Correction to “Identification of genetic polymorphisms of CYP2W1 in the three main Chinese ethnicities: Han, Tibetan, and Uighur”

Yanwei Li and Huijuan Wang, the senior authors of the above article [Li YW, Kang X, Yang G, Dai PG, Chen C, and Wang HJ (2016) Drug Metab Dispos 44(9):1510-5; DOI: https://doi.org/10.1124/dmd.115.069153] have recently identified two errors in data entry and analysis.

1. In the second row of Table 1, the SNP “v01, -717C>A” has a rsID “rs111601777”. Therefore, the SNP “v01, -717C>A” would not be considered as “novel”. Accordingly, the description in the text of “Results” should be corrected as: “A total of 26 different genetic variations including two novel ones in the CYP2W1 gene were detected in 150 individuals (Table 1). The two novel variants were all SNPs, located in the intron 7 (v21), and 3’ untranslated region (3’UTR) (v25), respectively.” In addition, in the third row of Table 1, the SNP “v02, -668G>A” has a rsID “rs1025780441”, instead of “rs6768161064”.

2. In Supplemental Table 2, the 10th, 11th, and 12th row of the first column, the superscript “b” should be replaced by “c”, i.e. these three data are from 1000 genomes (c).

The authors apologize for any inconvenience caused.