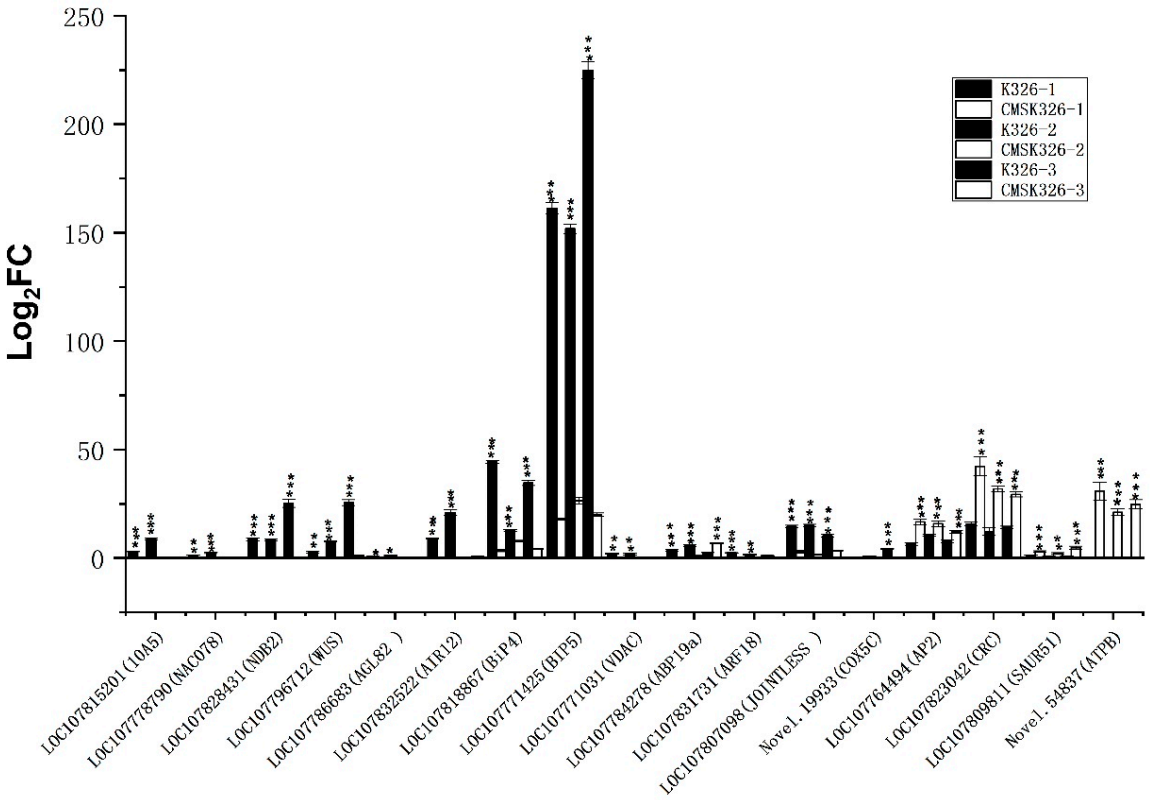
Table S1. Genes and their primers used in RT-PCR.

Gene ID	Forward primer	Reverse primer	Gene Name
LOC107815201	GGTCCTTTGGTCCTCCCTTG	GGAAAGCTCATTCATGGCAACA	*10A5*
LOC107778790	TCACGGCTTCCTAGTGTTCG	CAGAAATCTCCCAGGGTGGC	*NAC078*
LOC107828431	CGCAACAGCAAAACTCAGGT	AGCGTTTACGAATTCTTACATGGAC	*NDB2*
LOC107796712	GGAAGCTGCCCAACAACAAA	ACCACCAACACCATCTTCAATAAT	*WUS*
LOC107786683	TGGTGAAAGGACAAATATTCCTAGT	TCTTCAACTCAAAACGGCCA	*AGL82*
LOC107832522	TGGCAAATTGGGCCAGTAGT	CATGAGTAGGGGCAATGGCA	*AIR12*
LOC107818867	AAGAAAGTTGATAAGTGGAACTGGT	ACGATCAAGGATGTGCGTCT	*BiP4*
LOC107771425	TCCTTGTGGACCTACGAGCTA	AACAAGCTCCCGAACAGGAC	*BiP5*
LOC107771031	TCTCTCACGGCTCGGTCTAA	ATGCCCCTGAGAGAGACTGT	*VDAC*
LOC107784278	TCAAGTGGCACTCAATTTGCTC	GCGACGCAGAAATCTTGGAC	*ABP19a*
LOC107831731	AGGCACTTCAACTTTCTTCACTT	GCAAAAGCATCTGAAAACCAGA	*ARF18*
LOC107807098	TACAACAACCCAGCAGCAGTAT	GGGCAACCCCAACTTGAGAG	*JOINTLESS*
Novel.19933	CCTCTCCCATGTCAACTCGG	ACGAAGTCAGTTCCACAACCA	*COX5C*
LOC107764494	CCCAAGGCTCTTTTGGTTGG	TGGACAAATGCGGAGGAGAC	*AP2*
LOC107823042	CTTGTCTCGTTCGCAACCTTC	GGATGCAACTTAAAACCAGGAGT	*CRC*
LOC107809811	TCACAAACAGCAATGTTGAAGCA	TGGAAGCCCTAATTGCTCGT	*SAUR51*
Novel.54837	TTTAGCCCCTTATCGCCGTG	CTTCCCGAGTACGTTCACCC	*ATPB*
LOC107804067	CCCCTCAATCGTGCATAGGA	GAGACCAGAACACAGTCCACAA	*Actin*

Figure S1. Validation of DEGs by RT-PCR.



Note: Significance was analyzed using Student's t test, and Error bars show the SD (n=3). All data are shown as means ± standard deviations (SDs) (n=3). ***significant at $p \leq 0.001$, **significant at $p \leq 0.01$, *significant at $p \leq 0.05$.