ERRATUM

Erratum to: Differential toxicity biomarkers for irinotecanand oxaliplatin- containing chemotherapy in colorectal cancer

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The author would like to correct the errors in the publication of the original article. The corrected details are given below for your reading.

In the abstract, the result section should read as:

Results In irinotecan-treated patients, T allele of ABCB1 C1236T SNP was associated with a lower risk of asthenia (OR = 0.047; 95 % CI = 0.004–0.493; P = 0.011) and T allele of ABCB1 C3435T SNP was associated with a lower risk of diarrhea (OR = 0.177; 95 % CI = 0.034–0.919; P = 0.039), and individuals with two copies of GSTT1 gene had a lower risk for asthenia (OR = 0.093; 95 % CI = 0.011–0.794; P = 0.030). In oxaliplatin-treated patients, carriers of one or two T variants of Asn118Asn ERCC1 SNP had a lower risk for neutropenia (OR = 0.205; 95 % CI = 0.061–0.690; P = 0.01).

In results section, last paragraph should read as:

All these associations were analyzed in a multivariate analysis by linear regression including other factors (sex,

performance status, and adjuvant or metastatic setting) as variables (Table 5). Three of the associations found in the sample of patients treated with irinotecan-based regimens were confirmed by this analysis: rs1128503 (ABCB1 1236) and CNV of GSTT1 with asthenia (CT/TT vs. CC: OR = 0.047; 95 % CI = 0.004-0.493; P = 0.011; SP =91.2 % and OR = 0.093; 95 % CI = 0.011-0.794; P = 0.030; SP = 85.7 %, respectively), and rs1045642 (ABCB1 3435) with diarrhea (CT/TT vs. CC: OR = 0.177; 95 % CI = 0.034-0.919; P = 0.039; SP = 65.7 %). In patients treated with oxaliplatin-containing regimens, the association between rs11615 (ERCC1) with neutropenia was confirmed in the multivariate analysis (CT/TT vs. CC: OR = 0.205; 95 % CI = 0.061-0.690; P = 0.010; SP = 79.2 %).

In discussion, first sentence of the fifth paragraph should read as:

The multivariate analysis also revealed a lower risk of severe asthenia in patients with two copies of *GSTT1* than in patients with the null genotype or one copy of the gene in patients receiving an irinotecan-based regime (OR = 0.093; 95 % CI = 0.011-0.794; P = 0.030).

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