Corrigendum: Transcriptional Time Course After Rotator Cuff Tear

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A Corrigendum on

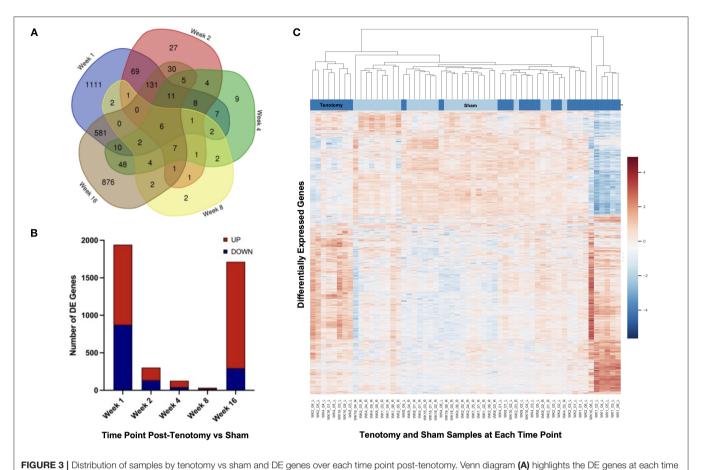**Transcriptional Time Course After Rotator Cuff Tear**

by Vasquez-Bolanos, L. S., Gibbons, M. C., Ruoss, S., Wu, I. T., Vargas-Vila, M., Hyman, S. A., Esparza, M. C., Fithian, D. C., Lane, J. G., Singh, A., Nasamran, C. A., Fisch, K. M., and Ward, S. R. (2021). Front. Physiol. 12:707116. doi: 10.3389/fphys.2021.707116

In the original article, there was a mistake in **Figure 3** as published. **An older version of the figure was published by mistake.** The corrected **Figure 3** appears below.

The authors apologize for this error 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 | Distribution of samples by tenotomy vs sham and DE genes over each time point post-tenotomy. Venn diagram **(A)** highlights the DE genes at each time point and the overlap with other time points. The bar chart **(B)** displays the number of DE genes which are up or down regulated at each timepoint. Data in the heatmap **(C)** is presented as normalized expression for each tenotomy and sham sample at each time point post-tenotomy with a z-score scale by rows and an average hierarchical clustering by columns.