CD137 Is Required for M Cell Functional Maturation but Not Lineage Commitment

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Mucosal immune surveillance depends on M cells that reside in the epithelium overlying Peyer’s patch and nasopharyngeal associated lymphoid tissue to transport particles to underlying lymphocytes. M cell development is associated with B lymphocytes in a basolateral pocket, but the interactions between these cells are poorly understood. In a cell culture model of M cell differentiation, we found lymphotoxin/tumor necrosis factor (TNF)/H9251 induction of CD137 (TNFRSF9) protein on intestinal epithelial cell lines, raising the possibility that CD137 on M cells in vivo might interact with CD137L expressed by B cells. Accordingly, while CD137-deficient mice produced UEA-1/H11545 putative M cell progenitors in nasopharyngeal associated lymphoid tissue and Peyer’s patch epithelium, they showed an abnormal morphology, including the absence of basolateral B cell pockets. More important, CD137-deficient nasopharyngeal associated lymphoid tissue M cells were defective in microparticle transcytosis. Bone marrow irradiation chimeras confirmed that while induction of UEA-1+ putative M cell precursors was not CD137-dependent, full M cell transcytosis function required expression of CD137 by radioresistant stromal cells as well as by bone marrow-derived cells. These results are consistent with a two-step model of M cell differentiation, with initial CD137-independent commitment to the M cell lineage followed by a CD137-CD137L interaction of M cells with CD137-activated B lymphocytes or dendritic cells for functional maturation. (Am J Pathol 2010, 177:666–676; DOI: 10.2353/ajpath.2010.090811)

The differentiation of lymphoid tissue stromal cells is dependent on complex inducing signals that lead to changes in specific patterns of gene expression among mesenchymal cells, endothelium, and epithelium. One particular apparent paradox in these developmental pathways is the finding that cytokines in the tumor necrosis factor (TNF)/lymphotoxin family are critical to both proinflammatory processes and to differentiation of lymphoid tissue stroma. Signaling by TNF/lymphotoxin superfamily receptors can activate nuclear factor κB (NF-κB) through both the classical (IKK-dependent) and non-classical (relB-dependent) pathways.1 Thus, there is no clear distinction between signals that lead to production of inflammatory cytokines versus those that lead to stable development of lymphoid tissue stromal cells such as high endothelial venules or lymphoid mesenchymal cells producing chemokines such as CCL21.2,3 Chronic production or presentation of TNF/lymphotoxin signals, as in transgenic mice or chronic autoimmune diabetes4 may lead to generation of lymphoid structures resembling secondary lymphoid tissue, but it is also possible that controlled combinations of factors may also specify differentiation versus inflammation.

In the case of mucosal lymphoid tissues such as Peyer’s patch (PP) and nasopharyngeal associated lymphoid tissue (NALT), in addition to the stromal cells associated with the scaffolding in the lymphoid follicle and high endothelial venule, specific inducing factors are required for the differentiation of the follicle associated epithelium (PFPAE). In the crypts adjacent to the PFPAE, crypt stem cells are induced by unknown factors to give rise to at least three or more distinct phenotypic subsets: the common follicle associated epithelial cell, occasional goblet cells, and M cells.5 The common follicle associ-
ated epithelial cell resembles the intestinal enterocyte by morphology (eg, tight junction, brush border microvilli), but recent analysis of gene expression profiling data\textsuperscript{6–11} reveal that these cells show a distinct pattern of gene expression, including expression of unusual extracellular matrix and extracellular matrix-interacting proteins. PPFAE have also been shown to be constitutively positive for the NF-κB gene reB,\textsuperscript{12} which suggests that these cells have persistent activation of NF-κB signaling, as previously described for dendritic cells which are also reB-positive.\textsuperscript{13} This may be through TNF/lymphotoxin signals provided by follicular lymphocytes; these factors have been implicated in differentiation of secondary and tertiary lymphoid tissue, relying on the alternative NF-κB pathway.\textsuperscript{14} Moreover, it has been reported that lymphotoxin signaling may be responsible for inducing expression of CCL20 in PPFAE.\textsuperscript{15,16} In this context, Katkai et al\textsuperscript{18} showed that stromal cell lines would initiate stromal cell like differentiation in the presence of TNFα or LTα, and that even more rapid differentiation would occur in the presence of both TNFα and LTβR agonist. Accordingly, we found that treatment of intestinal epithelial cell lines (Caco-2BBbe and IEC-6) with LTβR agonist and TNFα induced expression of both PPFAE and M cell-associated genes.\textsuperscript{17}

Specification of M cell development is likely to be even more complex than for common PPFAE. It has been suggested that M cells are generated from specialized crypt cell precursors,\textsuperscript{18} but differentiation also requires an association with B lymphocytes that become embedded within a basolateral pocket in the M cell.\textsuperscript{19–21} So whether or not M cell precursors are predestined to their phenotype, additional signals from the association with B cells appear to complete the sequence toward final M cell development. What signals are involved remain unclear; an elegant study by Chervonsky and colleagues\textsuperscript{22} have shown that while B cells are required for M cell development, lymphotoxin signals do not have to be directly provided by them, so other, as yet unidentified, B cell factors may be involved. The failure in M cell development in CCR6-deficient mice\textsuperscript{23,24} also suggest that CCR6-expressing cells, likely recruited by PPFAE expression of CCL20, are important though it remains unclear whether the critical cells are B cells or myeloid cells.

Considering the likely role of lymphotoxin/TNF and NF-κB signaling on PPFAE development, we examined the effect of lymphotoxin/TNF treatment on intestinal epithelial cells. The colon carcinoma cell line Caco-2 has been used in co-culture studies as a model for M cell differentiation.\textsuperscript{19,25,26} Though in these experiments, the role of lymphotoxin/TNF signaling on gene expression has only recently been studied.\textsuperscript{19} Thus, we studied cytokine effects on Caco-2BBbe (a subclone of Caco-2\textsuperscript{27}) and the rat IEC-6\textsuperscript{28} cell lines for the induction of PPFAE and M cell-associated gene expression.\textsuperscript{17} Our data showed up to 60-fold induction of transcription of the TNF receptor superfamily member CD137/4-1BB (TNFRSF9).\textsuperscript{17} This induction by the combination of lymphotoxin and TNF was higher than the induction by either cytokine stimulus alone, suggesting at least an additive effect. We also were able to confirm the report of Rumbo et al,\textsuperscript{15} that under specific conditions the chemokine CCL20, strongly associated with PPFAE, could be induced by LTβR agonist antibody and/or TNFα. Most important, the treatment with the combination of TNFα and LTβR agonist showed the strongest induction of CD137 transcripts that persisted as long as 72 hours.\textsuperscript{17} This persistent induction of specific genes by combinations of lymphotoxin/TNF ligands may be an important discriminator between their acute proinflammatory effects versus the chronic induction of secondary lymphoid stromal cell development.

Here, we report that the CD137 (4-1BB) protein is also expressed by intestinal epithelial cells in response to TNF/lymphotoxin signaling. In CD137-deficient mice, while the development of mucosal lymphoid tissue appears normal,\textsuperscript{29,30} M cell functional differentiation is strikingly abnormal, with defects in particle transcytosis by NALT epithelium. Our studies suggest that the development of M cells in these mice fails at a relatively late stage dependent on the association of immature M cells and basolateral pocket B lymphocytes. This model may provide us with a useful model to study M cell differentiation in which early and late developmental stages may be separated by the expression of and signaling through CD137.

**Materials and Methods**

**Cell Lines**

Caco-2BBbe (ATCC no. CRL-2102) colon carcinoma cells were grown in advanced Dulbecco’s modified Eagle’s medium (Gibco) with the addition fetal bovine serum to a final concentration of 10%, penicillin-streptomycin-glutamine, and 15 mmol/L HEPES buffer. T84 (ATCC no. CCL-248) colon carcinoma cell lines were grown in a 1:1 mixture of advanced Dulbecco’s modified Eagle’s medium and F12 nutrient mixture HAM (Gibco) with fetal bovine serum (5%), penicillin-streptomycin-glutamine, and 15 mmol/L HEPES. Cell treatments included addition of goat anti-LTβR agonist antibody at 5 μg/ml (R&D Systems), TNFα at 100 ng/ml (PeproTech, Inc.) or both. Cytokine response studies indicated optimal LTβR agonist effects (induction of gene expression\textsuperscript{17}) at doses at or above 2.5 μg/ml, and TNFα responses at or above 50 ng/ml.

**Cultures on Transwells**

Cells were cultured on Transwells containing polystyrene supports with a pore size of 0.4 μm (Corning) at 10–20,000 cells per well, in a final volume of 600 μl. Cytokine treatments were done using the concentrations described above. Filters were removed and fixed with PBS/4% paraformaldehyde for 10 minutes and then rehydrated with PBS for 10 minutes. Casein/PBS (Fisher) with a final concentration of 0.1% Triton X was then used for blocking and for dilutions of antibodies. TO-PRO-3 or DAPI was used as a nuclear stain. The slides were post fixed for 5 minutes with 4% paraformaldehyde, and mounted using ProLong Gold (Molecular Probes). Cells were analyzed using a BD CARV II spinning disk confocal
microscope, using iPLab software. Image deconvolution and analysis was performed using Volocity software (PerkinElmer).

Animals

CD137-deficient mice on the BALB/c background were provided by Dr. B.S. Kwon and bred in the UC Riverside vivarium under SPF conditions. In each experiment, wild-type BALB/c age-matched mice were used as controls for CD137-deficient mice. Mice were handled in accordance with institutional IACUC and NIH guidelines.

Bone Marrow Irradiation Chimeric Mice

Recipient mice were irradiated on day 1 with 1100 rads and were injected with $6 \times 10^6$ purified bone marrow cells within 20 hours after irradiation. All bone marrow recipient mice received two additional intravenous injections of $40 \times 10^6$ purified lymphocytes per injection from PP, LN, and spleen on days 6 and 13 (in proportions of approximately 45% to 60% spleen cells, 35% to 50% LN, and 1% to 5% PP). All bone marrow chimera mice were studied for particle uptake and tissue histology on day 19.

Electron Microscopy

For Scanning EM, tissues were dissected and fixed in 4% paraformaldehyde/1% glutaraldehyde (Electron Microscopy Sciences) and then prepared by critical point drying and gold sputter coating. Samples were examined with a FEI XL30 field emission scanning electron microscopy (SEM) at 5 kV. For transmission electron microscopy (TEM), specimens were fixed in 2.5% glutaraldehyde in 0.1 mol/L sodium phosphate, pH 7.2–7.4, on ice. Tissues were then postfixed with 1% OsO$_4$ in 0.1 mol/L PBS. After gradient ethanol dehydration, samples were infiltrated with Spurr’s resin. Ultrathin slices were mounted on Formvar-coated grids and stained with uranyl acetate followed by lead citrate. After air drying, grids were analyzed on an FEI Tecnai-12.

Whole Tissue Confocal Microscopy and Microparticle Uptake Studies

Dissected NALT or PP tissue was fixed in 1 to 4% paraformaldehyde/30% sucrose/PBS, and flash-frozen for cryostat sections, which were then stained according to the protocol described for cell cultures. For whole tissue mounts, dissected tissues were fixed in 1% paraformaldehyde/30% sucrose/PBS and then stained in whole tissue fragments. Tissues were mounted in wells built from reinforcement rings on microscope slides for confocal microscopy studies. For microparticle uptake studies, 0.2-micron 505/515 (yellow-green) Neutravidin Fluospheres (Molecular Probes/Invitrogen) were used uncoated (ie, without biotinylated peptide), coated with biotinylated P13 scramble sequence peptide (ie, a Clostridium perfringens enterotoxin (CPE) 30-related sequence that was scrambled), or coated with biotinylated CPE30 peptide. Under anesthesia, mice were given $6 \times 10^8$ beads in 40 $\mu$m of PBS intranasally (20 $\mu$m/nostril), and after 10 minutes, the NALT was dissected for microscopy. For latex bead (Fluoresbrite YG 0.2-micron microspheres; Polysciences) uptake studies in the NALT, mice were given $6 \times 10^8$ beads in 40 $\mu$m of PBS intranasally (20 microliters/nostril) approximately 1–5 minutes before sacrificing mice for dissection. Tissues were analyzed using a BD CARV II spinning disc confocal microscope, with Volocity software (PerkinElmer) for deconvolution and image analysis. Bead uptake was measured from z-projection images from UEA-1-stained NALT taking 40 $\times$ 40 $\mu$m areas where UEA-1-positive cells were evident. Each experimental group consisted of at least three individual mice (two NALT per mouse); three non-overlapping confocal z-stacks were taken from each NALT, and three to five non-overlapping 40 $\times$ 40 $\mu$m image “stamps” were counted from each NALT image. Bead counts were performed using Volocity software. M cell density counts were performed in similar manner. Statistical analysis of bead counts and M cell density were performed by Student’s t-test and also by Mann-Whitney (Prism, GraphPad Software), with similar results.

Results

TNF/Lymphotoxin Signals and Induction of CD137 in Intestinal Epithelium

To follow on our observation that CD137 (TNFRSF9/4-1BB) gene transcripts were induced in Caco-2BBe cells by LTβR agonist antibody plus TNFα, we confirmed that this cytokine treatment also induced expression of CD137 protein as revealed by staining of cytokine treated Caco-2BBe cells both in chamber slides and Transwell cultures (Figure 1). Surprisingly, the CD137 was present in a vesicular distribution within the cytoplasm, and flow

Figure 1. Induction of CD137 expression in cytokine treated Caco-2BBe cells. Caco-2BBe cells were cultured alone or with a LTβR agonist antibody plus TNFα and stained for CD137 (green) and fibronectin (red). The inset shows co-localization of cytoplasmic CD137 with fibronectin (arrows).
cytometry analysis could not detect significant levels of cell surface antigen (not shown). The cytoplasmic vesicles with CD137 did not co-localize with the tight junction protein Claudin 4, a protein that we previously reported to be associated with particle endocytosis in cytokine-treated cells. However, we did find that most of the cytoplasmic CD137 co-localized with the extracellular matrix protein fibronectin (Figure 1), though it was not associated with fibronectin at the surface of the cells. The cytoplasmic associations may also account for the apparent decrease in cell-associated fibronectin in the cytokine treated cells; this pattern suggests that fibronectin synthesized by the epithelial cells was endocytosed in association with CD137 either during synthesis or soon after both proteins appeared at the surface of the cells. Interestingly, while CD137 has generally been considered in the context of its expression on lymphocytes and interaction with CD137L expressed on B cells and myeloid cells, it may be more important in the connection of cells to the basement membrane during cytokine-dependent changes, including M cell differentiation. Whether this interaction also induces signaling through CD137 is not clear.

We attempted histological staining to detect CD137 expression in mouse PPFAE in vivo. While the antibody against human CD137 was able to readily detect the induced protein in the Caco-2BBe cells (Figure 1), with the anti-mouse CD137 antibody we were unable to detect expression in any mouse tissues (not shown). While available antibodies against mouse CD137 could detect the protein on T cells by flow cytometry, they were not sensitive enough to detect CD137 on either T cells or M cells in cryostat sections. Given the much lower level induction in the rat IEC-6 cell line as compared with the human Caco-2BBe line, it is possible that CD137 expression on mouse mucosal epithelium in vivo is too low to detect or the available reagents are not sensitive enough. Indeed, we have found no published studies showing detection of mouse CD137 by staining mouse tissue sections.

Mucosal Lymphoid Tissues in CD137 Knockout Mice

Electron Microscopy Studies

M cells are most easily found in mucosal lymphoid tissues such as the NALT and PPFAE. Therefore, to assess the importance of CD137 on mucosal M cell development, we examined NALT and PP from mice with a CD137 gene knockout, which had been backcrossed to the BALB/c background. In previous reports, CD137 knockout mice were not noted to have any unusual developmental defects in any lymphoid tissue, and lymphoid cell populations were noted to be relatively normal. Immune response deficiencies in cytolytic T cells, NK, and NKT cells have been reported, and there have also been reports of alterations in T lymphocyte homeostasis and regulation, including enhanced CD4 T cell responses in vitro, but many aspects of immune responses including antibody production have been reported to be normal, depending on the stimulus. Yet while CD137-deficient mice appear to generate good T and B cell responses to systemic or intraperitoneal immunization, specific studies on mucosal immune function have not been reported. In PPFAE, SEM reveals that most of the epithelial cells have a well defined brush border with tightly packed microvilli uniformly extending less than a micron from the apical surface (Figure 2A). Here, M cells lacking brush border microvilli can be identified across the surface of PP follicle epithelium as gaps in the brush border. The apical surface PP M cells have small projections from the apical cilia (asterisk), but with basolateral membranes (arrows) extending around basolateral pocket lymphocyte (L). TEM of CD137-deficient NALT, showing narrow cross-section and absence of apical cilia (asterisk) of M cells (M). Most M cells lack basolateral lymphocytes; the M cell at upper left may be associated with a lymphocyte, although membrane extensions contacting the lymphocyte are not evident.

Figure 2. Electron microscopy analysis of PP and NALT. PP and NALT from wild-type (BALB/c) and CD137-deficient mice were prepared for SEM and TEM to assess the morphology of M cells in mucosal lymphoid tissues. A: SEM of wild-type PP M cells. B: SEM of CD137-deficient PP M cells, showing abnormally narrowed apical surfaces of M cells (arrows). C: SEM of wild-type NALT. D: SEM of CD137-deficient NALT. E: TEM of wild-type NALT, showing M cell (M) lacking apical cilia (asterisk), but with basolateral membranes (arrows) extending around basolateral pocket lymphocyte (L). F: TEM of CD137-deficient NALT, showing narrow cross-section and absence of apical cilia (asterisk) of M cells (M). Most M cells lack basolateral lymphocytes; the M cell at upper left may be associated with a lymphocyte, although membrane extensions contacting the lymphocyte are not evident.
distinguished from the neighboring microvilli of FAE enterocytes (Figure 2A). By contrast, M cells were more difficult to identify in CD137-deficient PP. Presumed M cells (identified provisionally by the lack of brush border microvilli) did not have a clearly flattened polygonal apical surface and instead presented a much smaller apical surface that was also more narrow in shape (Figure 2B).

In the NALT, most of the epithelium displays a dense cover of cilia 2 to 3 \( \mu m \) long, so M cells from wild-type mice were identifiable as cells lacking cilia but having smaller projections from the apical surface (Figure 2C). M cells could be found across NALT from both control and CD137-deficient mice (Figure 2D), but due to the larger number of ciliated cells in the NALT epithelium, differences in the number of distribution of M cells were not easy to establish. Morphological differences were also difficult to establish, though many of the M cells appeared narrower than in wild-type controls. However, TEM of the NALT from wild-type and control mice showed some differences. Putative M cells in wild-type NALT showed an absence of apical cilia, and in most cases, the basolateral membrane appeared to be intimately associated with an adjacent lymphocyte (Figure 2E, arrows). In NALT from CD137-deficient mice, cells that lacked apical cilia were more difficult to find, with the few cells identified appearing narrow, with most lacking evidence for an associated basolateral lymphocyte (Figure 2F).

Confocal Studies of UEA-1 Lectin Binding

The distribution of M cells across the face of the PPFAE can also be visualized using whole mount staining of the follicles with the lectin UEA-1. Although goblet cells can also bind UEA-1, they are rare in the PPFAE and NALT epithelium, and they also present a distinctive goblet-shaped morphology easily distinguished from conventional M cells. In control PP, UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology (Figure 3B). UEA-1 staining of wild-type adjacent crypts (Figure 3A) showed pattern consistent with migration of M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of wild-type adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of wild-type adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of wild-type adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of wild-type adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of wild-type adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of CD137-deficient adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration). UEA-1 staining of wild-type adjacent crypts (asterisks and arrows as in C) showed a similar radial pattern, but with a clearly narrowed apical morphology. M cells from crypts (asterisk, with arrows indicating apparent course of migration).
Since M cells in the PPFAE are generated from crypt cell precursors alongside the other PPFAE, we can follow the progression of cells from the crypt of normal and CD137-deficient mice. The pattern of UEA-1-positive cells arrayed in lines extending from the marginal crypt indicates that a crypt cell progenitor has been committed to produce M cells (Figure 3C); the presence of similar radial lines of UEA-1-positive cells in CD137-deficient mice (Figure 3D) suggests that this commitment is normal despite the unusual morphology of cells generated. This phenotype is distinct from the situation in CCR6-deficient mice, in which UEA-1-positive cells were nearly entirely absent; thus, the CD137 knockout appears to affect a later stage in M cell development than the CCR6 knockout, after commitment of the crypt stem cell to the M cell lineage. Importantly, the absence of CD137 also did not appear to affect their survival in the PPFAE.

**Defective Formation of M cell/B Lymphocyte Complexes in CD137-Deficient Mice**

The induction of CD137 on intestinal epithelial cell lines was surprising, since it had previously been described mainly on bone marrow-derived cells such as activated T cells and myeloid cells, though other reports on CD137 expression detected it specifically on vascular endothelial cells in the lung associated with tumors, or associated with atherosclerosis. The main known ligand for CD137 is the molecule 4-1BBL expressed on B cells and a variety of myeloid cells. Since B cells are known to be important in the induction and maintenance of PPFAE M cells, it was possible that CD137 on developing M cells is involved in the crosstalk between M cells and B cells in the basolateral pocket.

We therefore assessed the morphology of M cells and associated cells in Peyer’s patch follicle epithelium. The narrow morphology of the UEA-1-positive M cells in CD137-deficient mice would appear to be inconsistent with the development of a basolateral pocket with B lymphocytes that is characteristic of both NALT and PP M cells. Staining for B cell specific markers CD19 and CD45/B220 revealed strong staining on follicular B cells but only very weak staining of the M cell basolateral pocket cells (Figure 3E, showing B220). By contrast, strong major histocompatibility complex (MHC) class II staining was consistent among B cells and putative dendritic cells in the subepithelial space. It is possible that the change in B220 and CD19 expression may be a phenotypic change in B cells activated to enter the basolateral pocket. Therefore, in cryostat sections of PP we stained for both UEA-1 to detect M cells and MHC class II antigens (I-A/d-E) to detect putative B cells as well as dendritic cells (Figure 3F). In wild-type BALB/c PP, MHC II-positive cells were found associated with most UEA-1-positive M cells, located in a central-basolateral pocket. The small rounded morphology and fine ring staining for MHC II (arrows) of the basolateral pocket cells was most consistent with their putative identity as B lymphocytes. However, with CD137-deficient mice, while MHC II-positive cells could be found in the follicle epithelium, few if any of these cells were clearly contained within any identifiable M cell basolateral pocket, and their morphology was more irregular (Figure 3G). Thus, it appears that in the absence of CD137, MHC class II-positive cells, whether B cells or dendritic cells, were less efficient in forming stable basolateral pockets or other complexes with UEA-1-positive M cells.

**NALT M Cells**

The mouse NALT epithelium is not generated from crypt cell precursors as in the intestine, so there is no radial pattern of UEA-1-positive M cells across the epithelium. Instead, the NALT epithelium of control mice shows a distribution of UEA-1-positive M cells across the NALT. Three-dimensional renderings from confocal z-series showed that these cells were large and more rounded in morphology (Figure 4A). By contrast, the UEA-1-positive M cells in CD137-deficient mice were more irregular in morphology (Figure 4B).

Using confocal z-series of UEA-1 staining cells, we could also construct three-dimensional surface renderings of these cells to assess their morphology. Thus, cells from wild-type NALT showed a roughly cuboid shape, with many displaying a clear basolateral indentation presumably to contain associated lymphocytes (Figure 4C).
By contrast, UEA-1-positive cells from the CD137 knock-out mice (Figure 4D) showed mainly narrow shaped cells, with no evidence for a basolateral indentation among these cells, consistent with the TEM studies (Figure 2F).

**M Cell Particle Uptake in NALT and PP**

M cells, in their role in mucosal immune surveillance, have a specialized capacity for uptake and transcytosis of antigens and microparticles. To test whether the CD137 deficiency had an effect on M cell particle uptake, we relied on the observation that M cells show regulation and redistribution of the tight junction claudin 4, suggesting of a role for claudin 4 in microparticle uptake. Thus, we took streptavidin-conjugated fluorescent microbeads (0.2 micron diameter) and used them uncoated, or coated them with a control biotinylated peptide, or with a biotinylated 30 amino acid C-terminal peptide from CPE known to bind to an external domain of claudin 4. We found that NALT M cells were particularly active in microparticle transcytosis; while intestinal PP M cells could be shown to take up particles after hours of lumenal instillation, NALT follicles were found to accumulate microparticles within minutes of administration of a suspension of targeted (CPE peptide-coated) fluorescent styrene microbeads. This transcytosis appears to be specifically dependent on M cells; in very brief exposure (<5 minutes) to targeted microparticle suspensions, particles found in the wild-type NALT epithelial layer were present within UEA-1+ M cells but many particles were already evident within the lymphoid follicle below. For more consistent quantitation of particle uptake, bead (both CPE-coated and control peptide-coated) suspensions were instilled into the nasal passages for ten minutes before dissection of the NALT for microscopic analysis of whole mounted tissue (Figure 5, A and B). Control peptide-coated microbeads were found only sparsely taken up in the NALT follicles of wild-type mice. By contrast, large numbers of beads were found taken into the NALT when the beads were coated with the claudin 4-binding peptide (Figure 5C).

In CD137-deficient mice, the uptake of CPE peptide-coated styrene microparticles was far less effective (Figure 5C). Minimal particle uptake was evident using beads coated with control peptide, but more importantly the uptake of beads with claudin 4-binding peptides remained at background levels, and significantly below the uptake by wild-type NALT (P < 0.0001; Figure 5C). This suggested that stromal cells such as mucosal epithelial cells can also be induced to express CD137. Thus, despite the presence of UEA-1-positive epithelial cells in the CD137-deficient NALT, microparticle uptake was minimal, suggesting that cells committed to M cell differentiation (evidenced by UEA-1 binding) failed to acquire the capacity to transcytose particles.

Mucosal M cell particle uptake is commonly assessed using latex microbeads; in contrast to protein-coated styrene beads, the apparent low surface charge of these uncoated synthetic particles enables uptake without the need for M cell targeting ligands (unpublished observations). Thus, we also administered suspensions of 0.2 micron diameter fluorescent latex beads and looked for uptake into the NALT. Here too, we found that uptake of the latex beads in wild-type NALT was significantly greater than uptake by CD137-deficient NALT (Figure 5D).

**CD137-Deficient Radiation-Resistant Stroma Supports M Cell Lineage Commitment but Not Function**

CD137 is known to be expressed by bone marrow-derived cells such as T lymphocytes, and our studies have suggested that stromal cells such as mucosal epithelial cells can also be induced to express CD137. Thus, M cell development may require CD137 on bone marrow-derived cells, mucosal epithelium, or both. To test this question directly, we made radiation chimeras in which wild-type and CD137-deficient mice were given lethal irradiation, and reconstituted with multiple injections of bone marrow and lymphoid cell populations (spleen, lymph nodes, Peyer’s patch) from wild-type or CD137-deficient donors (“wild-type→wild-type” and “wild-type→CD137KO” chimeras, and “CD137KO→wild-type” and “CD137KO→CD137KO” chimeras). Thus, in these mice the major difference is the CD137 genotype.
of the bone marrow-derived cells such as lymphocytes and dendritic cells, versus the radioresistant stromal cells, including mucosal epithelium. If CD137 was only required on bone marrow-derived cells (e.g., activated T cells and dendritic cells) then only wild-type wild-type and wild-type CD137KO chimeras would be expected to have normal M cell development and function. If CD137 is only required on stromal cells such as mucosal epithelium, then only wild-type—wild-type and CD137KO—wild-type would have normal M cells. A third possibility is that CD137 has two roles, in bone marrow-derived cells and mucosal epithelium; here, only the wild-type—wild-type chimeras would have normal M cells.

NALT epithelium in all types of chimeras showed similar development of UEA-1/H11001 M cells (Figure 6, A–D), despite the fact that non-irradiated CD137-deficient NALT showed more irregular M cells (Figure 3); this may be an effect of the irradiation on the epithelium. Fortunately, when tested for latex microparticle uptake (Figure 6E), the wild-type—wild-type chimeras still showed normal particle uptake, indicating that M cell function was present even after irradiation. By contrast, the wild-type—CD137KO, CD137KO—wild-type, and CD137KO—CD137KO chimeras all showed very poor to nearly absent microparticle uptake, similar to both unmanipulated CD137-deficient mice and RAG-1-deficient mice (not shown). The difference in the uptake of particle in the NALT of wild-type—wild-type versus wild-type—CD137KO and other chimeras was highly significant (P < 0.0001). Thus, CD137-deficient bone marrow-derived cells failed to support M cell function even in wild-type mucosal epithelium. CD137-deficient mice have been reported to have some T cell functional defects due to a lack of CD137 signaling, so lymphocyte defects could be responsible in part for the M cell abnormalities. Lymphocyte expression of CD137 is associated with T cell costimulation, so it may be that the connection to M cells is through CD4 T cell interactions with B lymphocytes, which are known to be critical to M cell development. Interestingly, however, CD137 expression was also required in the radioresistant stromal cells, as wild-type bone marrow failed to induce normal M cell function in the CD137-deficient recipients. This would be consistent with a second role for CD137 in the interactions between B cells and M cells.

As noted in the unmanipulated CD137-deficient mice, both NALT and PP epithelium showed the development of UEA-1+ M cells (Figure 6, A–D), but had abnormal morphology. In the irradiation chimeras, we also saw UEA-1+ presumptive cells in the NALT epithelium. The morphology of the UEA-1+ cells was not strikingly different among the different types of chimeras (Figure 6, A–D); however, the radiation did seem to cause more variable morphology among all of the UEA-1+ cells. Similarly, intestinal Peyer’s patches were small and more difficult to examine, likely due to the residual effects of the lethal irradiation. When we counted the distribution of M cells across the NALT epithelium, all chimeras showed a similar density of M cells per surface area of NALT (not significantly different by t-test; Figure 6F). Together, these results are consistent with the notion that CD137 is not required for M cell lineage commitment in mucosal epithelium, but it remains critical for the development of M cell transcytosis function.
M cells have been historically difficult to study in detail due to the difficulty in isolating sufficient numbers of cells. Reports on co-cultures of B lymphocytes and Caco-2 cells suggested that this cell culture model could help provide more material for study, but this approach has not been widely used, perhaps due to variations among specific cell line clones available. Moreover, as Kerneis et al. and Kanaya et al. have reported, the induction of M cell-like function in vitro is fairly short-lived (only a few days), and from those studies it is generally not clear what proportion of the cultured cells actually developed M cell-like function. This perhaps also illustrates the main present limitation in M cell studies: a clear molecular definition of the M cell phenotype is only now beginning to be formulated by the collected studies of several groups. Despite these issues, we were able to use the Caco-2BBe and IEC-6 cell lines to assess their response to cytokine induction, and identified genes relating to M cell-like function. This may also provide some additional clues to the stages in M cell differentiation in mucosal lymphoid tissues. Although lymphotixin/TNF ligands are known to be important in the development of secondary and tertiary lymphoid tissues, these cytokines are apparently not by themselves sufficient to induce full maturation and function of M cells in vivo; the physical association with B cells (or subepithelial dendritic cells) may provide additional important signals. A report suggests that signaling through intestinal epithelium expression of the TNF receptor superfamily member RANK (TNFRSF11a) is also important in M cell differentiation, like through interaction with the ligand RANKL (TNFSF11) on stromal cells. Our results suggest that CD137/CD137L signaling between M cell progenitors and basolateral pocket B cells may provide at least one of these important signals that induce final functional maturation of developing M cells.

In the radiation chimera studies, results suggest that M cell function is dependent on CD137 expression by both bone marrow-derived cells and radioresistant stromal cells (presumably epithelial precursors of M cells). The nature of any lymphocyte-dependent defect (if present) is not yet clear, and lymphoid tissue inducer cells (LTi) are probably normal; the lymphoid tissues of CD137-deficient mice showed overall relatively normal structure and organization, and the development of UEA-1-expressing M cell lineage appeared unimpaired despite their abnormal morphology. The CD137-dependent lymphoid cell contributions to M cell function may be limited to induction of function after lineage commitment. For example, it may be possible that CD137- CD4 T cells are involved in inducing B cells to become basolateral pocket lymphocytes in mucosal lymphoid tissues; the basolateral pocket B cells in turn may be necessary to induce M cell transcytosis function. In preliminary studies, we found that CD137L triggering increased CCR6 expression in the RajiB cell line (not shown). CCR6 is important in the homing of lymphocytes toward Peyer’s patch follicle epithelium expressing the ligand CCL20,9,15,17 and CCR6- deficient mice also have been reported to have M cell defects.23,24 so effects on CD137-induced CCR6 may partly affect M cell development in the CD137-deficient mice. The requirement for CD137 in radioresistant stroma also suggests a second step in which the recruited CCR6+ CD137L+ basolateral pocket B cells interact with CD137 on M cells.

Our histological studies, by both TEM and confocal microscopy, suggested that the CD137-deficient M cells in the NALT and Peyer’s patch present a smaller luminal surface than normal M cells. The particle uptake defect in these cells could be attributed at least in part to a simple reduction in this apical surface area. However, these morphological differences may be insufficient to explain the functional defect, especially in the radiation chimeras. It may be simpler to suggest that a primary functional defect and failure to form a significant basolateral pocket may also be secondarily associated with the reduced apical surface.

In view of our results, a rough outline of a model might be developed as a working model for M cell differentiation from crypt cell precursors: First, cytokines from lymphoid tissue inducer cells or other follicle associated cells induce nearby crypt cells to convert to M cell progenitors instead of conventional enterocytes. At this point, the committed progenitors produce the fucose moiety that binds to UEA-1 lectin, and they would also begin to express CD137. Second, a CCR6+ B lymphocyte (perhaps induced by a CD137+ CD4 T cell) migrating to the CCL20+ follicle epithelium encounters the CD137 expressing M cell progenitor, and cross-signaling by CD137/CD137L establishes a bond between the M cell and B cell. Third, the presence of the B cell induces cytoskeletal changes in the M cell, including formation of the basolateral pocket, and induction of particle transcytosis activity by the M cell. The B cells in the basolateral pocket might only be for the induction and maintenance of M cell function, since transcytosed antigens and particles may be more dependent on underlying dendritic cells for presentation to T cells. Thus, basolateral pocket B cells may be a supplementary source of cytokine ligands including lymphotixin, TNFa/LT, and CD137L, and together these may be responsible for the maintenance of the M cell functional phenotype. Additionally, the geometry of the basolateral pocket may provide directional clues for transcytosis of particles to the basolateral side of the M cells. CD137 interactions with extracellular matrix proteins may also be important for stabilizing the M cell contact with the basement membrane, especially since much of the basolateral surface is lost to forming the B cell pocket. It is interesting to note that the failure of UEA-1+ cells to associate with B cells in CD137-deficient mice does not appear to cause the death of these cells, as they persist across the face of the follicle epithelium dome; thus, B cells do not seem to be needed for any kind of survival signal. The specific effects of CD137-mediated signaling in M cell development is not clear; signaling pathways in CD137+ T cells might be quite different from those mucosal epithelium. However, recent studies have suggested that other non-hemopoietic cells such as vascular endothelium in vivo and
HUVEC in vitro can be induced to express CD137 by TNFα, and direct triggering of CD137 on HUVEC can induce expression of ICAM-1 and VCAM-1. Similar CD137-induced expression of such adhesion molecules by M cells would help reinforce interactions between M cells and their basolateral pocket B cells expressing receptors such as LFA-1 and VLA-4.

With the specific defect in function by UEA-1-positive cells in CD137-deficient mice, we now have an in vivo system for studying M cell differentiation in which crypt cells or NALT M cell precursors are able to receive the initial signals that give rise to "immature" M cells (identified by UEA-1 binding), but differentiation fails to proceed past that point. The defect in CD137-deficient M cells also provides us with a tool for dissecting the role of M cell particle uptake in mucosal immunity.

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