ERRATUM

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Erratum to: Meta-analytic support vector machine for integrating multiple omics data

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Erratum

After publication of the original article [1], the author noticed errors to Table 1. The errors occurred in the fourth and seventh columns labelled 'Youden'. The correct version of Table 1 is included in this erratum.

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References
1. Kim SH, Jhong JH, Lee JJ, Koo JY. Meta-analytic support vector machine for integrating multiple omics data. BioData Mining. 2017;10:2. doi:10.1186/s13040-017-0126-8.

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Table 1 Shown are the results of experimental studies to compare the meta-logistic model with the meta-analytic SVM

Variance (R)	Meta-SVM			Meta-logistic regression		
	Sensitivity (SE)	Specificity (SE)	Youden	Sensitivity (SE)	Specificity (SE)	Youden
	No inclusion of random study					
0.1	0.828 (0.001)	0.9843 (0)	0.812	0.1073 (0)	1 (0)	0.107
0.3	0.8127 (0.002)	0.8707 (0.001)	0.683	0.2087 (0.001)	0.996 (0)	0.205
0.5	0.76 (0.002)	0.867 (0.001)	0.627	0.2633 (0.001)	0.9123 (0.001)	0.176
	Inclusion of one random study					
0.1	0.8007 (0.011)	0.9837 (0.002)	0.784	0.102 (0.004)	0.997 (0.001)	0.099
0.3	0.6673 (0.013)	0.8497 (0.009)	0.517	0.2113 (0.009)	0.966 (0.005)	0.177
0.5	0.6013 (0.017)	0.852 (0.009)	0.453	0.2527 (0.01)	0.8667 (0.008)	0.119
	Inclusion of two random studies					
0.1	0.624 (0.016)	0.9737 (0.009)	0.598	0.0847 (0.005)	0.994 (0.001)	0.079
0.3	0.51 (0.016)	0.8433 (0.006)	0.353	0.1727 (0.011)	0.9317 (0.005)	0.104
0.5	0.4167 (0.012)	0.85 (0.009)	0.267	0.256 (0.012)	0.8193 (0.006)	0.075