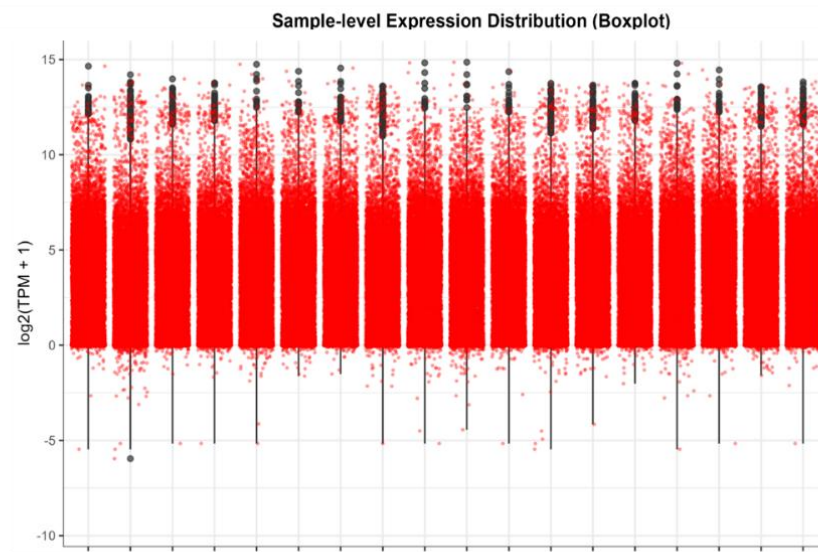
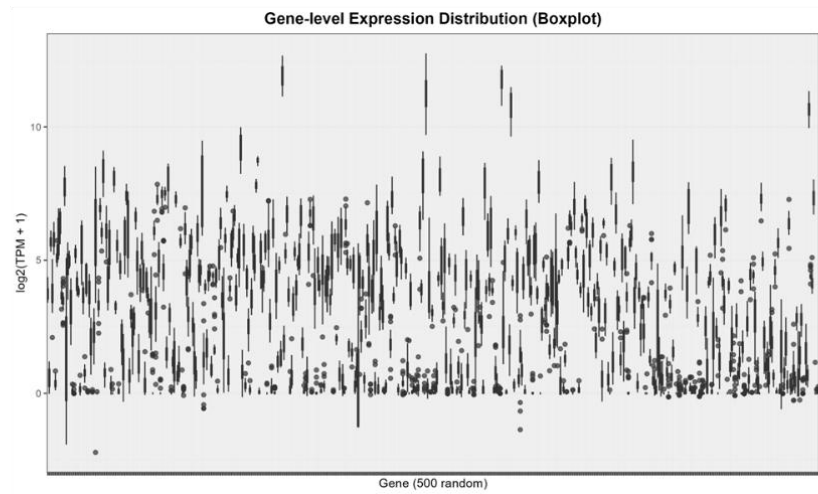


**Figure S1**

**A**



**B**



**Figure S1** The sample-level (A) and gene-level (B) expression distribution after data clean.

# Figure S2

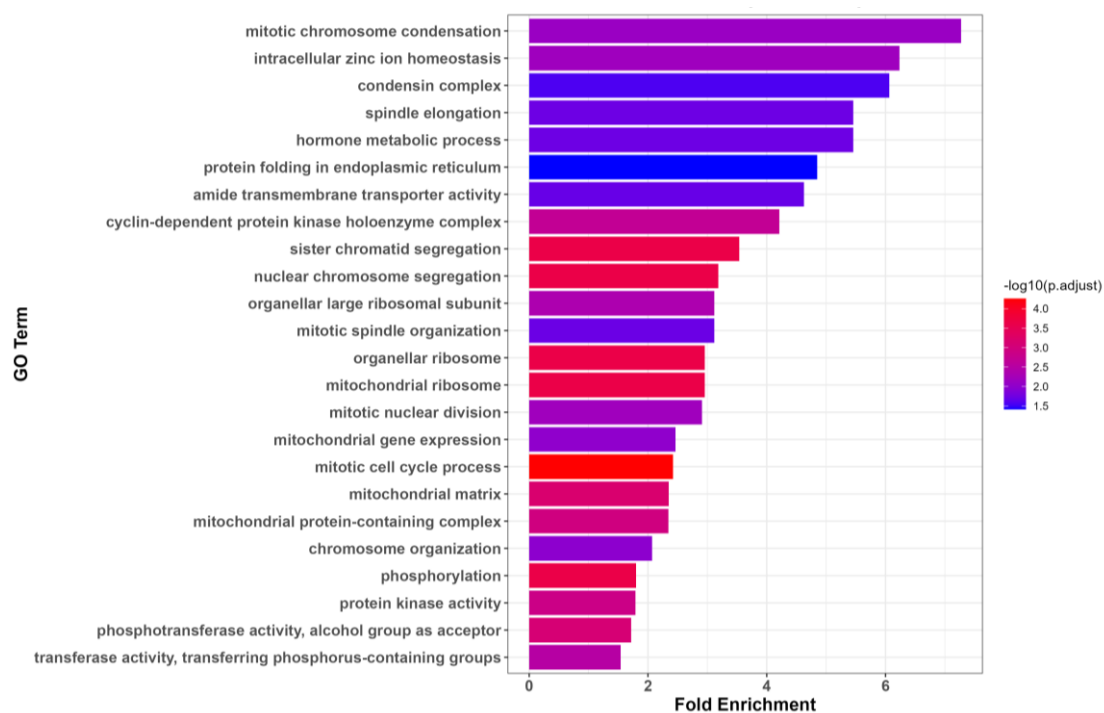


Figure S2 GO enrichment analysis of the consistently differentially expressed genes across all three time points

Figure S3

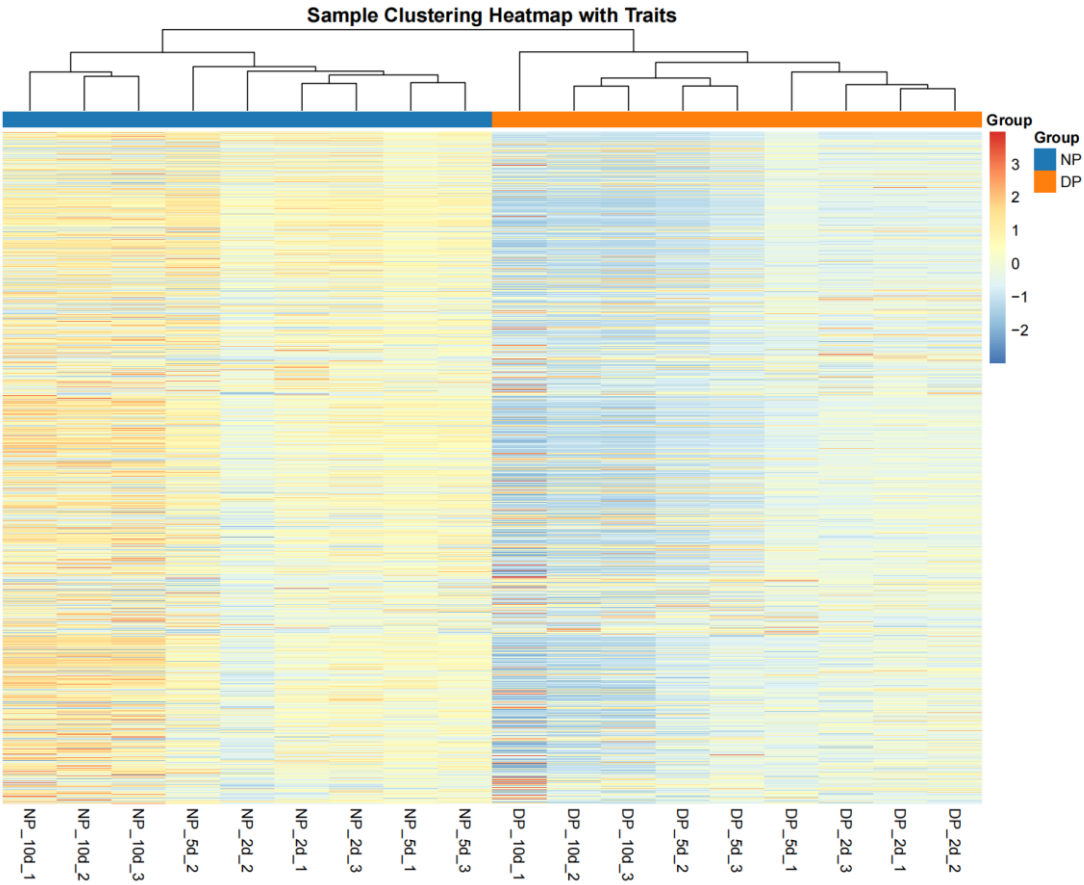


Figure S3 The sample clustering heatmap with traits.

## Table.S1

**Table.S1 Primers used in this work**

|          |                                 |
|----------|---------------------------------|
| TAF5L -F | 5'-GGTAAAAAATGAGCTTCTTGGATT-3'  |
| TAF5L -R | 5'-GATTGCCGCTTTCAGGTCCTGTAA-3'  |
| PLK4-F   | 5'-GAAGATGCACATTATGTTTATTTAG-3' |
| PLK4-R   | 5'-TCCATTGTACAAGAAGGGATGTTCT-3' |
| DDX5-F   | 5'-GAAGAAGGAGGGTTTCCAGATTA-3'   |
| DDX5-R   | 5'-GCCAGTGGTGTGGCGATGACTAT-3'   |
| Actin-F: | 5'-CAACGGCATCCACGAGACCAC-3'     |
| Actin-R: | 5'-GGGCTGTGATCTCCTTCTGC-3'      |