According to the published sequences in Amaranthaceae or Caryophyllales, the partial sequences of the six candidate reference genes were obtained from homology cloning. These sequences have not been added to the GenBank database.

(1)*Sa18SrRNA*\_1\_\_(T150120-0216)RV-M\_A77616\_D06\_1504063341J.seq (256bp)

GCGGCTTAATTTGACTCAACACGGGGAAACTTACCAGGTCCAGACATAGTAAGGATTGACAGACTGAGAGCTCTTTCTTGATTCTATGGGTGGTGGTGCATGGCCGTTCTTAGTTGGTGGAGCGATTTGTCTGGTTAATTCCGTTAACGAACGAGACCTCAGCCTGCTAACTAGCTATGCGGAGGTAGTCCTTCGCAGCTAGCTTCTTAGAGGGACTATGGCCTTTTAGGCCACGGAAGTTTGAGGCAATAACAGG

(2)*Sa28SrRNA*\_4\_\_(T150120-0216)RV-M\_A77616\_D04\_1505062346J.seq (154bp)

GCCGACCCTGATCTTCTGTGAAGGGTTCGAGTGAGAGCATACCTGTCGGGACCCGAAAGATGGTGAACTATGCCTGAGCGGGGCGAAGCCAGAGGAAACTCTGGTGGAGGCCCGAAGCGATACTGACGTGCAAATCGTTC GTCTGACTTGGGTA

(3)*SaACTIN*\_3\_\_(T150120-0216)RV-M\_A77616\_H03\_1505062342J.seq (149bp)

CCAAAGGCCAACAGAGAGAAGATGACCCAGATCATGTTTGAAACCTTCAATGTTCCAGCCATGTACGTTGCCATCCAGGCTGTTCTTTCTCTTTACGCTAGTGGTCGTACCACAGGTATTGTGCTTGATTCTGGTGATGGTGTGTCTCA

(4)*SaTUB*\_2\_\_(T150120-0216)RV-M\_A77616\_F05\_1504092476J.seq (368bp)

GGACTCTCAAGCTAACTACACCAAGCTTTGGTGATTTGAACCATTTGATATCGGCCACCATGTCTGGAGTAACATGTTGTCTGAGGTTCCCAGGTCAATTAAACTCTGACCTTAGGAAACTTGCAGTCAACCTTATTCCATTCCCCAGGCTTCATTTCTTCATGGTGGGTTTCGCCCCACTAACATCTCGTGGGTCTCAGCAATACCGCTCCTTGACAGTCCCTGAGCTAACCCAGCAAATGTGGGATTCCAAGAACATGATGTGTGCCGCAGACCCTAGACATGGGCGTTACTTGACGGCTTCTGCCATGTTCCGAGGAAAGATGAGCACAAAGGAGGTTGATGAGCAGATGATCAACGTGCAGAAC

(5)*SaGAPDH*\_10\_\_(T150120-0216)RV-M\_A77616\_H05\_1504092478J.seq (808bp)

GAGCTTGTTGCTGTTAACGATCCTTTCATCACCACTGACTACATGACATACATGTTTAAGTACGACAGTGTTCACGGTCAATGGAAACACCATGAGTTGAAGGTTAAGGATGAGAAGACTCTTCTTTTTGGTGACAAGCCAGTTACCGTATTCGGTTGCAGGAACCCAGAGGATATCCCATGGAGTTCGACTGGAGCTGACTTTGTTGTTGAGTCTACTGGTGTTTTCACTGACAAGGACAAGGCTGCTGCTCATTTGAAGGGTGGTGCCAAGAAGGTTGTCATATCTGCTCCTAGCAAGGATGCTCCCATGTTCGTCGTTGGTGTCAATGAGCATGAGTACAAGTCGGATCTTCACATTGTCTCCAATGCCAGTTGTACCACCAATTGCCTTGCTCCTTTGGCCAAGGTTATAAATGACAAGTTTGGCATCGTTGAGGGTCTTATGACCACTGTCCACTCCATCACTGCCACCCAAAAGACTGTTGACGGTCCATCAGCTAAGGACTGGCGAGGTGGAAGGGCTGCCTCATTTAACATCATTCCCAGCAGCACTGGAGCAGCTAAGGCTGTGGGAAAGGTTTTGCCTGCCCTCAATGGAAAATTGACTGGAATGGCTTTCCGTGTTCCAACTGTTGATGTTTCTGTTGTTGATCTTACTGTTAGACTTGAGAAGGCTGCTTCCTATGATCAGATTAAGGCTGCCATCAAGGAGGAGTCAGAGGGCAAGTTGAAGGGAGTTTTGGGTTACACCGACGAGGATGTTGTTTCCACTGATTTCATTGGTGACAACAGATCAAGTATTTTTG

(6)*SaUBQ*\_A10\_\_(T150120-0216)RV-M\_A77616\_D03\_1505062338J.seq (162bp)

GCTAAGATTCAAGACAAGGAGGGTATCCCACCAGACCAGCAGAGGCTTATTTTTGCTGGGAAACAGCTAGAGGAAGGTAGAACACTGGCAGACTATAACATTCAGAAAGAATCAACCCTCCACCTGGTTCTTCGCCTTAGGGGAGGTATGCAGATTTTCGTG