Check for updates

OPEN ACCESS

EDITED AND REVIEWED BY Wouter H. Van Geffen, Medical Center Leeuwarden, Netherlands

*CORRESPONDENCE Armin Frille armin.frille@medizin.uni-leipzig.de Maximilian von Laffert maximilian.von-laffert@medizin.uni-

[†]These authors have contributed equally to this work

RECEIVED 30 July 2024 ACCEPTED 22 August 2024 PUBLISHED 11 September 2024

CITATION

leipzig.de

Frille A, Boeschen M, Wirtz H, Stiller M, Bläker H and von Laffert M (2024) Corrigendum: *TP53* co-mutations in advanced lung adenocarcinoma: comparative bioinformatic analyses suggest ambivalent character on overall survival alongside *KRAS*, *STK11* and *KEAP1* mutations. *Front. Oncol.* 14:1473239. doi: 10.3389/fonc.2024.1473239

COPYRIGHT

© 2024 Frille, Boeschen, Wirtz, Stiller, Bläker and von Laffert. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms. Corrigendum: *TP53* comutations in advanced lung adenocarcinoma: comparative bioinformatic analyses suggest ambivalent character on overall survival alongside *KRAS*, *STK11* and *KEAP1* mutations

Armin Frille^{1*†}, Myriam Boeschen^{2†}, Hubert Wirtz¹, Mathias Stiller², Hendrik Bläker² and Maximilian von Laffert^{2*}

¹Department of Respiratory Medicine, Leipzig University, Leipzig, Germany, ²Institute of Pathology, Leipzig University, Leipzig, Germany

KEYWORDS

NSCLC, lung adenocarcinoma, KRAS, STK11, KEAP1, TP53, co-mutations, survival

A Corrigendum on

TP53 co-mutations in advanced lung adenocarcinoma: comparative bioinformatic analyses suggest ambivalent character on overall survival alongside *KRAS*, *STK11* and *KEAP1* mutations

By Frille A, Boeschen M, Wirtz H, Stiller M, Bläker H and von Laffert M (2024). Front. Oncol. 14:1357583. doi: 10.3389/fonc.2024.1357583.

In the published article, there was a mistake in the **Funding** statement. The original text of the **Funding** statement was unspecific, since financial support was received for the publication costs only.

The first two sentences of the original Funding statement read:

"The author(s) declare financial support was received for the research, authorship, and/ or publication of this article. This publication was funded by the Open Access Publishing Fund of Leipzig University, supported by the German Research Foundation (DFG)."

The correct Funding statement appears below.

Funding

The authors declare that financial support was received for the publication of this article, which was funded by the Open Access Publishing Fund of Leipzig University, supported by the German Research Foundation (DFG). The funder was not involved in the study design, collection, analysis, interpretation of data, the writing of this article or the decision to submit it for publication.

In the published article, there was an error in the Abstract section.

The words "in patients with lung cancer" were added after the term "(mOS)".

Abstract, Background. This sentence previously stated:

"Recently, we could show that the co-mutations of *KRAS* + *KEAP1*, *STK11* + *KEAP1* and *KRAS* + *STK11* + *KEAP1* lead to a significantly shorter median overall survival (mOS) across treatments by analyzing multiple datasets."

The corrected sentence appears below:

"Recently, we could show that the co-mutations of *KRAS* + *KEAP1*, *STK11* + *KEAP1* and *KRAS* + *STK11* + *KEAP1* lead to a significantly shorter median overall survival (mOS) in patients with lung cancer across treatments by analyzing multiple datasets."

In the published article, there was an error in the **Abstract** section. The word "*STK1*" was replaced with "*STK11*" at the end of the sentence.

Abstract, Results. This sentence previously stated:

"Whereas TP53-mutations seem to have protective effects in the context of further *KEAP1*- and *KRAS* + *KEAP1*-alterations (improved mOS), their role seems contrary if acquired in an already existing combination of mutations as *KRAS* + *STK11*, *KRAS* + *STK11* + *KEAP1* and *STK1* + *KEAP1*."

The corrected sentence appears below:

"Whereas TP53-mutations seem to have protective effects in the context of further *KEAP1*- and *KRAS* + *KEAP1*-alterations (improved mOS), their role seems contrary if acquired in an already existing combination of mutations as *KRAS* + *STK11*, *KRAS* + *STK11* + *KEAP1* and *STK11* + *KEAP1*."

In the published article, there was an error in the **Abstract** section. The word "co-mutationshad" was replaced with "co-mutations had" in the last sentence of the results section.

Abstract, Results. This sentence previously stated:

"TP53 co-mutationshad a negative influence on KRAS- only mutated LUAD (mOS reduced significantly by more than 30%)."

The corrected sentence appears below:

"TP53 co-mutations had a negative influence on KRAS- only mutated LUAD (mOS reduced significantly by more than 30%)."

In the published article, there was an error in the **Introduction** section.

The word "KRAS" was italicized.

1 Introduction, Paragraph 5. This sentence previously stated: "Whereas, *TP53* mutations alone had no impact on response and survival, a subgroup (12 patients) with KRAS *G12C* + *TP53* comutations defined long-term responders to immunotherapy (9)."

The corrected sentence appears below:

"Whereas, *TP53* mutations alone had no impact on response and survival, a subgroup (12 patients) with *KRAS G12C* + *TP53* comutations defined long-term responders to immunotherapy (9)." In the published article, there was an error in the **Material and methods** section.

The word "compromise" was replaced with "comprised".

2 Materials and methods, Paragraph 1. This sentence previously stated:

"All data compromise patients with advanced tumor stages (mainly stage IV)."

The corrected sentence appears below:

"All data comprised patients with advanced tumor stages (mainly stage IV)."

In the published article, there was an error in the **Results** section.

The symbol ">" was replaced with "<".

3 Results, *3.1 Demographics and incidences of (co-)mutations in NSCLC and LUAD*, Paragraph 3. This sentence previously stated:

"While mutations in *KRAS*, *STK11* and *KEAP1* significantly co- occurred among themselves (q > 0.05), there was neither a significant co-occurrence nor a significant mutual exclusivity between mutations in one of the three genes and *TP53* mutations."

The corrected sentence appears below:

"While mutations in *KRAS*, *STK11* and *KEAP1* significantly cooccurred among themselves (q < 0.05), there was neither a significant co-occurrence nor a significant mutual exclusivity between mutations in one of the three genes and *TP53* mutations."

In the published article, there was an error in the **Discussion** section.

The word "mutation" was replaced with "mutational pattern".

4 Discussion, Paragraph 1. This sentence previously stated:

"By employing this database approach, we were able to show that *TP53* mutations had an influence on mOS for better or worse depending on the concurrent mutation."

The corrected sentence appears below:

"By employing this database approach, we were able to show that *TP53* mutations had an influence on mOS for better or worse depending on the concurrent mutational pattern."

The authors apologize for these errors and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.