Author Correction: A SARS-like cluster of circulating bat coronaviruses shows potential for human emergence

Vineet D. Menachery, Boyd L. Yount Jr, Kari Debbink, Sudhakar Agnihothram, Lisa E. Gralinski, Jessica A. Plante, Rachel L. Graham, Trevor Scobey, Xing-Yi Ge, Eric F. Donaldson, Scott H. Randell, Antonio Lanzavecchia, Wayne A. Marasco, Zhengli-Li Shi and Ralph S. Baric

Correction to: *Nature Medicine* <https://doi.org/10.1038/nm.3985>, published online 9 November 2015.

In the version of this article initially published, the sequence of the mouse adapted SHC014-MA15 virus had not been deposited in GenBank. The sequence has now been deposited in GenBank under accession number [MT308984](https://www.ncbi.nlm.nih.gov/nuclseq/MT308984).

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Author Correction: Caring for patients with cancer in the COVID-19 era

Joris van de Haar, Louisa R. Hoes, Charlotte E. Coles, Kenneth Seamon, Stefan Fröhling, Dirk Jäger, Franco Valenza, Filippo de Braud, Luigi De Petris, Jonas Bergh, Ingemar Ernberg, Benjamin Besse, Fabrice Barlesi, Elena Garralda, Alejandro Piris-Giménez, Michael Baumann, Giovanni Apolone, Jean Charles Soria, Josep Tabernero , Carlos Caldas and Emile E. Voest

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-020-0874-8>, published online 16 April 2020.

In the version of this article initially published, the CCE Contributors Appendix in the Supplementary Note was missing the name 'Paul D.P. Pharoah'. The correct text should read as follows: "...and Paul D.P. Pharoah, of the Cancer Research UK Cancer Centre...". An updated Supplementary Note is now provided.

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OPEN

Author Correction: Integrative molecular and clinical modeling of clinical outcomes to PD1 blockade in patients with metastatic melanoma

David Liu, Bastian Schilling , Derek Liu, Antje Sucker, Elisabeth Livingstone, Livnat Jerby-Arnon, Lisa Zimmer, Ralf Gutzmer, Imke Satzger, Carmen Loquai, Stephan Grabbe, Natalie Vokes, Claire A. Margolis, Jake Conway, Meng Xiao He, Haitham Elmarakeby, Felix Dietlein, Diana Miao , Adam Tracy, Helen Gogas, Simone M. Goldinger, Jochen Utikal , Christian U. Blank , Ricarda Rauschenberg, Dagmar von Bubnoff, Angela Krackhardt, Benjamin Weide, Sebastian Haferkamp, Felix Kiecker, Ben Izar, Levi Garraway, Aviv Regev , Keith Flaherty , Annette Paschen, Eliezer M. Van Allen and Dirk Schadendorf

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-019-0654-5>, published online 2 December 2019.

In the version of this article initially published, the sixth author's surname (Jerby-Amon) was incorrect. The correct surname is 'Jerby-Arnon'. The error has been corrected in the HTML and PDF versions of the article.

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Publisher Correction: Pulmonary venous circulating tumor cell dissemination before tumor resection and disease relapse

Francesca Chemi, Dominic G. Rothwell, Nicholas McGranahan, Sakshi Gulati, Chris Abbosh, Simon P. Pearce , Cong Zhou, Gareth A. Wilson, Mariam Jamal-Hanjani, Nicolai Birkbak, Jackie Pierce, Chang Sik Kim, Saba Ferdous, Deborah J. Burt, Daniel Slane-Tan, Fabio Gomes, David Moore, Rajesh Shah, Maise Al Bakir, Crispin Hiley, Selvaraju Veeriah, Yvonne Summers, Philip Crosbie, Sophia Ward, Barbara Mesquita, Marek Dynowski, Dhruva Biswas , Jonathan Tugwood, Fiona Blackhall, Crispin Miller, Allan Hackshaw, Ged Brady , Charles Swanton , Caroline Dive and TRACERx Consortium

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-019-0593-1>, published online 7 October 2019.

In the version of this article initially published, the list of members and affiliations for the TRACERx Consortium was provided as a Supplementary Note. The list should have been provided in the online version. Also, some of the numbers in the second row (Non-adenocarcinoma) of the Tumor histology section of Table 1 were incorrect ($n = 43$; 20 (47%); 23 (53%)). The correct numbers are as follows (respectively): $n = 41$; 20 (49%); 21 (51%). Finally, the HR value for Extended Data Fig. 1b (HR = 2.78) was incorrect in the final sentence of the fourth paragraph of the main text (it is correct in Extended Data Fig. 1b itself). The correct value is 'HR = 2.89'. The errors have been corrected in the HTML and PDF versions of the article.

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Publisher Correction: Spatial heterogeneity of the T cell receptor repertoire reflects the mutational landscape in lung cancer

Kroopa Joshi, Marc Robert de Massy , Mazlina Ismail, James L. Reading, Imran Uddin , Annemarie Woolston , Emine Hatipoglu, Theres Oakes, Rachel Rosenthal, Thomas Peacock , Tahl Ronel, Mahdad Noursadeghi , Virginia Turati, Andrew J. S. Furness, Andrew Georgiou, Yien Ning Sophia Wong, Assma Ben Aissa, Mariana Werner Sunderland, Mariam Jamal-Hanjani, Selvaraju Veeriah, Nicolai J. Birkbak, Gareth A. Wilson, Crispin T. Hiley, Ehsan Ghorani , José Afonso Guerra-Assunção , Javier Herrero , Tariq Enver, Sine R. Hadrup , Allan Hackshaw, Karl S. Peggs, Nicholas McGranahan, Charles Swanton , TRACERx consortium, Sergio A. Quezada and Benny Chain

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-019-0592-2>, published online 7 October 2019.

In the version of this article initially published, the list of members and affiliations for the TRACERx Consortium was provided as a Supplementary Note. The list should have been provided in the online version. The error has been corrected in the HTML and PDF versions of the article.

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Publisher Correction: A clonal expression biomarker associates with lung cancer mortality

Dhruva Biswas, Nicolai J. Birkbak , Rachel Rosenthal, Crispin T. Hiley, Emilia L. Lim, Krisztian Papp , Stefan Boeing, Marcin Krzystanek, Dijana Djureinovic, Linnea La Fleur, Maria Greco, Balázs Döme, János Fillinger, Hans Brunnström , Yin Wu, David A. Moore, Marcin Skrzypski, Christopher Abbosh, Kevin Litchfield, Maise Al Bakir, Thomas B. K. Watkins, Selvaraju Veeriah, Gareth A. Wilson, Mariam Jamal-Hanjani, Judit Moldvay, Johan Botling, Arul M. Chinnaiyan, Patrick Micke, Allan Hackshaw, Jiri Bartek, Istvan Csabai, Zoltan Szallasi, Javier Herrero , Nicholas McGranahan , Charles Swanton and TRACERx Consortium

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-019-0595-z>, published online 7 October 2019.

In the version of this article initially published, the list of members and affiliations for The TRACERx Consortium was provided as a Supplementary Note. The list should have been provided in the online version. Also, an incorrect version of Fig. 2b was published. The correct version is provided below. The errors have been corrected in the HTML and PDF versions of the article.

Fig. 2 | Original and Corrected.

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Publisher Correction: Safety and feasibility of CRISPR-edited T cells in patients with refractory non-small-cell lung cancer

You Lu , Jianxin Xue, Tao Deng, Xiaojuan Zhou, Kun Yu, Lei Deng, Meijuan Huang, Xin Yi, Maozhi Liang, Yu Wang, Haige Shen, Ruizhan Tong, Wenbo Wang, Li Li, Jin Song, Jing Li, Xiaoxing Su, Zhenyu Ding, Youling Gong, Jiang Zhu, Yongsheng Wang, Bingwen Zou, Yan Zhang, Yanying Li, Lin Zhou, Yongmei Liu, Min Yu, Yuqi Wang, Xuanwei Zhang, Limei Yin, Xuefeng Xia, Yong Zeng, Qiao Zhou, Binwu Ying, Chong Chen, Yuquan Wei, Weimin Li and Tony Mok

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-020-0840-5>, published online 27 April 2020.

In the version of this article initially published, the label ‘TPS1 $\geq$ 1%’ in the top left box of Fig. 1a was incorrect. The correct label is ‘TPS $\geq$ 1%’. The error has been corrected in the HTML and PDF versions of the article.

Published online: 18 June 2020

<https://doi.org/10.1038/s41591-020-0973-6>

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Publisher Correction: China needs a national intelligent syndromic surveillance system

Peng Jia and Shujuan Yang

Correction to: *Nature Medicine* <https://doi.org/10.1038/s41591-020-0921-5>, published online 20 May 2020.

In the version of this article initially published, the numbering and order of the second and third affiliations (²West China School of Public Health and West China Fourth Hospital, Sichuan University, Chengdu, China. ³International Initiative on Spatial Lifecourse Epidemiology, Hong Kong, China.) was incorrect. The correct order is as follows: ²International Initiative on Spatial Lifecourse Epidemiology, Hong Kong, China. ³West China School of Public Health and West China Fourth Hospital, Sichuan University, Chengdu, China. Also, the second half of the first time frame (“under 2 days”) in the first sentence of the third paragraph was incorrect. The correct phrasing is “under 2 hours.” The errors have been corrected in the HTML and PDF versions of the article.

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Addendum: Estimating clinical severity of COVID-19 from the transmission dynamics in Wuhan, China

Joseph T. Wu , Kathy Leung, Mary Bushman, Nishant Kishore, Rene Niehus, Pablo M. de Salazar, Benjamin J. Cowling , Marc Lipsitch and Gabriel M. Leung

Addendum to: *Nature Medicine* <https://doi.org/10.1038/s41591-020-0822-7>; published online 19 March 2020

On 17 April 2020, the Wuhan Municipal Health Commission revised the official COVID-19 death counts in Wuhan up by 1,290 to 3,869. No new information on the age distribution of COVID-19 deaths in China has been published since February 2020. In light of the changes in the number of fatalities in Wuhan, we are providing a revised estimate of clinical severity as follows, assuming that the age distribution of COVID-19 deaths remains the same as before.

We assumed that the abovementioned revision of the total number of COVID-19 deaths in Wuhan would increase the number of confirmed COVID-19 deaths during our study period by the same proportion; i.e., from 2,169 to $2,169 \times 3,869 / (3,869 - 1,290) = 3,254$. The resulting age-specific symptomatic case-fatality risks (sCFRs) are shown in Fig. 1 in this Addendum. The overall sCFR would increase to 2.2% (95% credible interval, 1.4–3.2%), 2.0% (1.2–3.4%) or 1.8% (1.0–2.8%) if the probability of developing symptoms after infection (P_{sym}) was 0.5, 0.75 or 0.95, respectively. If $P_{\text{sym}} = 0.5$, the sCFRs are 0.5% (0.2–1.0%), 0.8% (0.5–1.2%) and 3.9% (2.6–5.9%) for those <30 years of age, 30–59 years of age and >59 years of age, respectively.

We will update our estimates again as and when new and more detailed information, including the total number and age distribution of COVID-19 deaths, are made available.

Fig. 1 | Estimates of age-specific sCFR values and susceptibility to symptomatic infection for COVID-19 in Wuhan. a, Estimates of age-specific sCFRs, assuming P_{sym} is 0.5 (red), 0.75 (green) or 0.95 (blue). **b,** Estimate of relative susceptibility to symptomatic infection by age, assuming P_{sym} is 0.50 (red), 0.75 (green) or 0.95 (blue). Symbols in **a,b** show the posterior means, and the bars show 95% credible intervals.

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