CORRECTION

Correction: Whole Blood Transcriptional Profiling Reveals Deregulation of Oxidative and Antioxidative Defence Genes in Myelofibrosis and Related Neoplasms. Potential Implications of Downregulation of Nrf2 for Genomic Instability and Disease Progression

The PLOS ONE Staff

The image quality of [Fig. 1](#) is poor, making interpretation difficult. Please see the correct [Fig. 1](#) here.

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Citation: The PLOS ONE Staff (2015) Correction: Whole Blood Transcriptional Profiling Reveals Deregulation of Oxidative and Antioxidative Defence Genes in Myelofibrosis and Related Neoplasms. Potential Implications of Downregulation of Nrf2 for Genomic Instability and Disease Progression. PLoS ONE 10(2): e0118049. doi:10.1371/journal.pone.0118049

Published: February 2, 2015

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Figure 1. ATOX1, DEFB122, GPX8, PRDX2, PRDX6, PTGS1, and SEPP1 were progressively and significantly upregulated in patients with ET, PV, and PMF (FDR <0.05). Fold changes for each gene are shown on the y-axis. doi:10.1371/journal.pone.0112786.g001

doi:10.1371/journal.pone.0118049.g001

Reference

1. Hasselbalch HC, Thomassen M, Hasselbalch Riley C, Kjær L, Stauffer Larsen T, Jensen MK et al. (2014) Whole Blood Transcriptional Profiling Reveals Deregulation of Oxidative and Antioxidative Defence Genes in Myelofibrosis and Related Neoplasms. Potential Implications of Downregulation of Nrf2 for Genomic Instability and Disease Progression. PLoS ONE 9(11): e112786. doi: [10.1371/journal.pone.0112786](https://doi.org/10.1371/journal.pone.0112786) PMID: [25397683](https://pubmed.ncbi.nlm.nih.gov/25397683/)