

CORRECTION

Correction: A Novel Quantitative Hemolytic Assay Coupled with Restriction Fragment Length Polymorphisms Analysis Enabled Early Diagnosis of Atypical Hemolytic Uremic Syndrome and Identified Unique Predisposing Mutations in Japan

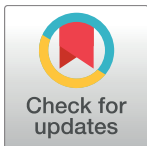
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There are two errors in the “Hemolytic assay using citrated human plasma and sheep RBCs” section of the Materials and Methods. The fourth sentence of the second paragraph should read: After incubation, the reaction was quenched by the addition of 1 ml of VBS-EDTA buffer (2.5 mM barbital, 1.5 mM sodium barbital and 144 mM NaCl, and 2 mM EDTA, pH 7.4).

The sixth sentence of the second paragraph should read: Plasma, diluted with AP-CFTD buffer containing 50 mM EDTA was treated in the same manner and used as a blank.

Reference

1. Yoshida Y, Miyata T, Matsumoto M, Shirotani-Ikejima H, Uchida Y, Ohyama Y, et al. (2015) A Novel Quantitative Hemolytic Assay Coupled with Restriction Fragment Length Polymorphisms Analysis Enabled Early Diagnosis of Atypical Hemolytic Uremic Syndrome and Identified Unique Predisposing Mutations in Japan. PLoS ONE 10(5): e0124655. doi:[10.1371/journal.pone.0124655](https://doi.org/10.1371/journal.pone.0124655) PMID: [25951460](https://pubmed.ncbi.nlm.nih.gov/25951460/)



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