



**Fig. S8. Effect of TGF- $\beta$  and oATP treatment on expression of genes interacting within the TGF- $\beta$  pathway (<http://www.netpath.org>).** The hierarchical cluster plot shows the median centralized relative expression data of differentially expressed genes resulting from the comparison of TGF- $\beta$ , TGF- $\beta$ +oATP and oATP treated with untreated T<sub>conv</sub> cells. Genes showing a similar expression profile are clustered together (left tree) as well as samples showing a similar expression behavior among the TGF- $\beta$  pathway (top tree).