Author Correction: Metagenomic analysis of colorectal cancer datasets identifies cross-cohort microbial diagnostic signatures and a link with choline degradation

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Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-019-0405-7>, published online 1 April 2019.

In the version of this article initially published, the affiliation for Chiara Pozzi, Sara Gandini and Davide Serrano was given as the European Institute of Oncology. The correct affiliation is IEO, European Institute of Oncology IRCCS. In addition, the data for age, BMI and sex were missing from Table 1 for the HanniganGD_2018 dataset, and consequently for the totals of all datasets. These data are given in Table 1 here. The errors have been corrected in the HTML and PDF versions of the article.

Table 1 | Size and characteristics of the large-scale CRC metagenomic datasets included in this study

Dataset	Groups (n)	Age (mean $\pm$ s.d.)	BMI (mean $\pm$ s.d.)	Sex F(%) / M(%)	Country	No. of reads ($\times 10^9$)
HanniganGD_2018 (ref. ³¹)	Control (28)	55.46 $\pm$ 9.4	26.63 $\pm$ 5.1	60.7/39.3	USA (54), Canada (28)	0.5
	Adenoma (27)	61.74 $\pm$ 10.8	27.36 $\pm$ 4.45	40.7/59.3		
	CRC (27)	58.6 $\pm$ 10.9	30.41 $\pm$ 7.5	25.9/74.1		
Total	Control (308)	62.01 $\pm$ 3.7	25.4 $\pm$ 1.4	43.5/56.5		45.6
	Adenoma (143)	63.88 $\pm$ 1.9	26.8 $\pm$ 1.1	40.5/59.5		
	CRC (313)	64.67 $\pm$ 3.8	25.8 $\pm$ 1.7	33.5/66.5		

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Author Correction: Reporting guidelines for clinical trials evaluating artificial intelligence interventions are needed

The CONSORT-AI and SPIRIT-AI Steering Group

Correction to: *Nature Medicine* <https://www.nature.com/articles/s41591-019-0603-3>, published online 24 September 2019.

In the version of this article originally published, the name of author Lavinia Ferrante di Ruffano was incorrectly spelled as Lavinia Ferrante Di Ruffano. The error has been corrected in the HTML and PDF versions of the article.

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