# CORRECTION Open Access

# Correction: Granulin as an important immune molecule involved in lamprey tissue repair and regeneration by promoting cell proliferation and migration



Ruixiang Sun<sup>1,2,3†</sup>, Dong Wang<sup>1,2,3†</sup>, Yuxuan Song<sup>1,2,3†</sup>, Qingwei Li<sup>1,2,3</sup>, Peng Su<sup>1,2,3\*</sup> and Yue Pang<sup>1,2,3\*</sup>

<sup>†</sup>Ruixiang Sun, Dong Wang and Yuxuan Song contributed equally to this work

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\*Correspondence: sp4046@163.com; pangyue01@163.com

<sup>1</sup> College of Life Sciences, Liaoning Normal University, Dalian 116081, China <sup>2</sup> Lamprey Research Center, Liaoning Normal University, Dalian 116081, China <sup>3</sup> Collaborative Innovation Center of Seafood Deep Processing, Dalian Polytechnic University, Dalian 116034, China

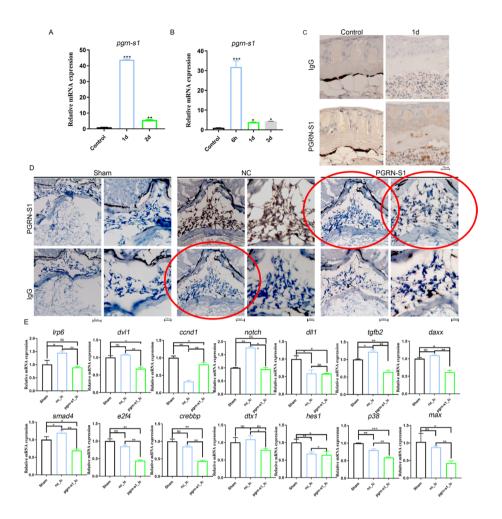
# Correction: Cellular & Molecular Biology Letters (2022) 27:64

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Following publication of the original article [1], the authors identified an error in Fig. 5. They reused the picture of NC group of Fig. 5D (published on line) in the PGRN-S1 group. Thus, they used the correct picture to replace it. The incorrect and the correct figure is given below.

The incorrect Fig. 5 is:

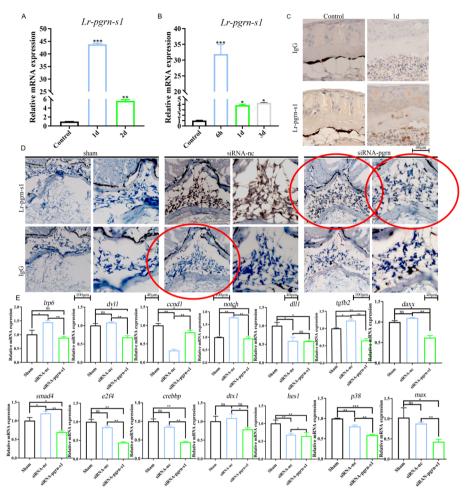




**Fig. 5** Expression of Lr-pgrn-s1 after lamprey skin and spinal cord injury. **A** Expression of Lr-pgrn-s1 in lamprey skin lesions 1d and 2d. **B** Expression of Lr-pgrn-s1 in lamprey spinal cord injury at 6 h, 1d and 3d by qRT-PCR. **C** Detection of tissue localization of Lr-PGRN-S1 in skin injury 1d by immunohistochemistry. Take pictures at  $40 \times$  magnification. **D** Immunohistochemistry of Lr-pgrn-s1 was knocked down 6 h after lamprey spinal cord injury. Results under  $20 \times$  and  $40 \times$  microscopy in Sham, NC (normal control) and Lr-PGRN-S1 groups. **E** The qRT-PCR analysis of genes in MAPK, Notch, Wnt and TGF-β signaling pathways. The statistical differences between experimental groups were detected by the Student's t test. All data were presented as the means  $\pm$  SDs based on three independent samples with three replicates per sample. ns non-significant, \*P < 0.05, \*\*P < 0.001, \*\*\*P < 0.0001.

The correct Fig. 5 is:

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**Fig. 5** Expression of Lr-pgrn-s1 after lamprey skin and spinal cord injury. **A** Expression of Lr-pgrn-s1 in lamprey skin lesions 1d and 2d. **B** Expression of Lr-pgrn-s1 in lamprey spinal cord injury at 6 h, 1d and 3d by qRT-PCR. **C** Detection of tissue localization of Lr-PGRN-S1 in skin injury 1d by immunohistochemistry. Take pictures at 40× magnification. **D** Immunohistochemistry of Lr-pgrn-s1 was knocked down 6 h after lamprey spinal cord injury. Results under 20× and 40× microscopy in Sham, NC (normal control) and Lr-PGRN-S1 groups. **E** The qRT-PCR analysis of genes in MAPK, Notch, Wnt and TGF-β signaling pathways. The statistical differences between experimental groups were detected by the Student's *t* test. All data were presented as the means  $\pm$  SDs based on three independent samples with three replicates per sample. ns: non-significant, \*P<0.005, \*\*P<0.001, \*\*\*P<0.0001

### Reference

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