



Corrigendum: Phenotypic, Transcriptomic, and Metabolomic Signatures of Root-Specifically Overexpressed *OsCKX2* in Rice

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1

A Corrigendum on

Phenotypic, Transcriptomic, and Metabolomic Signatures of Root-Specifically Overexpressed *OsCKX2* in Rice

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In the original article, there was a mistake in Figure 1 as published. The corrected Figure 1 appears below.

In the original article, there was a mistake in **Supplementary Table 2** as published. The corrected **Supplementary Table 2** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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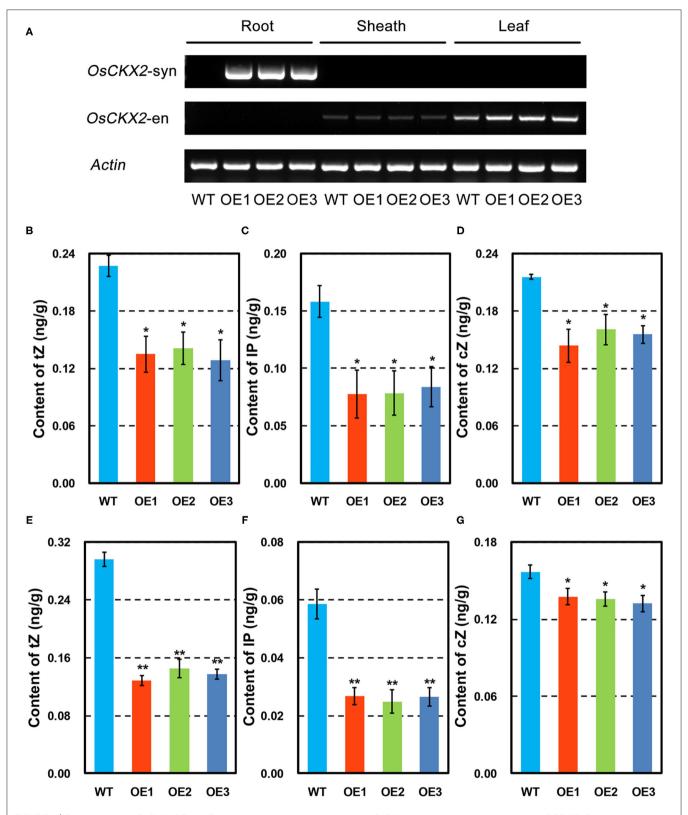


FIGURE 1 | Characteristics of OsCKX2 OE rice. Expression of synthetic and endogenous OsCKX2 in root, sheath, and leaf by semi-qPCR **(A)**. Contents of cytokinins in different forms of tZ, IP, and cZ at seedling stage **(B-D)** and mature stage **(E-G)**. Data represents means \pm SEM (n=3). $^*P < 0.05$, $^*P < 0.05$.

SUPPLEMENTARY TABLE 2 | List of primers used in this study.

Name	Forward primer (5'-3')	Reverse primer (5'-3')
OsCKX2-syn	GAGCAGTGCGTCCT	GCTTGCCCTCAAAAGTGC
OsCKX2-en	TTCGTCCGCCTCCTTCCT	CGACGCGCAGCAGCGC
Actin	GGAAGTACAGTGTCTGGATTGGAG	TCTTGGCTTAGCATTCTTGGGT
OsPSK4	GCCGTGCTGATTTTC	GTGATGCTGCTGGGTGTAGA
DsPUP6	TCCCTGATGCAGCTCACGTT	TCGCCCTTCTTGTACCCGTC
BD11-1	CCAGAACCAAGTCTCCCAGC	TCCACATGGACTCTTTCTTGAGG
DsPsbR2	GACCGAACCTGAAAGACGGT	CCAGCCAGCAGAATTCCAGA
DsMST1	TGACGTTCTCGGTGGTCATC	TAGACGCAGTACTCGTTCCC
DsPME22	ACTCGCTTCGCCAGTTCTAC	CTGGGGTGTATCAGGCTGTC
DsPGL21	AACTTGGCAGGGAGGTTCAG	CATGCAAAATGGGCAGGCTT
DsAP25	CCGAACTACACGTTCGGGTG	AGCGAGCCGGAGAAGTAGTA
DsSub31	GCTTACAGCGCCATGGAAAG	CCGGAGTAGTTCTTGCCGTT
OsBGal1	AGGAACCATCCGTCAACGAC	AGTCACAGTTGGATCGGCAG