CORRECTION

Correction: Validation of Novel Biomarkers for Prostate Cancer Progression by the Combination of Bioinformatics, Clinical and Functional Studies

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The gene LAMB1 appears incorrectly throughout the article. The correct gene name should be LMNB1. In [Fig 8](#), the label for [Fig 8c](#) is incorrect. Please see the corrected [Fig 8](#) here.

OPEN ACCESS

Citation: Alinezhad S, Väänänen R-M, Mattsson J, Li Y, Tallgrén T, Tong Ochoa N, et al. (2016) Correction: Validation of Novel Biomarkers for Prostate Cancer Progression by the Combination of Bioinformatics, Clinical and Functional Studies. PLoS ONE 11(6): e0158255. doi:10.1371/journal.pone.0158255

Published: June 21, 2016

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Fig 8. Images of 3D organotypic cell cultures of PC3 cells, embed in Matrigel, after segmentation and subsequent image analysis using the AMIDA software package A. Untreated cells form large organoids with overt invasive processes. B. Silencing of PLA2G7 and C. laminin beta 1 (LMNB1) result in well-rounded, poorly invasive organoids, with higher cell density. D. In contrast, silencing of the TDRD1 gene results in significant induction of invasive properties, further loss of the structural organization or maturation of organoids, and decreased cell density.

doi:10.1371/journal.pone.0158255.g001

Reference

1. Alinezhad S, Väänänen R-M, Mattsson J, Li Y, Tallgrén T, Tong Ochoa N, et al. (2016) Validation of Novel Biomarkers for Prostate Cancer Progression by the Combination of Bioinformatics, Clinical and Functional Studies. PLoS ONE 11(5): e0155901. doi: [10.1371/journal.pone.0155901](https://doi.org/10.1371/journal.pone.0155901) PMID: [27196083](https://pubmed.ncbi.nlm.nih.gov/27196083/)