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CORRECTION

Correction: Construction of transcriptome atlas of white yak hair follicle during anagen and catagen using single-cell RNA sequencing

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Following the publication of the original article [1], the authors identified an error in Fig. 6. The correct figure is given below.

The original article [1] has been corrected.

The original article can be found online at https://doi.org/10.1186/s12864-022-09003-8.

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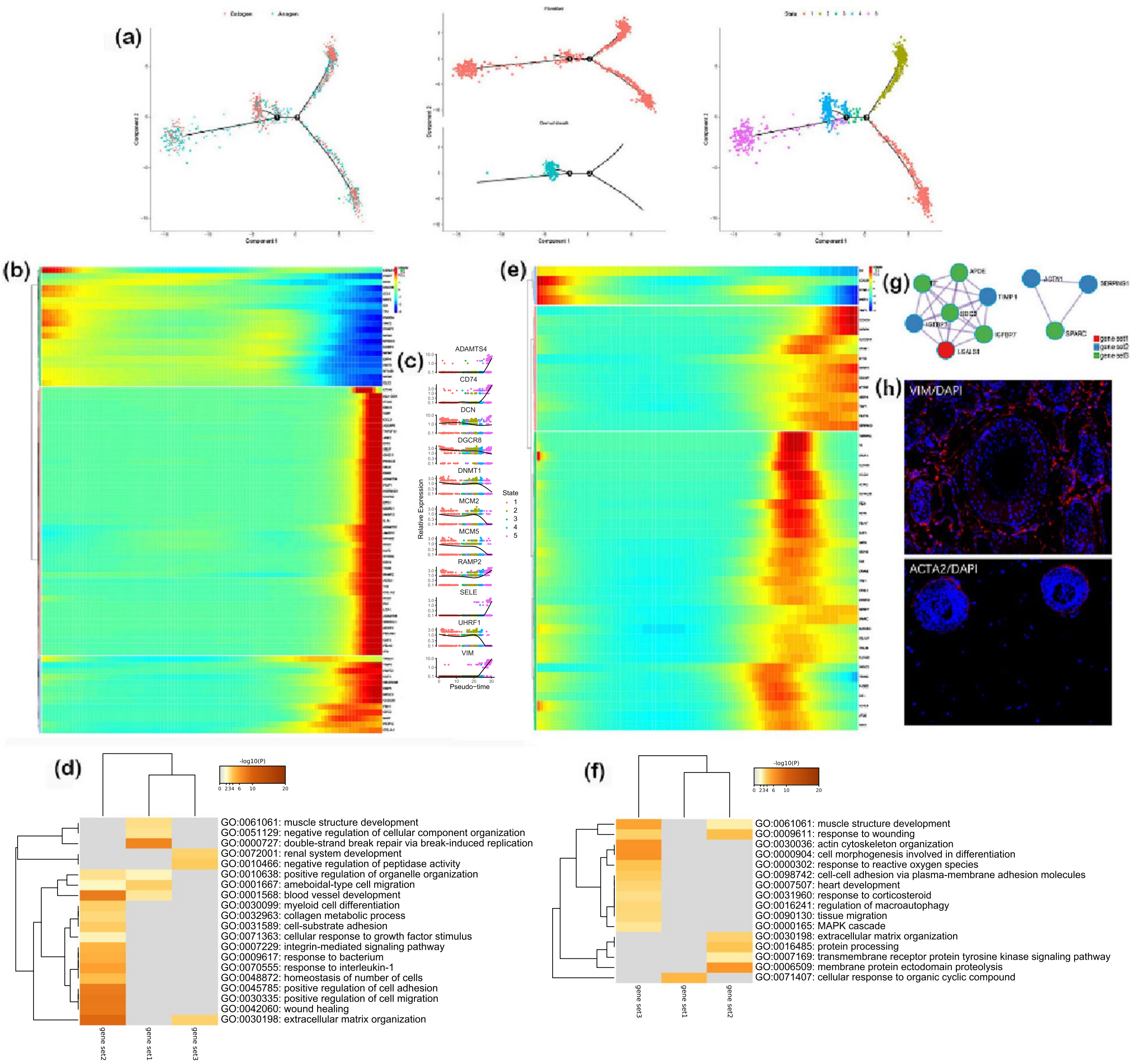

Fig. 6 Dynamic changes in the gene expression during specialization of hair follicle fibroblasts and DS. **a** Construction of the pseudotime differentiation trajectory between the fibroblasts and DS; **b** Gene expression during fibroblast specialization; **c** Expression of the characteristic genes in different stages of fibroblast; **d** GO enrichment analysis of fibroblast characteristic genes; **e** Gene expression during DS specialization; **f** GO enrichment analysis of the characteristic genes in DS; **g** Interaction analysis of the characteristic genes in DS; **h** Immunofluorescence analysis of the hair follicles

Reference

1. Zheng Q, Ye N, Bao P, et al. Construction of transcriptome atlas of white yak hair follicle during anagen and catagen using single-cell RNA sequencing. BMC Genomics. 2022;23:813. https://doi.org/10.1186/s12864-022-09003-8.