

## Supplementary Material

### Contribution of Mesenchymal Stem Cells from Obese Adipose Tissue to PD-L1 Over-Expression and Breast Cancer Progression through Pathogenic Th17 Cell Activation

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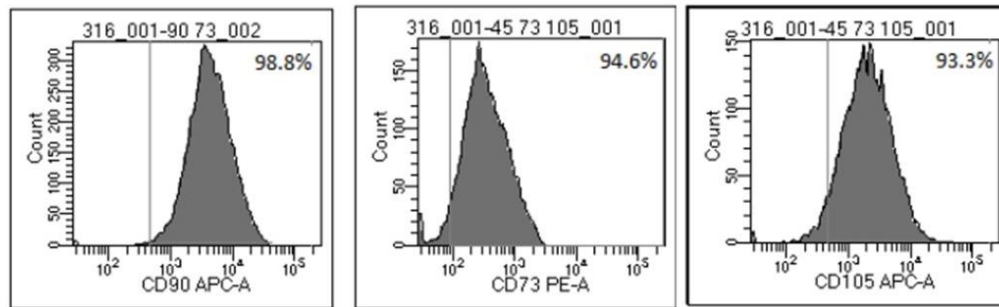
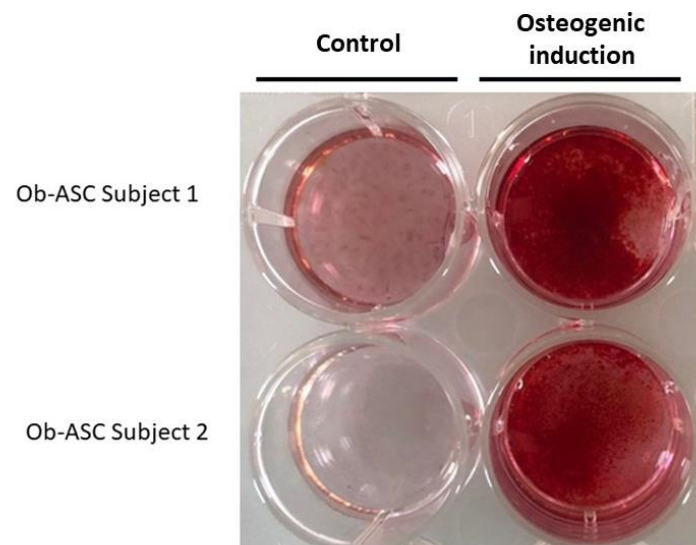
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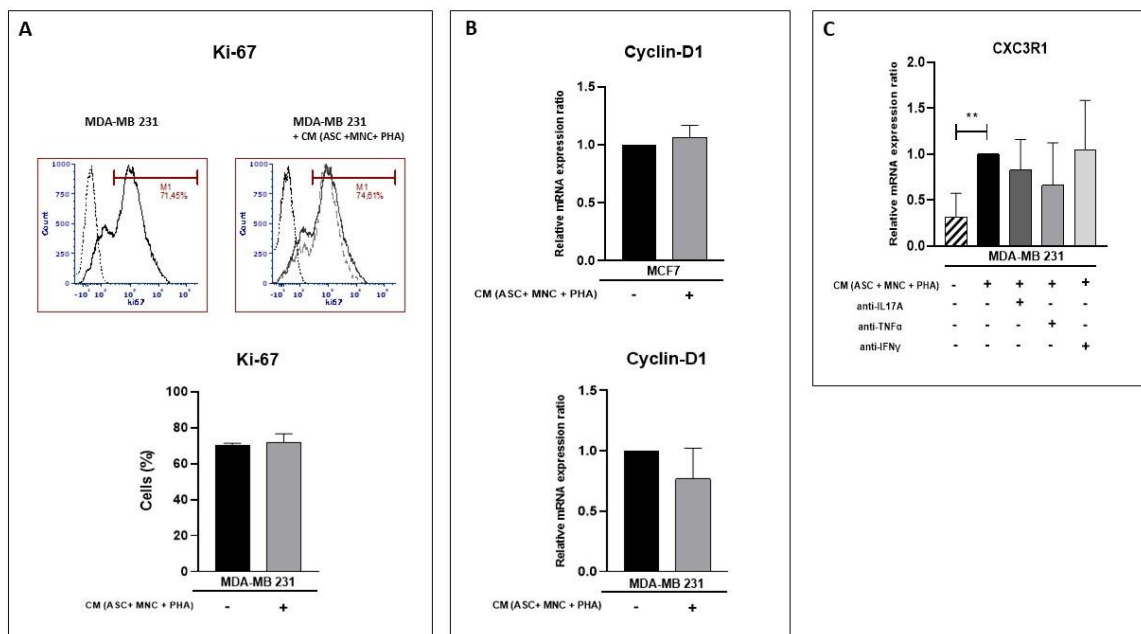
**Table S1.** Primers for Human transcripts used in real-time PCR

Gene name	5'-3' sequence	Amplicon size, bp	RefSeq number
PPIF	F : GGCTACAAAGGCTCCACCTTC R : GAAAGCGGCTTCCGTAGATG	112	NM_005729
IL-6	F : AGCCCTGAGAAAGGAGACATGTAACAAG R : TTCTGCAGGAAGTGGATCAGGACTTT	273	NM_000600
IL-1 $\beta$	F : GGCAATGAGGATGACTTGTT R : TGTAGTGGTGGTCGGAGATT	124	NM_000576
IL-8	F : TTCTGCAGCTCTGTGTGAAG R : ACAGAGCTCTCTCCATCAG	187	NM_000584
VEGF-A	F : AGTACCCTGATGAGATCGAG R : TATGTGCTGGCCTTGGTGAG	161	NM_001025366
MMP-9	F : GGGAAGATGCTGCTGTTCA R : TTCAACTCACTCCGGGAAGT	203	NM_004994
CD-274	F : CTGCAGGGCATTCCAGAAAAG R : GTTCAGCAAATGCCAGTAGG	71	NM_001314029
CX3CR1	F : AGAGGCTGGTTCTTACGATG R : CAGGTTGGCAGTAGCATGAA	170	NM_001171172
CyclinD1	F : CTTCTCTCCAAAATGCCAG R : AGGGCGGATTGGAAATGAAC	113	NM_053056

**A****B**

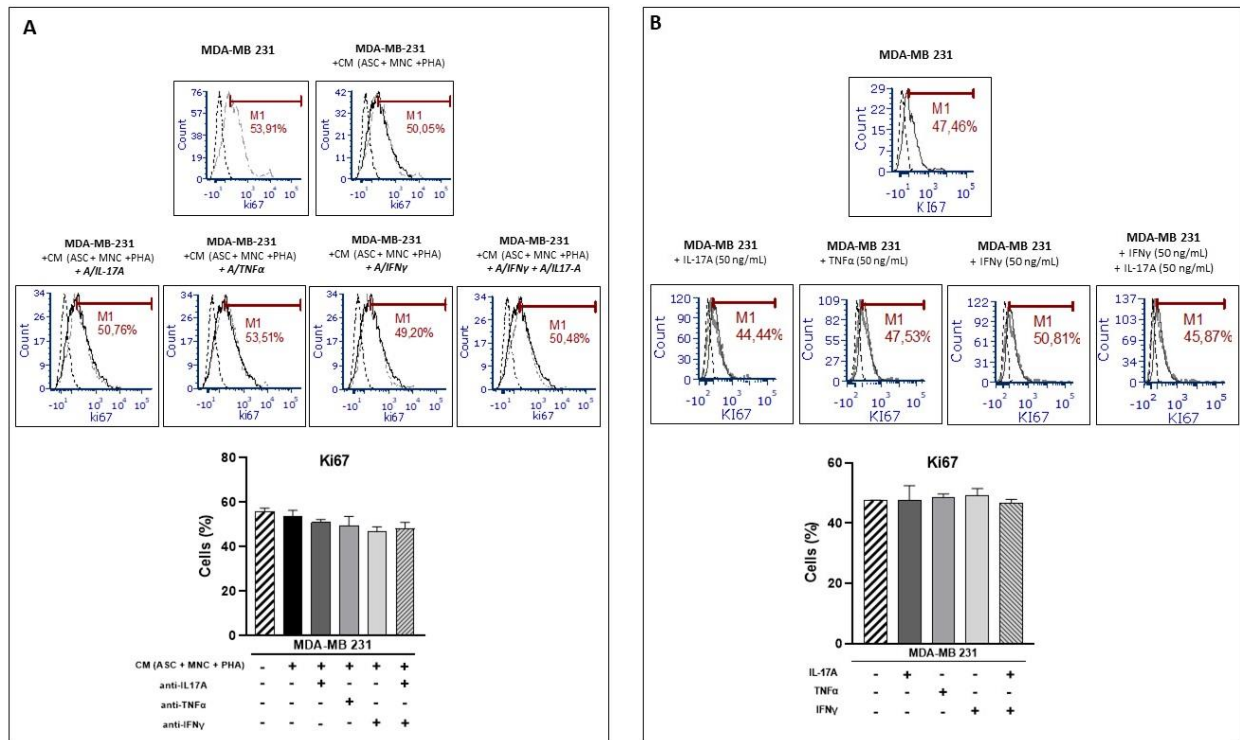
**Supplementary Figure S1. Characterization of Adipose tissue-derived-mesenchymal stem cells (ASC).**

**(A)** ASC were stained for CD90, CD73 and CD105. The expression of these surface molecules was measured by flow cytometry. This figure is representative of 3 experiments. **(B)** ASC were differentiated or not (as control) into osteoblasts using osteoblast differentiation medium for 14 days and stained with alizarin red (calcium deposits).



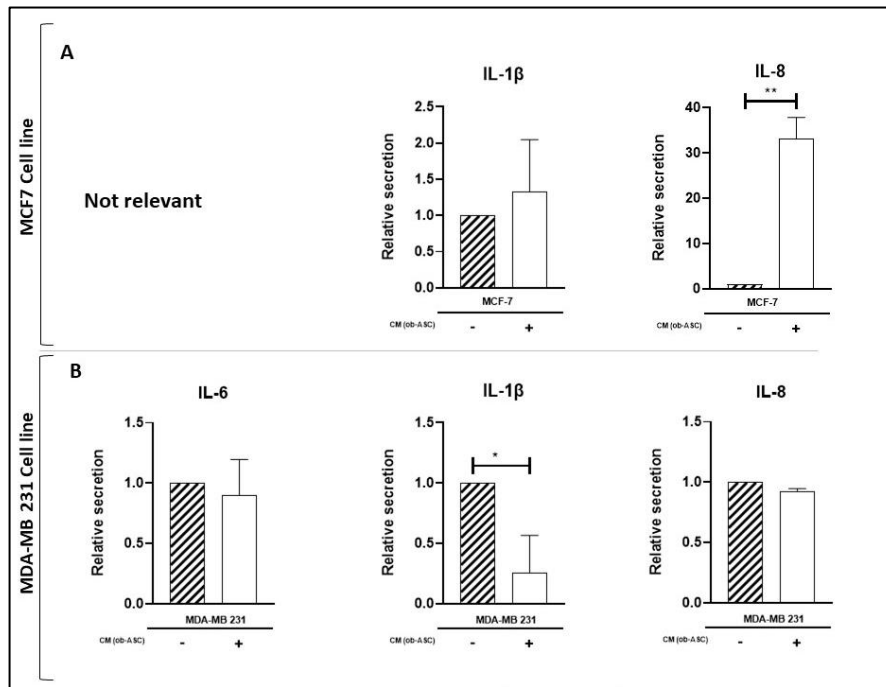
**Supplementary Figure S2. CM from activated co-cultures does not influence BCCL proliferation , but migration**

**(A)** MDA-MB-231 cells ( $0.3 \times 10^5$  cells) were cultured for 24 hours with or without CM harvested from with PHA activated ob-ASC/MNC co-cultures and Ki67 expression were measured by flow cytometry. Overlays correspond to isotypic control (hatched-black) versus Ki67 in MDA-MB231(solid-black) or MDA-MB231 plus CM (hatched-grey). Statistical analysis of 3 experiments is shown. **(B)** CyclinD1 mRNA expression levels were measured in each BCCL by qRT-PCR and results were normalized to PPIF. **(C)** MDA-MB 231 were cultured with CM harvested from PHA-ob-ASC/MNC cocultures plus or minus (i) A/IL-17A, (ii) A/ TNFα or (iii) A IFNγ. CX3CR1 expression levels were measured by qRT-PCR and results were normalized to PPIF. \* represent  $p < 0.05$  as obtained by the one way ANOVA followed by Fisher LSD multiple comparison tests.



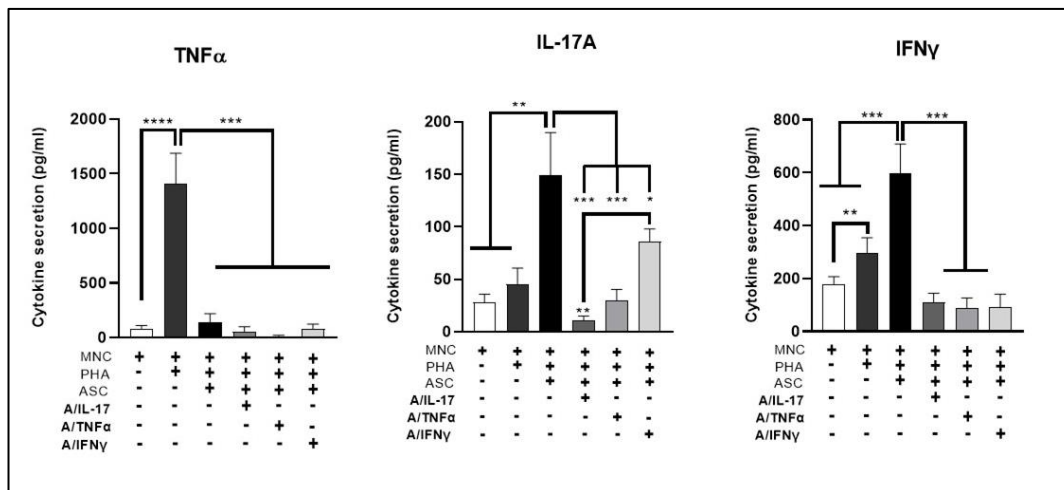
**Supplementary Figure S3: MDA-MB 231 cell proliferation is not influenced by pathogenic Th17cell cytokines**

**(A)** MDA-MB-231 cells ( $0.3 \times 10^5$  cells) were cultured for 24 hours with or without CM harvested from with PHA activated ob-ASC/MNC co-cultures plus or minus A/IL-17A, (ii) A/ TNF-  $\alpha$ , (iii) A/ IFN- $\gamma$  mAbs or (iv) combination of A/IL-17A and A/IFN- $\gamma$  mAbs and Ki67 expression were measured by flow cytometry. **(B)** MDA-MB-231 cells ( $0.3 \times 10^5$  cells) were cultured for 24 hours with or without recombinant human (i)IL-17A (50 ng/mL), (ii) TNF $\alpha$  (50 ng/mL), (iii) IFN $\gamma$  (50 ng/mL) and (iv) combination of IL-17A (50 ng/mL) and IFN $\gamma$  (50 ng/mL) Ki67 expression were measured by flow cytometry. Overlays correspond to isotypic control (hatched-black) versus Ki67 in MDA-MB 231(solid-black) or MDA-MB231 plus cytokines (grey). Statistical analysis of 3 experiments is shown.



**Supplementary Figure S4 : BCCL cytokine secretion profile following culture with Ob--ASC CM**

MCF7 and MDA-MB-231 cells ( $1 \times 10^5$  cells/well) cells were cultured for 24h with CM harvested from ob-ASC alone. **(A,B)** Cell culture supernatants were analyzed by ELISA. Error bars represent standard deviations (SD) from  $n = 3$  independent experiments. \*, \*\* represent  $p < 0.05$ ,  $p < 0.01$ , respectively, as obtained by the one-way ANOVA, followed by Fisher LSD multiple comparison tests.



**Supplementary Figure S5. Neutralization of T cell cytokines in ob-ASC/MNC cocultures, and residual cytokine levels among CM**  
MNC from healthy donors were activated or not with PHA (5 $\mu$ g/mL) and co-cultured with ob-ASC (5:1 ratio) (i) plus or minus A/IL-17A, (ii) A/ TNF $\alpha$  or (iii) A/ IFN $\gamma$  mAbs. 48h later, cytokine secretion was analyzed by ELISA for TNF $\alpha$ , IL-17A and IFN $\gamma$  secretion. .  
\*, \*\*, \*\*\*, \*\*\*\* represent  $p < 0.05$ ,  $p < 0.01$ ,  $p < 0.001$ ,  $p < 0.0001$ , respectively, as obtained by the one way ANOVA followed by Fisher LSD multiple comparison tests.