

CORRECTION

Correction: Gene expression and network-based analysis reveals a novel role for hsa-miR-9 and drug control over the p38 network in glioblastoma multiforme progression

Rotem Ben-Hamo and Sol Efroni*

Correction

It has come to our attention that there is an error in the Acknowledgements section of our article [1]. The Acknowledgements section should read:

SE is supported by the European Union through the IRG program. The results published here are in whole or part based upon data generated by The Cancer Genome Atlas pilot project established by the NCI and NHGRI. Information about TCGA and the investigators and institutions who constitute the TCGA research network can be found at the project website [2].

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References

1. Ben-Hamo R, Efroni S: Gene expression and network-based analysis reveals a novel role for hsa-miR-9 and drug control over the p38 network in glioblastoma multiforme progression. *Genome Medicine* 2011, **3**:77.
2. The Cancer Genome Atlas [<http://cancergenome.nih.gov/>].

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*Correspondence: sol.efroni@biu.ac.il

The Mina and Everard Goodman Faculty of Life Science, Bar Ilan University,
1 Keren-Hayesod St, Ramat-Gan, 52900, Israel