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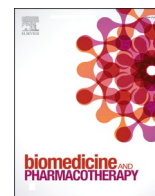
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Corrigendum to “A phenotype-driven data and drug repurposing strategy used to identify potential treatments targeting chondrocyte hypertrophy in osteoarthritis” [Biomed. Pharmacother. 193 (December) (2025) 118773]

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The authors regret that the original data availability statement of our manuscript was not included in the final submission and this oversight was not identified during proofreading. In the context of open-access publication, it is essential that the correct data availability information is provided.

The correct statement is as follows:

The data generated and analysed during the current study can be found in the Supplementary Data file with the exception of the single cell

RNA sequencing data. The count matrices derived from the raw single cell RNA sequencing data have been uploaded to Gene Expression Omnibus under accession number GSE290973. Due to privacy considerations, the raw sequencing files have not been deposited in the Gene Expression Omnibus but can be obtained from the corresponding author upon reasonable request and completion of a Data Use Agreement.

The authors would like to apologise for any inconvenience caused.

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