

CORRECTION

Correction: Comprehensive proteomics and functional annotation of mouse brown adipose tissue

The PLOS ONE Staff

The image for Fig 1 is incorrect. There is also an error in the caption for Fig 1. Please see the complete, correct Fig 1 and Fig 1 caption here. The publisher apologizes for the error.

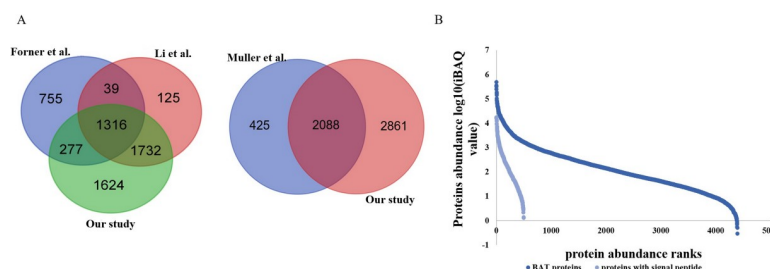
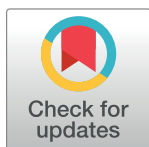


Fig 1. BAT proteome profile analysis. (A) A Venn diagram comparing BAT proteome large-scale studies. (B) The quantitative protein abundance ranges in BAT samples and the proteins that included a secreted signal peptide according to the iBAQ algorithm.

<https://doi.org/10.1371/journal.pone.0244359.g001>

Reference

1. Li J, Li J, Zhao W-G, Sun H-D, Guo Z-G, Liu X-Y, et al. (2020) Comprehensive proteomics and functional annotation of mouse brown adipose tissue. PLoS ONE 15(5): e0232084. <https://doi.org/10.1371/journal.pone.0232084> PMID: 32374735



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