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Author Correction: Genome-wide identification of *Arabidopsis* long noncoding RNAs in response to the blue light

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This Article contains errors in the Reference list. References 28 and 29 are incorrectly listed as references 30 and 31 respectively. As a result, references 30 and 31 are listed incorrectly as references 28 and 29. The correct references 28–31 are listed below as references 1–4 respectively.

1. Kong, L. et al. Cpc: Assess the Protein-Coding Potential of Transcripts Using Sequence Features and Support Vector Machine. *Nucleic Acids Res.* **35**, W345–W349 (2007).
2. Sun, K. et al. Iseerna: Identification of Long Intergenic Non-Coding Rna Transcripts from Transcriptome Sequencing Data. *BMC Genomics.* **14**(Suppl 2), S7 (2013).
3. Kim, D., Langmead, B. & Salzberg, S.L. Hisat: A Fast Spliced Aligner with Low Memory Requirements. *Nat. Methods.* **12**, 357–360 (2015).
4. Pollier, J., Rombauts, S. & Goossens, A. Analysis of Rna-Seq Data with Tophat and Cufflinks for Genome-Wide Expression Analysis of Jasmonate-Treated Plants and Plant Cultures. *Methods Mol Biol.* **1011**, 305–315 (2013).



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