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## Human tolerogenic Dendritic cells produce IL-35 in the absence of other IL-12 family members.

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## Abstract

IL-35 is a novel cytokine of the IL-12 family, existing as a heterodimer of IL-12p35 and EBV-induced gene 3 (Ebi3). IL-35 has anti-inflammatory properties and has been described to be produced by regulatory T cells in humans and mice, where it is required for optimal suppression. Distinct from other IL-12 family members, expression of IL-35 has not been clearly described in antigen presenting cells. In view of its immune-regulatory properties, we investigated the expression, regulation and function of IL-12p35 and Ebi3 in human monocyte derived dendritic cells (DCs) and tolerogenic DCs (tolDCs), generated in the presence of dexamethasone. These toIDCs did not produce detectable levels of bioactive IL-12p70 or the homodimer IL-12p40. In line with this, we demonstrate by Q-PCR that tolDCs completely lack transcriptional expression of IL-12p40. However, toIDCs maintain mRNA expression of IL-12p35 and Ebi3. Using intracellular flow cytometry and western blot we show that tolDCs are able to produce Ebi3 and IL-12p35, and both can be further enhanced upon stimulation with IFNy, LPS or CD40 ligation. Supernatant of tolDCs has the capacity to suppress T cell activation. Using IL-12A silencing we demonstrate that IL-12p35 is required for tolDCs to reach their full suppressive potential. Taken together, our results indicate that toIDCs produce IL-35, providing an additional novel mechanism by which these cells elicit their tolerogenic potential.

## Introduction

Dendritic cells (DCs) represent a heterogeneous family of professional APCs that have many functions in both the initiation and maintenance of immunity and immunological tolerance<sup>1,2</sup>. Several factors, including IL-10, TGF-β, Vitamin D<sub>3</sub> and corticosteroids have been shown to modulate DC function in vitro and in vivo, and lead to the generation of tolerogenic DCs (tolDCs)<sup>3-6</sup>. These tolDCs represent an altered status in DC differentiation and are currently considered in the design of therapeutic strategies to treat patients suffering from autoimmune disease<sup>7,8</sup> and those undergoing transplantation<sup>9,10</sup>. Impaired expression of IL-12 in tolDCs is thought to be a crucial requirement for these cells to elicit their tolerogenic potential, and indeed many tolDC populations have been shown to lack IL-12<sup>3,5</sup>. However the IL-12 family is ever expanding and little is known about the regulation and expression of other IL-12 family members in toIDCs. The IL-12 family is composed of four heterodimeric complexes each consisting of an alpha chain (IL-12p35, IL-23p19 and IL-27p28) and a beta chain (Ebi3 and IL-12p40)<sup>11</sup>. The IL-12p40 subunit can pair with either IL-12p35 or IL-23p19 to yield IL-12 and IL-23 respectively<sup>12,13</sup>. The second beta chain subunit Ebi3, can pair with 2 other alpha chain subunits, IL-27p28 and IL-12p35, to yield IL-27 and IL-35 respectively. Although there is a large degree of chain sharing within

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this family, each cytokine possesses a unique and distinct function. IL-12 and IL-23 have primarily pro-inflammatory properties and play an important role in the development of Th1 and Th17 T helper cell subsets respectively. IL-27 possess dual roles playing a protective role in tumour immunity<sup>14,15</sup>, but as an immunoregulatory cytokine can induce Tr1 cells<sup>16,17</sup> and lead to the generation of a tolerogenic DC expressing elevated levels of CD39<sup>18</sup>. The most recently characterised member of this family is IL-35, a potent anti-inflammatory cytokine, which has been shown to be required for the optimal suppressive activity of regulatory T cell populations in both mice and man<sup>19-22</sup>. DCs are known producers of three family members, IL-12, IL-23 and IL-27. In toIDCs little is known about the expression of these family members aside from the characteristic loss of IL-12 production and limited transcriptional data on IL-23 and IL-27<sup>23</sup>. For IL-35, production by myeloid cells in human or mouse has not yet been demonstrated. In this study we addressed the expression profile and regulation of all IL-12 family members in human DCs and tolDCs with a particular emphasis on the subunits which combined make up IL-35. We show that toIDCs did not produce detectable levels of bioactive IL-12p70, the homodimer IL-12p40 or express IL-12p40 transcripts. However toIDCs maintain mRNA and protein expression of IL-12p35 and Ebi3. We show that supernatants of toIDCs are sufficient to suppress T cell proliferation, and that supernatants of cells silenced for IL-12p35 are hampered in their suppressive potential, thereby implying a role for IL-35 in toIDC function.

### **Materials and Methods**

#### Generation of human monocyte derived DCs

Human monocytes were isolated from buffy coats obtained from healthy donors using Ficoll density gradient centrifugation followed by positive selection using anti-CD14 MACS microbeads (Miltenyi Biotech GmBH, Bergisch Gladbach, Germany). DCs were generated and cultured in RPMI supplemented with 10% heat inactivated FCS, 90 U/ml penicillin and 90 U/ml streptomycin (Gibco/ Life Technologies, Bleiswijk, The Netherlands), supplemented with 5 ng/ml GM-CSF and 10 ng/ml IL-4 (Invitrogen/ Life Technologies, Bleiswijk, The Netherlands) as described before<sup>3</sup>. ToIDCs were generated by the addition of Dexamethasone (10<sup>-6</sup>M Dex) (Pharmacy, LUMC, Leiden, The Netherlands), only at the start of culture (day 0). Cultures were refreshed with medium containing cytokines on day 3. For differentiation into mature DCs, cells were harvested, washed and stimulated on day 6 with 200 ng/mL LPS (E.Coli 0127:B8 Sigma-Aldrich, Zwijndrecht), 20µg/ml Poly I:C, 10 µg/ml PGN (InvivoGen) or 100 ng/ml IFN $\gamma$  (Peprotech, Germany). CD40L-activation was performed with CD40L transfected L cells (L-CD40L) in a DC: L cell ratio of 5:1. Non-transfected L cells served as control cells (L-Orient).

#### DC-T cell co culture

Total CD4<sup>+</sup> lymphocytes were negatively selected from PBMCs using CD4<sup>+</sup> T cell isolation kit II (Miltenyi Biotech GmBH, Bergisch Gladbach, Germany), and subsequently labelled with carboxyfluorescein diacetate succinamidyl ester (CFSE) (Molecular Probes, Europe BV Leiden, The Netherlands). In short, T cells were

suspended in PBS and incubated for 15 minutes at 37°C with 5  $\mu$ M CFSE. The reaction was quenched by washing the cells in medium containing 10% FCS before resuspending at 1×10<sup>6</sup>/ml in total RPMI culture medium. Labelled CD4<sup>+</sup> T cells were co-cultured with allogeneic DCs at a 10:1 T cell/DC ratio for 5 days. The T cells were then harvested and proliferation was assessed by CFSE dilution.

#### Surface and Intracellular flow cytometry

For cell surface flow cytometric analysis, cells were stimulated overnight, harvested, washed, and stained for 30 minutes at 4°C in FACS buffer (PBS, 0.5% heat inactivated NHS, 1% BSA, 0.02% NaN<sub>3</sub>) with anti-CD86, anti-CD14 MΦ P9 (BD biosciences, San Diego, CA, USA), anti-B7H1, anti-B7DC, anti-ILT3 (Biolegend), anti-HLADR (B8.11.2, ATCC), anti-Mannose Receptor (clone D547, msIgG1 in house) or anti-DC-SIGN (R&D Systems, Wiesbaden, Germany). Nonconjugated antibodies were detected with PE-conjugated goat-anti-mouse Ig (Dako, Glostrup, Denmark). Isotype matched control antibodies were used to determine the level of background staining. For intracellular staining cells were stimulated overnight, followed by incubation in the presence of Brefeldin-A (Sigma-Aldrich, Zwijndrecht) for a further 5 hours. The cells were then harvested, washed and fixed in PBS containing 4% formaldehyde and 1% heat inactivated FCS, washed with PBS containing 10% heat inactivated FCS and permeabilised with perm buffer (PBS, 0.5% saponin, 1% heat inactivated FCS) for 10 minutes. Cells were stained with anti-Ebi3-APC or anti-IL-12p35-APC for 30 minutes at 4°C in perm buffer. APC-conjugated isotype matched control antibodies were used to determine the level of background staining. The fluorescence was measured on a FACS Calibur flow cytometer, and data were analysed with Cell Quest Software (BD Biosciences, San Diego, CA, USA) and FlowJo Software (Tree star, USA).

#### mRNA isolation, cDNA synthesis, and real-time PCR

Day 6 DCs were plated in a 12-well plate at a density of 1x10<sup>6</sup>/ml and harvested after indicated time points. mRNA was isolated using an RNeasy kit (Qiagen, Hilden, Germany) following the manufacturer's instructions. Digestion of genomic DNA was performed by using the on-column RNase-free DNase set. Reverse transcriptase system kit (Promega, Leiden, The Netherlands) was used to synthesise cDNA conforming to the manufacture's instruction and was stored at -20°C for further analysis. Specific primers (Table I) for human *IL12A, IL12B, IL23A, IL27A, IL27B, IL10* and *GAPDH* were designed using the computer software Oligo explorer and synthesised at Biolegio. Quantitative PCR was performed using SYBR Green qPCR master mix (Bio-Rad, Veenendaal, The Netherlands). *GAPDH* was used as the endogenous reference gene. Data analysis was performed using Bio-Rad CFX Manager Software (Bio-Rad). For each sample, the relative abundance of target mRNA was calculated from the obtained Ct values for the target gene and expressed relative to the endogenous reference gene *GAPDH*.

Gene	NCBI ID	Protein ID	Forward Sequence (5'-3')	Reverse Sequence (5`-3`)
IL12B	3593	IL-12p40	CAGCAGCTTCTTCATCAGGG	GAGTACTCCAGGTGTCAGG
IL12A	3592	IL-12p35	CCAGAGTCCCGGGAAAGTC	ACCAGGGTAGCCACAAGG
IL27B	10148	Ebi3	TGGATCCGTTACAAGCGTC	AGTTCCCCGTAGTCTGTG
IL27A	246778	IL-27p28	GAGGGAGTTCACAGTCAGC	GCAGGAGGTACAGGTTCAC
IL23A	51561	IL-23p19	CCAAGGACTCAGGGACAAC	CTGAGGCTTGGAATCTGCTG
Ш10	3586	IL-10	GCGCTGTCATCGATTTCTTCC	GTAGATGCCTTTCTCTTGGAGCTTA
GAPDH	2597	GAPDH	TTCCAGGAGCGAGATCCCT	CACCCATGACGAACATGGG

#### Table I: Real Time PCR Oligonucleotide sequences

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#### Cytokine production

DCs were plated at  $1x10^{6}$ /ml and stimulated accordingly. Cell culture supernatants were harvested after 48 hours and frozen at  $-80^{\circ}$ C until analysis. Subsequently they were tested for the presence of IL-12p70, IL-12p40, (Biolegend) IL-10 (Sanquin, Amsterdam, The Netherlands) and IL-27 (eBioscience) by ELISA according to manufacturer's instructions. IFN $\gamma$  levels were determined in the cellular supernatants of T cells from indicated time points (eBioscience).

#### Western Blotting

DCs were plated at  $1 \times 10^6$  /ml, stimulated for 16 hours, followed by the addition of Brefeldin A for a further 5 hours. Cells were harvested, washed in ice-cold phosphate-buffered saline and lysed in medium stringency lysis buffer for 30 minutes. The lysates were centrifuged at 13,000 rpm, the supernatants harvested and protein concentration was determined by a Pierce assay. The samples were diluted in 1x SDS loading buffer, boiled for 5 minutes in reducing conditions, followed by separation on 10% SDS-PAGE gel. Gels were transferred to nitrocellulose membranes, blocked with TBS plus 0.1% Tween-20 (TBS-T) and 3% BSA or milk and probed with indicated primary antibodies; purified anti-IL-12p40, anti-IL-12p35 (Santa Cruz Heidelberg, Germany) or purified anti-Ebi3 (Abnova, Huissen The Netherlands). After incubation with HRP-conjugated secondary antibody, proteins were detected with super signal ECL system. Blots were stripped and  $\beta$ -actin was used as a loading control using HRP conjugated anti- $\beta$  actin (Abcam, Cambridge, UK).

#### Immunofluorescence microscopy

Cells were cultured on 8 well chamber slides (NUNC) in the presence of stimuli and Brefeldin-A. The cells were subsequently washed and fixed in PBS containing 4% formaldehyde, 1% heat inactivated FCS followed by washing with PBS containing 10% heat inactivated FCS and permeabilised with perm buffer (PBS, 0.5% saponin, 1% heat inactivated FCS) for 10 minutes. The DCs were then stained using IL-12p35-APC or purified anti-Ebi3 (Abnova), followed by goat anti-mouse-Alexa488 (Molecular Probes). Hoechst was used for nuclear staining.

#### **RNA** interference

DCs were transfected with 50 nM siRNA through the use of the transfection reagent Lipofectamine 2000 (Life Technologies) and were used for experiments 24 hours after transfection. The following SMARTpool siRNAs were used (Dharmacon, Fisher Scientific, Netherlands): IL-12A and nontargeting siRNA as a control. Silencing of expression was verified by real-time PCR in toIDC and IL-12p40/p70 ELISAs in DC.

#### T cell proliferation assays

Naïve CD4<sup>+</sup> T cells were isolated by negative selection using naïve CD4<sup>+</sup> T cell isolation kit II (Miltenyi Biotech), from cord blood derived PBMCs. Cells were stimulated as described previously<sup>20</sup>. Briefly cells were plated at a density of 25,000/well in 96 well round bottom plates in the presence of  $\alpha$ CD3/28 coated beads. Supernatants from stimulated toIDCs were harvested and incubated with the T cells at a dilution of 1:2. T cell proliferation was assessed by the addition of <sup>3</sup>[H]-thymidine (0,5 µCi/well) for the last 8 hours. Percentage suppression was calculated by 1-(cpm T cells + toIDC sup / cpm T cells alone) times 100.

#### Statistical analysis

Statistical analysis was performed with Graph Pad Prism (Graph Pad Software, San Diego, CA) using a one-tailed t-test. P-values  $\leq 0.05$  were considered statistically significant.

## Results

## DCs generated in the presence of dexamethasone display typical features of tolerogenic DCs with low expression of CD86 but high B7H1 and B7DC.

We have previously demonstrated that Dex treated DCs display reduced levels of co-stimulatory molecules<sup>3</sup>. To further investigate the phenotypical characteristics of DCs and tolDCs, freshly isolated monocytes were cultured in the presence of GM-CSF and IL-4, with the addition of Dexamethasone to generate toIDCs. Both cell types expressed characteristic markers of myeloid DCs including DC-SIGN (CD209) and Mannose Receptor (CD206), while toIDCs preserved the expression of CD14 upon differentiation from monocyte to immature DC (data not shown). In an immature state both DCs expressed comparable levels of CD86 and immunoregulatory molecules B7H1 (CD274) and ILT3 (CD85k), while HLA-DR and B7DC (CD273) were more abundantly expressed on tolDCs (Fig.1). Stimulation using IFNy, LPS or PGN resulted in the generation of mature DCs as shown by increased expression of co-stimulatory molecules including CD86 and HLA-DR (Fig.1A,B). tolDCs were significantly impeded in their ability to upregulate CD86 or HLA-DR upon stimulation with any agents used (Fig.1A, B). In contrast toIDC upregulated B7H1 (Fig.1C) and to a lesser degree B7DC and ILT3 (Fig.1D,E). To fully ascertain the allostimulatory capabilities of tolDCs, immature and mature DC/tolDCs were co-cultured with allogenic CD4<sup>+</sup> T lymphocytes for 5 days, after which the T cells were analysed for CFSE dilution (Fig.1F). DCs were potent inducers of T cell proliferation which was further enhanced when the DCs were first matured for 24 hours with LPS or PGN. Conversely toIDCs were hampered in their ability to stimulate allogenic T cells regardless of maturation stimulus used (Fig. 1G). In line with the proliferation data DCs induced a strong IFNy production by the T cells, whereas tolDCs induced almost negligible amounts of IFNy production (Fig.1G). In summary the tolDCs used in this study display typical characteristics of myeloid tolerogenic DCs.

## ToIDC maintain expression of IL-12p35 in the absence of IL-12p40.

A key characteristic of toIDCs is their ability to preserve production of immunoregulatory cytokines while maintaining low to absent production of inflammatory cytokines. Upon activation, DCs showed strong production of IL-12p40 and IL-12p70, whereas these cytokines were completely absent in supernatants of activated toIDCs (Fig.2A,B). In contrast, toIDCs produced significantly higher levels of IL-10 upon stimulation as compared to DC (Fig.2C). To further explore the expression of IL-12 in DC and toIDC we assessed by RT-PCR the expression of *IL12A* (IL-12p35) and *IL12B* (IL-12p40), together making IL-12p70. In line with our ELISA data DCs showed a strong *IL12B* 

expression upon stimulation while toIDCs were greatly inhibited in this ability (Fig.2D). However, toIDCs maintained their expression of *IL12A* and expressed significantly higher basal levels compared to DC (Fig.2E). Upon stimulation, where *IL12B* remains absent (Fig.2D), *IL12A* expression is significantly increased in toIDCs to comparable and even higher levels than DC (Fig.2E).



**Figure 1:** DCs generated in the presence of dexamethasone display a tolerogenic phenotype and have a reduced allostimulatory capacity. Monocytes were isolated from PBMCs and cultured for 6 days in the presence of IL-4 and GM-CSF to obtain immature DC. toIDC were generated by the addition of 10<sup>-6</sup>M Dexamethasone to the moDC cultures at day 0. Cells were activated overnight with IFNγ, LPS or PGN. The expression of (A) CD86, (B) HLA-DR, (C) B7H1, (D) B7DC and (E)

ILT3 and on DCs and toIDCs was analysed using flow cytometry. Dashed line indicates the mean MFI of isotype controls used. (F) DC and toIDC were harvested on day 6 and either unstimulated or stimulated with IFN $\gamma$ , LPS or PGN for 24 hours. The DCs were then harvested and co cultured with CFSE labelled allogenic CD4<sup>+</sup>T cells at a ratio of 1:10. On day 5 the T cells were harvested and the CFSE staining was analysed using flow cytometry. Data shown is representative of 3 independent experiments. Data shown in (G) is the mean ± SD of those 3 experiments. Cytokine production was measured using ELISA. T cell culture supernatants were harvested on day 5 and analysed for (H) IFN $\gamma$  production. Data shown is the mean ± SD of 3 experiments.

### Differential kinetics of LPS induced IL-12 family members in DC and toIDC.

The IL-12 family is composed of 4 related yet distinct heterodimeric cytokines and chain sharing has become a quintessential feature of this cytokine family<sup>24,</sup> <sup>25</sup>. We demonstrated that toIDCs lack expression of *IL12B* (IL-12p40) yet maintain expression of *IL12A* (IL-12p35) (Fig.2). We investigated the expression of all family members in immature unstimulated DCs and toIDCs and found that although toIDCs expressed lower levels of *IL12B* compared to DC, they maintained expression of all other family members and *IL10* (Fig.3A).

Notably both *IL27B* (Ebi3) and *IL12A* expression was even higher in toIDC compared to DC. We further investigated the regulation of the IL-12 family in DC and toIDC in LPS triggered cells. We found that DCs displayed a pronounced upregulation of *IL12A* and *IL27A* (IL-27p28) with the expression of both subunits peaking at 6 hours (Fig.3C,F). This was distinct from *IL10* and *IL23A* (IL-23p19) which showed the greatest expression at 2 hours (Fig.3G,E), and *IL27B* which gradually increased over time and reached its highest level at 48 hours



**Figure 2: ToIDC maintain expression of IL-12p35 in the absence of IL-12p40.** Dendritic cells were harvested after 6 days of culture and stimulated with irradiated CD40L cells at a ratio of 1:5 dendritic cells, or a combination of IFN $\gamma$  + LPS. After 48 hours the supernatants were harvested and (A) IL-12p40, (B) IL-12p70 and (C) IL-10 were measured. Data shown is the mean ±SD

of 6 independent experiments. Cells were stimulated for 6 hours after which mRNA was isolated followed by cDNA synthesis. The transcript levels of (D) *IL12B* (E) *IL12A* were determined by RT-PCR. *GAPDH* mRNA expression from the same samples was used as an endogenous reference gene (relative mRNA expression). Data shown is mean  $\pm$  SD of 4 independent experiments. Untransfected L cells were used as a control for CD40L cells and yielded results in line with medium conditions (data not shown).

(Fig.3D). While *IL12B* was rapidly induced in DCs, expression remained very low in tolDCs (Fig.3B). tolDCs demonstrated only a minor increase in *IL12A* and *IL27B* despite possessing up to 100 fold higher basal level of expression compared to DC (Fig.3C,D). *IL23A* expression was also higher in tolDC at time 0 but gradually decreased over time before returning to the initial expression level (Fig.3E). *IL27A* expression was comparable between the two cell types up until 6 hours, after which a drop off was noted for tolDC (Fig.3F). Finally although *IL10* started off with a higher level of expression in tolDC the pattern of upregulation



Figure 3: Kinetic expression of IL-12 family members and IL-10 mRNA in LPS-treated DCs and tolDCs. Cells were harvested on day 6, mRNA was isolated followed by cDNA synthesis. The transcript levels of (A) *IL12B, IL12A, IL27B, IL23A, IL27A* and *IL10* were determined by RT-PCR in unstimulated cells. Cells were stimulated with LPS for indicated time points; mRNA was isolated followed by cDNA synthesis. The transcript levels of (B) *IL12B, (C) IL12A, (D) IL27B* (E) *IL23A,* (F) *IL27A* and (G) *IL10* were determined. *GAPDH* mRNA expression from the same samples was used as an endogenous reference gene (relative mRNA expression). Data shown is mean  $\pm$  SD of 2 independent experiments.

was comparable between the two cell types with a transient upregulation at 2 hours followed by a gradual decrease over time (Fig.3G). The relative abundance of both *IL27B* and *IL12A* transcripts in unstimulated toIDC compared to DC, in contrast to other IL-12 family subunits, is intriguing and may indicate some inherent regulation of these subunits within tolerogenic DCs.

## Stimulation of toIDCs favours upregulation of the immunoregulatory chains IL-12p35, Ebi3 and IL-27p28.

Although toIDCs specifically maintained the expression of subunits involved in immunoregulatory processes (Fig.3), they remained relatively refractory to LPS stimulation in terms of IL-12 family expression. We investigated if this profile was intrinsic for these cells or was activation dependent.

We observed that although DCs expressed relatively abundant levels of *IL12B*, particularly upon stimulation with LPS, IFN $\gamma$  + LPS or Poly I:C, transcripts of this subunit were significantly reduced in toIDC (Fig.4A).

We demonstrate that *IL12A* is maintained in toIDCs and expressed comparably to DC across the different stimuli used. Notably treatment with a combination of IFN $\gamma$  + LPS yielded the highest *IL12A* expression in toIDCs (Fig.4B). IFN $\gamma$ , and a combination of IFN $\gamma$  +LPS also displayed the greatest increase in *IL27B* expression when comparing DC to toIDCs (Fig.4C), with CD40 ligation showing the next highest expression in toIDC. *IL27A* levels were comparable between both DCs but IFN $\gamma$  yielded the strongest induction of *IL27A* transcripts in toIDCs (Fig.4D). In summary toIDCs do not express *IL12B* but do maintain expression of *IL12A*, *IL27A* and *IL27B* which can be further induced upon activation with IFN $\gamma$ , IFN $\gamma$  + LPS and CD40 ligation.



**Figure 4: Stimulation of tolDCs favours upregulation of the immunoregulatory chains IL-12p35, Ebi3 and IL-27p28.** Dendritic cells were stimulated for 6 hours with IFNγ, LPS, PGN,

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Poly (I: C) or a combination of both IFN  $\gamma$  + LPS. Irradiated CD40L cells were added at a ratio of 1:5. The transcript levels of (A) *IL12B*, (B) *IL12A*, (C) *IL27B*, (D) *IL27A* were determined by RT-PCR. *GAPDH* expression from the same samples was used as an endogenous reference gene (relative mRNA expression). Data shown is mean  $\pm$  SD of 3-5 independent experiments. Untransfected L cells were used as a control for CD40L cells and yielded results in line with medium conditions (data not shown).

# ToIDC produce low levels of IL-27 but maintain production of IL-35 related chains.

We investigated if our observations regarding transcriptional expression of IL-27p28, Ebi3 and IL-12p35 were translated into protein by assessing production in DC and toIDCs. Using an IL-27 ELISA we found that DCs, but not toIDCs, produced IL-27 upon IFN $\gamma$  + LPS stimulation, although we did find considerable variation across the 6 donors analysed (Fig.5A). Notably a combination of IFN $\gamma$  + LPS or CD40L with either IFN $\gamma$  or LPS, were the best inducers of IL-27 production in DCs while the individual stimuli alone were poor inducers (Fig.5B). Under all conditions toIDCs were significantly hampered in their IL-27 production (Fig.5B).

Ebi3 and IL-12p35 can dimerise to form IL-35; however there are no reliable methods for the detection of IL-35 in human cellular supernatants. We established by intracellular fluorescent microscopy that toIDCs do express the individual chains, Ebi3 and IL-12p35, and that production of both can be increased upon stimulation with IFN $\gamma$  or IFN $\gamma$  + LPS (Fig.5C). To quantify this, we measured Ebi3 and IL-12p35 by intracellular flow cytometric staining in both DC and toIDC. We found that although toIDCs had lower basal levels of Ebi3 compared to DC, they showed a stronger induction upon stimulation (Fig.5D). The relative increase in Ebi3 expression upon stimulation with IFN $\gamma$  or IFN $\gamma$  + LPS in tolDCs was 6 fold or higher (Fig.5D). Although toIDC have a lower basal level of IL-12p35 compared to DC, they increase expression comparably to DC upon stimulation with IFN $\gamma$  or IFN $\gamma$  + LPS (Fig.5E). To further reinstate our findings we performed a western blot for IL-12p35 and Ebi3 on whole cell lysates. In line with our flow cytometry and microscopy data, tolDCs do express both IL-12p35 and Ebi3, and both appear to increase upon stimulation. Importantly, in contrast to DC, toIDC do not express IL-12p40 using any stimuli assessed (Fig.5F).

## ToIDCs require expression of IL-12p35 to elicit their full tolerogenic potential.

To determine the functional significance of IL-12p35 and Ebi3 produced by toIDC, we investigated the suppressive activity of toIDC supernatants on  $\alpha$ CD3/CD28 activated T cells. Supernatants from CD40 ligated toIDCs demonstrated a pronounced inhibition of T cell proliferation which was not observed with mock activated toIDCs (Fig.6A). Using different toIDC and T cell donors, we found



Figure 5: ToIDC maintain and can enhance their protein expression of IL-12p35 and Ebi3 in the absence of IL-12p40. (A) DCs and toIDCs were harvested on day 6, replated and stimulated with IFN $\gamma$  + LPS; IL-27 levels were assessed by ELISA. (B) Cells were stimulated with IFN $\gamma$ + LPS, IFN $\gamma$ +CD40L, LPS+CD40L or CD40L for 48 hours followed by detection of IL-27 by ELISA. Data shown is mean ± SD of 3-6 independent experiments. (C) toIDCs were cultured on 8 well chamber slides and either untreated, or stimulated with IFN $\gamma$  or IFN $\gamma$  + LPS followed by overnight incubation with Brefeldin A. The cells were then fixed and permeabilised, followed by incubation with purified anti-huEbi3 or IL-12p35-APC. Ebi3 was detected using GaM-Alexa<sup>488</sup>. For flow cytometry DCs and toIDCs were either untreated or stimulated for 16 hours with IFN $\gamma$ alone or IFN $\gamma$  + LPS, followed by culture for a further 5 hours in the presence of 10µg/ml Brefeldin A. The cells were then harvested, fixed and permeabilised, followed by incubation with (D) Ebi3-APC or (E) IL-12p35-APC. Data shown is representative of 3 independent experiments, with graphed data showing the mean ± SD of those 3 experiments. (F) DCs were treated similarly for the generation of cellular lysates. Samples were loaded on 10% SDS gel followed by transfer to

nitrocellulose membranes. Blots were probed with indicated primary antibodies and detected with super signal ECL system. (data not shown).

that supernatant from CD40L activated toIDC consistently induced suppression ranging from 30-80% (median 60%) (Fig.6B).

To establish the role of IL-35 production by tolDC in T cell suppression, we performed RNA interference of *IL12A*. By mRNA analysis we found at least a tenfold inhibition of *IL12A* expression in activated tolDCs (Fig.6C). Importantly *IL27B* and *IL10* remained unaffected by *IL12A* silencing, and use of a non-specific siRNA did not affect *IL12A* expression (Fig.6C).



Figure 6: ToIDCs require IL-12p35 in order to elicit their full tolerogenic potential. toIDCs were harvested on day 6 and stimulated with irradiated CD40L or mock transfected cells at a ratio of 1:5 for 48 hours. Supernatants were harvested and incubated with  $\alpha$ CD3/28 stimulated naïve CD4<sup>+</sup> T cells at a dilution of 1:2 for 72 hours. <sup>3</sup>H incorporation was determined during the last 8-12 hours of culture. Results shown in (A) are from a representative experiment (mean ± SD of CPM from triplicate cultures). Data shown in (B) is the mean ± SEM percentage suppression from 14 independent experiments. (C) *IL12A* was silenced in toIDCs demonstrating up to 90% interference in mRNA transcripts of *IL12A*. IL-12A silenced or control treated toIDCs were harvested and stimulated as described above for 48 hours. The supernatants were then incubation with  $\alpha$ CD3/28 stimulated naïve CD4<sup>+</sup> T cells at a dilution of 1:2 for 72 hours. <sup>3</sup>H incorporation was determined during the last 8-12 hours of culture. CPM shown in (D) are from a representative experiment of

3 independent experiments performed. Percentage inhibition shown in (E) is the mean  $\pm$  SD of those 3 experiments. DCs were harvested on day 6 and were either mock (-), IL-12A, or Non-target siRNA treated, followed by stimulation with IFN $\gamma$  + LPS. Cell supernatants were harvested and assessed for production of (F) IL-12p70 and IL-12p40 by ELISA, data shown is mean  $\pm$  SD of 4 independent experiments. siRNA treated DCs were stimulated with CD40L, supernatants were harvested after 48 hours and incubated with  $\alpha$ CD3/28 activated total CD4<sup>+</sup> T cells for 3 days. (G) IFN $\gamma$  levels were determined in the T cell supernatants by ELISA.

Using supernatants from CD40L stimulated toIDCs we found that *IL12A* silenced supernatants were significantly impaired in their suppressive capacity (Fig 6D, 6E), thereby indicating that IL-12p35 is required by toIDCs to elicit their full tolerogenic potential. To confirm that the IL12A siRNA directly targeted IL-12p35 protein we demonstrated, in activated control DCs, that IL-12p70 was up to 60% inhibited while IL-12p40 was unaffected (Fig 6F). As a consequence, the capacity of DC supernatant to induce IFNy production in activated T cells was significantly reduced (Fig 6G).

## Discussion.

The IL-12 family is gaining increasing attention as being a key regulator of immunity and immunological tolerance<sup>25,26</sup>. A common feature of tolerogenic DCs is the lack of IL-12, however little is known about the expression, regulation or function of the other family members in human tolDCs. In this study we focused on the IL-12/IL-35 axis and found that tolDCs maintain, and can increase expression of Ebi3 and IL-12p35, and that IL-12p35 is required for their full tolerogenic potential.

Despite the remarkable degree of chain sharing between the four members of the IL-12 family, each one has its own unique biological function. IL-12 and IL-23 are pro-inflammatory cytokines crucial for the development of Th1 and memory Th17 cells respectively<sup>27,28</sup>. On the other hand IL-27 and IL-35 are inhibitory and immunoregulatory cytokines<sup>20,29-31</sup>. IL-35 although a relatively new player within this family has already been shown to induce proliferation of iTr35, a regulatory T cell subset which possesses a profound anti-inflammatory phenotype with the ability to reduce inflammation in animal models of IBD and EAE<sup>20</sup>. With the striking immunoregulatory functions of IL-35 it became evident to us that the possible secretion of IL-35 from tolerogenic DCs could be a new previously unexplored mechanism into how these cells can suppress T cell responses. To study this we used a well characterised human toIDC and first established that they displayed typical features associated with regulatory DC populations -low CD86, high B7H1<sup>32</sup> and poor ability to stimulate allogenic T cells<sup>8,33</sup>. In addition we observed that these cells express high levels of the endocytic receptor CD206 as well as the immunoregulatory molecules B7DC<sup>34,35</sup> and ILT3<sup>36</sup>. Our focus was

on the cytokine repertoire of these cells and one of the more intriguing qualities of toIDCs is the profound lack of IL-12p40/70 production regardless of stimulation used. We have shown that toIDCs completely lack IL-12p40 expression yet preserve their transcriptional expression of the other IL-12 family members, in particular those who, upon dimerisation are associated with anti-inflammatory properties, namely Ebi3, IL-12p35 and IL-27p28.

Analysing LPS triggered DCs, we observed that IL-12p35/p40 were transient and superimposable with both subunits reaching their highest level at 6 hours in DCs. A similar peak in expression was noted for IL-27p28, though the increase in expression appears to occur more rapidly than for p40/p35. Conversely Ebi3 does not peak until 24-48 hours post activation in DCs. This differential pattern of expression has been likewise noted in previous studies<sup>37,38</sup> and has fuelled speculation that Ebi3 may have a unique biological function independent of its typical pairing subunits. Recent evidence using murine cells showed that type 2 anti-inflammatory macrophages express high level of Ebi3<sup>39</sup>. We have also observed elevated Ebi3 levels in M-CSF generated human M2 compared to GM-CSF generated M1 macrophages (unpublished data). Intriguingly, the authors show that intracellular Ebi3 expression can block LPS induced M1 to M2 transition and IL-12p70 production<sup>39</sup>, suggesting that Ebi3 may play a role in maintaining the regulatory functions of M2 macrophages. Further work is needed to fully extrapolate this data to human macrophage subsets.

In our study with dendritic cells we not only observed consistently higher levels of Ebi3 in tolDCs, but also IL-12p35. We observed minimal upregulation when the toIDCs were stimulated with LPS so we addressed whether this was an intrinsic characteristic of these cells or was activation dependent. Notably IFNy seemed to have the greatest effect on toIDC in terms of Ebi3/p35/p28 expression with just a moderate increase noted in DCs. This may be explained by the finding that toIDCs express higher levels of IFN $\gamma R1/2$  and STAT1 compared to DC (data not shown). Although IFNy is well acknowledged as an inflammatory mediator, in recent years it has also been appreciated as a key coordinator of immunity with many anti-inflammatory molecules also under its control, including B7H1<sup>40</sup> and IDO41,42. More recently IFNy has even been demonstrated to lead to the generation of a tolerogenic DC population in vitro<sup>43</sup>. In fact, Ebi3 and IFNy have been demonstrated to be important to limit allograft rejection in mouse, whereby IFNy upregulation of Ebi3 in autologous tolerogenic DCs was found to be a crucial mechanism in graft maintainance<sup>44</sup>. It is intriguing that in our current study, using human cells, Ebi3 was also strongly upregulated by IFNy in tolDCs. As described earlier Ebi3 can dimerise with either IL-27p28 or IL-12p35 to yield IL-27 or IL-35 respectively. In view of the expression of IL-27p28 in toIDCs we investigated IL-27 production in both DC and toIDC. IL-27 has been shown to

be produced by DCs stimulated with Poly I:C<sup>45</sup> or macrophages stimulated with IFN $\gamma$  + LPS<sup>46</sup>. We did not find any striking effect with Poly I:C in our study but did find that IFN $\gamma$  + LPS induced a strong IL-27 production in DCs. In addition we show that CD40 ligation alone or in particular when combined with IFN $\gamma$  or LPS can also induce IL-27 production. In contrast toIDCs had low levels of IL-27 in their supernatants with only a combination of IFN $\gamma$  with either LPS or CD40L showing any detectable levels. This may be explained by the rapid loss of IL-27p28 transcripts in toIDC after 6 hours which we observed using LPS induced kinetics. Further work is required to evaluate whether IL-27p28 transcripts are less stable in toIDC compared to DC or are possibly targeted for degradation.

Unfortunately there are no reliable methods to determine IL-35 in cellular supernatants. However we did show that tolDCs do express the individual chains required to form IL-35. We established the functional significance of our findings in toIDCs by performing a T cell proliferation assay in the presence of supernatants from activated toIDC and IL-12A silenced toIDC. Using the supernatants from control silenced toIDCs we demonstrated a robust inhibition of T cell proliferation while supernatants from IL-12A silenced toIDCs were significantly impaired in this ability, indicating that toIDCs require IL-12p35 to elicit their full tolerogenic potential. There has been much speculation as to whether DCs, particularly tolerogenic DCs could produce IL-35<sup>26</sup>. It has been reported that IL-12p35 needs to be part of a heterodimer in order to be secreted<sup>47</sup>. IL-12p35 can heterodimerise with IL-12p40 or Ebi3 and the toIDCs used in this study do not produce IL-12p40. Although we cannot exclude that there may be a yet unidentified additional chain which IL-12p35 may pair with it, it seems likely considering the abundance of IL-12p35 and Ebi3 in tolDCs, that the partial loss of suppressive capacity in IL-12A silenced toIDCs may be a reflection of the loss of IL-35. Taken together, our results suggest that human toIDCs produce IL-35, an exciting new immunoregulatory cytokine, providing an additional novel mechanism by which these cells can elicit their tolerogenic potential.

#### **Conflict of interest**

D.A.A.V. has submitted patents covering IL-35 that are pending and is entitled to a share in net income generated from licensing of these patent rights for commercial development.

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