

CORRECTION

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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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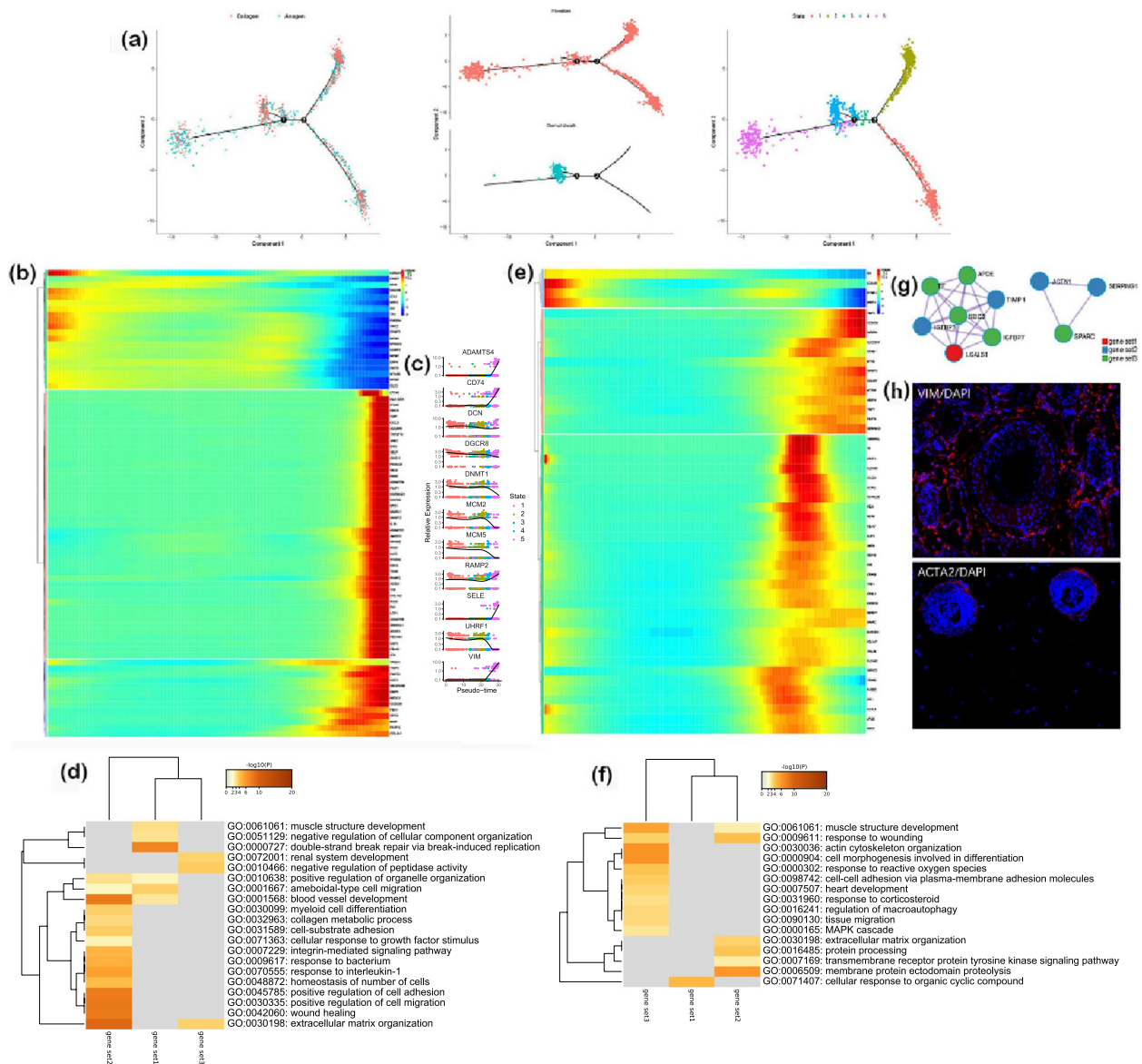


Fig. 6 Dynamic changes in the gene expression during specialization of hair follicle fibroblasts and DS. **a** Construction of the pseudotime differentiation trajectory between the fibroblasts and DS; **b** Gene expression during fibroblast specialization; **c** Expression of the characteristic genes 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

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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>.