

Functional and phenotypic characterization of distinct porcine dendritic cells derived from peripheral blood monocytes

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SUMMARY

Dendritic cells (DCs) are bone marrow-derived antigen-presenting cells that have an exquisite capacity to interact with T cells and modulate their responses. Little is known about porcine DCs despite the fact that they represent an important target in strategies that are aimed at modulating resistance to infection in pigs and may be of major importance in transplantation biology. We generated immature monocyte-derived porcine dendritic cells (MoDCs) directly from adherent peripheral blood cells treated with porcine granulocyte-macrophage colony-stimulating factor (GM-CSF) and interleukin-4 (IL-4). The cells were observed via electron microscopy and their phenotype was characterized using monoclonal antibodies. The functionality of the porcine MoDCs was demonstrated showing that the cells were capable of different specialized functions relevant to antigen capture and were potent stimulators in a primary allo-mixed leucocyte reaction. Treatment of the MoDCs with porcine cell line-derived necrotic factors resulted in the phenotypic and functional maturation of MoDCs. We confirmed also that monocyte-derived DCs were differentially regulated by cytokines, showing that transforming growth factor- β 1 (TGF- β 1) is able to redirect monocytic precursors into the differentiation pathway of Langerhans' cells presenting typical Birbeck granules. Interestingly, and in contrast to the human and murine model, we showed that the monocyte-derived porcine Langerhans'-type cells (MoLCs) were much more potent activators of allogeneic T cells than MoDCs obtained without TGF- β 1.

INTRODUCTION

Dendritic cells (DCs) are professional antigen-presenting cells (APC) that are responsible for the activation of naive T cells and the generation of primary T-cell responses.¹ Their specific role is to capture, process and present antigens to T cells. Immune and inflammatory signals are responsible for the migration of DCs from tissues to lymphoid organs where they initiate an immune response. These processes induce the maturation of DCs, which results in a shift from a processing to a presenting stage.

DCs are characterized by their unusual dendritic morphology, by their potent antigen-presenting capacity and by the lack of T cells, B cells, natural killer (NK) cells and monocyte-specific markers. Nevertheless, considerable functional and

phenotypic heterogeneity has been documented within the DC system. It is still not clear whether the different DCs represent different stages of maturation of unique DC lineage or whether they stem from different progenitors.²

DCs are found in virtually all tissues of the body and circulate in the peripheral blood, but large-scale preparation of DCs is hampered by their very low frequency. Previous reports have described the *in vitro* generation of human DC using the CD34⁺ progenitor pool in the cord blood or by using monocyte/macrophage CD14⁺ blood cells.^{3–6} Cord blood CD34⁺ haematopoietic progenitor cells treated with granulocyte-macrophage colony-stimulating factor (GM-CSF) and tumour necrosis factor- α (TNF- α) differentiate into two distinct DC lineages: Langerhans' cell (LC)-type DCs and CD14⁺-derived DCs. Monocytes treated with GM-CSF and interleukin-4 (IL-4) rapidly become non-adherent, acquire DC morphology and down-regulate CD14 expression. LCs represent a discrete population among DCs, and the current hypothesis is that LCs arise from a specific precursor.^{4,7} However, it has been shown recently that LCs may originate from monocytes in response to GM-CSF, IL-4 and transforming growth factor- β 1 (TGF- β 1).⁸

Little is known about porcine DCs despite the fact that they represent an important target in strategies that are aimed at

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Abbreviations: APC, antigen-presenting cells; DC, dendritic cell; DX, dextran; MLR, mixed leucocyte reaction; MoDC, monocyte-derived DC; MoLC, monocyte-derived Langerhans' cell.

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modulating resistance to infection in pigs and should have major importance in transplantation biology.

As monocytes are directly accessible from peripheral blood and seem to represent a cell population with a high developmental flexibility towards DCs, we chose to generate monocyte-derived porcine DCs (MoDCs). In the present study, we characterized the phenotype and the functionality of cells generated from adherent peripheral blood mononuclear cells (PBMC) using porcine GM-CSF and IL-4. Furthermore, we confirmed that MoDCs are differentially regulated by cytokines, showing that TGF- β 1 is able to redirect monocytic precursors into the differentiation pathway of LCs.

MATERIALS AND METHODS

Animals

Blood samples were taken from Large White/Landrace hybrid pigs (10 to 13 weeks of age) of both sexes. Pigs were maintained in conventional conditions on a commercial diet.

Cytokines, maturation factors and cell line

Recombinant swine GM-CSF and IL-4 were prepared in our laboratory from CHO-K1 cells (ECACC no. 85051005, European Collection of Cell Culture, Salisbury, UK) transiently transfected with GM-CSF- or IL-4-coding plasmids. The supernatant from transfected cells was harvested after 24 hr, centrifuged (4800 g) and stored at -20° . The functionality of these two cytokines has been tested previously using bioassays (data not shown): IL-4-containing supernatant was shown to induce porcine lymphoblast proliferation up to a dilution of 1:2000, and GM-CSF-containing supernatant was shown to induce the differentiation of granulomacrophagic colonies (CFU-GM) from porcine bone marrow cell culture *in vitro*.

Porcine TNF- α and TGF- β 1 were obtained from Endogen (Woburn, MA). Lipopolysaccharide (LPS) (from *Salmonella abortus equi*) was purchased from Sigma (St Louis, MO). Necrotic factors were obtained following four freeze-thaw cycles of the IRP porcine cell line. The IRP cell line originates from porcine kidney and has been established in-house.

Cell isolation and cultures

PBMC were isolated by using Pancoll ($d=1.077$; D. Dutscher, Strasbourg, France) density-gradient centrifugation (1100 g for 30 min) of heparinized blood obtained from healthy pigs. These PBMC were plated (1×10^7 cells/3 ml per well) into 6-well plates (Corning, Brumath France) in medium (RPMI-1640 supplemented with 10% heat-inactivated fetal calf serum [FCS], 50 μ mol/l of β -mercaptoethanol and antibiotics; Life Technologies, Cergy Pontoise, France) supplemented with 1% GM-CSF- and IL-4-containing supernatants. Non-adherent cells were removed after 16 hr of incubation at 39° ; the adherent cells were maintained in the same culture conditions. The DC cultures were fed every 3 days with fresh medium and 1% GM-CSF- and IL-4-containing supernatants. In some cultures, 10 ng/ml of TGF- β 1 was added at the time of culture initiation.

Electron microscopy

Cultured cells (sampled after 7 days of culture) were fixed in 2% glutaraldehyde, postfixed in 1% osmium tetroxide (Euromedex, Souffelweyersheim, France) and embedded in Epon 812

(Sigma). Ultrathin sections were then cut, stained with uranyl acetate and lead citrate, and examined using an electron microscope (Philips CM10; Lyon 1 Microscopy Center, Lyon, France).

Immunofluorescence analysis

Immunofluorescence analysis was performed on 5×10^5 cells. Monoclonal antibodies (mAbs) to pig CD1 (clone 76-7-4), human (hu) CD36 (CB38) and huCD68 (Y1/82 A), were obtained from PharMingen (La Jolla, CA). mAbs to pig CD11a/18 (MUC76A), CD14 (M-M9), CD11b/c (PGBL18A) and major histocompatibility complex (MHC) class I (H58A) were obtained from VMRD Inc. (Pullman, WA). mAbs to CD25 (231.3B2), CD45Ra (MIL13) and MHC class II (274 3G8) were obtained from Serotec (Oxford, UK). mAb to hup55 Fascin (55K-2) was obtained from Dako (Carpinteria, CA). In indirect assays, reactivity was detected using either fluorescein isothiocyanate (FITC)- or phycoerythrin (PE)-conjugated anti-mouse immunoglobulin G (IgG) and immunoglobulin M (IgM) (Interchim, Montlucon France). Intracellular staining was performed using the Cytofix/Cytoperm plus kit (PharMingen). Cells were analysed using a fluorescence-activated flow cytometer (FACScalibur flow cytometer; Becton-Dickinson, Mountain View, CA) calibrated with Calbrite beads (Becton-Dickinson) for FITC and PE. The distribution of debris, dead cells and any contaminating red blood cells was assessed on the basis of forward and right-angle scatter before proceeding with the analysis. A total of 10 000 events were examined using a 488-nm wavelength excitation. Acquired events were analysed using CELL QUEST software (Becton-Dickinson).

Endocytosis

Cells were washed in phosphate-buffered saline (PBS) and incubated with 0.1 mg/ml of FITC-dextran (DX-FITC; 40 000 molecular weight) or with 1 mg/ml of bovine serum albumin (BSA)-FITC (Sigma Chemical Co.) at 39° or 0° for different periods of time. Uptake was stopped by adding ice-cold fluorescence-activated cell sorter (FACS) buffer (PBS, 5% BSA, 0.02% sodium azide) containing 1% formol.

Formation of immune complexes (ICs)

ICs were formed by incubating 5 μ g of β -galactosidase (Sigma) with anti- β -galactosidase serum from Sigma (1:100). Free β -galactosidase was eliminated by heat denaturation at 62.5° . The resulting immune complexes were then assayed by detection of the remaining β -galactosidase enzyme activity protected by antibodies, using FDG reagent (Sigma). DCs were fed with the β -galactosidase ICs or non-denatured β -galactosidase protein (as a control) for 4 hr. Incorporated β -galactosidase was finally revealed by flow cytometry using the FDG substrate.

Mixed leucocyte reaction (MLR)

APCs were treated with mitomycin C (50 μ g/ml; Sigma) for 20 min and washed three times in PBS. PBMC (5×10^4) were stimulated at different APC/effector-cell ratios for 5 days. Cultures were performed in triplicate (in 96-well plates) in 200 μ l of culture medium. During the final 8 hr of incubation, each culture was labelled with 0.5 μ Ci of [3 H]thymidine (Amersham, Les Ulis France). Cells were harvested onto glass fibre disks using a multiple cell harvester, and thymidine

incorporation was determined (in counts per minute [c.p.m.]) in a liquid scintillation counter (MicroBeta counter; EG & G Wallac, Evry France). The results were expressed as index of proliferation:

$$\text{Index of proliferation} = \frac{(\text{mean c.p.m. of stimulated wells})}{(\text{mean c.p.m. of control wells})} \times 100$$

RESULTS

Generation of porcine DCs from cultured adherent PBMC

DCs were derived from adherent PBMC cultured in medium supplemented with porcine GM-CSF and IL-4. Adherent aggregates were visible on day 3. After 7 days, cultures consisted mainly of single and aggregated floating cells, presenting a typical veiled morphology. The cells obtained were evaluated by transmission electron microscopy, which revealed that the cells had a large diameter, showed pronounced protrusions and microvillous projections of their plasma membrane, and contained abundant multivacuolar and multilamellar vesicles (Fig. 1). Overall, the adherent PBMC-derived porcine cells showed ultrastructural features characteristic of DCs.

The surface phenotype of this DC-enriched cell population was analysed. The cells did not express T-, B-, or NK-cell markers (data not shown). Expression of the monocyte/macrophage marker, CD14, was assayed on days 3 and 7 after the start of the culture (Fig. 2). A marked down-regulation of its surface expression was observed, indicating the differentiation of the adherent monocytes into DCs. To assess more specifically the phenotype of these cells, several mAbs directed against surface markers were used. The results presented in Fig. 2 showed that the generated cells were MHC class I⁺, MHC class II⁺, CD1⁺, CD11a/18⁺, CD11b/c⁺, CD25⁻, CD36⁺, CD68⁺ and p55 fascin positive. Although none of the markers tested is specific for the dendritic lineage, they all have been reported to be expressed by DCs. The cells expressing the CD1 antigen are recognized by the mAb 76-7-4. This antibody was recently described as being specific to a new member of the CD1 family expressed in porcine cells and shares a high homology with the human CD1a antigen.⁹ The porcine cells strongly expressed β 2-integrins (CD11b/c and CD18), which are important in cell-cell interactions and represent binding molecules for bacteria and LPS.¹⁰ Using an anti-human CD68 mAb, we found that the porcine DCs expressed the putative intracellular and surface CD68, which is a lysosome-associated membrane glycoprotein abundant in MoDCs.¹¹ Surface CD68 expressed by DCs has been reported to possibly act as a ligand for E-selectin.¹² The actin-bundling p55 fascin protein, which causes aggregation of F-actin into bundles in membrane protrusions,¹³ was also detected in the porcine DCs, using an anti-human p55 mAb. p55 fascin has been reported to be expressed and restricted to mature DCs.⁵ On the other hand, these DCs express a putative CD1a molecule and low levels of CD14, which represent a phenotype compatible with an immature stage of differentiation.^{14,15}

Taken together, these results suggest that adherent porcine PBMC cultured with GM-CSF and IL-4 generate cells displaying all the characteristics of MoDCs. Further functional

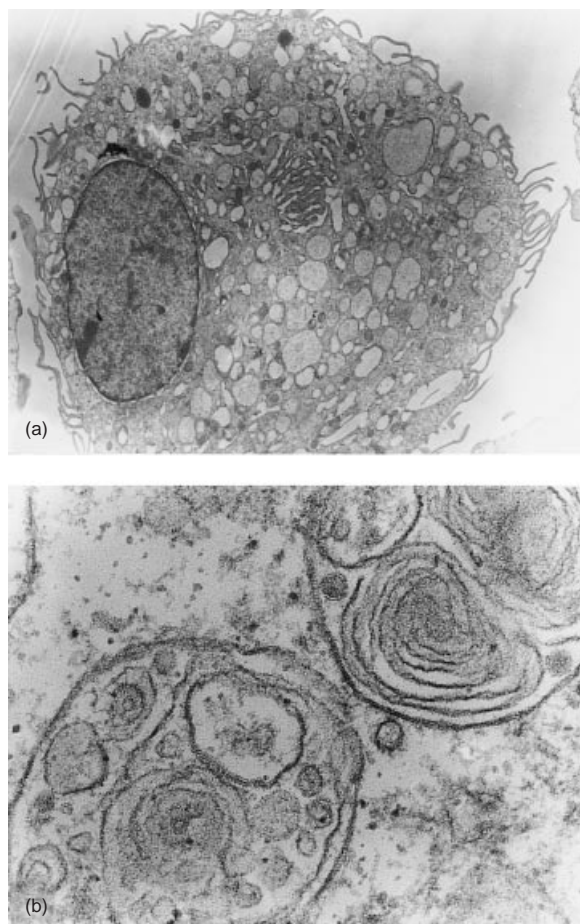


Figure 1. Transmission electron microscopy was used to evaluate the cellular morphological characteristics of adherent peripheral blood mononuclear cells (PBMC) after 7 days of culture in the presence of porcine granulocyte-macrophage colony-stimulating factor (GM-CSF) and interleukin-4 (IL-4). (a) Electron micrograph showing a cell with a large diameter and microvillous projection of the plasma membrane (magnification $\times 2600$). (b) Details of the multivacuolar and multilamellar intracytoplasmic organelles (magnification $\times 46\,000$).

analyses are needed to define the stage of differentiation of the derived porcine MoDCs.

Porcine MoDCs are capable of several antigen-capture mechanisms

Immature DCs usually possess different specialized functions relevant to antigen uptake, including receptor-mediated endocytosis and macropinocytosis.¹ We investigated whether the *in vitro*-derived porcine MoDCs were able to take up FITC-conjugated ligands. Figure 3 shows that porcine MoDCs were efficient at the uptake of both DX-FITC and BSA-FITC. This property was abolished when the experiments were performed at 4°.

The kinetic experiments performed at short time intervals indicated that 15 min is sufficient to load the majority of DCs with DX-FITC or BSA-FITC. The fluorescent signal obtained after 15 min of DX-FITC loading was maximal and did not

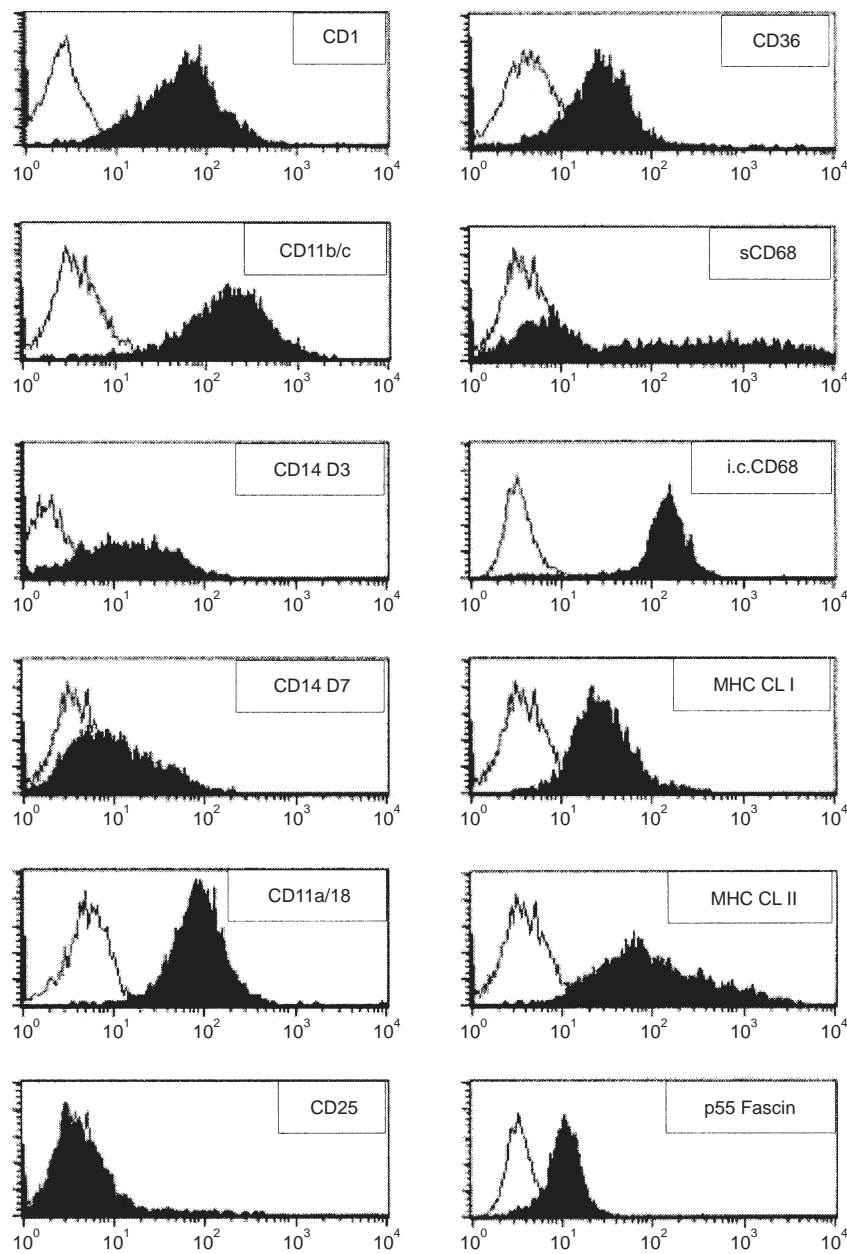


Figure 2. Flow cytometric phenotyping of the porcine dendritic-type cell. Cells were stained with control isotypes (open histogram) or with the indicated anti-human or anti-porcine monoclonal antibody (mAb) against cell surface markers (solid histogram). Intracellular (i.c.) staining was performed for the human (hu) p55 Fascin. Both i.c. and surface (s) staining were performed for CD68. Data are representative of at least three experiments performed.

increase when the incubation time was extended to 2 hr (Fig. 3). In contrast, the signal intensity obtained after incubation with BSA-FITC was increased after 120 min. This observation suggests that the two ligands are taken up by different mechanisms. DX-FITC uptake has been shown to be mannose-receptor mediated and the saturated signal probably represents saturation of the mannose receptors.³ In contrast, BSA-FITC uptake is not saturated over time, suggesting that its uptake is mediated by macropinocytosis, independently of membrane receptors. Uptake of both DX-FITC and BSA-FITC was

significantly increased when the incubations were carried out for a longer time-period (6–8 hr; data not shown).

Another pathway for antigen uptake by DCs is mediated through binding of ICs to Fc γ R receptors.¹⁶ MoDCs were fed with β -galactosidase ICs for 4 hr and intracellular β -galactosidase activity was revealed using the fluorescent FDG substrate. Figure 3 shows that β -galactosidase ICs were efficiently internalized into the MoDCs. Therefore, the porcine MoDCs appeared to be efficient at both receptor- and non-receptor-mediated endocytosis.

MoDCs are efficient APC in primary allo-MLR

It is well established that the ability of DCs to stimulate a potent allo-MLR distinguishes them from other leucocytes.¹⁷ Immature porcine MoDCs were used on day 7 as stimulators of unprimed heterologous porcine PBMCs. As shown in Fig. 4, MoDCs were much more potent activators of allogeneic T cells

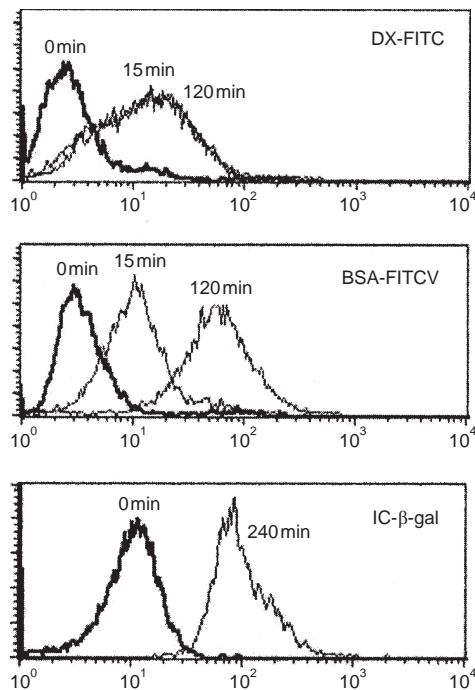


Figure 3. Uptake of dextran-fluorescein isothiocyanate (DX-FITC), bovine serum albumin (BSA)-FITC and β -galactosidase (β -gal) immune complexes (ICs). Monocyte-derived dendritic cells (MoDCs) were allowed to take up the different ligands for the indicated times at 39° (—) or on ice (---). The resulting fluorescence was read by flow cytometry. Results are representative of at least three experiments.

than were PBMCs from the same animal. Under the same conditions, autologous MoDCs did not induce proliferation (data not shown).

Phenotypic and functional maturation of the porcine MoDCs can be induced *in vitro*

LPS, TNF- α and necrotic factors were tested for their ability to induce the maturation of MoDCs. Figure 5(a) shows the down-regulation of the CD1 and CD14 surface markers and the up-regulation of MHC class II molecules observed after 3 days of treatment with the necrotic factors. The down-modulation of CD1a and CD14 markers has been reported to occur during DC maturation.^{14,15} We showed here that the CD1 molecule recognized by the 76-7-4 mAb seemed to behave like the CD1a molecule expressed by human DCs.

The down-modulation of receptor-mediated endocytosis of DCs after treatment of immature DCs with maturation stimuli has been well documented.¹⁸ To evaluate whether the necrotic factor-induced phenotypic maturation of MoDCs was accompanied with down-regulation of antigen uptake, we measured DX-FITC uptake by untreated and necrotic factor-treated MoDCs. In contrast to untreated MoDCs, DX-FITC uptake was markedly down-regulated in necrotic factor-treated MoDCs (Fig. 5a). The same phenotypic and functional alterations were obtained using LPS or TNF- α as maturation factors, but the strongest effect was mediated by the necrotic factors (data not shown).

Finally, the stimulatory properties in a primary allo-MLR of the necrotic factor-treated MoDCs were compared with those of the untreated MoDCs. Untreated MoDCs were clearly weaker in their stimulatory capacity (Fig. 5b). Therefore, treatment with necrotic factors induced MoDCs that were much more potent activators of allogeneic T cells as compared to non-treated cells.

Our data clearly demonstrate that porcine MoDCs can mature *in vitro* and suggest that MoDCs derived in the presence of GM-CSF and IL-4, and without the addition of other factors, give rise to immature porcine MoDCs.

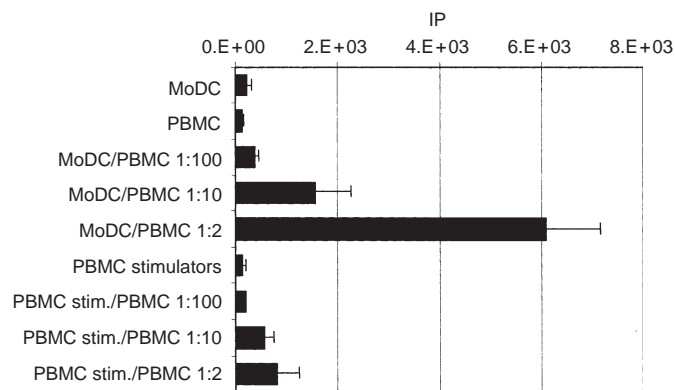


Figure 4. Five-day primary mixed leucocyte reaction (MLR) showing allogeneic T-lymphocyte activation in response to monocyte-derived dendritic cells (MoDCs). Proliferation was determined by [³H]thymidine incorporation. Represented are the means and SD of triplicate wells containing mitomycin-treated allogeneic MoDCs or peripheral blood mononuclear cells (PBMC) and 2×10^5 fresh responding PBMC, at the indicated ratios. MoDCs or PBMC alone were included as controls. Data shown are representative of four experiments. IP, index of proliferation (see Materials and methods).

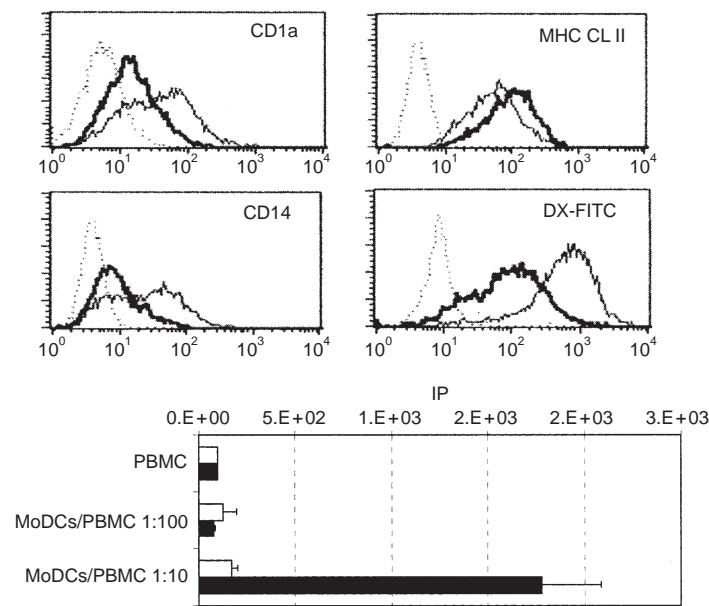


Figure 5. Phenotypic and functional modification induced by treatment of the monocyte-derived dendritic cells (MoDCs) with IRP necrotic factors. (a) Five-day-old MoDCs were cultured for 4 days with (thick line) or without (thin line) IRP necrotic factor (1:2 dilution). Untreated (thin line) and necrotic factor-treated (thick line) MoDCs were stained for CD1a, CD14 and major histocompatibility complex (MHC) class II molecules and control isotypes (dotted line). Untreated (thin line) and necrotic factor-treated (thick line) MoDCs were incubated for 4 hr with dextran-fluorescein isothiocyanate (DX-FITC) at 39° or on ice (dotted line). (b) Proliferative responses of 2×10^5 porcine PBMC in response to allogeneic untreated (open histogram) or necrotic factor-treated (solid histogram) MoDCs. IP, index of proliferation (see Materials and methods).

Addition of TGF- β 1 induces Langerhans²-type cell derivation

Several observations suggest that monocytes can differentiate into LCs when TGF- β 1 is added to the culture medium together with GM-CSF and IL-4.⁸ We tested whether the addition of TGF- β 1 could induce porcine monocytes to differentiate into LC. After 7 days of culture in the presence of GM-CSF, IL-4 and TGF- β 1, the generated cells were observed by electron microscopy. Dendritic-shaped cells with a moderate amount of Birbeck granules (resembling tennis racket-like structures) were observed (Fig. 6a, 6b), indicating a typical LC morphology. These MoLCs contained moderate amounts of Birbeck granules compared to LCs observed *in vivo*. We observed that the cell morphology of MoLCs was slightly different from the MoDCs generated in the absence of TGF- β , displaying shorter and thicker dendrites (Fig. 6a).

Relative efficiency of MoDCs and MoLCs to stimulate a primary allo-MLR

It has been shown previously that DCs grown in the presence of TGF- β 1 exhibit characteristics of fully immature or early DCs.^{19–21} Moreover, inhibition of acquisition of mature DC features by cultured LCs has been consistently observed and appears to be mediated by TGF- β 1. In the present study, no phenotypic differences were found between the two types of cells by flow cytometry analysis using the mAbs described above (data not shown). When matured with necrotic factors, LPS or TNF- α , the MoLCs behaved like the MoDCs, showing down-regulated expression of CD1 and CD14 surface markers,

a down-regulated DX-FITC uptake ability and an enhanced MLR activator potency (data not shown).

Surprisingly, when MoDCs and MoLCs were compared for their capacity to stimulate allogeneic PBMCs in a primary allo-MLR, it was found that porcine MoLCs were significantly more efficient for primary T-cell activation (Fig. 6c). This result suggests that TGF- β 1 regulates, in a positive manner, the antigen-presenting capability of the porcine MoLCs generated.

DISCUSSION

We have shown in the present work that adherent porcine PBMC cultured *in vitro* in the presence of porcine GM-CSF and IL-4 differentiate into cells displaying the morphology, the phenotype and the functions of MoDCs. In addition, phenotypic and functional modifications of the MoDCs in response to maturation stimuli allow us to predict the differentiation stage of the generated cells. We showed that the matured MoDCs presented a down-regulated CD14 and CD1 surface expression, while MHC class II molecules were up-regulated in comparison to the untreated MoDCs. The loss of CD14 and CD1a surface markers has been shown to correlate with human DC maturation.^{14,15} The porcine CD1 molecule (recognized by the 76-7-4 mAb in this study) is encoded by the recently cloned pCD1.1 gene and represents a new member of the CD1 family.⁹ It was shown (see ref. 9) that the pCD1.1 nucleotide and deduced amino-acid sequences present the highest homology with the human CD1a molecule. The results reported in the present study suggest that the expression of this porcine CD1 molecule on DCs also may be regulated like the CD1a human molecule. It is known that up-regulation of MHC class II

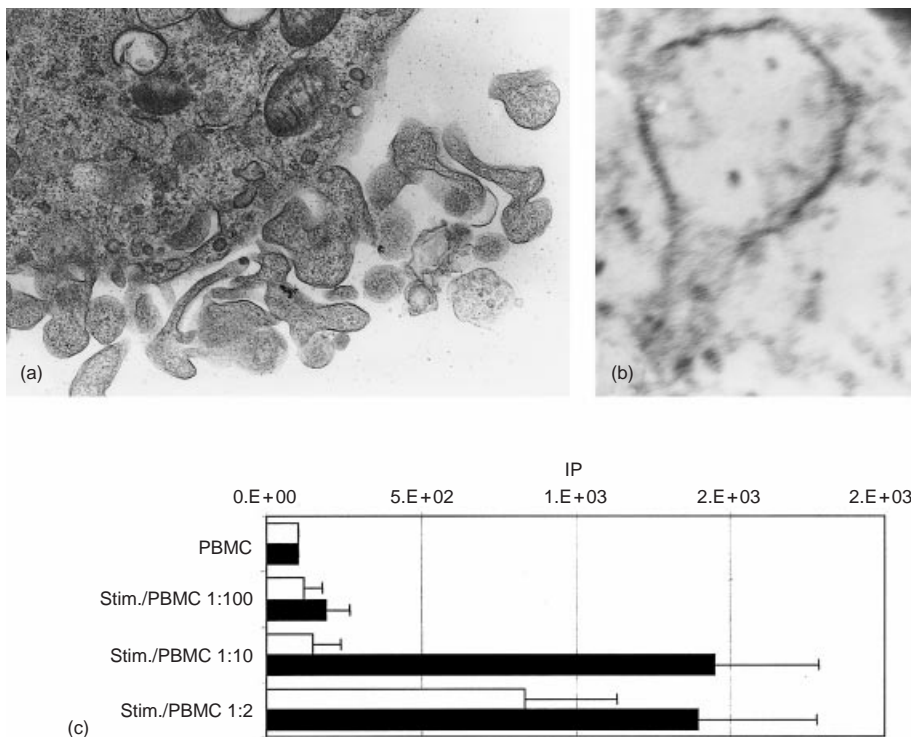


Figure 6. Addition of transforming growth factor- β 1 (TGF- β 1) induces Langerhans'-type cell derivation. Electron micrographs are presented of monocyte-derived Langerhans' cells (MoLCs) showing (a) dendrite details (magnification $\times 10\,500$) and (b) a Birbeck granule (magnification $\times 92\,000$). (c) Proliferative responses of 2×10^5 porcine peripheral blood mononuclear cells (PBMC) to allogeneic mitomycin-treated monocyte-derived dendritic cells (MoDCs) (open histogram) or MoLCs (solid histogram) after 5 days of culture. Data shown are representative of three experiments. IP, index of proliferation (see Materials and methods).

molecules correlates with DC maturation and results from the redistribution of molecules from the intracellular vesicular compartments to the cell surface.²² Moreover, we showed that the phenotypic maturation was accompanied with a down-regulation of the receptor-mediated uptake activity and an increase of the MLR stimulatory capacity. These functional changes correlate with the shift from the processing stage to the presenting stage, which occurs during DC maturation, and have been widely characterized in the literature.^{18,22} In a recent report published by Basta *et al.*, analysing the changes occurring during porcine monocyte differentiation into macrophage, the uptake of DX-FITC was used as a pH-sensitive probe.²³ It was shown that during the maturation process, DX-FITC fluorescence, which is lost in the acidic endosomal vesicles, is inversely correlated with the endocytic efficiency. In contrast, in our experiment the intensity of the fluorescence signal in DX-FITC-loaded DCs seemed to be directly correlated with the uptake efficiency. This difference suggests that the DX-FITC intracellular trafficking is different between the two types of cells and that DX-FITC might escape acidic endosomal vesicles in DCs, as shown by Rodriguez *et al.* in a mouse DC line.²⁴

It has been described in detail that, during the maturation process, DCs increase their level of surface costimulatory molecules such as CD40, CD80 and CD86, which provide the additional signals required to initiate T-cell activation.²² Unfortunately, we were not able to investigate the expression and up-regulation of the costimulatory molecules in the

absence of porcine-specific or cross-reacting antibodies directed against these molecules. The expression of the costimulatory B7 molecules on bone-marrow-derived porcine DCs has been reported by West *et al.* using the human cytotoxic T-lymphocyte antigen-4 (CTLA-4)-immunoglobulin fusion protein.²⁵ Overall, the porcine DCs appeared to behave in a similar manner to the well-characterized murine or human MoDCs and we predict that the regulation of costimulatory molecule expression on porcine DCs should be equivalent to what has been described for mouse and human models. To our knowledge, the work reported here is the first to describe both immature and mature stages of porcine DCs derived *in vitro*.

Necrotic cell lines have been recently described as maturation factors for DCs.^{26,27} Among the different maturation factors tested, the necrotic factors obtained from the IRP porcine cell line seemed to be the most effective. Despite the fact that immature MoDCs expressed low levels of CD14 and strongly expressed β 2 integrin receptors, LPS seemed to represent a weaker maturation signal, suggesting that the supernatant of the necrotic IRP cell line may contain a combination of several factors involved in DC maturation. It is of interest that the role of necrotic factors in DC maturation has been challenged and may, in fact, be related to the presence of mycoplasma in the cell lines.²⁸ We cannot rule out this possibility as the IRP cell line used in the present work was not assayed for mycoplasma contamination.

LCs are DCs that are present in the epidermis, bronchi and mucosae. Little is known about their lineage of origin. TGF- β 1

has been shown to play an important role in their biology,^{29,30} and *in vitro* studies of human or murine haematopoietic progenitor cells revealed that precursor LCs generated in response to TGF- β 1 stimulation expressed monocyte-lineage marker molecules.^{31,32} Recently, Geissmann *et al.* demonstrated that TGF- β 1 in the presence of GM-CSF and IL-4 could induce the differentiation of human monocytes into Langerhans'-type cells.⁸ We showed in the present work that, in the same manner, porcine TGF- β 1 added to porcine GM-CSF and IL-4 drove *in vitro* differentiation of pig monocytes into Langerhans'-type cells that displayed moderate amounts of Birbeck granules.

It has been reported that the addition of TGF- β 1 *in vitro* promotes the differentiation of fully immature DCs and inhibits acquisition of mature DC features.^{19–21} Geissmann *et al.* have shown that Langerhans'-type cells derived from monocytes in the presence of TGF- β 1 express almost exclusively intracellular class II antigens and a low level of costimulatory molecules.¹⁹ Furthermore, these Langerhans'-type cells did not mature in response to inflammatory stimuli but needed cognate stimulus induced by CD40 to fully mature. In the present study, the phenotype of the porcine MoLCs was not distinguishable from the phenotype of the MoDCs, when studied using our panel of mAbs. Treatment of the MoLCs with maturation factors resulted in the same phenotype and functional modifications to the ones reported here for the MoDCs. We cannot exclude that a more detailed analysis of intracellular MHC class II localization, costimulatory molecule expression and cytokine production by our MoLCs would reveal some differences that distinguished them from the MoDCs. Surprisingly, we observed that the MoLCs were more potent activators in primary allo-MLR than the MoDCs. This observation suggests that TGF- β 1 promotes the differentiation of MoLCs with enhanced antigen-processing or costimulatory functions. Additionally, Riedl *et al.* have shown that TGF- β 1 reduced TNF- α -induced Fas expression on LCs derived from CD34⁺ progenitors, protecting them against Fas-mediated apoptosis.³³ We can hypothesize that a reduced susceptibility to Fas-mediated apoptosis of the stimulator MoLCs cells would account for the observed result. Additional work is required to clarify the effect of TGF- β 1 on the differentiation of porcine MoLCs.

Despite the fact that pigs represent an important food-animal group and, owing to their similarities with humans, constitute an animal model in medical research, little is known about the cellular immune responses in this species. In particular, access to the porcine CD8 T-cell-mediated immune response would be of great help in vaccine design and xenotransplantation studies. Moreover, the pig presents a original T-lymphocyte compartment containing sizeable, unconventional DP (CD4⁺ CD8⁺), DN (CD4⁻ CD8⁻) and $\gamma\delta$ T-cell subsets,^{34–37} rendering this species a suitable model for elucidating the potential role of these T cells in immune defences. Analysis of the *in vitro* pig CD8 T-cell response is hampered, in part, by the difficulty of generating APC in an outbred context. We showed in this report that DCs can be easily generated from individual pigs. Moreover, we demonstrated that these porcine DCs can be antigen loaded in different ways. It is now widely accepted that DCs are endowed with cross-presentation capacity^{38,39} and can therefore present exogenous antigen in combination with class I molecules.

We emphasize that these porcine DCs represent an important tool for *in vitro* studies.

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