



Corrigendum: Macrophage Polarization: Different Gene Signatures in M1(LPS+) vs. Classically and M2(LPS-) vs. Alternatively Activated Macrophages

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A Corrigendum on

Macrophage Polarization: Different Gene Signatures in M1(LPS+) vs. Classically and M2(LPS-) vs. Alternatively Activated Macrophages

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In the original article, there was a mistake in **Figure 3** as published. Three gene names were misspelled in the previous version. In Figure 3A, M1(=LPS+) panel, the correct gene name for EIFE3 is EIF4E3. In Figure 3B, alternatively activated (IL-4) panel, the correct gene name for WRD4 is WDR4. Also in Figure 3B, the correct gene name for POLRR3K is POLR3K. The corrected **Figure 3** appears below.

Additionally, in the original article, there was a mistake in the legend for **Figure 3** as published. "M1(=LPS+)" was written as "M1(LPS+)", and "M2(=LPS-)" was written as "M2(LPS-)". The correct legend for **Figure 3** appears below.

The authors apologize for these errors and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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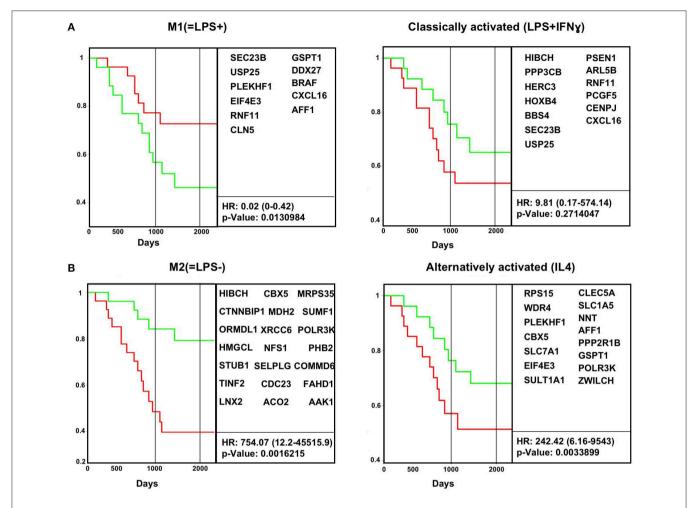


FIGURE 3 | In vivo macrophage signatures predict survival in osteosarcoma cancer biopsy transcriptomes. Survival data for human osteosarcoma cancer biopsies (GSE21257) were analyzed for the impact of M1(=LPS+) and classically activated (LPS+|FN-γ) (A) and M2(=LPS-) and alternatively activated (IL-4) (B) gene expression signatures in the tumor biopsy transcriptome. Kaplan-Meier curves were plotted using ProggeneV2, divided by the median of the mean expression of a tumor-specific gene list (in boxes). Hazard ratio (HR, cox proportional hazard analysis) and significance (log rank P-value) are shown. Red, green curves indicate high, low expression of the respective signature genes. The two vertical black lines indicates 3 and 5 years, respectively.