

Fig. S1. Flowering phenotypes of Nipponbare and the *OsFLZ*2 transgenic plants under natural long-day conditions (NLD, from mid-April to July, Guangzhou).

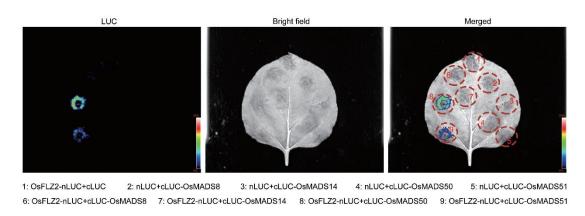


Fig. S2. The LCI assay showing the interaction of OsFLZ2 and OsMADS8/14/50/51 in the leaves of *N. benthamiana*.

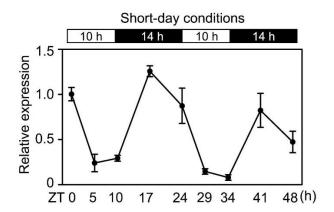


Fig. S3. qRT-PCR analysis of the diurnal expression pattern of OsFLZ2 under short-day conditions (10 h light). Penultimate leaves were collected at the indicated time points during a 48-h period of 14-d-old *Nipponbare* plants. *Ubiquitin* gene was used as an internal control. The expression level was set as 1.0 at ZT0. Values are means $\pm SD$ (n = 3). The white bars and black bars at the top represent the light and dark periods, respectively. ZT, Zeitgeber time.

 Table S1. Complete list of potential interactors of OsFLZ2 identified in IP-MS assay.

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Table S2. Primers used in this study

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