



# Corrigendum: Transcriptomics Identifies Modules of Differentially Expressed Genes and Novel Cyclotides in *Viola pubescens*

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## OPEN ACCESS

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## A Corrigendum on

### Transcriptomics Identifies Modules of Differentially Expressed Genes and Novel Cyclotides in *Viola pubescens*

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In our original article, there was a mistake in the legend for **Figure 1** as published. The descriptions of the figure labels (A) and (B) were switched. The correct legend appears below.

“*Viola pubescens* var. scabriuscula bearing (A) cleistogamous and (B) chasmogamous flowers. Photographs were taken over native populations located in Sells Park, Athens County, Ohio, 45701 (39°20′40.6″N 82°04′31.9″W).”

Additionally, there was a mistake in **Table 1** as published. In the first column, “# of scaffolds > 10 kbp” should be “# of scaffolds > 1 K nt.” The corrected **Table 1** appears below.

The authors apologize for these errors and state that they do not change the scientific conclusions of the article in any way. The original article has been updated.

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**TABLE 1** | Summary statistics of the *V. pubescens* genome assembly via ABySS and CEGMA and BUSCO assessments of genome completeness.

|   |   |
|---|---|
| # of scaffolds  | 157,722   |
| Total size of scaffolds (bp)                                  | 318,370,682   |
| Longest scaffold (bp)   | 86,685  |
| # of scaffolds > 1 K nt                                       | 80,885  |
| N50 length (bp)   | 3,500   |
| Average length of break (>25 N's) between contigs in scaffold | 45  |
| Scaffold %N   | 0.06  |
| Percent in scaffolded contigs                                 | 3.2   |
| Percent in unscaffolded contigs                               | 96.8  |
| CEGMA Partial (%)   | 94, <i>n</i> = 248  |
| CEGMA Complete (%)  | 76, <i>n</i> = 248  |
| BUSCO   | C: 79.7% [D:13.5%], F: 8.9%,<br>M: 11.4%, <i>n</i> = 2121 |

C, complete; D, duplicated; F, fragmented; M, missing; *n*, gene number.