



## Correction to: Menatretrenone facilitates hematopoietic cell generation in a manner that is dependent on human bone marrow mesenchymal stromal/stem cells

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In the original publication of the article, the figures 4 C, F and 5 B, C were published with unexpected appearance of dots. The corrected Figs. 4, 5 are given in this correction.

In addition, Y.M.'s grant should be “18K08323/Grants-in-Aid from the Ministry of Education, Culture, Sports, Science, and Technology in Japan”.

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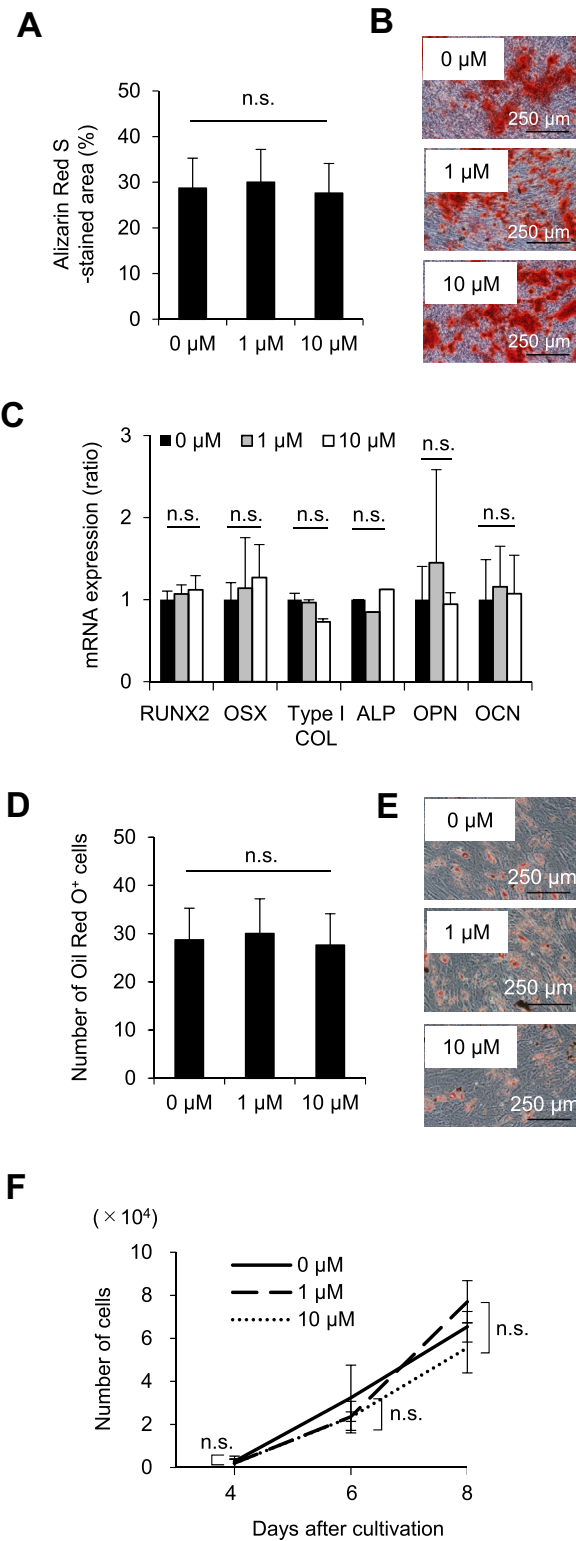
The original article can be found online at <https://doi.org/10.1007/s12185-020-02916-8>.

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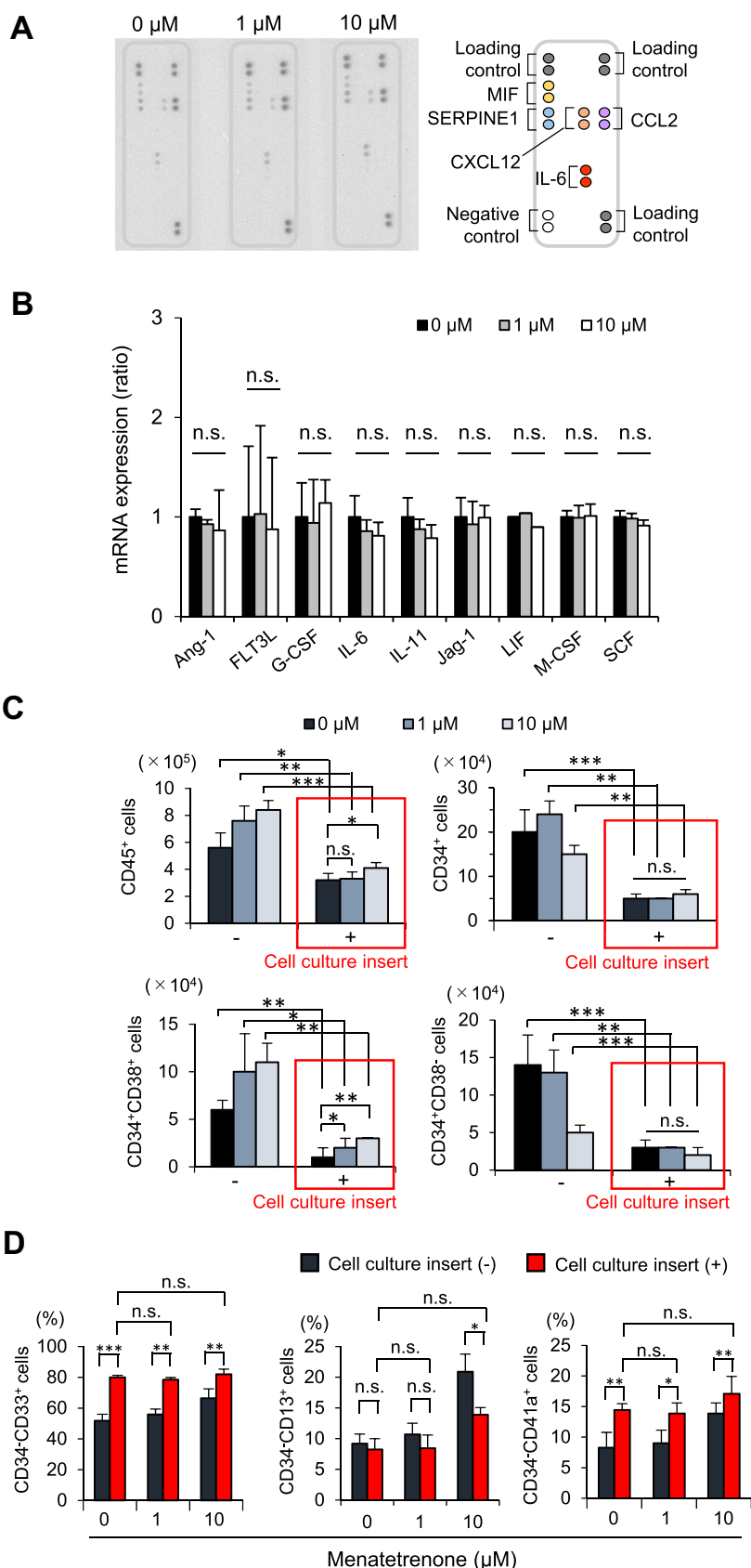
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**Fig. 4** Menatretrenone does not affect the differentiation or proliferation capabilities of BM-MSCs. **a, b** Osteogenic differentiation of BM-MSCs treated with or without menatretrenone, as assessed by Alizarin Red S staining (**a**). Red nodules indicate mineralization (**b**). Representative images are shown. **c** Expression levels of osteogenesis-associated genes in BM-MSCs treated with or without menatretrenone, as assessed by qRT-PCR. *RUNX2* runt-related transcription factor 2, *OSX* osterix, *COL* collagen, *ALP* alkaline phosphatase, *OPN* osteopontin, *OCN* osteocalcin. **d, e** Adipogenic differentiation of BM-MSCs treated with or without menatretrenone, as assessed by Oil Red O staining (**d**). Red-staining indicates fat-laden cells (**e**). Representative images are shown. **f** The proliferation of BM-MSCs treated with or without menatretrenone. **a, c, d, f** Data are represented as the mean  $\pm$  SD;  $n=5$  per group; *n.s.* not significant



**Fig. 5** Direct cell–cell interactions, but not soluble factors, mediate enhanced HPC expansion in co-cultures with menatretrenone-treated human BM-MSCs. **a** Cytokine arrays of supernatants from cultures of BM-MSCs treated with 0, 1, or 10  $\mu$ M menatretrenone. *CCL2* C–C motif chemokine ligand 2, *CXCL12* C–X–C motif chemokine ligand 12, *MIF* macrophage migration inhibitory factor, *SERPINE1* serpin family E member 1, *IL-6* interleukin-6. **b** The mRNA expression levels of multiple hematopoiesis-associated genes in BM-MSCs treated with 0  $\mu$ M (black bars), 1  $\mu$ M (gray bars), or 10  $\mu$ M (white bars) menatretrenone, as assessed by qRT-PCR;  $n=5$  per group. *Ang-1* angiopoietin-1, *FLT3L* fms-related tyrosine kinase 3 ligand, *G-CSF* granulocyte colony-stimulating factor, *IL-6* interleukin-6, *IL-11* interleukin-11, *Jag-1* jagged-1, *LIF* leukemia inhibitory factor, *M-CSF* macrophage colony-stimulating factor, *SCF* stem cell factor. **c** Co-culture of CD34<sup>+</sup> HSPCs with BM-MSCs treated with 0 (black bars), 1 (blue bars), or 10  $\mu$ M (gray bars) menatretrenone in the presence (+, red squares) or absence (–) of cell culture inserts ( $n=4$  per group). Flow cytometry analyses showing the numbers of CD45<sup>+</sup> cells, CD34<sup>+</sup> cells, CD34<sup>+</sup>CD38<sup>–</sup> cells, and CD34<sup>+</sup>CD38<sup>+</sup> cells. **d** Co-culture of CD34<sup>+</sup> HSPCs with BM-MSCs treated with 0, 1, or 10  $\mu$ M menatretrenone in the presence (red bars) or absence (black bars) of cell culture inserts ( $n=4$  per group). Flow cytometry plots showing the percentages of CD34<sup>–</sup>CD33<sup>+</sup> cells, CD34<sup>–</sup>CD13<sup>+</sup> cells, and CD34<sup>–</sup>CD41a<sup>+</sup> cells. **b–d** Data are presented as the mean  $\pm$  SD; \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ; *n.s.* not significant



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