

ERRATUM

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Erratum to: Estimating the population abundance of tissue-infiltrating immune and stromal cell populations using gene expression

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Erratum

After the publication of this work [1] it was noticed that the legend of Fig. 1a was incorrect. The correct legend should read: Flowchart of MCP-counter's development and validation. It was also noted that in Fig. 2c the name of the last two genes was partially removed. The horizontal lines that were delineating the specificity of the displayed genes are missing. The corrected Fig. 2c is noted below.

The equal contributors symbol was missing for authors Aurélien de Reyniès and Wolf H. Fridman. This has now been corrected.

In the PDF version of the article, equation 5 in the section 'Correlation profiles of TM in microenvironment and tumor datasets' is incorrect. Please see the correct equation below:

$$\pi_{k,j} = \frac{f_{S,j}}{f_k} = \frac{g_{S,j}}{g_k} \quad (5)$$

The publisher apologises for these errors.

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Reference

1. Becht E, Giraldo NA, Lacroix L, Buttard B, Elarouci N, Petitprez F, et al. Estimating the population abundance of tissue-infiltrating immune and stromal cell populations using gene expression. *Genome Biol.* 2016;17:218.

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