




Correction



Life Science Alliance

Correction: The intracellular pathogen *Francisella* escapes from adaptive immunity by metabolic adaptation

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We would like to apologize for an incorrect description in our manuscript entitled “The intracellular pathogen *Francisella tularensis* escapes from adaptive immunity by metabolic adaptation” (PMID: 35667686, PMCID: PMC9170078, DOI: [10.26508/lisa.202201441](https://doi.org/10.26508/lisa.202201441)). We recently found that, during the revision process, a sentence of the results section in the published manuscript was mistakenly inserted. We think that the deletion does not affect the conclusion of the manuscript.

In the Results section

Where it reads:

Four of the substitutions are located in the zinc-binding region of the cytidine and deoxycytidylate deaminase domain and one substitution is in the bacterial bifunctional deaminase-reductase domain (InterPro database, <https://www.ebi.ac.uk/interpro/>) (Fig 2C). No substitutions were found in ribAB, ribH and ribC. Among the five substitutions in ribD, H80C and Q254R were shared with seven other virulent strains, based on genome sequences deposited in the KEGG database (Fig 2D).

It should read:

Four of the substitutions are located in the zinc-binding region of the cytidine and deoxycytidylate deaminase domain and one substitution is in the bacterial bifunctional deaminase-reductase domain (InterPro database, <https://www.ebi.ac.uk/interpro/>) (Fig 2C). Among the five substitutions in ribD, H80C and Q254R were shared with seven other virulent strains, based on genome sequences deposited in the KEGG database (Fig 2D).



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