**Supplemental Table**

Supplemental Table 1 Prime names and sequences

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| Primer name | DNA sequences and restriction sites |
| NAC13F1 | 5’-CCAAGAAGCAATTGTGCGCTAAGC-3’ |
| NAC13R1 | 5’-GCATCTAATCCTCTAGCAACCCAG-3’ |
| NAC13F2 | 5’GGGTCGACTGACTAGTATGGGAGTGCAAGAAACAGACC-3’ |
| NAC13R2 | 5’-TGCTCACCATACTAGTCTGATTAAACCCGAACCCAC-3’ |
| NAC13F3 | 5’-GAATTCATGGGAGTGCAAGAAACAGACC-3’, |
| NAC13R3 | 5’- GTCGACTTACTGATTAAACCCGAACCCA-3’ |
| NAC13F4 | 5’GCGTCTAGAGGTGGCGACGACTCCTGGAGCCCGCCAAGAAGCAATTGTGCGCTAAGC-3’ |
| NAC13R4 | 5’GCGGAGCTCCATTACCAGTTGGTCTGGTGTCAAGCATCTAATCCTCTAGCAACCCAG-3’ |
| NAC13aF | 5’-GAATTCATGGGAGTGCAAGAAACAGACC-3’, |
| NAC13aR | 5’-GTCGACCATGTTTGCGAGAGGATTCAAT-3’, |
| NAC13bF | 5’-GAATTCATTGAATCCTCTCGCAAACATG-3’, |
| NAC13bR | 5’-GTCGACTTACTGATTAAACCCGAACCCA-3’ |

The primers of NAC13 are used to amplify the coding sequence of *NAC13* gene. NAC13F1 and NAC13R1 are used to clone the ORF sequence of *NAC13* gene; NAC13F2 and NAC13R2 are used to generate the recombinant construct 35S::NAC13-GFP, and the underlined area is the restriction site *Spe* I; NAC13F3 and NAC13R3, NAC13aF and NAC13aR, NAC13bF and NAC13bR are used to generate bait vector pGBKT7-NAC13, pGBKT7-NAC13a and pGBKT7-NAC13b, respectively. The underlined areas are *EcoR* I and *Sal* I restriction sites; NAC13F4 and NAC13R4 are used to amplify the complete cDNA sequence of *NAC13* gene and introduce into the plant binary vector pBI121. The underlined areas are restriction sites *Xba* I and *Sac* I.