Oncogenesis www.nature.com/oncsis

CORRECTION OPEN



Correction: Human colon cancer cells highly express myoferlin to maintain a fit mitochondrial network and escape p53-driven apoptosis

Gilles Rademaker, Brunella Costanza, Justine Bellier, Michael Herfs, Raphaël Peiffer, Ferman Agirman, Naïma Maloujahmoum, Yvette Habraken, Philippe Delvenne, Akeila Bellahcène, Vincent Castronovo 🗈 and Olivier Peulen 🕞

© The Author(s) 2023

Oncogenesis (2023)12:11; https://doi.org/10.1038/s41389-023-00455-5

Correction to: Oncogenesis https://doi.org/10.1038/s41389-019-0130-6, published online 08 March 2019

During figure preparation, the same protein samples were used in the western blots depicted in figures 4 and 5. For the sake of completeness, the same myoferlin western blots were included in figures 4 and 5. It has been noted that the western blot duplication can be misleading. Consequently, the duplicated myoferlin western blots have been removed from figures 4 and 5. The corrected figures are presented below.

Published online: 02 March 2023

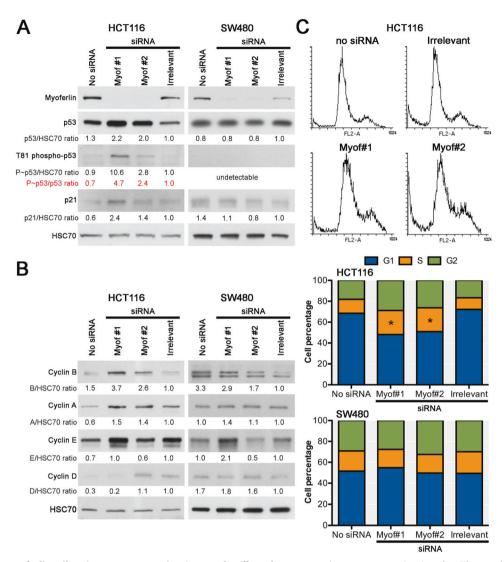


Fig. 4 Effects of myoferlin silencing on p53 activation and cell cycle progression. a p53 activation by Thr81 phosphorylation and subsequent p21 abundance were evaluated in HCT116 and SW480 48h after myoferlin silencing. **b** Cyclin abundance was evaluated by western-blot in HCT116 and SW480 48h after myoferlin silencing. Total protein extracts (10 μg) were subjected to SDS-PAGE followed by western blot analysis with specific antibodies. HSC-70 was used as a loading control. **c** Cell cycle was analyzed by flow cytometry after propidium iodide incorporation in HCT116 and SW480 48h after myoferlin silencing. Distribution of FL2 fluorescence (propidium iodide) was shown in HCT116. Proportion of cells in G1, S or G2 was shown in HCT116 and SW480. One representative experiment out of three is illustrated. **P* < 0.05.

SPRINGER NATURE Oncogenesis (2023)12:11

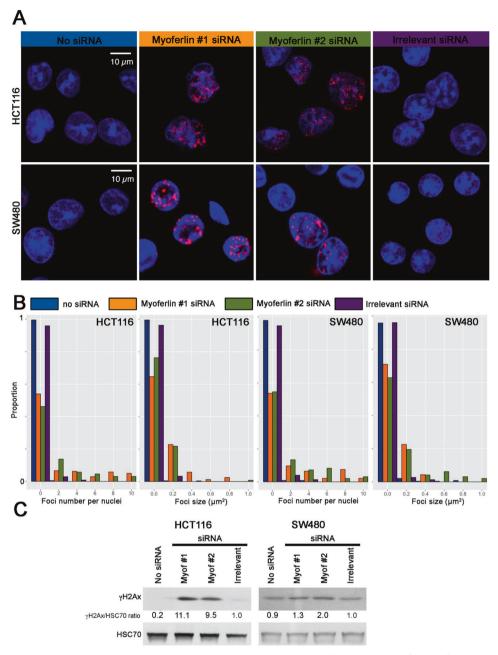


Fig. 5 Myoferlin-silencing induces a DNA damage response. a HCT116 and SW480 cell lines, silenced for myoferlin during 48h, were stained for γ H2Ax and observed under a confocal microscope. b γ H2Ax foci number and size were quantified using ImageJ. Number and size distributions were established (n > 210 nuclei). c γ H2Ax abundance was evaluated by western-blot in HCT116 and SW480 48h after myoferlin silencing. Total protein extracts (10 μ g) were subjected to SDS-PAGE followed by western blot analysis with specific antibodies. HSC-70 was used as a loading control.

Open Access This article is licensed under a Creative Commons on International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit http://creativecommons.org/licenses/by/4.0/.

© The Author(s) 2023

Oncogenesis (2023)12:11 SPRINGER NATURE