1*^{S173A}-*CFP*/Col-0 grown on medium containing 100 nM eBL and indicated concentrations of sucrose. (B) Immunoblot analysis of BZR1^{S173A}-CFP protein in seedlings in (A). Histone H3 was probed as a loading control in the immunoblot analysis. (TIF)

S5 Fig. Loss of BIN2 function enhances BR-induced BZR1 dephosphorylation in the presence of sugar. Immunoblot analysis of BZR1 protein in wild type and the *bin2-triple* mutant grown on media containing 90 mM sucrose and 0 (-BL) or 10 nM BL (+BL) for six days. The ratio of dephosphorylated BZR1 to phosphorylated BZR1 (BZR1/P-BZR1) was measured. (TIF)

S1 Table. List of primers used in this study. (XLSX)

S1 Data. Raw data for all quantitative assays shown in the manuscript. (XLSX)

Acknowledgments

We thank Dr. Jen Sheen for the mutant *gin2-1* and *tor-es*. We thank Marica Margis-Pinheiro and Tina Tingting Wang for editing the manuscript.

Author Contributions

Conceptualization: Zhenzhen Zhang. Data curation: Zhenzhen Zhang, Zhi-Yong Wang. Formal analysis: Zhenzhen Zhang. Funding acquisition: Zhenzhen Zhang, Wenfei Wang, Zhi-Yong Wang. Investigation: Zhenzhen Zhang, Xue Jiang, Wenfei Wang, Zhi-Yong Wang. Methodology: Zhenzhen Zhang, Wenfei Wang, Zhi-Yong Wang.

Project administration: Zhi-Yong Wang.

Visualization: Zhenzhen Zhang.

Writing - original draft: Zhenzhen Zhang, Zhi-Yong Wang.

Writing – review & editing: Zhenzhen Zhang, Ying Sun, Xue Jiang, Wenfei Wang, Zhi-Yong Wang.

References

- Wu Y, Shi L, Li L, Fu L, Liu Y, Xiong Y, et al. Integration of nutrient, energy, light, and hormone signalling via TOR in plants. J Exp Bot. 2019; 70(8):2227–38. Epub 2019/02/05. <u>https://doi.org/10.1093/jxb/</u> erz028 PMID: 30715492; PubMed Central PMCID: PMC6463029.
- Zhang Z, Zhu JY, Roh J, Marchive C, Kim SK, Meyer C, et al. TOR Signaling Promotes Accumulation of BZR1 to Balance Growth with Carbon Availability in Arabidopsis. Curr Biol. 2016; 26(14):1854–60. https://doi.org/10.1016/j.cub.2016.05.005 PMID: 27345161; PubMed Central PMCID: PMC5126233.
- Paparelli E, Parlanti S, Gonzali S, Novi G, Mariotti L, Ceccarelli N, et al. Nighttime sugar starvation orchestrates gibberellin biosynthesis and plant growth in Arabidopsis. Plant Cell. 2013; 25(10):3760–9. Epub 2013/10/08. https://doi.org/10.1105/tpc.113.115519 PMID: 24096343; PubMed Central PMCID: PMC3877829.
- Gupta A, Singh M, Laxmi A. Interaction between glucose and brassinosteroid during the regulation of lateral root development in Arabidopsis. Plant physiology. 2015; 168(1):307–20. <u>https://doi.org/10.1104/pp.114.256313 PMID: 25810094</u>; PubMed Central PMCID: PMC4424020.
- Mishra BS, Singh M, Aggrawal P, Laxmi A. Glucose and auxin signaling interaction in controlling Arabidopsis thaliana seedlings root growth and development. PloS one. 2009; 4(2):e4502. Epub 2009/02/19. https://doi.org/10.1371/journal.pone.0004502 PMID: <u>19223973</u>; PubMed Central PMCID: PMC2637607.
- Li J, Nagpal P, Vitart V, McMorris TC, Chory J. A role for brassinosteroids in light-dependent development of Arabidopsis. Science. 1996; 272(5260):398–401. Epub 1996/04/19. <u>https://doi.org/10.1126/science.272.5260.398</u> PMID: 8602526.
- Symons GM, Smith JJ, Nomura T, Davies NW, Yokota T, Reid JB. The hormonal regulation of de-etiolation. Planta. 2008; 227(5):1115–25. Epub 2008/01/25. https://doi.org/10.1007/s00425-007-0685-x PMID: 18214530.
- Oh E, Zhu JY, Wang ZY. Interaction between BZR1 and PIF4 integrates brassinosteroid and environmental responses. Nature cell biology. 2012; 14(8):802–9. https://doi.org/10.1038/ncb2545 PMID: 22820378; PubMed Central PMCID: PMC3703456.
- Liang T, Mei S, Shi C, Yang Y, Peng Y, Ma L, et al. UVR8 Interacts with BES1 and BIM1 to Regulate Transcription and Photomorphogenesis in Arabidopsis. Dev Cell. 2018; 44(4):512–23 e5. Epub 2018/ 02/06. https://doi.org/10.1016/j.devcel.2017.12.028 PMID: 29398622.
- Zhang Y, Liu Z, Wang J, Chen Y, Bi Y, He J. Brassinosteroid is required for sugar promotion of hypocotyl elongation in Arabidopsis in darkness. Planta. 2015; 242(4):881–93. https://doi.org/10.1007/ s00425-015-2328-y PMID: 25998528
- Kim TW, Wang ZY. Brassinosteroid signal transduction from receptor kinases to transcription factors. Annu Rev Plant Biol. 2010; 61:681–704. https://doi.org/10.1146/annurev.arplant.043008.092057 PMID: 20192752.
- Wang ZY, Nakano T, Gendron J, He J, Chen M, Vafeados D, et al. Nuclear-localized BZR1 mediates brassinosteroid-induced growth and feedback suppression of brassinosteroid biosynthesis. Dev Cell. 2002; 2(4):505–13. https://doi.org/10.1016/s1534-5807(02)00153-3 PMID: 11970900.
- Yin Y, Wang ZY, Mora-Garcia S, Li J, Yoshida S, Asami T, et al. BES1 accumulates in the nucleus in response to brassinosteroids to regulate gene expression and promote stem elongation. Cell. 2002; 109(2):181–91. Epub 2002/05/15. https://doi.org/10.1016/s0092-8674(02)00721-3 PMID: 12007405.
- Wang ZY, Bai MY, Oh E, Zhu JY. Brassinosteroid signaling network and regulation of photomorphogenesis. Annu Rev Genet. 2012; 46:701–24. https://doi.org/10.1146/annurev-genet-102209-163450 PMID: 23020777.
- Vert G, Chory J. Downstream nuclear events in brassinosteroid signalling. Nature. 2006; 441 (7089):96–100. https://doi.org/10.1038/nature04681 PMID: 16672972.

- Gampala SS, Kim TW, He JX, Tang W, Deng Z, Bai MY, et al. An essential role for 14-3-3 proteins in brassinosteroid signal transduction in Arabidopsis. Dev Cell. 2007; 13(2):177–89. https://doi.org/10. 1016/j.devcel.2007.06.009 PMID: 17681130; PubMed Central PMCID: PMC2000337.
- Kim TW, Guan S, Sun Y, Deng Z, Tang W, Shang JX, et al. Brassinosteroid signal transduction from cell-surface receptor kinases to nuclear transcription factors. Nature cell biology. 2009; 11(10):1254– 60. https://doi.org/10.1038/ncb1970 PMID: 19734888; PubMed Central PMCID: PMC2910619.
- Zhu JY, Li Y, Cao DM, Yang H, Oh E, Bi Y, et al. The F-box Protein KIB1 Mediates Brassinosteroid-Induced Inactivation and Degradation of GSK3-like Kinases in Arabidopsis. Mol Cell. 2017; 66(5):648– 57.e4. Epub 2017/06/03. https://doi.org/10.1016/j.molcel.2017.05.012 PMID: 28575660; PubMed Central PMCID: PMC5935450.
- Tang W, Yuan M, Wang R, Yang Y, Wang C, Oses-Prieto JA, et al. PP2A activates brassinosteroidresponsive gene expression and plant growth by dephosphorylating BZR1. Nature cell biology. 2011; 13(2):124–31. Epub 2011/01/25. <u>https://doi.org/10.1038/ncb2151</u> PMID: <u>21258370</u>; PubMed Central PMCID: PMC3077550.
- Oh E, Zhu JY, Ryu H, Hwang I, Wang ZY. TOPLESS mediates brassinosteroid-induced transcriptional repression through interaction with BZR1. Nat Commun. 2014; 5:4140. <u>https://doi.org/10.1038/ncomms5140</u> PMID: 24938363; PubMed Central PMCID: PMC4232713.
- Kim H, Shim D, Moon S, Lee J, Bae W, Choi H, et al. Transcriptional network regulation of the brassinosteroid signaling pathway by the BES1-TPL-HDA19 co-repressor complex. Planta. 2019; 250 (4):1371–7. Epub 2019/07/08. https://doi.org/10.1007/s00425-019-03233-z PMID: 31280329.
- Oh E, Zhu JY, Bai MY, Arenhart RA, Sun Y, Wang ZY. Cell elongation is regulated through a central circuit of interacting transcription factors in the Arabidopsis hypocotyl. Elife. 2014;3. <u>https://doi.org/10.7554/eLife.03031</u> PMID: 24867218; PubMed Central PMCID: PMC4075450.
- Bai MY, Shang JX, Oh E, Fan M, Bai Y, Zentella R, et al. Brassinosteroid, gibberellin and phytochrome impinge on a common transcription module in Arabidopsis. Nature cell biology. 2012; 14(8):810–7. https://doi.org/10.1038/ncb2546 PMID: 22820377; PubMed Central PMCID: PMC3606816.
- Chaiwanon J, Wang W, Zhu JY, Oh E, Wang ZY. Information Integration and Communication in Plant Growth Regulation. Cell. 2016; 164(6):1257–68. https://doi.org/10.1016/j.cell.2016.01.044 PMID: 26967291; PubMed Central PMCID: PMC5126258.
- Moore B, Zhou L, Rolland F, Hall Q, Cheng WH, Liu YX, et al. Role of the Arabidopsis glucose sensor HXK1 in nutrient, light, and hormonal signaling. Science. 2003; 300(5617):332–6. <u>https://doi.org/10.1126/science.1080585</u> PMID: 12690200.
- 26. Nolan TM, Brennan B, Yang M, Chen J, Zhang M, Li Z, et al. Selective Autophagy of BES1 Mediated by DSK2 Balances Plant Growth and Survival. Dev Cell. 2017; 41(1):33–46.e7. Epub 2017/04/12. https://doi.org/10.1016/j.devcel.2017.03.013 PMID: 28399398; PubMed Central PMCID: PMC5720862.
- Kim EJ, Lee SH, Park CH, Kim SH, Hsu CC, Xu S, et al. Plant U-Box40 Mediates Degradation of the Brassinosteroid-Responsive Transcription Factor BZR1 in Arabidopsis Roots. Plant Cell. 2019; 31 (4):791–808. Epub 2019/03/01. https://doi.org/10.1105/tpc.18.00941 PMID: 30814258; PubMed Central PMCID: PMC6501603.
- Yang M, Li C, Cai Z, Hu Y, Nolan T, Yu F, et al. SINAT E3 Ligases Control the Light-Mediated Stability of the Brassinosteroid-Activated Transcription Factor BES1 in Arabidopsis. Dev Cell. 2017; 41(1):47– 58.e4. Epub 2017/04/12. https://doi.org/10.1016/j.devcel.2017.03.014 PMID: 28399399; PubMed Central PMCID: PMC6283279.
- Srivastava M, Srivastava AK, Orosa-Puente B, Campanaro A, Zhang C, Sadanandom A. SUMO Conjugation to BZR1 Enables Brassinosteroid Signaling to Integrate Environmental Cues to Shape Plant Growth. Curr Biol. 2020. Epub 2020/02/29. https://doi.org/10.1016/j.cub.2020.01.089 PMID: 32109396.
- He JX, Gendron JM, Sun Y, Gampala SS, Gendron N, Sun CQ, et al. BZR1 is a transcriptional repressor with dual roles in brassinosteroid homeostasis and growth responses. Science. 2005; 307 (5715):1634–8. https://doi.org/10.1126/science.1107580 PMID: 15681342; PubMed Central PMCID: PMC2925132.
- Moorhead G, Douglas P, Cotelle V, Harthill J, Morrice N, Meek S, et al. Phosphorylation-dependent interactions between enzymes of plant metabolism and 14-3-3 proteins. Plant J. 1999; 18(1):1–12. https://doi.org/10.1046/j.1365-313x.1999.00417.x PMID: 10341439.
- Cotelle V, Meek SE, Provan F, Milne FC, Morrice N, MacKintosh C. 14-3-3s regulate global cleavage of their diverse binding partners in sugar-starved Arabidopsis cells. EMBO J. 2000; 19(12):2869–76. Epub 2000/06/17. https://doi.org/10.1093/emboj/19.12.2869 PMID: 10856232; PubMed Central PMCID: PMC203364.
- Peng P, Zhao J, Zhu Y, Asami T, Li J. A direct docking mechanism for a plant GSK3-like kinase to phosphorylate its substrates. J Biol Chem. 2010; 285(32):24646–53. https://doi.org/10.1074/jbc.M110. 142547 PMID: 20522560; PubMed Central PMCID: PMC2915701.

- Xiong Y, McCormack M, Li L, Hall Q, Xiang C, Sheen J. Glucose-TOR signalling reprograms the transcriptome and activates meristems. Nature. 2013; 496(7444):181–6. https://doi.org/10.1038/ nature12030 PMID: 23542588; PubMed Central PMCID: PMC4140196.
- Baena-Gonzalez E, Rolland F, Thevelein JM, Sheen J. A central integrator of transcription networks in plant stress and energy signalling. Nature. 2007; 448(7156):938–42. <u>https://doi.org/10.1038/</u> nature06069 PMID: 17671505.
- 36. Johnston CA, Taylor JP, Gao Y, Kimple AJ, Grigston JC, Chen JG, et al. GTPase acceleration as the rate-limiting step in Arabidopsis G protein-coupled sugar signaling. Proceedings of the National Academy of Sciences of the United States of America. 2007; 104(44):17317–22. https://doi.org/10.1073/pnas.0704751104 PMID: 17951432; PubMed Central PMCID: PMC2077254.
- Chen D, Juarez S, Hartweck L, Alamillo JM, Simon-Mateo C, Perez JJ, et al. Identification of secret agent as the O-GlcNAc transferase that participates in Plum pox virus infection. J Virol. 2005; 79 (15):9381–7. Epub 2005/07/15. https://doi.org/10.1128/JVI.79.15.9381-9387.2005 PMID: 16014901; PubMed Central PMCID: PMC1181581.
- Urano D, Phan N, Jones JC, Yang J, Huang J, Grigston J, et al. Endocytosis of the seven-transmembrane RGS1 protein activates G-protein-coupled signalling in Arabidopsis. Nature cell biology. 2012; 14 (10):1079–88. https://doi.org/10.1038/ncb2568 PMID: 22940907; PubMed Central PMCID: PMC3463750.
- Fu Y, Lim S, Urano D, Tunc-Ozdemir M, Phan NG, Elston TC, et al. Reciprocal encoding of signal intensity and duration in a glucose-sensing circuit. Cell. 2014; 156(5):1084–95. https://doi.org/10.1016/j.cell. 2014.01.013 PMID: 24581502; PubMed Central PMCID: PMC4364031.
- Tunc-Ozdemir M UD, Jaiswal D K, et al. Direct modulation of heterotrimeric G protein-coupled signaling by a receptor kinase complex. Journal of Biological Chemistry. 2016; 291(27):13918–25. <u>https://doi.org/ 10.1074/jbc.C116.736702</u> PMID: 27235398
- Tunc-Ozdemir M, Jones AM. BRL3 and AtRGS1 cooperate to fine tune growth inhibition and ROS activation. PloS one. 2017; 12(5):e0177400. https://doi.org/10.1371/journal.pone.0177400 PMID: 28545052; PubMed Central PMCID: PMC5436702.
- Tunc-Ozdemir M, Jones AM. Ligand-induced dynamics of heterotrimeric G protein-coupled receptorlike kinase complexes. PloS one. 2017; 12(2):e0171854. https://doi.org/10.1371/journal.pone.0171854 PMID: 28187200; PubMed Central PMCID: PMC5302818.
- Peng Y, Chen L, Li S, Zhang Y, Xu R, Liu Z, et al. BRI1 and BAK1 interact with G proteins and regulate sugar-responsive growth and development in Arabidopsis. Nat Commun. 2018; 9(1):1522. Epub 2018/ 04/20. https://doi.org/10.1038/s41467-018-03884-8 PMID: 29670153; PubMed Central PMCID: PMC5906681.
- Zentella R, Sui N, Barnhill B, Hsieh WP, Hu J, Shabanowitz J, et al. The Arabidopsis O-fucosyltransferase SPINDLY activates nuclear growth repressor DELLA. Nat Chem Biol. 2017; 13(5):479–85. Epub 2017/03/01. https://doi.org/10.1038/nchembio.2320 PMID: 28244988; PubMed Central PMCID: PMC5391292.
- 45. Xu SL, Chalkley RJ, Maynard JC, Wang W, Ni W, Jiang X, et al. Proteomic analysis reveals O-GlcNAc modification on proteins with key regulatory functions in Arabidopsis. Proceedings of the National Academy of Sciences of the United States of America. 2017; 114(8):E1536–e43. Epub 2017/02/06. <u>https://doi.org/10.1073/pnas.1610452114</u> PMID: 28154133; PubMed Central PMCID: PMC5338445.
- 46. Wang W, Lu X, Li L, Lian H, Mao Z, Xu P, et al. Photoexcited CRYPTOCHROME1 Interacts with Dephosphorylated BES1 to Regulate Brassinosteroid Signaling and Photomorphogenesis in Arabidopsis. Plant Cell. 2018; 30(9):1989–2005. Epub 2018/08/23. https://doi.org/10.1105/tpc.17.00994 PMID: 30131420; PubMed Central PMCID: PMC6181010.
- He G, Liu J, Dong H, Sun J. The Blue-Light Receptor CRY1 Interacts with BZR1 and BIN2 to Modulate the Phosphorylation and Nuclear Function of BZR1 in Repressing BR Signaling in Arabidopsis. Mol Plant. 2019; 12(5):689–703. Epub 2019/02/15. https://doi.org/10.1016/j.molp.2019.02.001 PMID: 30763615.
- Wu J, Wang W, Xu P, Pan J, Zhang T, Li Y, et al. phyB Interacts with BES1 to Regulate Brassinosteroid Signaling in Arabidopsis. Plant Cell Physiol. 2019; 60(2):353–66. Epub 2018/11/06. <u>https://doi.org/10.1093/pcp/pcy212</u> PMID: 30388258.
- Dong H, Liu J, He G, Liu P, Sun J. Photoexcited phytochrome B interacts with brassinazole resistant 1 to repress brassinosteroid signaling in Arabidopsis. J Integr Plant Biol. 2020; 62(5):652–67. Epub 2019/ 05/14. https://doi.org/10.1111/jipb.12822 PMID: 31081597.
- Xiong Y, Sheen J. Rapamycin and glucose-target of rapamycin (TOR) protein signaling in plants. J Biol Chem. 2012; 287(4):2836–42. https://doi.org/10.1074/jbc.M111.300749 PMID: 22134914; PubMed Central PMCID: PMC3268441.