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Correction to: A small-molecule/cytokine combination enhances hematopoietic stem cell proliferation via inhibition of cell differentiation

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Correction to: Stem Cell Research & Therapy (2017) 8:169 https://doi.org/10.1186/s13287-017-0625-z

Following the publication of the original article [1], the authors identified an error in Fig. 4b. The authors noticed that the gel bands of Runx1 were accidentally used again

as the ones of Bmi1 during the layout of figures. It has been corrected after they double checked the original data. The results and conclusion concluded in this paper are still valid.

The original article can be found online at https://doi.org/10.1186/s13287-017-0625-z.

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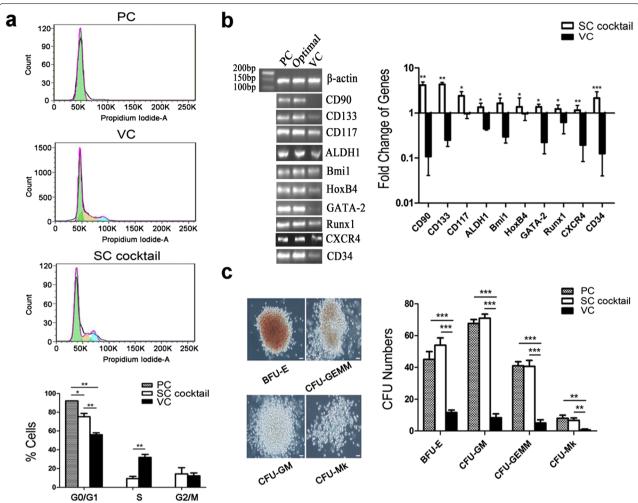


Fig. 4 Invitro functional assessments of the SC cocktail-expanded CD34+ cells. **a** Cell cycle analysis. One of threerepresentative experiments is shown, and the percentage of different phases issummarized in the histogram. **b**HSC-specific gene expression. Results of qualitative RNA-PCR are shown on theleft for one representative of three independent RNA samples analyzed pergroup, and the results of quantitative PCR are shown in the right. **c** The morphology and numbers ofcolony-forming units (*CFUs*).Morphology (20× objective, scale bar = 50 μ m) and colony numbers were recorded nday 14 after cell seeding. All data are shown as means \pm SD, n = 3. *p < 0.05, **p < 0.01, ***p < 0.01. *p < 0.01. *

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Published online: 17 August 2021

Reference

 Wang L, et al. A small-molecule/cytokine combination enhances hematopoietic stem cell proliferation via inhibition of cell differentiation. Stem Cell Res Ther. 2017;8:169. https://doi.org/10.1186/s13287-017-0625-z.

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