



Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active.

Correction

Compartmentalization-aided interaction screening reveals extensive high-order complexes within the SARS-CoV-2 proteome

Weifan Xu, Gaofeng Pei, Hongrui Liu, Xiaohui Ju, Jing Wang, Qiang Ding, and Pilog Li*

*Correspondence: pilogli@mail.tsinghua.edu.cn

<https://doi.org/10.1016/j.celrep.2021.109778>

(Cell Reports 36, 109482-1–109482-18.e1–e7; August 3, 2021)

Due to a technical error, two small images were erroneously converted to gray squares in the original Figure S5A. The original and corrected Figure S5A are included here, and the corrected Figure S5A can now be found with the article online.

The production team apologizes for the error.

A

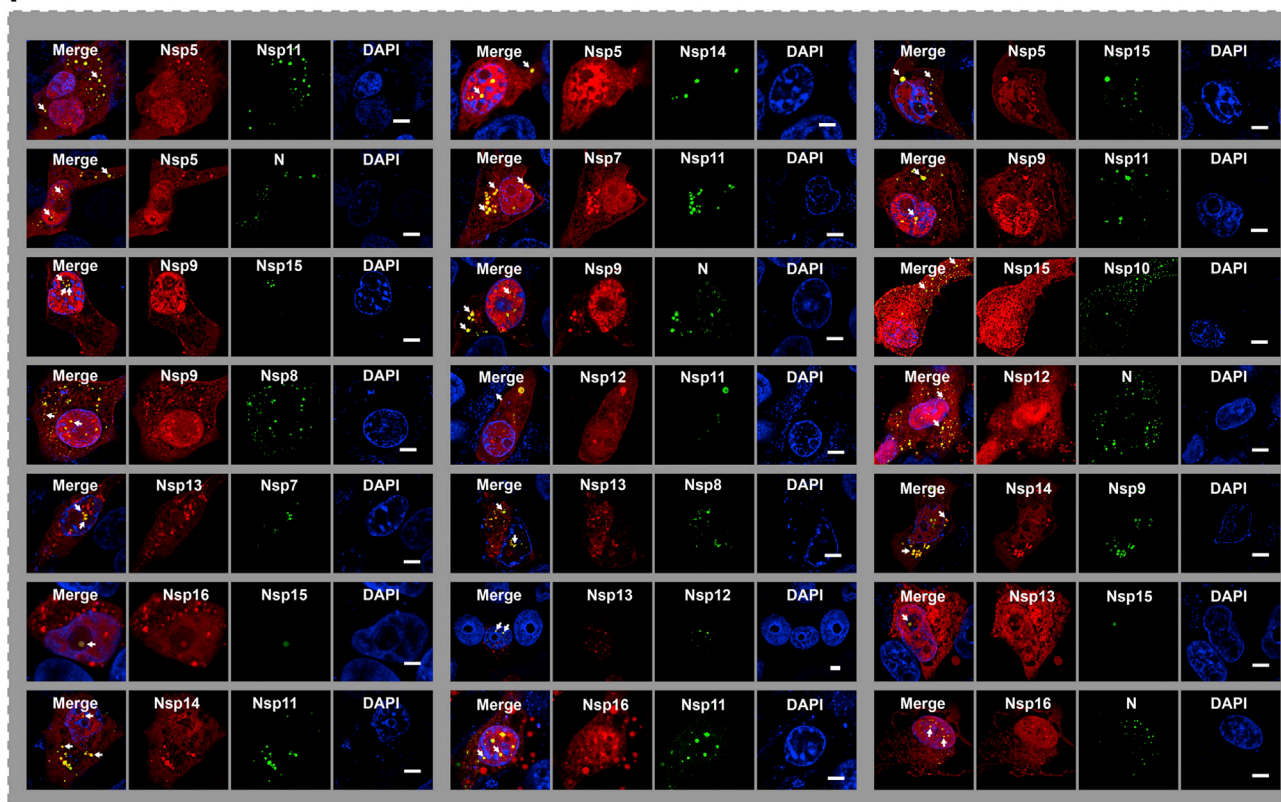


Figure S5A. Positive pairwise (unidirectional-) interactions of SARS-CoV-2 identified by CoPIC screening (corrected)



A

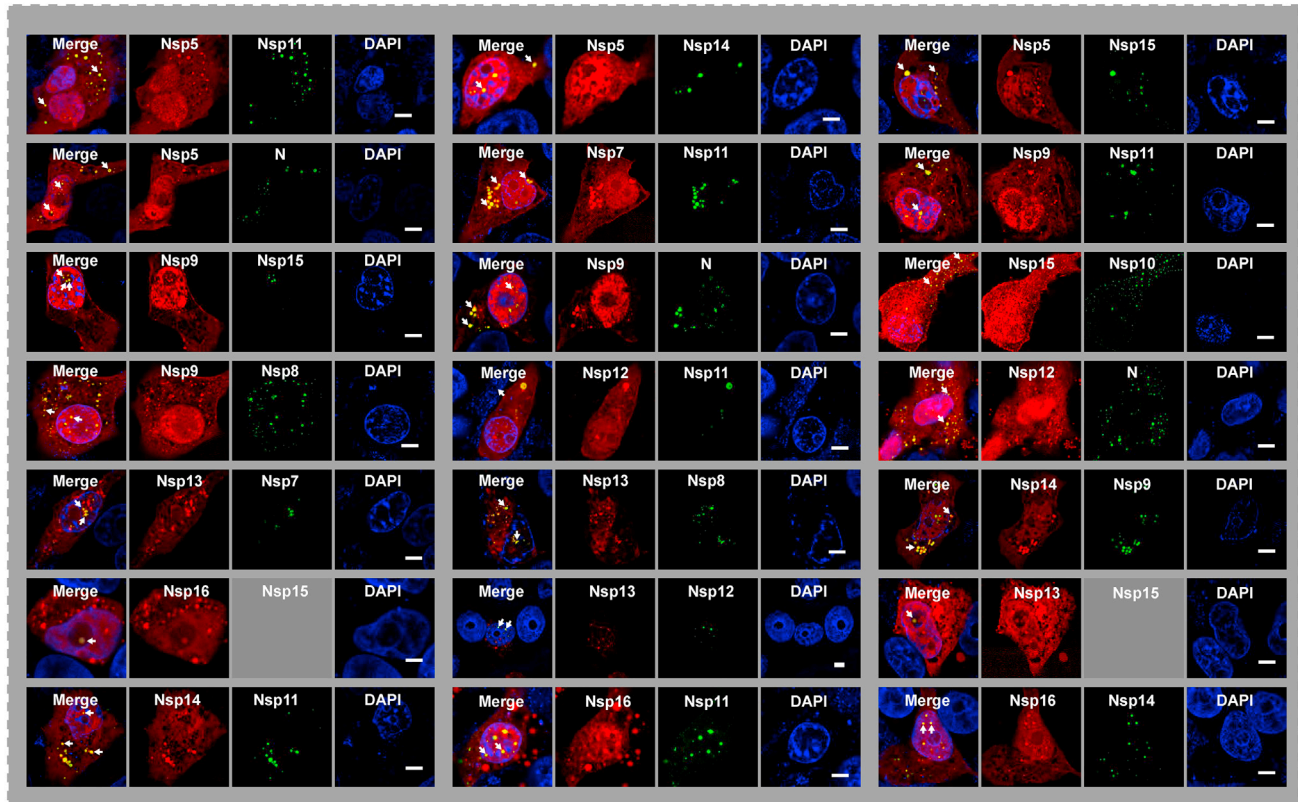


Figure S5A. Positive pairwise (unidirectional-) interactions of SARS-CoV-2 identified by CoPIC screening (original)