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We did not use any statistical method to determine the sample sizes prior to performing experiments. However, our sample sizes are in line with previous publications. Each experiment was repeated at least twice to confirm reproducibility of findings.

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Each experiment was repeated at least twice to confirm reproducibility of findings. Raw sequencing reads for the bulk and single cell RNA-seq have been deposited in the NCBI BioProject database under accession numbers PRJNA415135 and PRJNA415187. Mass spec protein identification data has been deposited in http://www.peptideatlas.org/PASS/PASS01443. The materials used in this study will be available from the corresponding authors upon request. The source data underlying all main and extended figures are provided as a Source Data file.

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* Indicate how samples were allocated into experimental groups (in the case of clinical studies, please specify allocation to treatment method); if randomization was used, please also state if restricted randomization was applied
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