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CORRECTION

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′-TTGGACTAGAAATCTCGTGCTGA-3′ reverse: 5′-GCTACATAGGGAGCCTTGTCCT-3′ probe: 5′-FAM-CAATCCACACAAACGCACGCGTA-TAMRA-3′	79
starch synthase (SSIIb-3)	forward: 5′-CCAATCCTTTGACATCTGCTCC-3′ reverse: 5′-GATCAGCTTTGGGTCCGGA-3′ probe: 5′-FAM-AGCAAAGTCAGAGCGCTGCAATGCA-TAMRA-3′	114
invertase (IVR)	forward: 5′-CGCTCTGTACAAGCGTGC-3′ reverse: 5′-GCAAAGTGTTGTGCTTGGACC-3′ probe: 5′-FAM-CACGTGAGAATTTCCGTCTACTCGAGCCT-TAMRA-3′	104
alcohol dehydrogenase (ADH)	forward: 5′-CGTCGTTTCCCATCTCTTCCTCC-3′ reverse: 5′-CCACTCCGAGACCCTCAGTC-3′ probe: 5′-FAM-AATCAGGGCTCATTTTCTCGCTCCTCA-TAMRA-3′	136
starch synthase (SSIIb-2)	forward: 5′-CTCCCAATCCTTTGACATCTGC-3′ reverse: 5′-TCGATTTCTCTCTTGGTGACAGG-3′ probe: 5′-FAM-AGCAAAGTCAGAGCGCTGCAATGCA-TAMRA-3′	151
zein	forward: 5′-GCCATTGGGTACCATGAACC-3′ reverse: 5′-AGGCCAACAGTTGCTGCAG-3′ probe: 5′-FAM-AGCTTGATGGCGTGTCCGTCCCT-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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