

Supplementary material: Potential Metabolite Markers for Pancreatic Cancer Identified by Metabolomic Analysis of Induced Cancer-Associated Fibroblasts

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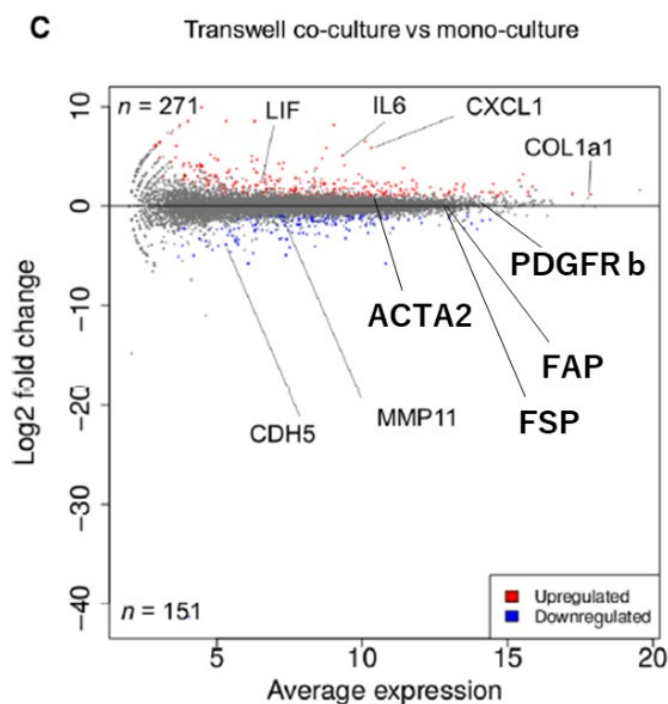


Figure S1: Global gene expression pattern showed that AD-MSCs were differentiated into CAF. MA plot, a scatter plot of log2 fold change versus the average expression, showing differentially expressed genes (adjusted p -value < 0.05 and \log_2 [fold change] ≥ 1) in transwell co-cultured AD-CAFs compared to those in monocultured AD-MSCs. Upregulated genes are shown in red, and downregulated genes are shown in blue.