

Evaluating Precision and Accuracy When Quantifying Different Endogenous Control Reference Genes in Maize Using Real-Time PCR [*J. Agric. Food Chem.* **2009**, *57*, 2903. DOI: 10.1021/jf803599t]. Tandace A. Scholdberg, Tim D. Norden, Daishia D. Nelson, AND G. Ronald Jenkins*

A corrected version of **Table 1** is given below.

Table 1. Primers and Probe Sequences for the Taxon-Specific Endogenous Control Reference Genes

| reference system name | endogenous control reference gene primer/probe sequence | amplicon (bp) |
|-----------------------------|--|---------------|
| high mobility group (HMG) | forward: 5'-TTGGA CTAGAAATCTCGTGCTGA-3' reverse: 5'-GCTACATAGGGAGCCTTGTCCT-3' probe: 5'-FAM-CAATCCACACAAACGCACGCGTA-TAMRA-3' | 79 |
| starch synthase (SSIIb-3) | forward: 5'-CCAATCCTTTGACATCTGTCC-3' reverse: 5'-GATCAGCTTTGGGTCCGGA-3' probe: 5'-FAM-AGCAAAGTCAGAGCGCTGCAATGCA-TAMRA-3' | 114 |
| invertase (IVR) | forward: 5'-CGCTCTGTACAAGCGTGC-3' reverse: 5'-GCAAAGTGTTGTGCTTGACC-3' probe: 5'-FAM-CACGTGAGAATTTCCGTCTACTCGAGCCT-TAMRA-3' | 104 |
| alcohol dehydrogenase (ADH) | forward: 5'-CGTCGTTTCCCATCTCTTCTCC-3' reverse: 5'-CCACTCCGAGACCCTCAGTC-3' probe: 5'-FAM-AATCAGGGCTCATTTTCTCGCTCCTCA-TAMRA-3' | 136 |
| starch synthase (SSIIb-2) | forward: 5'-CTCCCAATCCTTTGACATCTGC-3' reverse: 5'-TCGATTTCTCTTGGTGACAGG-3' probe: 5'-FAM-AGCAAAGTCAGAGCGCTGCAATGCA-TAMRA-3' | 151 |
| zein | forward: 5'-GCCATTGGGTACCATGAACC-3' reverse: 5'-AGGCCAACAGTTGCTGCAG-3' probe: 5'-FAM-AGCTTGATGGCGTGCCGTCCCT-TAMRA-3' | 104 |

On page 2905, ref 12 should be added to the sentence to read “Primers and probe sequences for the taxon-specific endogenous control reference genes used in this study are shown in **Table 1** (12, 28)”.

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