

## **Endogenous retrovirus-encoded Syncytin-2 contributes to exosome-mediated immunosuppression of T cells**

Running title: Immunosuppressive function of Syncytin-2 on T cell

Summary sentence: Human endogenous retroviral Syncytin-2 negatively downmodulates Th1 response via its immunosuppressive domain and acts through its association to exosomes

Keywords: Syncytin-2, Th1 cytokines, syncytiotrophoblast, placenta, exosomes, immunosuppression

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### Abstract

Modulation of the activation status of immune cell populations during pregnancy depends on placental villous cytotrophoblast (VCT) cells and the syncytiotrophoblast (STB) layer. Failure in the establishment of this immunoregulatory function leads to pregnancy complications. Our laboratory has been studying Syncytin-2 (Syn-2), an endogenous retroviral protein expressed in placenta and on the surface of placental exosomes. This protein plays an important role in STB formation through its fusogenic properties, but also possesses an active immunosuppressive domain (ISD). Considering that Syn-2 expression is importantly reduced in preeclamptic placentas, we were interested in addressing its possible immunoregulatory effects on T cells. Activated Jurkat T cells and peripheral blood mononuclear cells (PBMCs) were treated with monomeric or dimerized version of a control or a Syn-2 ISD peptide. Change in phosphorylation levels of ERK1/2 MAP kinases was selectively noted in Jurkat cells treated with the dimerized ISD peptide. Upon incubation with the dimerized Syn-2 ISD peptide, significant reduction in Th1 cytokine production was further demonstrated by ELISA and Human Th1/Th2 Panel Multi-Analyte Flow Assay. To determine if exosome-associated Syn-2 could also have an immunosuppressive effect, placental exosomes were incubated with activated Jurkat T cells and PBMCs. Upon quantification of Th1 cytokines in the supernatants, T cell activation was severely reduced. Interestingly, exosomes from Syn-2-silenced VCT incubated with PBMCs were less suppressive when compared to exosome derived from VCT transfected with control siRNA. Our results suggest that Syn-2 could be an important immune regulator both locally and at the systemic level, via its association with placental exosomes.

## Introduction

During normal pregnancy, the maternal immune system has been thought to be only modulated in order for the immunologically foreign foetus to be protected from rejection [1].

30 Recent studies have highlighted that placenta-mediated modulation of the activity of the immune responses is more complex and involves pro-inflammatory and anti-inflammatory phases, in which placental trophoblasts play an essential role [2]. Several immune cell populations are targeted by trophoblasts at different stages of pregnancy and include monocytes/macrophages, Treg cells, NKs, and dendritic cells, which are recruited and modulated by various secreted  
35 chemokines and cytokines [3]. CD4+ and CD8+ T cells also importantly participate in the immune response through their presence in circulating maternal blood [4]

Placenta development involves the differentiation of trophoblasts into extravillous and villous cytotrophoblasts [5]. Unlike extravillous cytotrophoblasts, villous cytotrophoblasts are non-invasive but fuse to form and maintain the structure of the peripheral syncytiotrophoblast  
40 layer through a well-regulated process. A number of studies have demonstrated that envelope genes from Human Endogenous Retrovirus (HERV), remnant of former ancestral retroviral infectious events, are implicated in this fusion through their fusogenic domain [6, 7]. Convincing data has clearly suggested that envelope (Env) Syncytin-1 and Syncytin-2 indeed participate in the formation of the syncytiotrophoblast layer by mediating fusion with underlying  
45 cytotrophoblast cells through interaction with their specific receptors [8].

Interestingly, HERV Env proteins have been reported to contain an immunosuppressive domain (ISD) in their sequence, which shared similarity to a previously described immunoregulatory region of envelope proteins from exogenous retroviruses [9-11]. We and

others have further provided evidence that these HERV Env ISD were functional [12-16].

50 Importantly, the potential immunomodulatory role of HERV Env proteins in pregnant women have been suggested to be altered in pregnancy disorders, such as pre-eclampsia [17-21]. In all these studies, including those with ISD region from exogenous retroviruses, a 17 amino acid-long peptide in a dimerized form was shown to be sufficient to mediate immunosuppression [16, 22]. Although the exact molecular mechanisms and interacting partners involved in the  
55 modulation of the immune response are not known, the ISD domain-induced cell signalling results in activation of ERK1/2 MAP kinases and other cellular effectors [22]. Studies on the Friend murine leukemia virus have further shown that the ISD affects both the innate and adaptive immunity [11].

A number of studies have suggested that the HERV Env-specific immunosuppressive  
60 activity might be attributed to their incorporation in extracellular microvesicles, such as exosomes. These various vesicles are released from the syncytiotrophoblast and have been shown to functionally alter several immune cell types [23, 24]. We and others have clearly demonstrated that Syncytin-1 is present on the surface of these extracellular vesicles and that, in this context, Syncytin-1 was suggested to be immunosuppressive [16, 25].

65 A recent study from our team has demonstrated that like Syncytin-1, Syncytin-2 is also incorporated in the surface of villous cytotrophoblast-derived extracellular vesicles. As limited studies have been aimed at the immunosuppressive activity of Syncytin-2, we were interested in studying the ISD activity of Syncytin-2 in the form of a dimerized peptides and in association to exosomes.

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## Materials and Methods

### Reagents, antibodies and peptides

75 The following activating agents were purchased from Millipore Sigma (Oakville, Canada):  
phorbol 12-myristate 13-acetate (PMA) (P1585) and ionomycin (I3909). In this study, the  
following antibodies were used (see also Supplementary Table 1): mouse anti-CD28 (#302902,  
Biolegend, San Diego CA), mouse anti-CD3 (#317318, Biolegend), rabbit anti-phospho-ERK1/2  
(#9101, Cell Signaling, Danvers MA), rabbit anti-ERK1/2 (#4695, Cell Signaling), mouse anti-  
80 TSG101 (ab83, Abcam, Cambridge MA) and horseradish peroxidase (HRP)-conjugated goat  
anti-rabbit/ anti-mouse antibodies (# 7074 and #7076; Cell Signaling). Antibodies against  
Syncytin-2 have been described previously and target the extracellular SU subunit [26]. Peptides  
were synthesized by solid phase peptide synthesis and were characterized by LC/MS-TOF.  
Dimerization was achieved by dimethyl sulfoxide oxidation. Amino acid sequence of the  
85 peptides were as follow: LQNRRLDMLTAAQGGI (Sync-2 ISD) and  
IGGQAATLMDLGRNQL (Rev Sync-2-ISD), the latter containing the same amino acid in  
reverse sequence (negative control).

### Cell line

90 The human Jurkat cell line was obtained from American Type Culture Collection (Manassas VA)  
and was maintained in RPMI 1640 medium (350-000-CL; Wisent, Montreal, Canada) and 10%  
exosome-free FBS (ultracentrifuged overnight at 100,000 g at 4°C; 12483-020; Life technologies,  
Burlington, Canada).

**Isolation of primary cytotrophoblast cells and peripheral blood mononuclear cells (PBMCs)**

This study was approved by the Ethic committees of both Université du Québec à Montréal and The Ste Justine Hospital. Venous blood was collected from four healthy male adult donors and PBMCs were prepared from Ficoll gradient, as previously described [27]. Human placentas of  
100 uncomplicated term pregnancies from 37 to 41 weeks were obtained after spontaneous vaginal delivery. All participating women signed an informed consent form. Primary human villous cytotrophoblasts were isolated through a previously published protocol [28]. Briefly, placental villi were cut, thoroughly washed to remove blood, and digested four times in Hanks' balanced salt solution containing trypsin (from  $9.6 \times 10^5$  to  $1.8 \times 10^6$  U) and DNase I (Millipore Sigma;  
105 from 15 to 30 mg per digestion) for 30 minutes at 37°C in a water bath under continuous shaking. Dispersed cells were layered on top of a discontinuous 5%-70% Percoll gradient and centrifuged for 23 min at 507 g. Intermediate layers, with a density between 1.048 and 1.062, containing cytotrophoblast cells, were collected and washed extensively. Cells were seeded at a density of  $1.5 \times 10^6$  cells/well in Dulbecco's modified Eagle's medium (Wisent, Montreal, Canada), 2 mM  
110 glutamine, 10% exosome-depleted FBS, and penicillin/streptomycin/neomycin (Invitrogen Canada, Inc., Burlington, Canada) and cultured for a maximum of 4 days. The purity of each cytotrophoblast preparation was evaluated by flow cytometry with FITC-conjugated monoclonal anti-cytokeratin 7 antibody (1:500) (CBL194F, Millipore Sigma), a specific marker of trophoblasts. Only preparations with a minimum of 96% of cytotrophoblast cells were used. All  
115 experiments with primary cytotrophoblast cells were conducted in triplicate with different placenta donors.

**120 Cell transfection**

Jurkat cells ( $1 \times 10^7$  cells) were transfected by the DEAE-Dextran protocol, as previously described [29]. Plasmids pNFAT-Luc and pNF- $\kappa$ B-Luc have been previously described and respectively contains NFAT and NF- $\kappa$ B-binding sites positioned upstream of a minimal promoter and the luciferase reporter gene [27]. Small interfering RNAs (siRNA) were transfected in human  
125 primary cytotrophoblast cells using the Microporator MP-100 apparatus (Digital Bio). All siRNAs were synthesized by Invitrogen Canada (Syncytin-2, s53886 and the negative control, AM4642). Freshly isolated primary cytotrophoblast cells ( $1.5 \times 10^6$ ) were transfected with 300 ng siRNA using the MicroPorator device with 1 pulse at 1300 V (30 ms.).

**130 Isolation of exosomes**

Supernatants of cultured human primary cytotrophoblast cells (day1 and day 2) were first cleared by successive centrifugations at 2000g for 15 min and at 10,000g (4°C) for 30 min. Following two subsequent 100,000g centrifugations (4°C), supernatants were discarded and pellets containing exosomes were resuspended in PBS (100  $\mu$ l). Exosome preparations were stored at -  
135 80°C in 50  $\mu$ l aliquots and analysed by Western blot.

**Incubation of Jurkat and PBMCs with peptides and exosomes**

In a first series of experiments, unstimulated Jurkat cells were incubated in serum-free medium for 2 h at 37 °C and treated with 0 to 60  $\mu$ M of monomeric or dimeric Sync-2-ISD or Rev Sync-  
140 2-ISD peptides between 2 and 60 min at 37°C. In subsequent experiments, PBMCs or Jurkat ( $5 \times 10^5$  cells) were cultured in sodium bicarbonate-free supplemented RPMI 1640 medium (20mM

Hepes, 2mM L-glutamine, 40 µg/mL gentamicin and 10% FBS). Cells were pre-treated for 2 h with Sync-2-ISD or Rev Sync-2-ISD peptides before activation with PMA (20 ng/ml)/Ionomycin (1 µM) or anti-CD3 (0.5 µg/ml) and anti-CD28 (1.5 µg/ml) and cultured for 18 to 48 h before  
145 harvesting the supernatant. For exosomes, PBMC or transfected Jurkat cells ( $5 \times 10^5$  cells) cultured in sodium bicarbonate-free supplemented RPMI 1640 medium (20mM Hepes, 2mM L-glutamine, 40 µg/ml gentamicin and 10% exosomes free FBS) were either left untreated or pre-treated for 24 h with different concentration of exosomes before activation with PMA/Ionomycin or anti-CD3/anti-CD28 antibodies. Treated cells were cultured at 37°C prior to analyses  
150 (luciferase activity, RT-PCR and harvesting of supernatant). Following incubation, cell viability was determined by Lactate Dehydrogenase Activity Assay (MAK066, Sigma-Aldrich), according to provided instructions (Promega).

#### Western blot analysis

155 Cells were resuspended in 2× SDS sample buffer containing 10% 2-mercaptoethanol, and a mixture of proteinase (Roche Diagnostics, Laval, Canada) and phosphatase inhibitors (Millipore Sigma) and incubated on ice for 15 min. Protein concentrations were evaluated using the bicinchoninic acid (BCA) protein assay (Thermo Fisher Scientific Inc, Rochester, New York). After boiling for 5 min., cell lysates were run on a 10-12% SDS-PAGE and transferred on  
160 polyvinylidene fluoride (PVDF) membranes (Millipore). Membranes were blocked in 5% powdered milk in Tris saline, pH 7.5, 0.15% Tween 20 (Millipore Sigma) for 1 h at room temperature. The following primary antibodies were then added: anti-phospho-ERK1/2 (1:1000), anti-ERK1/2 (1:1000), anti-Syncytin-2 (1:5000), and anti-TSG101 (1:1000). After overnight incubation at 4°C, membranes were washed three times in PBS-Tween 0.05% and incubated with

165 horseradish peroxidase (HRP)-conjugated goat anti-rabbit/anti-mouse antibodies (1:5000) for 1 h at room temperature. After three washes, signals were then revealed with the ECL Western blotting reagent (RPN2106, Millipore Sigma). Membranes were scanned with the Fusion FX5 system (Montreal Biotech Inc., Dorval, Canada).

#### 170 **RT-PCR analysis**

Total cellular RNA was isolated from cells using RNeasy mini kit (#74106, Qiagen, Mississauga, Canada) according to the supplier's protocol. Briefly, RNA (0.5 µg) was incubated in the presence of oligo(dT) (25 ng/µl), 10 mM DTT, 100 µM dNTP, 10 U SuperScript II reverse transcriptase (18064-014, Thermofisher Scientific, Waltham MA), and 20 U SUPERase-In (AM2696, Invitrogen Canada) at 37°C for 45 min. The resulting cDNAs (1 µl) were then PCR amplified in the presence of 1 U Taq DNA polymerase (Thermofisher Scientific), 1X Taq buffer, 100µM dNTP, and 0.5 µM of each primer. PCR conditions were as follows: a first step of denaturation at 95°C for 1 min, followed by 30 cycles of denaturation (30 s at 95°C), annealing (30 s from 55°C), and extension (30 s at 72°C) with a final cycle of 10 min at 72°C. The following primers were used: 5'-AGCCCATGTTGTAGCAAACC-3' (forward TNF-α), 5'-TGAGGTACAGGCCCTCTGAT-3' (reverse TNF- α), 5'-GAAGGTGAAGGTCGGAGTCAA-3' (forward GAPDH) and 5'-GGAAGATGGTGATGGGATTTTC-3' (reverse GAPDH).

#### **Quantification of Th1 and Th2 cytokine in Jurkat and PBMC cells**

185 Jurkat supernatants were assayed for TNF-α with the commercial ELISA MAX™ Deluxe kit (#430204, Biolegend, San Diego, California). PBMC supernatants were assayed for Th1 and Th2 cytokines using commercial LEGENDplex™ Human Th1/Th2 Panel (8-plex) (#740013,

Biolegend, San Diego, California). The sensitivity of the assays was 0.8 pg/ml for IL-13, 1pg/ml for IL-2, IFN- $\gamma$ , TNF- $\alpha$  and 1.1 pg/ml for IL-5, IL-6 and IL-10.

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### Statistics

All experiments were performed in triplicates, except for certain experiments using LEGENDplex™, which were representative of two independent experiments (indicated in Figure legends). Error bars in the graphic data represent means  $\pm$  S.E.M. Data were analysed by a 1-  
195 tailed or paired Student's t test. Statistical analyses were performed with the GraphPad Prism software (San Diego, CA).

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## Results

**The dimerized Syncytin-2 ISD peptide induces ERK1 and ERK 2 phosphorylation.** Previous studies reported that Syncytin-1 and -2 harbour a functional ISD domain [13, 16]. Based on former analyses of this domain, we were interested in more closely examining the inhibitory function of the Syncytin-2 ISD. Since it was reported that CKS-17, a 17 amino acid-long immunosuppressive peptide corresponding to the equivalent domain of Feline and Murine Leukemia Viruses, triggered activation of ERK 1 and ERK 2 MAP kinases in the monocytic THP-1 cell line [10], we first tested if comparable effects were induced in Jurkat CD4+ T cells. As previously reported, the peptide was tested as a monomer, termed Sync-2-ISD, or as its dimerized form, i.e. (Sync-2-ISD)<sub>2</sub>. Jurkat cells were cultured in serum starved-medium for 2 h and stimulated with different concentrations of these peptides and for different time lengths (Figure 1). We first assessed the impact of the (Sync-2-ISD)<sub>2</sub> dimeric peptide on Jurkat cells at a fixed concentration and for different time lengths (Figure 1A). Kinetic experiments showed that (Sync-2-ISD)<sub>2</sub> induced ERK1/2 phosphorylation in a time-dependent manner, as demonstrated by Western blot. In sharp contrast, no induction (or very minimal levels) of ERK1/2 phosphorylation was noted in Jurkat cells treated with monomeric/dimeric version of Rev Sync-2-ISD or the Sync-2-ISD monomer, demonstrating the specificity of the signalling potential of the Syncytin-2 ISD region. Levels of total ERK remained generally stable in these experiments. Furthermore, resulting densitometry analyses of the phosphorylation status of ERK1/2 normalised for protein levels further confirmed the important induction of phosphorylation in response to treatment with the (Sync-2-ISD)<sub>2</sub> peptide (Figure 1B). A dose response experiment was next performed and revealed a gradual increment of ERK1/2 phosphorylation in (Sync-2-ISD)<sub>2</sub>-treated Jurkat cells,

which levelled off at 30  $\mu$ M (Figure 1C). No similar induction was noted in Jurkat cells treated with the control non-dimerized form of the peptide (data not shown).

225           These results thereby indicated that the dimerized representative peptide of the ISD domain of Syncytin-2 specifically induced ERK1/2 phosphorylation in Jurkat cells in a comparable manner to the induction observed with ISD domain of envelope proteins from different retroviruses.

230   **(Sync-2-ISD)<sub>2</sub> inhibits TNF- $\alpha$  production in Jurkat cells.** In order to determine if the previously reported immunosuppressive capacity of Syncytin-2 could also affect activated T cells [13], monomeric/dimeric forms of Sync-2-ISD or Rev Sync-2-ISD were incubated in the presence of Jurkat T cells. After addition of PMA/Ionomycin, cell supernatants were quantified for TNF- $\alpha$  production. As presented in Figure 2, (Sync-2-ISD)<sub>2</sub> inhibited TNF- $\alpha$  production in  
235   response to PMA/Ionomycin stimulation of Jurkat cells, whereas its monomeric form or Reverse peptide controls did not show a similar important reduction. Of note, the dimerized Reverse Sync2-ISD did show significant reduction at 24 h post-treatment, but at a lower extent than (Sync-2-ISD)<sub>2</sub>-treated Jurkat cells and not at the 48 h time point. As expected TNF- $\alpha$  levels were undetectable in Jurkat cells treated with peptides in absence of PMA/Ionomycin (data not  
240   shown). Importantly, these treatments had no impact on cell viability, based on measurement of lactate dehydrogenase activity (data not shown).

          These data thereby confirmed that the ISD domain of Syncytin-2 demonstrated a specific inhibitory effect on T cell activation and that only the dimeric form was capable of inducing this inhibition.

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**(Sync-2-ISD)<sub>2</sub> inhibits cytokine production in PMA-Ionomycin-activated PBMCs.** In order to address the immunosuppressive potential of the Syncytin-2 ISD in a more physiologically relevant manner and to evaluate a wider range of cytokines, we next tested whether the (Sync-2-ISD)<sub>2</sub> peptide modulates the cytokine profile of PBMCs. Freshly isolated PBMCs were thus stimulated or not with PMA/Ionomycin following pre-incubation with dimerized Syncytin-2 ISD peptide or respective controls. Th1 and Th2 cytokine levels were next quantified in supernatants from PBMCs through the Biolegend multiplex analysis kit. This assay relies on the use of beads of two different sizes and with variation in their fluorescence. Each set of beads are bound to antibodies against a specific cytokine and, upon binding to their specific cytokine, can be detected and analysed following the addition of a pool of biotinylated antibodies recognizing all targeted cytokines. Using this assay, we first compared untreated vs. activated PBMCs (24 h) for the production of Th1 and Th2 cytokines (Figure 3A). Results showed that Th1 cytokines, IL-2, TNF- $\alpha$  and IFN- $\gamma$  were responsive to activation as depicted in our flow cytometry data. As opposed to this induction, only weak stimulation of Th2 cytokines was observed at 24 h post-stimulation. When PBMCs were incubated in the presence of (Sync-2-ISD)<sub>2</sub>, an important inhibition of the induced IL-2, IFN- $\gamma$  and TNF- $\alpha$  levels was noted, while no such strong impact was apparent in Rev (Sync-2-ISD)<sub>2</sub>-treated PBMCs (Figure 3B-C). Similar analyses were conducted on PBMCs stimulated for 48 h (Figure 4). Th1 cytokine production was again importantly induced, although Th2 cytokines showed more important increment in their levels in PBMC supernatant, most notably for IL-13 (Figure 4A). Importantly, induced cytokines were severely reduced when cells were incubated with (Sync-2-ISD)<sub>2</sub>, while a limited impact was observed in (Rev Sync-2-ISD)<sub>2</sub>-treated cells.

Through internal controls and standard curves provided in the assay, values of cytokine concentration for data presented in Figures 3 and 4 were estimated and are depicted in Figure 5.

270 Again, as demonstrated above, the dimerized Syncytin-2 ISD peptide was clearly shown to confer an inhibitory capacity toward Th1 cytokine production in activated PBMCs. Only limited inhibition by the control dimeric peptide was noted in treated PBMCs. In these experiments, no reduction in cell viability was noted in any of the tested treatment (data not shown).

These data thereby further supported the immunosuppressive function of the Syncytin-2  
275 ISD region and indicated that Th1 cytokines were consequently blocked in their production.

**(Sync-2-ISD)<sub>2</sub> inhibits cytokine production in anti-CD3/anti-CD28-activated PBMCs** To corroborate the above results and to test more representative T-cell-specific activators, PBMCs were treated with a combination of anti-CD3/anti-CD28 antibodies prior to their exposure to the peptides. As shown in Figure 6A and B, analyses of supernatant of PBMCs harvested at 24 and  
280 48 h after activation by the multiplex analysis kit again revealed that Th1 cytokines (solely depicted by IFN- $\gamma$  and TNF- $\alpha$  in these graphs) were again strongly induced but severely hampered upon the addition of the (Sync-2-ISD)<sub>2</sub> peptide. In contrast, incubation with (Rev Sync-2-ISD)<sub>2</sub> led to a modest impact on levels of Th1 cytokines. Estimated concentrations of Th1  
285 cytokines, including IL-2 (Figure 7) further confirmed this specific inhibitory action of (Sync-2-ISD)<sub>2</sub> in comparison to the control dimerized peptide.

Using two different combinations of activators, our data thus demonstrated that the ISD domain of Syncytin-2 possessed immunosuppressive activity on activated PBMCs, mainly toward a Th1 response.

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**Placental exosomes down-regulate TNF- $\alpha$  expression and activation of Jurkat cells.** Given the immunosuppressive potential of the Syncytin-2 ISD domain, we next examined the contribution of Syncytin-2 to the inhibitory capacity of placental exosomes on T cell activation. We first confirmed that indeed these exosomes were capable of negatively modulating T cell activation. Jurkat cells were thus incubated in the presence of increasing quantities of exosomes isolated from cultured villous cytotrophoblasts and then stimulated with the PMA/Ionomycin combination (Figure 8). Following analyses of by RT-PCR and normalisation with GAPDH, TNF- $\alpha$  expression was induced following PMA/Ionomycin treatment, while addition of trophoblast exosomes inhibited this induction (Figure 8A). Quantification of supernatant-associated TNF- $\alpha$  confirmed that placental exosomes reduced TNF- $\alpha$  secretion in Jurkat cells (Figure 8B). In order to further demonstrate the inhibitory capacity of cytotrophoblast-derived exosomes on T cells, Jurkat cells were transfected with constructs containing luciferase reporter gene under the regulation of either NF- $\kappa$ B or NFAT, two transcription factors of high importance in activation of T cells. As shown in Figure 8C-D, activation of both transcription factors by PMA/Ionomycin was severely affected following incubation with exosomes. This effect was further demonstrated to be dose-dependent (Figure 8D).

These results hence corroborated former studies showing that placental exosomes, in this case produced for villous cytotrophoblasts, act negatively on the activation of T cells.

**Impact of exosome-associated Syncytin-2 on the immunosuppressive capacity of placental exosomes.** Since we and others have previously demonstrated that both Syncytin-1 and Syncytin-2 are integrated at the surface of placental extracellular vesicles (including exosomes) [16, 25, 30], we were thus interested in determining the contribution of these proteins in exosome-

mediated immunosuppression. Using the FACS-based approach described above, we first  
315 quantified production of IFN- $\gamma$  and TNF- $\alpha$  (vs. IL-4) by PBMCs activated by the addition of anti-  
CD3/anti-CD28 antibodies and incubated in the presence of placental exosomes. Placental  
exosomes indeed showed an important inhibitory effect on the induced production of Th1  
cytokines (Figure 9; 10A). The addition of exosomes to PBMCs did not lead to higher cell death  
in comparison to untreated cells (data not shown).

320 Based on these results, we focussed on Syncytin-2 for its contribution in the  
immunosuppressive capacity of placental exosomes. To selectively deplete Syncytin-2 from these  
exosomes, Syncytin-2-specific siRNAs were tested, as we had previously noted efficient  
repression of expression by this approach [26, 30]. As expected, when isolated villous  
cytotrophoblasts were transfected with Syncytin-2 siRNAs, after 24 and 48 h, isolated exosomes  
325 showed limited levels of Syncytin-2 as compared to exosomes from control (scrambled) siRNA-  
transfected cytotrophoblasts (Figure 10B).

We then compared normal vs. Syncytin-2-deficient exosomes for their  
immunosuppressive potential on anti-CD3/anti-CD28-activated PBMCs. As depicted in Figure  
10A, exosomes derived from villous cytotrophoblasts transfected with control (scrambled)  
330 siRNAs showed strong immunosuppressive abilities, as demonstrated by IFN- $\gamma$ , IL-2 and TNF- $\alpha$   
levels present in the supernatant of treated PBMCs. However, importantly, Syncytin-2-depleted  
exosomes harvested at two different days (day 1 and 2) of culture of transfected villous  
cytotrophoblasts demonstrated a lesser inhibitory effect on IFN- $\gamma$ , IL-2 and TNF- $\alpha$  production.  
Results of the quantification of the different Th1 cytokines from these FACS analyses are  
335 summarized in Figure 11 for both 24 and 48 h conditions. These data indeed strongly argued that

placental exosomes showed an immunosuppressive property on Th1 cytokine secretion and that depletion of Syncytin-2 from their surface hampered this inhibitory potential.

Overall, these data demonstrated that Syncytin-2 contributed to the immunosuppression mediated by placental exosomes on activated PBMCs by reducing levels of secreted Th1 cytokines.

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## Discussion

345 In this study, we have analysed the immunosuppressive properties of Syncytin-2. It is now established that retroviral infections often promote severe immunosuppression in both animals and humans through a retroviral immunosuppression domain (ISD) [22, 31]. Based on our results, we have demonstrated that Syncytin-2 ISD, similarly to peptide CKS-17 and the Syncytin-1 ISD, activates Mitogen Activated Protein Kinase (MAP-kinase) and leads to inhibition of cytokine production in PBMCs activated by PMA/Ionomycin or by anti-CD3/anti-CD28 [10, 16]. Furthermore, an equal inhibition was demonstrated in Jurkat T cells, in which 350 TNF- $\alpha$  synthesis was specifically reduced following exposure to (Sync-2-ISD)<sub>2</sub>.

During normal pregnancy, the second trimester is characterized by a shift toward production of Th2 cytokines and inhibition of the Th1 immune response [2]. Our results 355 demonstrated that the Syncytin-2 ISD was able to inhibit the production of Th1 cytokines (TNF- $\alpha$ , IFN- $\gamma$ , IL-2 and IL-6), which extends the observation of the impact of Syncytin-1 ISD on the inhibition of Th1 cytokine secretion [16]. These results concur with many studies showing that HERV envelope proteins (like for several exogenous retroviruses and ERV from other species) harbour functional ISD domain that allowed viruses to achieve infection in its hosts before being 360 co-opted [32]. Interestingly, a previous study had indicated that the ISD region of HERV-K Envelope inhibited the proliferation of activated PBMCs and that this same peptide increased IL-10 production [14]. We have not been able to demonstrate consistent induction of IL-10 secretion in ISD-treated unstimulated or stimulated PBMCs and this could be attributed to concentration of the peptide and variation in time kinetics used to measure cytokine production. Clearly, however, 365 an early impact on Th1 cytokine is apparent and does not seem to rely on IL-10 production,

although this cytokine might intervene at later time points in regulating Th1 response. The function of HERV Env ISD needs further investigation but, in the context of pregnancy, numerous studies including our own, provide strong evidence for its role in regulating the immune response [33]. As pregnancy-associated immunity has been compared to immunity related to cancer development [2], it can also be argued that tumours associated with high expression levels of HERV Env might also benefit from a functional ISD in escaping the immune response, although other roles attributed to overexpressed HERV Env proteins have been suggested to contribute to the development of cancer [34].

The placenta, most specifically the syncytiotrophoblast layer, produces an important amount of extracellular vesicles, which, through their capacity to mediate intercellular communication and to carry miRNA, proteins and diverse molecules, are thought to act in different physiological and pathological processes [23]. Several reports have suggested that different components of placental extracellular microvesicles promote foeto-maternal tolerance [24, 35, 36]. In this study, we have confirmed that exosomes inhibited the production of Th1 cytokines (TNF- $\alpha$ , IFN- $\gamma$ , IL-2) and that Syn-2-negative exosomes lost this Th1 cytokine-suppressing capacity. These results are also very similar to those obtained for the Syncytin-2 peptide and suggest a similar mode of action. They also provide a mechanism by which Syncytin-2 through its incorporation into extracellular microvesicles could modulate the immune response at a distance and in different regions surrounding the placenta. Previous studies on the immunosuppressive function of retroviral envelope proteins have relied on an *in vivo* tumour rejection assay in which tumour cells expressing different Env proteins acquired the capacity to proliferate in resistant mice [11-13, 37, 38]. This immunosuppressive function of HERV Env proteins in this context might depend on the action of Env-containing exosomes released from tumour cells. In this view, it is also tempting to speculate that, given that exosomes share certain

390 features with retroviruses, such as size and biogenesis, they could also act similarly to  
retroviruses by generating an immunosuppressive state via surface-associated proteins, including  
Syncytin-2. In fact, it is highly likely that the distribution of Syncytin-2 on the surface of  
exosome mimics typical trimeric complexes formed by Env proteins at the surface of exogenous  
retroviruses, and suggested for Syncytin-1 and Syncytin-2 [39-41]. Such a Syncytin-2 trimer  
395 should also adopt the typical six-helix bundle preceding the fusion step. These different  
structures, i.e. trimers and six-helix bundles, might optimize the inhibitory potential of Syncytin-  
2 and would thereby be displayed appropriately on the surface of exosomes. Such a mechanism  
of action might in fact explain why only dimerized forms of ISD peptides are inhibitory, as  
opposed to their monomeric equivalent.

400 A recent study by Eksmond *et al.* (2017) had suggested that mutations known to affect  
ISD function was also severely impacting Friend-Murine Leukemia Virus replication and that the  
role played by the ISD domain during infection might not be related to an immunosuppressive  
function [42]. However, in our system, we have used a Syncytin-2 ISD peptide and exosome-  
associated Syncytin-2 to address the immunosuppressive capacity of this HERV Env protein,  
405 which therefore allowed us to avoid being constrained by a possible effect of the ISD region on  
viral replication. The current mechanisms of action of the ISD domain should involve cell surface  
factor, which are likely different from the identified receptors needed for HERV envelope-  
mediated fusion. More mechanistic studies are needed to better understand how this Envelope  
region can modulated immune response and which immune cell populations are targeted by the  
410 HERV immunosuppressive domain.

In conclusion, our data suggest that Syncytin-2 via its ISD domain, and particularly  
through its association to exosomes, contribute in the modulation of the microenvironment  
surrounding the foetus by acting on the Th1 response. Since placenta-derived-extracellular

vesicles of pre-eclamptic patients are altered in their content and number in comparison to those  
415 of normal pregnant women, it has been suggested that they might contribute to the development  
of the disorder [24]. Reduced Syncytin-2 levels in exosomes from pre-eclamptic patients [30]  
might therefore play a determinant role in altering immunomodulatory capacity of these vesicles  
and could be important in the appearance of the symptoms related to pre-eclampsia. More studies  
will be needed to address this possibility.

420

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**Figure legends**

560 **Figure 1: The dimerized Syncytin-2 ISD peptide induces ERK1/2 phosphorylation in Jurkat cells. A-B.** Jurkat cells were either left untreated or treated with monomeric Sync-2-ISD, dimeric (Sync-2-ISD)<sub>2</sub>, monomeric Reverse Sync-2-ISD or dimeric Reverse (Sync-2-ISD)<sub>2</sub> (60 μM) from 2 to 60 min. Cell lysates were prepared and the phosphorylation of MAP kinase was determined by Western blot using antibodies against phosphorylated ERK1 and ERK2 (pERK 42/44) (upper panels) or total ERK1/2 (ERK 42/44) (lower panels) (**A**). The phosphorylation of MAP kinase in this time course experiment was quantified by scanning densitometry for dimeric (Sync-2-ISD)<sub>2</sub>-treated cells (**B**). Data represents fold increase of pERK1/2 (normalized with signals for total ERK1/2) compared to non-stimulated control (set at a value of 1). **C.** Jurkat cells were either left untreated or treated with different concentrations (5, 10, 30 and 60 μM) of dimerized ISD (Sync-2-ISD)<sub>2</sub> for 1 h. Western blot analysis was performed as indicated in **A-B**. Results are representative of three independent experiments.

**Figure 2: (Sync-2-ISD)<sub>2</sub> inhibits TNF-α production in Jurkat T cells.** Jurkat cells were incubated with 60 μM of monomeric or dimerized Sync-2-ISD vs. control Rev Sync-2-ISD peptides. After 2 h, cells were stimulated with PMA (20ng/ml)/Ionomycin (1 μM) and TNF-α was measured in the supernatant after 24 h (**A**) and 48hrs (**B**) by ELISA. Means ± SE were calculated from triplicates and results are representative of three independent experiments. \* p < 0.05 and \*\*\* p < 0.001.

580 **Figure 3: (Sync-2-ISD)<sub>2</sub> inhibits Th1 cytokine production in PMA/Ionomycin-activated**  
**PBMC.** PBMCs were left untreated (A) or incubated with 60  $\mu$ M Rev (Sync-2-ISD)<sub>2</sub> (B) or  
(Sync-2-ISD)<sub>2</sub> (C). After 2 h, cells were stimulated with PMA (20ng/ml)/Ionomycin (1  $\mu$ M) and  
cytokines were measured in supernatants after 24 h through the LEGENDplex™ Human  
Th1/Th2 Panel kit. Data are depicted as the distribution of signals corresponding to each detected  
585 bead and are presented in terms of intensity for each measured cytokine.

**Figure 4: (Sync-2-ISD)<sub>2</sub> inhibits Th1 cytokine production in PMA/Ionomycin-activated**  
**PBMC.** PBMCs were left untreated (A) or incubated with 60  $\mu$ M Rev (Sync-2-ISD)<sub>2</sub> (B) or  
(Sync-2-ISD)<sub>2</sub> (C). After 2 h, cells were stimulated with PMA (20ng/ml)/Ionomycin (1  $\mu$ M) and  
590 cytokines were measured in supernatants after 48 h through the LEGENDplex™ Human  
Th1/Th2 Panel kit.

**Figure 5: Quantification of Th1 cytokine levels in PMA/Ionomycin-stimulated PBMCs**  
**following treatment with (Sync-2-ISD)<sub>2</sub>.** PBMCs were incubated with (Sync-2-ISD)<sub>2</sub> vs. control  
595 Rev (Sync-2-ISD)<sub>2</sub> (60  $\mu$ M). After 2 h, cells were stimulated with PMA (20ng/ml)/Ionomycin (1  
 $\mu$ M) and cytokines were measured. Quantification of cytokine levels from Figure 3 and 4 are  
presented in pg/ml and are based on standard curves generated for each cytokine. Means  $\pm$  SE  
were calculated for each measured cytokine and results are representative of two independent  
experiments. \*\*\*  $p < 0.001$ .

600 **Figure 6: (Sync-2-ISD)<sub>2</sub> inhibits Th1 cytokine production in anti-CD3/anti-CD28-activated**  
**PBMCs.** PBMCs were incubated with 60  $\mu$ M (Sync-2-ISD)<sub>2</sub> or Rev (Sync-2-ISD)<sub>2</sub>. After 2 h,

cells were stimulated anti-CD3 (0.5 µg/ml) and anti-CD28 (1.5 µg/ml) antibodies and cytokines were measured in supernatants after 24 (A) and 48 h. (B) through the LEGENDplex™ Human Th1/Th2 Panel kit.

**Figure 7: Quantification of cytokine levels in anti-CD3/anti-CD28-stimulated PBMCs following treatment with (Sync-2-ISD)<sub>2</sub>.** PBMCs were incubated with (Sync-2-ISD)<sub>2</sub> vs. Rev (Sync-2-ISD)<sub>2</sub> (60 µM). After 2 h, cells were stimulated with anti-CD3 (0.5 µg/ml) and anti-CD28 (1.5 µg/ml) antibodies and cytokines were measured in supernatants after 24 and 48 h. Quantification of cytokine levels from Figure 6 are presented in pg/ml. Means ± SE were calculated for each measured cytokine and results are representative of two independent experiments. \*\* p<0.01 and \*\*\* p < 0.001.

**Figure 8: Inhibition of TNF-α production in Jurkat T cells by trophoblast-derived exosomes.** Jurkat cells were incubated with different concentrations of exosomes (0, 5 or 10 µg/ml) in triplicates. After 24 h, cells were stimulated with PMA (20ng/ml)/Ionomycin (1 µM) and analysed for TNF-α expression by RT-PCR (6 h post-treatment, A) and TNF-α secretion by ELISA in harvested supernatants (18 h post-treatment, B). For RT-PCR analysis, GAPDH signals were amplified in parallel and signals were scanned for densitometric analyses. Results -are presented following normalisation with GAPDH. Results are representative of three different experiments. C-D. Jurkat cells transfected with pNF-κB-Luc (C) or pNFAT-Luc (C-D) were stimulated with PMA (20ng/ml)/Ionomycin (1 µM) for 8 h and incubated in the presence of fixed (50 µg/ml) (C) or varying concentrations (D) of isolated exosomes. Luciferase activity was

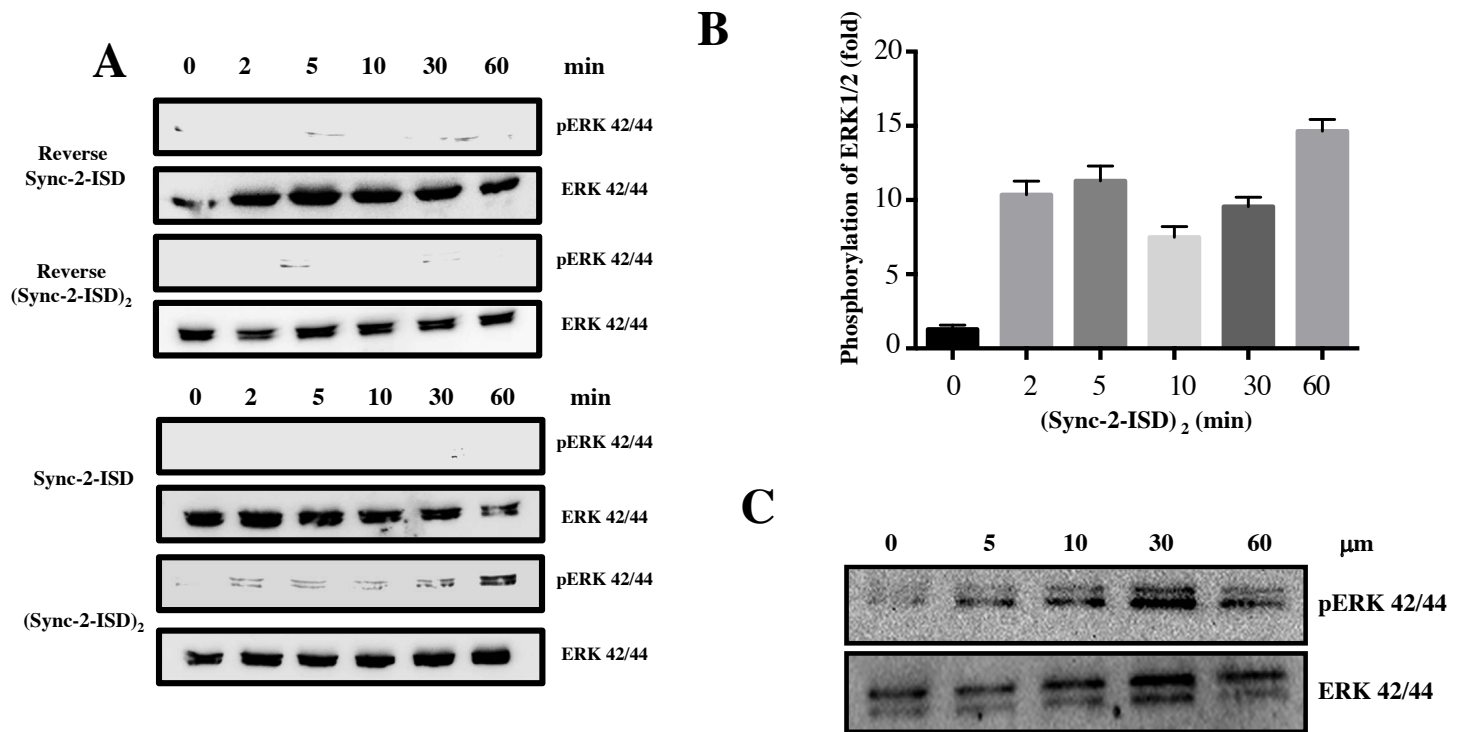
625 measured in three independent transfection samples and fold induction are presented as a mean  $\pm$   
SE. \*  $p < 0.05$ , \*\*  $p < 0.01$ , and \*\*\*\*  $p < 0.0001$ .

**Figure 9: Villous cytotrophoblast-derived exosomes inhibit Th1 cytokine production in anti-CD3/anti-CD28-activated PBMCs.** PBMCs were incubated in the presence of 50  $\mu\text{g/ml}$  of  
630 exosomes isolated from freshly purified day 1 villous cytotrophoblasts (CT-DE). After 24 h, cells  
were stimulated in the presence of anti-CD3 (0.5  $\mu\text{g/ml}$ ) and anti-CD28 (1.5  $\mu\text{g/ml}$ ) antibodies  
and cytokines were measured in supernatant after 24 h through the LEGENDplex™ Human  
Th1/Th2 Panel kit.

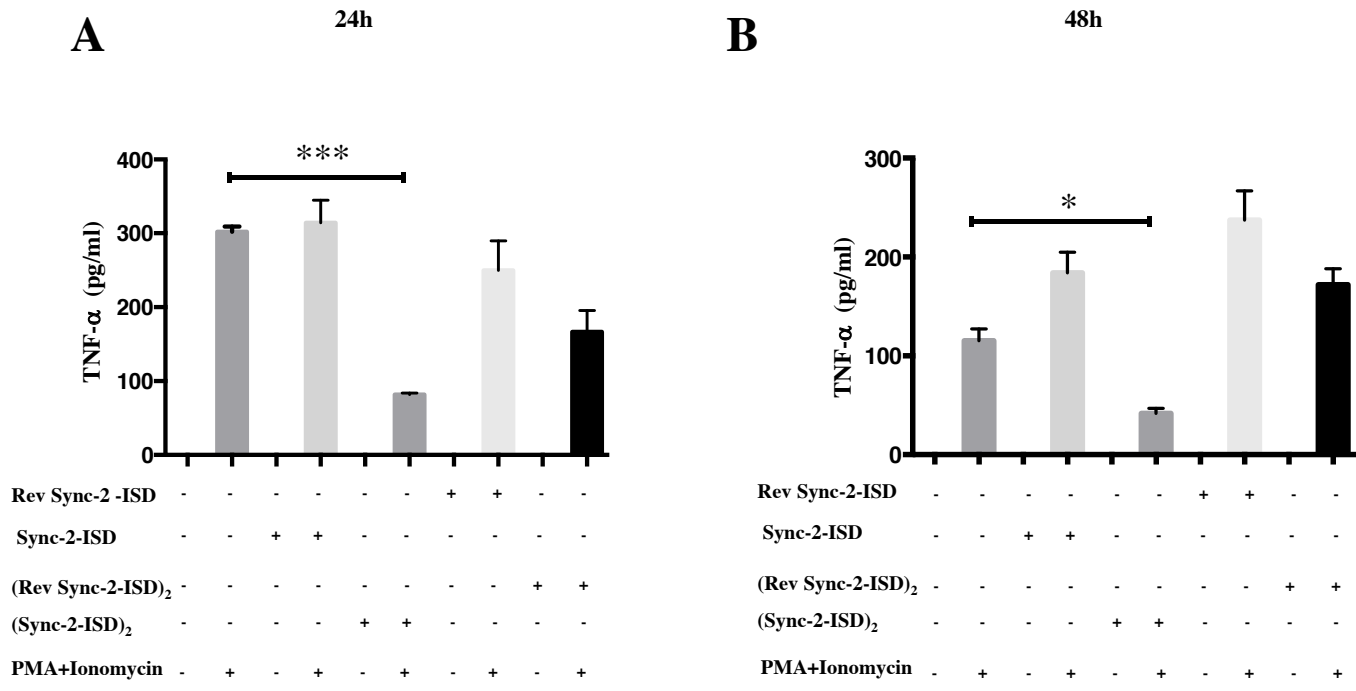
635 **Figure 10: Reduced levels of Syncytin-2 on the surface of cytotrophoblast-derived exosomes  
decrease their inhibitory capacity over Th1 cytokine production.** PBMCs were incubated in  
the presence of 50  $\mu\text{g/ml}$  of exosomes (CT-DE) isolated from untransfected primary  
cytotrophoblast cells or from primary cytotrophoblast cells transfected with Syncytin-2-specific  
siRNA (siSync2) or scrambled control siRNA (siScr.) (one or two-day cultures: D1- D2). After  
640 24 h, cells were stimulated with anti-CD3 (0.5  $\mu\text{g/ml}$ ) and anti-CD28 (1.5  $\mu\text{g/ml}$ ) and cytokines  
were measured in harvested supernatants after 24 h through the LEGENDplex™ Human  
Th1/Th2 Panel kit (A). Western blot analyses were performed on extracts from cytotrophoblast-  
derived exosomes from each transfection condition using anti-Syncytin-2 and anti-TSG101  
antibodies at 24 and 48 h after transfection (B). These results are representative of two  
645 independent experiments.

**Figure 11: Quantification of cytokine levels in anti-CD3/anti-CD28-stimulated PBMCs following exposure to cytotrophoblast-derived Syncytin-2- vs. Syncytin-2+ exosomes.** PBMCs

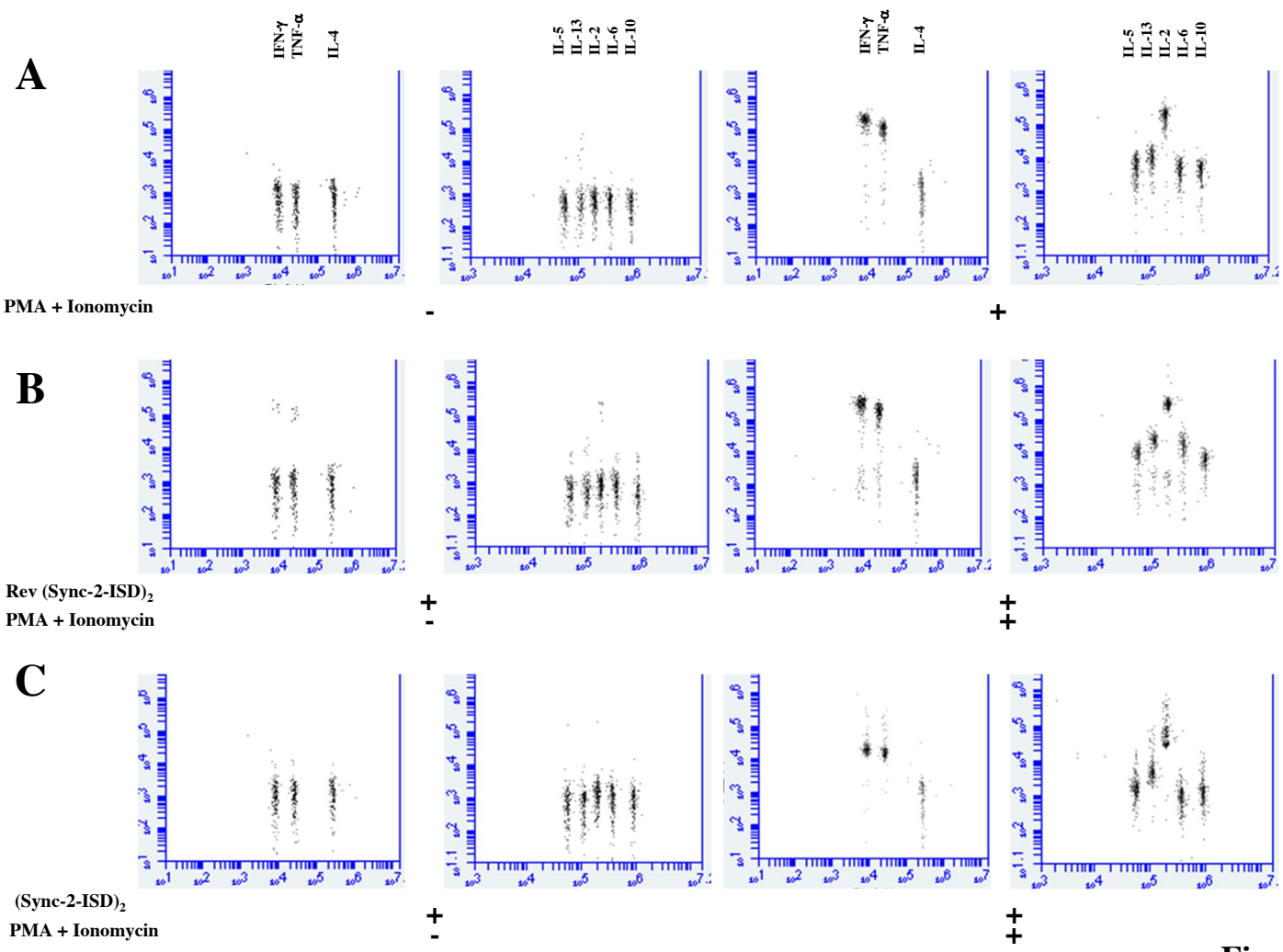
650 were incubated in the presence of 50 µg/ml of exosomes (CT-DE) isolated from untransfected primary cytotrophoblast cells or from primary cytotrophoblast cells transfected with Syncytin-2-specific siRNA (siSync2) or scrambled control siRNA (siScr.) (one or two-day cultures: D1- D2). After 24 h, cells were stimulated anti-CD3 (0.5 µg/ml) and anti-CD28 (1.5 µg/ml) and cytokines were measured in supernatants after 24 and 48 h through the LEGENDplex™ Human Th1/Th2  
655 Panel kit. Quantification of cytokine levels from Figure 10 are presented in pg/ml. Means ± SE were calculated for each measured cytokine and results are representative of two independent experiments. \*p<0.05, \*\* p < 0.01 and \*\*\* p < 0.001.



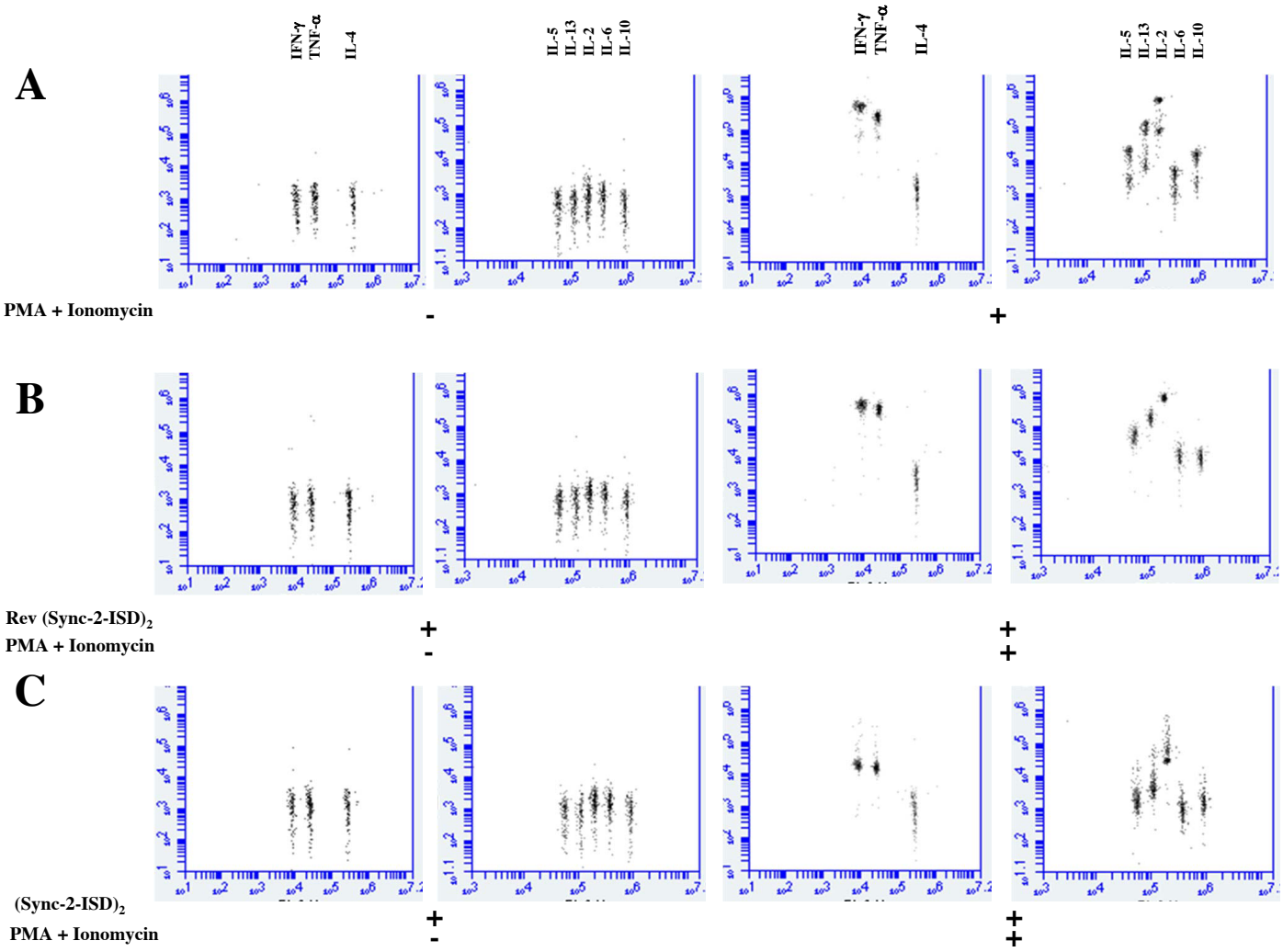
**Figure 1**



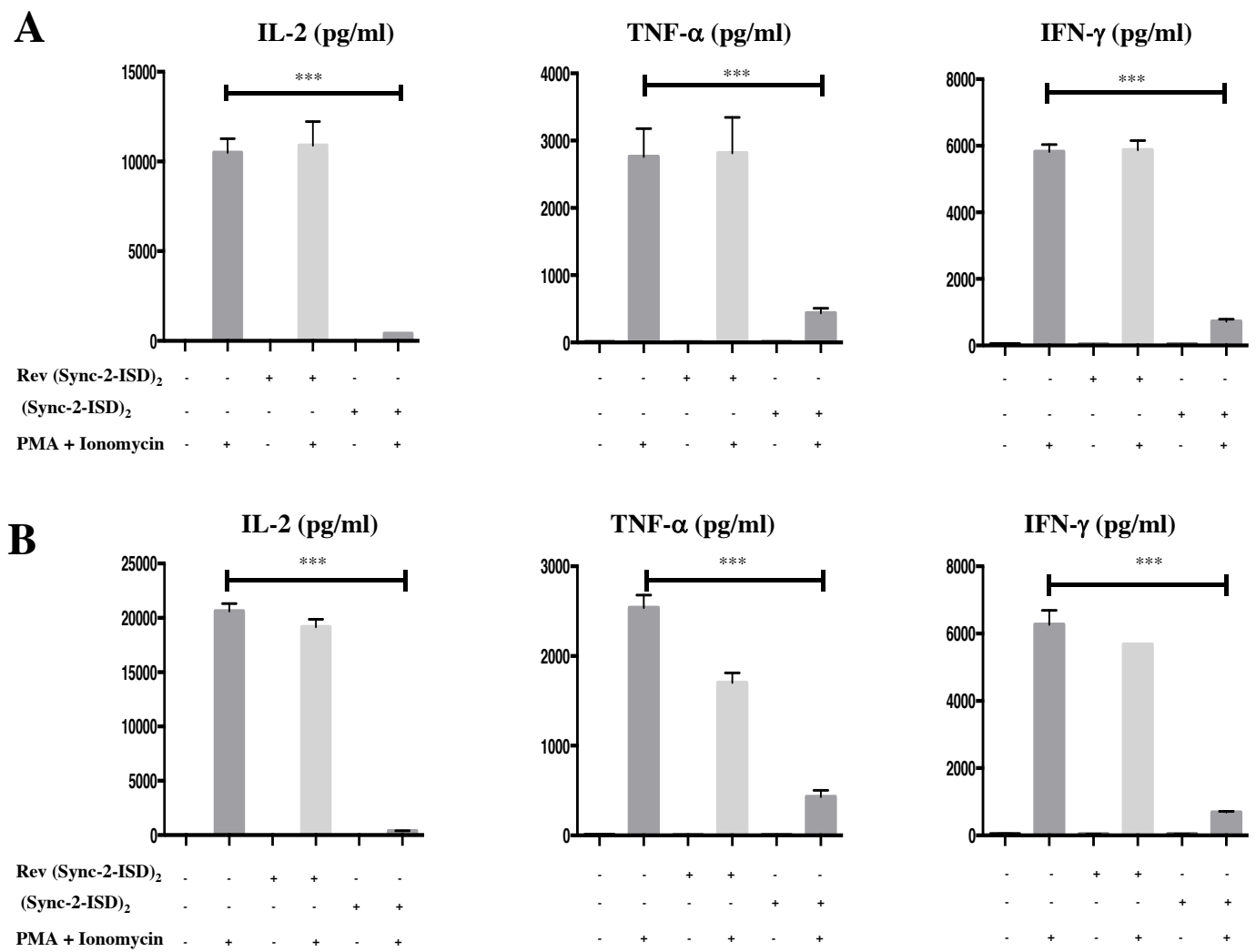
**Figure 2**



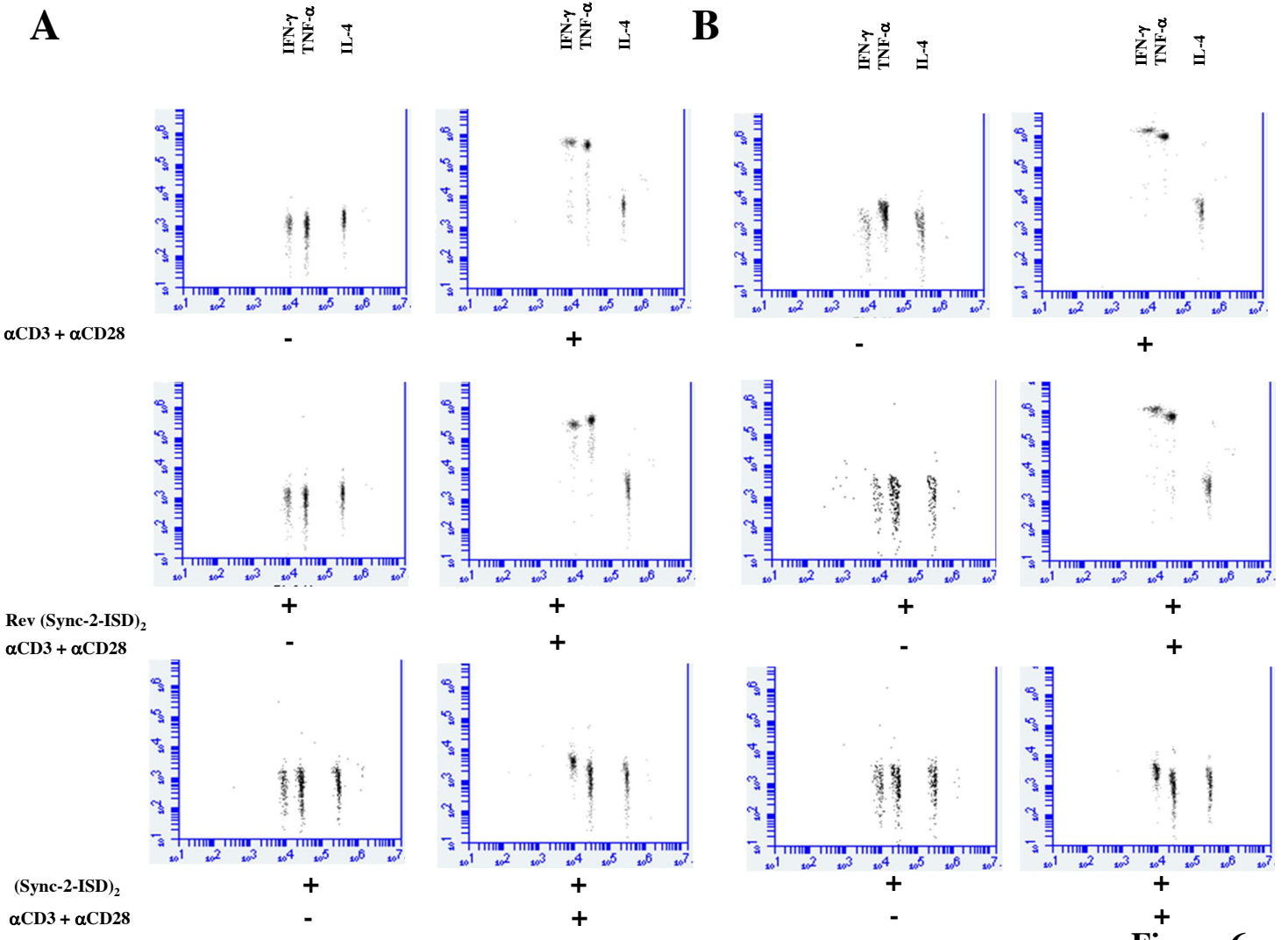
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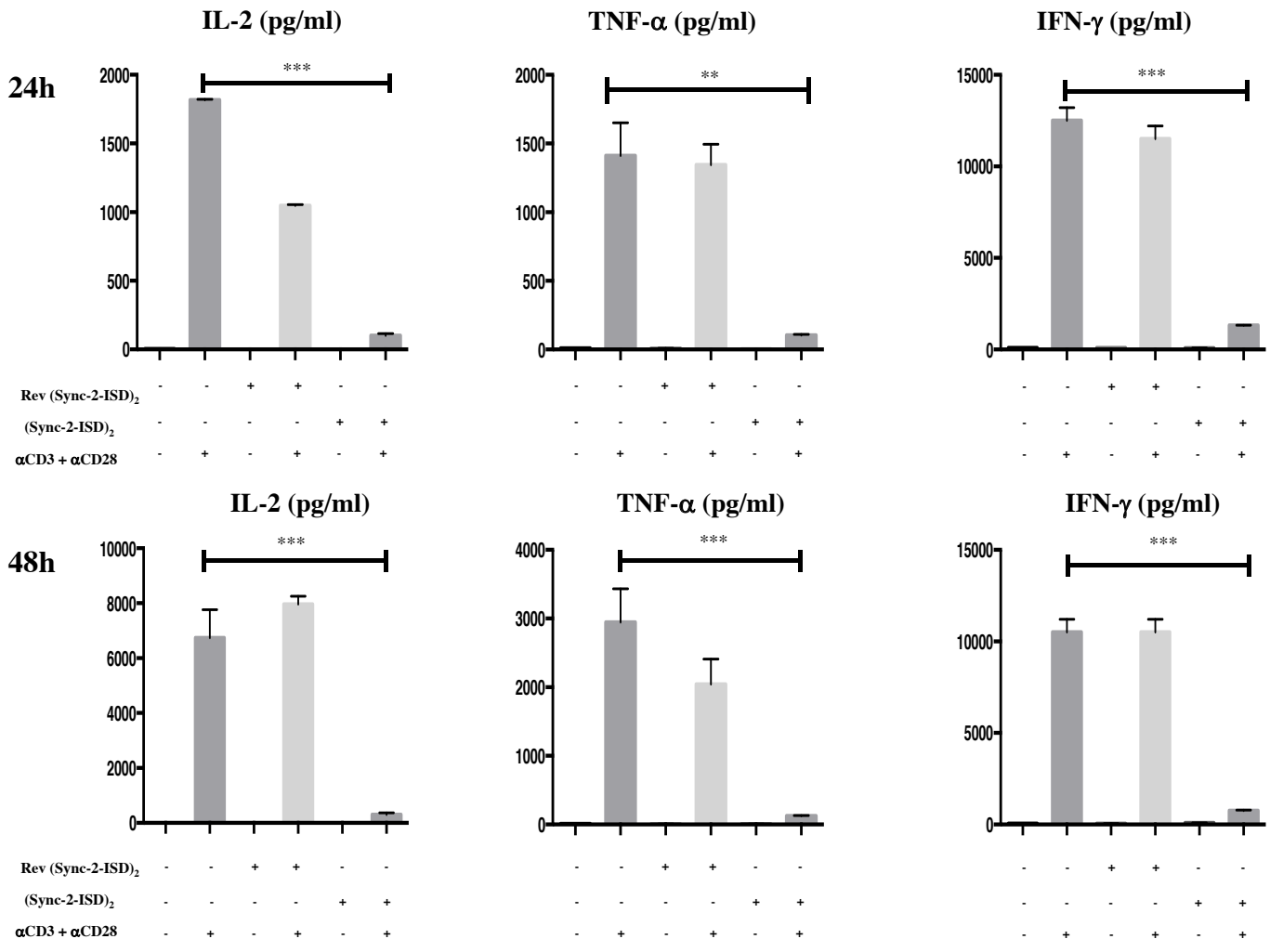
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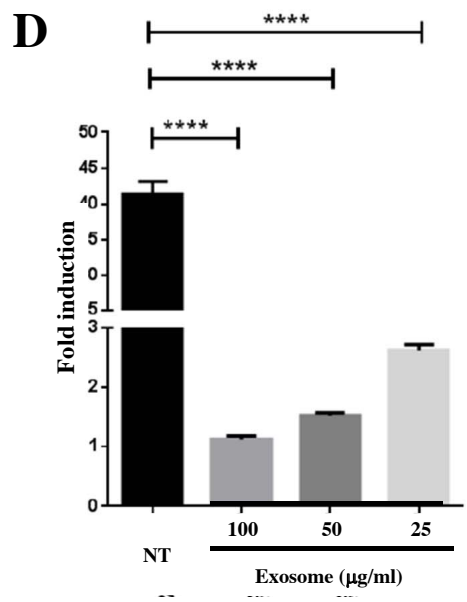
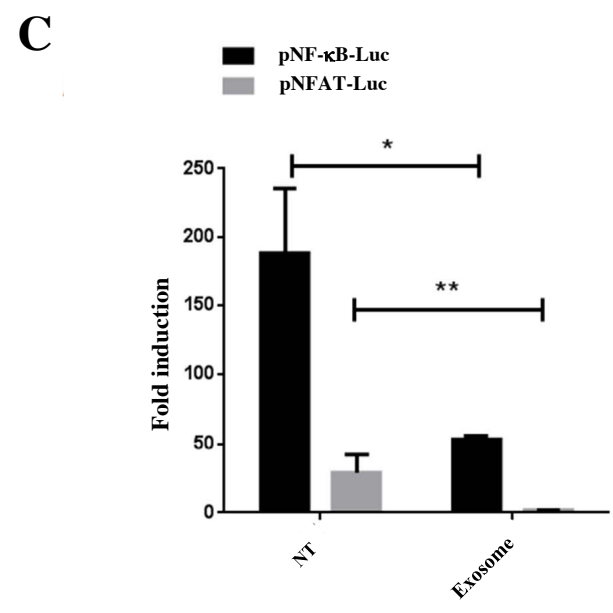
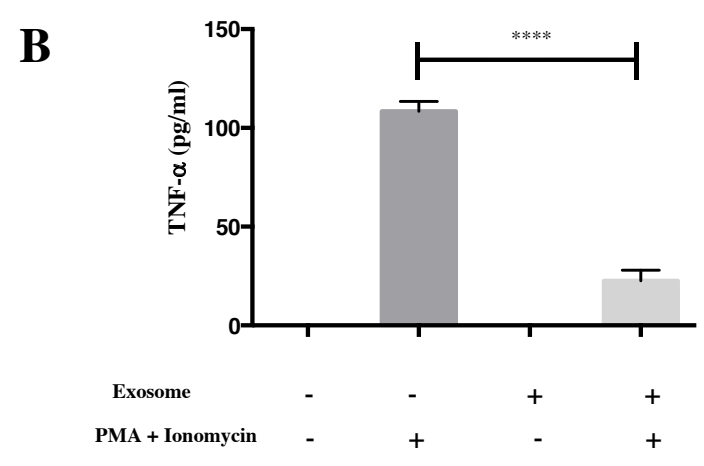
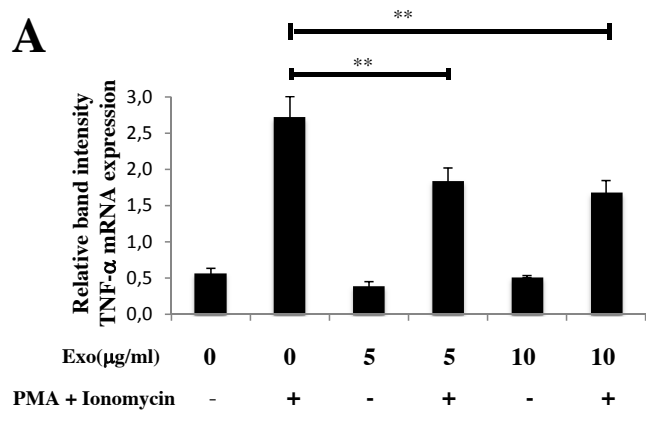
**Figure 5**



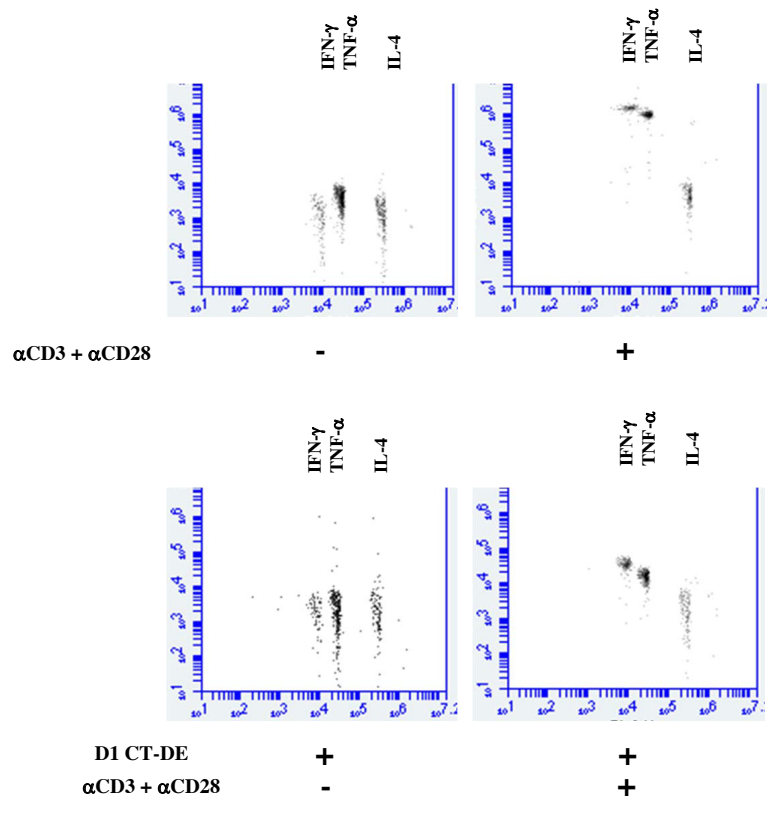
**Figure 6**



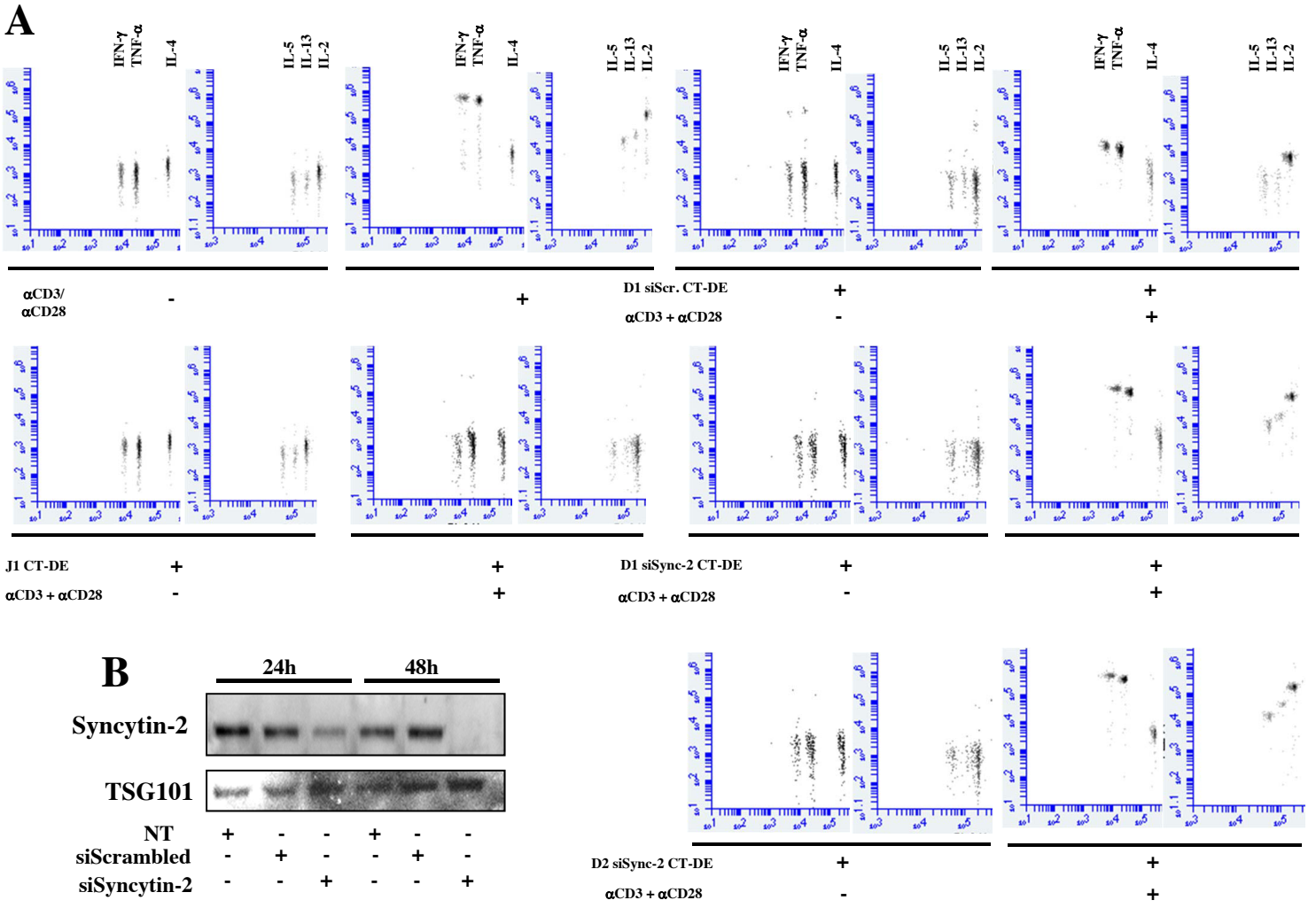
**Figure 7**



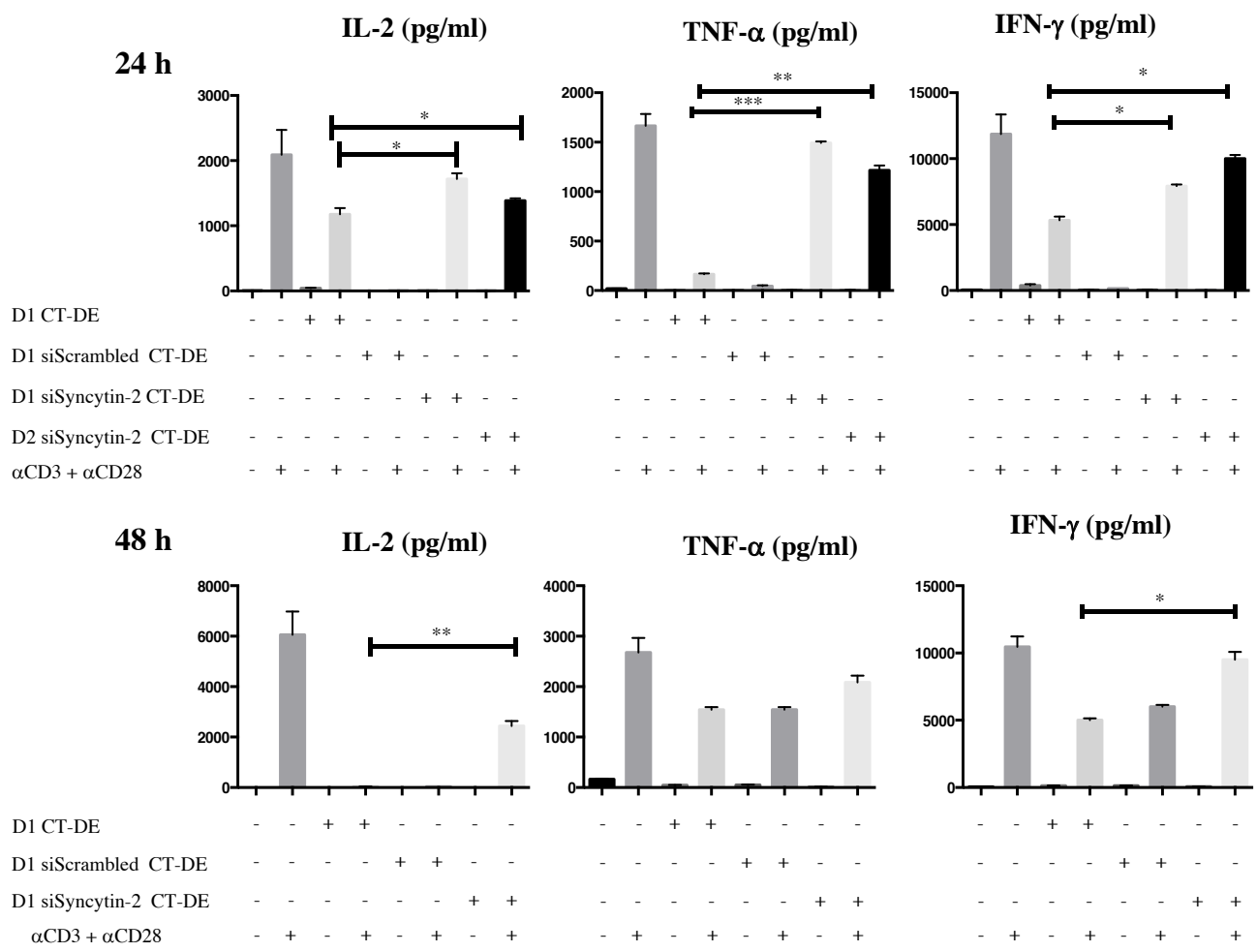
**Figure 8**



**Figure 9**



**Figure 10**



**Figure 11**