

CORRECTION

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Correction to: Comprehensive gene expression meta-analysis identifies signature genes that distinguish microglia from peripheral monocytes/macrophages in health and glioma

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Correction to: *Acta neuropathol commun* (2019) 7: 20
<https://doi.org/10.1186/s40478-019-0665-y>

The original publication of this article [1] contained 3 minor errors in Figs. 1, 3 and 5. In this correction article the updated figures are published. The figure captions describe the updated information in these figures.

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Published online: 08 January 2020

Reference

1. Haage V, Semtner M, Vidal RO et al (2019) Comprehensive gene expression meta-analysis identifies signature genes that distinguish microglia from peripheral monocytes/macrophages in health and glioma. *Acta Neuropathol Commun* 7:20. <https://doi.org/10.1186/s40478-019-0665-y>

The original article can be found online at <https://doi.org/10.1186/s40478-019-0665-y>

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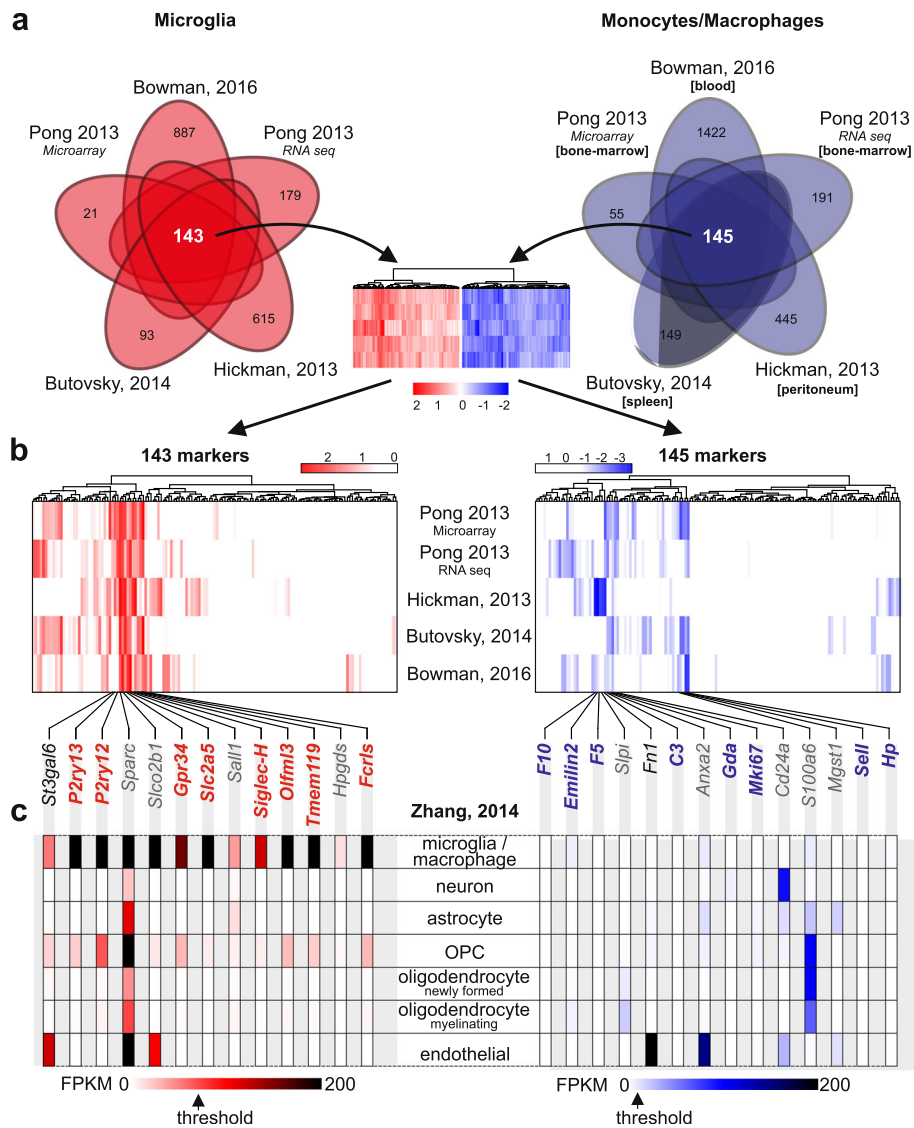


Fig. 1 The threshold FPKM expression value for excluding genes as microglia markers is 70 (panel c)

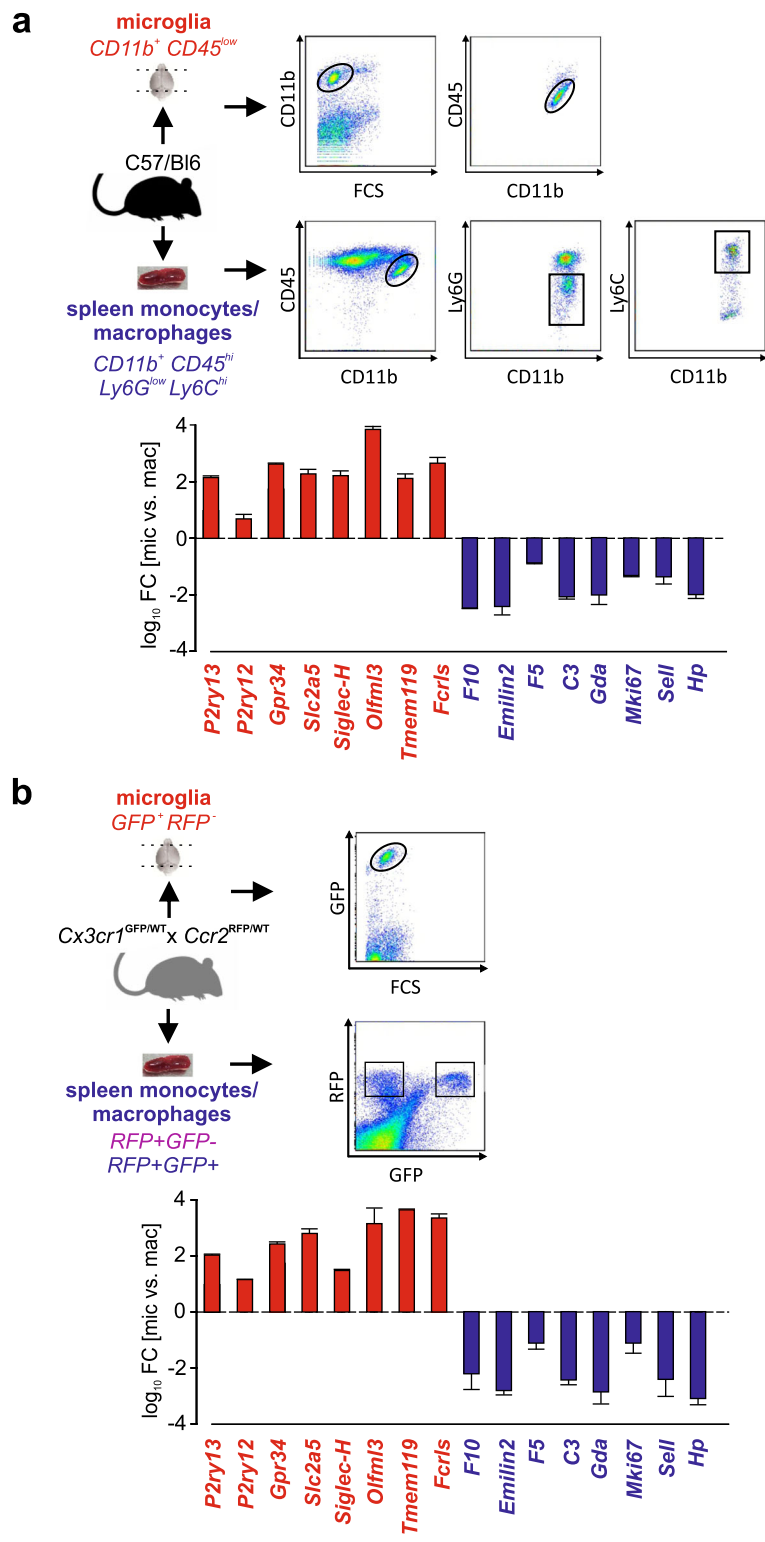


Fig. 3 Bar graphs represent the log_{10} fold change expression (panel **a** and **b**)

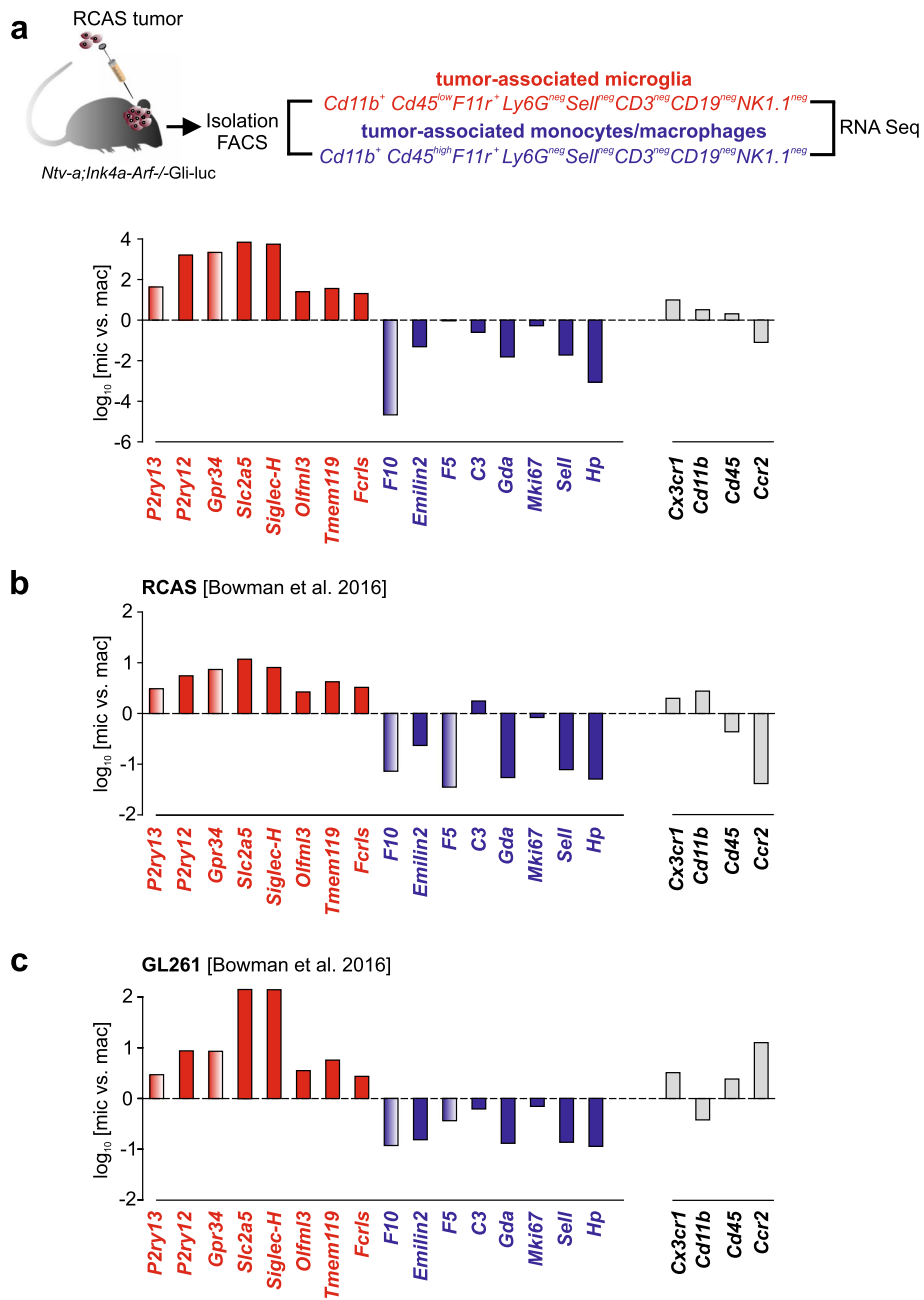


Fig. 5 Bar graphs represent the log₁₀ fold change expression (panel **a**, **b** and **c**)