

Fig. 1. Unique trafficking pathways for IELs in the intestinal compartments. In the antigen-primed IEL trafficking pathway, DCs in PPs take up antigens transported through M cells and present them to T cells in an MHC-dependent manner. Simultaneously, DCs produce retinoic acid, which causes antigen-primed T cells to express $\alpha 4\beta 7$ integrin and CCR9. $\alpha 4\beta 7$ integrin interacts with intestinal high endothelial venules (HEV) expressing MAdCAM-1, while CCR9 allows specific migration to the CCL25, a chemokine produced by intestinal ECs.

In the other pathway, RTEs migrate into the intestinal epithelium without antigen priming at PPs and MLNs. Alternatively, RTEs have already expressed $\alpha 4\beta 7$ integrin, CCR9, and αE integrin in the thymus. The other population of IELs may originate from the CPs, but their trafficking pathway has not been fully elucidated. Upon migration into the intestinal epithelium, IELs express several unique molecules capable of tethering ECs (e.g. αE integrin, Ep-CAM, and occludin) or of recognizing stressed or infected ECs (e.g. $\alpha \beta$ TCR, $\gamma \delta$ TCR, NKG2D, and CD8).

results obtained by the Study (32), one can surmise that almost half of the naive IELs in the small intestine are derived from RTEs, while PPs and MLNs contain a naive population of RTEs with much less frequency (less than 3%). With their diverse TCR repertoires, RTEs seem likely to play an important role in maintaining TCR diversity in the intestine. At present, however, it remains unknown whether or not RTEs are present in the large intestinal epithelium.

Differing compositions of naive and activated/memory IELs in the small and large intestine

Not surprisingly given the different environmental circumstances faced in the small and large intestine, IELs in the two environments differ with regard to the ratio between naive and activated/memory IELs. It is well established that small intestinal IELs are functionally characterized as antigen-experienced activated/memory cells expressing CD69 and that they contain few naive cells (less than 5%) (34–36). In contrast, the large intestine contains large numbers of naive-type cells

expressing CD62L^{high} and CD44^{int} (more than 40%) but a far less abundant supply of activated/memory cells (34–36).

A correlation may exist between the abundant presence of activated/memory IELs and of CD8 $\alpha\alpha$ IELs in the small intestine. As mentioned above, the small intestine contains a large number of type b IELs expressing CD8 $\alpha\alpha$ as well as type a IELs expressing CD8 $\alpha\alpha$ together with CD4 or CD8 $\alpha\beta$ (Table 1). CD8 $\alpha\alpha$, which can be expressed on activated conventional T cells of the lymph nodes and spleen, plays a key role in mediating the survival of those effector cells that further differentiate into mature memory T cells (37, 38). These surviving conventional T cells acquire the ability to migrate to and reside long-term in non-lymphoid tissues, including the gut, and they readily reinduce CD8 $\alpha\alpha$ expression after secondary restimulation. These observations indicate that the expression of CD8 $\alpha\alpha$ by the various subsets of mucosal T cells is not a marker to identify their origin or the specificity of their TCRs, but rather this expression reflects their effector/memory phenotype and plays a unique

role in maintaining immune homeostasis within the intestine.

Though both show a memory phenotype, type a and type b IELs represent distinct types of memory cells. Type a IELs resemble conventional memory T cells found in the periphery and lamina propria regions (39), and given that TCR β clonotypes were comparable for type a IELs and lamina propria T cells (39), it is possible that they differentiate via the same pathway. In contrast, CD8 $\alpha\alpha$ ($\alpha\beta$ IELs) are thought to be natural memory T cells. CD8 $\alpha\alpha$ $\alpha\beta$ TCR IELs show oligoclonal TCR repertoires that are different from those of type a IELs (40), indicating that CD8 $\alpha\alpha$ $\alpha\beta$ IELs differentiate via a distinct pathway. The accumulation of autoreactive TCRs in CD8 $\alpha\alpha$ $\alpha\beta$ IELs (41), along with their early appearance in neonatal mice (42) and human fetal intestine (43), when exposure to exogenous antigen is minimal, provides evidence for the self-specificity of these T cells. In addition, most human fetal IELs already express an effector/memory cell phenotype (43), indicating that a differentiation pathway other than exposure to peripheral exogenous antigen resulted in the acquisition by these cells of an activated memory-like phenotype. However, unlike the CD8 $\alpha\alpha$ $\gamma\delta$ TCR IELs ($\gamma\delta$ IELs), the $\alpha\beta$ IELs do not appear in the small intestine of germ-free mice lacking gut flora, although bacterial colonization of these animals restores the population of CD8 $\alpha\alpha$ $\alpha\beta$ IELs (44). These observations indicate that local stimulation from the luminal microflora is required for generation and maintenance of the CD8 $\alpha\alpha$ $\alpha\beta$ IELs.

Antigen recognition by IELs

Type a IELs are reactive to classical MHC-restricted antigen presentation

Similar to peripheral T cells, type a IELs express $\alpha\beta$ TCRs together with the coreceptor CD8 $\alpha\beta$, which recognizes MHC class I, or the coreceptor CD4, which recognizes MHC class II. In general, MHC class I presents peptide generated from cytoplasmic protein (endogenous antigen) (13), while antigens proteolytically processed after endocytosis/phagocytosis are presented by MHC class II molecules (45). Although both MHC class I and class II molecules express self- and non-self-antigen, type a IELs are specific for non-self-antigen because of the positive and negative selection in the thymus.

Antigen recognition by type b CD8 $\alpha\alpha$ $\alpha\beta$ IELs

The MHC restriction of type b CD8 $\alpha\alpha$ $\alpha\beta$ IELs is still enigmatic. CD8 $\alpha\alpha$ $\alpha\beta$ IELs are drastically reduced in number in β 2-microglobulin-deficient mice (46–49), but they are present in mice deficient either in the transporter associated with antigen

processing (TAP) (49) or in the classical class I molecules (K^b-/-D^b-/-) (50, 51). This finding suggests that type b IELs require a TAP-independent non-classical MHC class Ib molecule.

It has been reported that mice deficient in Qa-2, a non-classical MHC class I molecule that is TAP dependent (52), possessed fewer CD8 $\alpha\alpha$ $\alpha\beta$ IELs, suggesting that some CD8 $\alpha\alpha$ $\alpha\beta$ IELs are reactive to Qa-2 (53). However, when they compared BALB/c mice with the Qa-2-null Bailey substrain of BALB/c mice, Cheroutre's group (54) were unable to confirm these findings. Furthermore, Qa-2 is broadly expressed by the IELs themselves (18), and its expression is mostly TAP dependent. Two other non-classical class I molecules, the thymic leukemia (TL) antigen (55) and CD1d (56), have been reported to be expressed by mouse intestinal ECs, although this is controversial in the case of mouse CD1d (57). Both TL and CD1d are TAP independent, but they require β 2-microglobulin expression (58–60). However, the TCRs of this IEL subset in mice have never been shown to recognize either of these Ib molecules. Recent structural data indicate that the TL antigen has a very narrow and closed groove that makes interacting with $\alpha\beta$ TCRs unlikely (61).

Researchers using several MHC class-II-restricted TCR transgenic systems (62, 63) showed that CD8 $\alpha\alpha$ IELs were generated when agonist self-peptides were present, just as in the MHC class-I-restricted systems. Despite this finding, peripheral T cells in these same MHC class-II-restricted TCR transgenic mice were CD4⁺. These observations indicate that CD8 $\alpha\alpha$ does not function as a TCR coreceptor and that expression of CD8 $\alpha\alpha$ on the $\alpha\beta$ IELs does not necessarily imply MHC class I restriction. From these results, it can be surmised that CD8 $\alpha\alpha$ $\alpha\beta$ IELs, unlike type a IELs, are self-reactive but are not selected for self-reactivity to a single antigen and that they may include cells reactive to classical class I molecules, non-classical class I molecules, and MHC class II molecules.

Specificity of $\gamma\delta$ TCR IELs ($\gamma\delta$ IELs)

It is still uncertain to what extent the $\gamma\delta$ TCRs require a selection event for their development, especially in mice. Murine $\gamma\delta$ IELs make predominant use of the V γ 5 (also known as V γ 7) gene segment together with several V δ genes (64). Their CDR3 regions are more diverse than the invariant $\gamma\delta$ TCRs expressed in the skin and reproductive tracts (65). Although the great majority of these cells express CD8 $\alpha\alpha$, their numbers are not reduced in β 2-microglobulin-deficient mice (47, 48). In humans, the $\gamma\delta$ TCRs expressed by IELs predominantly use V δ 1 (66), and they have an oligoclonal CDR3 repertoire that can be sustained over a period of many months (67). These TCRs recognize MICA and

MICB, class-I-like molecules that are induced by heat shock or stress on ECs (26, 68, 69). Like TL antigen in mice, MIC molecules are not capable of presenting peptides or other ligands (61). As mentioned above, MICA is also recognized by NKG2D. Although mice do not have a functional MIC gene ortholog, they do have NKG2D receptors that recognize class-I-like molecules such as H60 (70) and members of the RAE class-I-like family (71). There are no reports, however, indicating that significant populations of $\gamma\delta$ IELs or other mouse $\gamma\delta$ T cells recognize these class-I-like molecules.

TL antigen and MICA for CD8 α IELs

Given the MHC class I or class II specificity of the TCRs of CD8 α $\alpha\beta$ IELs together with their requirement for β 2-microglobulin, an MHC class I-molecule-mediated function other than peptide presentation must be required for CD8 α $\alpha\beta$ IEL development and/or maintenance. In this regard, Leishman et al. (20) have shown that the mouse TAP-independent, β 2-microglobulin-dependent, non-classical MHC class I molecule TL is a unique ligand for CD8 α . TL is constitutively and abundantly expressed by the ECs of the small intestine and so is in close proximity to the CD8 α IELs. The structural analysis of the molecule indicates the absence of an obvious antigen-binding groove (61). The

specific interaction of TL with CD8 α on activated CD8 α $\alpha\beta$ IELs modifies the TCR-activation signals to allow for regulated immune responses that are compatible with the gut environment (20). Unlike CD8 $\alpha\beta$, CD8 α is not internalized with the TCR from the surface of activated T cells (72), and associated p56^{lck} is inefficiently provided to the TCR complex, in part because of the inability of CD8 α to locate effectively into lipid rafts (73). It is, therefore, possible that the interaction of TL with CD8 α might sequester CD8-associated p56^{lck} away from the TCR complex, consequently reducing activation signals while promoting survival. In contrast to the abundant expression of TL in the small intestinal epithelium, TL was only weakly expressed in the large intestinal epithelium (74), which might be consistent with the small numbers of CD8 α IELs in the large intestine.

Our previous study constructed a transgenic mouse with gut-specific MICA expression driven by a T3b promoter, and we surprisingly observed a clonal expansion of CD4⁺ T cells also possessing CD8 α in the small intestine (75), suggesting that MICA positively regulates the expression of CD8 α (Fig. 2). Furthermore, studies in the inflammatory bowel disease model showed that transgenic MICA was able to attenuate the acute colitis induced by dextran sodium sulfate administration

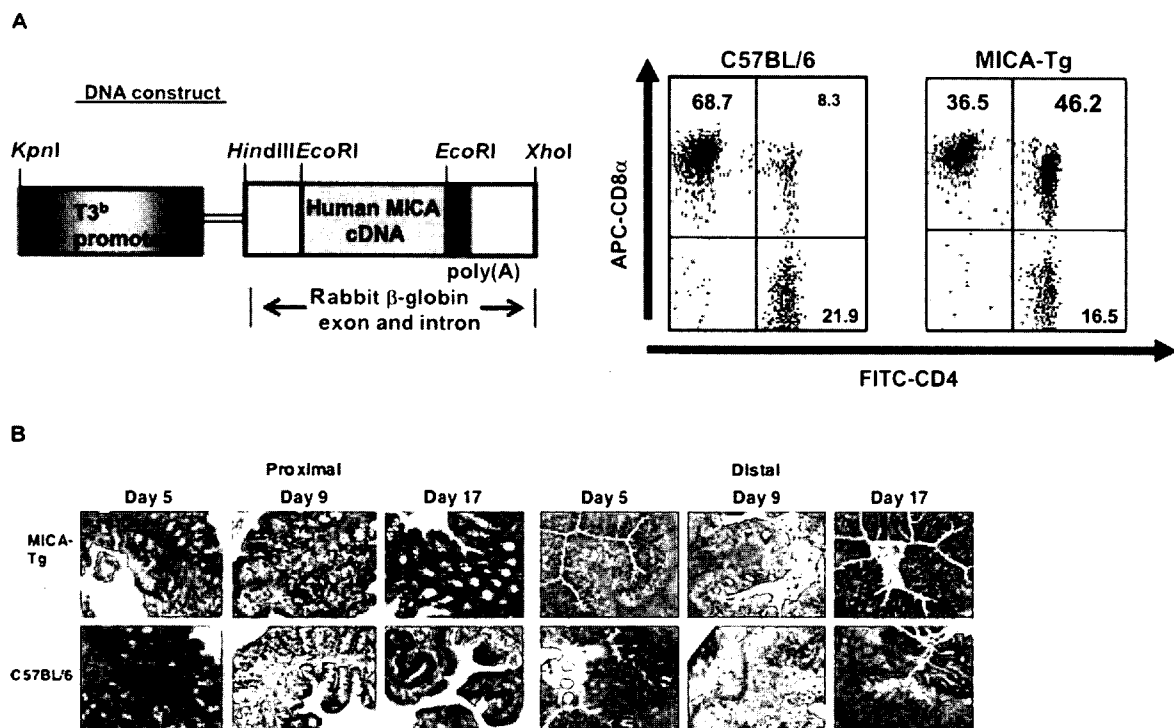


Fig. 2. CD8 α expression by mucosal T cells plays a unique role in maintaining immune homeostasis within the intestine. A transgenic mouse with gut-specific MICA expression driven by a T3b promoter was constructed. Surprisingly, a clonal expansion of CD4⁺ T cells also possessing CD8 α was observed in the small

intestine, suggesting that MICA positively regulates the expression of CD8 α (A). Furthermore, studies in the inflammatory bowel disease model show that transgenic MICA was able to attenuate the acute colitis induced by dextran sodium sulfate administration (B).

(Fig. 2). In this regard, it should be noted that MICA and TL, both of which belong to MHC class Ib molecules, are preferentially expressed in the intestinal ECs (55, 76). Thus, although MICA is both β 2-microglobulin and TAP independent, while TL is TAP independent but β 2-microglobulin dependent, it is likely that both molecules play a similar role in the development of IELs. Several years ago, Madakamutil et al. (38) demonstrated that TCR ligation and activation of CD8 $\alpha\alpha$ IELs by TL resulted in increased expression of the anti-apoptotic proteins Bcl-2 and Bcl-xL, suggesting that TL may abrogate antigen-induced cell death.

Unique signal transduction through TCR and Fc ϵ RI γ in type b IELs

Type b IELs in both the small and large intestine exclusively express homodimers of Fc ϵ RI γ or heterodimers of Fc ϵ RI γ with CD3 ζ (77–79), while type a IELs and peripheral T cells predominantly express homodimers of CD3 ζ , which can form a complex with TCR that is important for signal transduction (77). Thus, most T cells from CD3 ζ -deficient mice showed abnormal development (78, 80). However, type b IELs could express TCR in the CD3 ζ -deficient mice because Fc ϵ RI γ compensated for the CD3 ζ function in the TCR–CD3 complex (78–80). Because CD8 $\alpha\alpha$ molecules are not able to transfer signaling through TCR (73), selective expression of Fc ϵ RI γ on type b IELs might act as an alternative pathway for type b IEL activation.

Uniqueness of IEL developmental pathways

Agonist selection in CD8 $\alpha\alpha$ IELs

Although controversy remains as to what extent the $\gamma\delta$ IELs require thymic selection (42, 81), much evidence exists in mice indicating that self-specific CD8 $\alpha\alpha$ $\alpha\beta$ IELs are actively and preferentially selected in the thymus under agonist selection conditions (63, 82). The outcome of thymic selection of conventional $\alpha\beta$ T cells is determined by the signal strength received through the TCR when interacting with MHC molecules presenting self-antigens. Intermediate TCR signal strength results in positive selection of conventional $\alpha\beta$ T cells, whereas strong TCR-mediated signals delete cells that express those TCR from the conventional T-cell repertoire. By contrast, CD8 $\alpha\alpha$ $\alpha\beta$ IELs preferentially accumulate under conditions that lead to the deletion of conventional $\alpha\beta$ T cells, and, as a consequence, autoreactive TCRs are enriched in the TCR repertoire of CD8 $\alpha\alpha$ $\alpha\beta$ IELs (41). The existence of an agonist-dependent selection of CD8 $\alpha\alpha$ $\alpha\beta$ IELs has been shown using mice that express a transgenic TCR and cognate antigen during thymic selection. The H-Y TCR is specific for a Y-chromosome-encoded peptide presented by H-2D^b, and H-Y TCR transgenic

mice provide a model for the self-based agonist selection of CD8 $\alpha\alpha$ $\alpha\beta$ IELs (63, 83). When the H-Y TCR transgenic mice were crossed onto a recombination-activating gene (*Rag*)^{-/-} background, CD8 $\alpha\beta$ thymocytes matured in the thymus and populated the peripheral tissues of H-Y antigen-negative female mice, whereas few CD8 $\alpha\beta$ IELs and no CD8 $\alpha\alpha$ IELs were detected in these female mice. By contrast, despite deletion of conventional CD8 $\alpha\beta$ H-Y TCR⁺ T cells in *Rag*^{-/-} male H-Y TCR transgenic mice, large numbers of H-Y TCR⁺ IELs largely expressing CD8 $\alpha\alpha$ developed (83).

Similar observations have been made in TCR transgenic mice that coexpress the endogenous cognate antigen, such as 2C TCR transgenic mice expressing the H-2L^d alloantigen recognized by the 2C TCR (84) and double-transgenic mice expressing both the TCR and the cognate antigen. Examples of the latter include OT-1 TCR transgenic/RIPmOVA mice [membrane-bound ovalbumin (OVA) antigen transgene under the control of the rat insulin promoter (RIP)] (63), influenza virus nucleoprotein (NP)-specific F5 TCR transgenic mice that express NP peptides (85), and P14 lymphocytic choriomeningitis virus (LCMV)-specific TCR transgenic mice that coexpress the cognate LCMV-specific epitope GP33-41 (86).

The selection of CD8 $\alpha\alpha$ $\alpha\beta$ IELs is most effective in the presence of high-affinity interactions between the TCR and the agonist self-antigens, as was shown using the clonotypic F5 TCR, which has a high affinity for the influenza virus epitope NP68. NP68/F5 TCR double-transgenic mice preferentially generated CD8 $\alpha\alpha$ $\alpha\beta$ IELs, whereas double-transgenic F5 TCR mice coexpressing the antagonist peptide NP34 did not (85). Similarly, CD8 $\alpha\alpha$ IELs expressing the 2C TCR were readily generated in mice expressing the H-2L^d high-affinity ligand, whereas CD8 $\alpha\beta$ T cells developed preferentially in the presence of the low-affinity ligand presented by H-2K^b (84). Similarly, H-Y TCR⁺ CD8 $\alpha\alpha$ IELs were more efficiently generated in H-2D^{b/b} homozygous male mice than in their H-2D^{b/d} heterozygous counterparts (87).

Similar to conventional $\alpha\beta$ T cells, CD8 $\alpha\alpha$ $\alpha\beta$ IELs require the expression of pre-T α during the initial steps of $\alpha\beta$ TCR rearrangements (88). The fact that the efficient agonist selection of CD8 $\alpha\alpha$ $\alpha\beta$ IELs also requires the expression of the TCR α chain connecting peptide domain (α -CPM) further suggests that the agonist and conventional selection processes have similar features and may have evolved simultaneously (63).

Agonistic selection in the thymus

The marked reduction in the numbers of CD8 $\alpha\alpha$ $\alpha\beta$ IELs in congenic athymic nude mice (89, 90) and in mice that had been neonatally thymectomized (91) indicates that the thymus is

required for the generation of agonist-selected CD8 $\alpha\alpha$ $\alpha\beta$ IELs. This requirement is further supported by the observation that thymus grafts from male H-Y TCR transgenic mice generated CD8 $\alpha\alpha$ H-Y TCR⁺ IELs when transplanted into female recipient mice (63), and wildtype thymus grafts efficiently generated CD8 $\alpha\alpha$ $\alpha\beta$ IELs when transplanted into recipient nude mice (42). In the latter case, it was shown that CD8 $\alpha\alpha$ IELs and conventional CD4 and CD8 $\alpha\beta$ IELs were generated with different kinetics. Fetal thymus grafts or grafts from mice up to weaning age mainly repopulated the intestine of the recipient mice with CD8 $\alpha\alpha$ $\alpha\beta$ IELs, whereas thymus grafts from older mice were more efficient in generating the conventional $\alpha\beta$ T-cell subsets (42). It should not be assumed, however, that the adult thymus had a reduced ability to generate CD8 $\alpha\alpha$ $\alpha\beta$ IELs as the adult thymus grafts from the male H-Y TCR transgenic *Rag*^{-/-} mice efficiently did so (63). It is possible that in the absence of conventional positive selection, as in the case of the male H-Y TCR transgenic mice on a *Rag*^{-/-} background, the generation of CD8 $\alpha\alpha$ $\alpha\beta$ IELs can continue in adult mice as long as a functional thymus is present.

Is it true that the type b IELs are developed in the thymus? Given that CD8 $\alpha\alpha$ $\alpha\beta$ IELs continued to be efficiently generated under conditions that caused deletion of CD8 $\alpha\beta$ T cells in the

thymus, some speculated that the precursors of these cells had not been submitted to thymic negative selection (41, 83). Taken together with observations that, in the absence of a thymus, these type b IELs nevertheless develop in lymphopenic mice grafted with normal bone marrow, fetal liver, or intestine (92, 93), this finding suggests that CD8 $\alpha\alpha$ $\alpha\beta$ IELs are likely the progeny of extrathymic progenitor cells localized within the intestine itself.

Kanamori et al. (94) demonstrated that approximately 1500 cryptopatch aggregates (CPs) are dispersed throughout the wall of the small and large intestine in mice (Fig. 3). Cells within the CP are composed of mostly lymphoid progenitors expressing stem cell factor receptor, or c-kit, and IL-7R α , but lacking the lineage markers (CD3, B220, Mac-1, Gr-1, and TER-119). They possess transcripts for germline TCR genes, messenger RNA (mRNA) for CD3 ϵ as well as proteins (i.e. RAG-2 and pre-T α) involved in TCR gene rearrangement (95), and are able to generate TCR⁺ IELs, type b CD8 $\alpha\alpha$ $\alpha\beta$ IELs, as well as $\gamma\delta$ IELs, albeit with a strong bias toward the generation of $\gamma\delta$ T cells, in irradiated severe combined immune-deficient mice. These findings demonstrated that c-kit⁺ CP cells are committed to the T-cell lineage and are competent for the generation of IELs (93, 96). Both IL-7R α ^{-/-} mice (94) and common cytokine receptor γ -chain (γ_c)-deficient mice (95) lack CP structure, but local

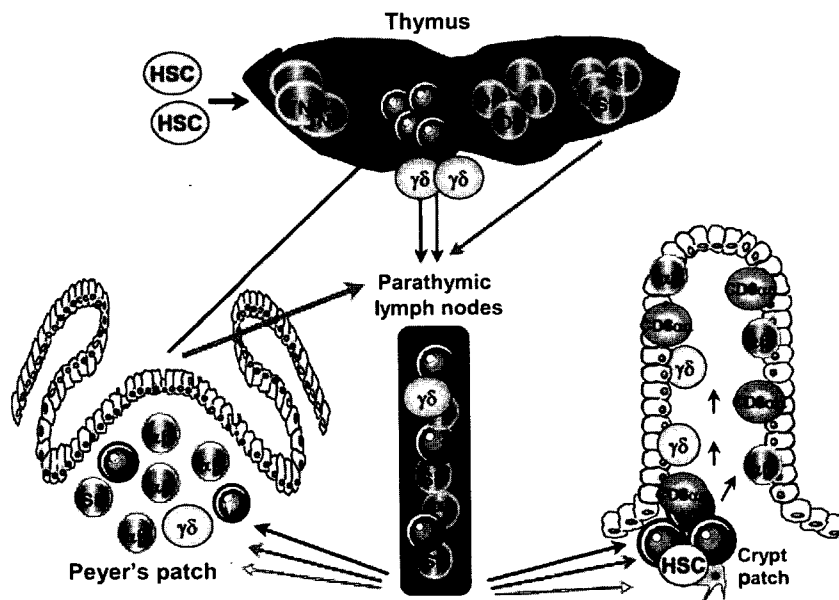


Fig. 3. Thymus-dependent versus thymus-independent IEL development. Bone marrow-derived or fetal-liver-derived hematopoietic stem cells (HSCs) enter the thymus as T-committed cells. Signals from the stroma and from other thymocytes (CD4⁺ CD8⁺ DP thymocytes?) induce differentiation toward type b unconventional T cells. Other immature CD4⁻ CD8⁻ CD3⁻ (TN) cells gradually switch off unconventional T-cell genes and differentiate toward conventional CD4⁺ or CD8⁺

single-positive (SP) $\alpha\beta$ T cells. Both immature and mature cells exit from the thymus. Unconventional T cells and their progenitors (TN2-3 cells) more readily populate the gut epithelium, possibly via CPs, where the progenitors may complete their differentiation into CD8 $\alpha\alpha$ IELs. Some TN2-3 progenitors may circulate back to the thymus. The gut epithelium could also be populated by bone marrow-derived HSCs that differentiate into type b $\gamma\delta$ T cells in CPs.

(fatty-acid-binding protein promoter) transgene expression of IL-7 by mature enterocytes of IL-7^{-/-} mice restored the development of CPs and $\gamma\delta$ IELs, whereas $\gamma\delta$ T cells remained absent from all other tissues (97), suggesting an important role for IL-7R signaling in their generation. These observations indicate a correlation between the presence of CP cells and lymphopoiesis in the gut, especially of $\gamma\delta$ IELs.

Lambolez et al. (88) reported difficulty identifying specific T-lineage-committed precursors among the CP cells. Using multiple lineage markers, they performed a detailed characterization of the progenitor cells isolated from CPs. They also transferred bone marrow from nude mice to irradiated thymectomized γ_c and RAG-2 double-deficient recipient mice to establish the sequence of appearance of these precursors during lymphoid development in the gut. This analysis provided little evidence that CP lineage (Lin)⁻ cells are T-cell committed, a finding consistent with data showing that several gene mutations that affect T-cell differentiation, including pre-T α , TCR α , and CD3 ϵ , had no effect on the appearance or ratio of CP precursor cells (88). However, CP Lin⁻ cells did express GATA-2, a transcription factor found in pluripotent hematopoietic progenitor cells, indicating that these CP cells might represent precursors of other hematopoietic cells. There is relatively little cell turnover in CPs, and very few CP cells express RAG-2 mRNA (88). In contrast, potential precursor cells isolated from the epithelium were more committed to the T-cell pathway, and TCR α mutation dramatically altered subtype ratios within the immature Lin⁻ cells of the epithelium (88).

In seeking to investigate which immune cells synthesize RAG-2, subsequent studies (98) have shown that RAG-2 production is detected exclusively in the thymus in normal mice with an intact thymus (euthymic), but RAG-2 is found in the MLNs of the gut and minimally in the PPs of nude or other athymic mice. Their findings suggest that extrathymic T-cell development is totally repressed in normal euthymic mice and becomes active only in cases of severe T-cell depletion. Whether in normal and athymic mice, neither CP cells nor IELs showed fluorescence-tagged RAG-2. To evaluate these findings, Nonaka et al. (99) generated athymic nude mice that lacked all lymph nodes and PPs by administering lymphotoxin- β receptor Ig and tumor necrosis factor (TNF)-R55-Ig fusion proteins into pregnant nu/+ mice and nu/nu *aly/aly* (alymphoplasia) mice that lacked all lymph nodes, PPs, as well as intestinal ILFs. These two kinds of mice, which harbored numerous DN as well as CD8 $\alpha\alpha$ $\gamma\delta$ IELs possessing TCR V γ 1, V γ 4, and V γ 7 segments in the small intestines, were found to retain CPs. These findings provide compelling evidence that in the lamina propria of the

intestine, at least some of the DN and CD8 $\alpha\alpha$ $\gamma\delta$ IELs developed within the tiny clusters, that is, CPs, that are filled with c-kit⁺IL-7R⁺Lin⁻ lymphopoietic cells.

Podd et al. (100) explored whether CD3 ϵ ⁺CP cells are linearly related to $\gamma\delta$ IELs using laser capture microdissection analysis. They observed that in-frame TCR V γ 5 rearrangements were detected in the CD3 ϵ ⁺CP cells that are shared with a subset of $\gamma\delta$ IELs. However, they did not detect mRNA for RAG-1 gene expression in the CP aggregates. Thus, their data lend support to the hypothesis that CPs are perhaps reservoirs for some antigen-experienced $\gamma\delta$ IELs in mice.

Eberl and Littman (101) performed interesting experiments in mice bearing two transgenes, a Cre recombinase transgene under the control of the retinoid-related orphan receptor γ t (ROR γ t) promoter and a green fluorescent protein (GFP) transgene controlled by a ubiquitous promoter that is only transcribed when a lox-flanked stop sequence is removed by Cre. ROR γ t, an orphan nuclear hormone receptor, was detected in immature DP thymocytes, fetal lymphoid-tissue-inducer cells, and c-kit⁺Lin⁻ cells in CPs. In these transgenic mice, all progeny of a Cre-expressing cell become constitutively GFP⁺ because the stop sequence of the GFP transgene has been deleted. These experiments, also performed with a CD4-Cre transgene, showed that GFP was expressed by all DP thymocytes and also by all single-positive T cells and $\alpha\beta$ IELs, including CD8 $\alpha\alpha$ $\alpha\beta$ IELs, but not by any other immune cells such as $\gamma\delta$ IELs, B cells, or NK cells. This finding elegantly demonstrates that all $\alpha\beta$ IELs are of thymic origin and must have been selected by the DP pathway. However, these findings did not exclude an extrathymic origin for $\gamma\delta$ and other IELs. They did suggest, however, that the contribution of intestinal precursors including c-kit⁺Lin⁻ cells is minimal in healthy mice but could become more substantial in lymphopenic mice or in mice in which large numbers of T cells have been mobilized in response to intestinal inflammation (101). However, Eberl and Littman (102) mentioned elsewhere that the principal function of the CP cells is to induce the formation of lymphoid follicles in the intestinal lamina propria in a manner similar to the induction of lymph nodes and PPs by the ROR γ t-expressing fetal lymphoid tissue inducer cells.

Resolution of the thymus-dependent versus thymus-independent controversy

Just when researchers seemed to hit an impasse regarding thymus involvement in IEL production, a truly incendiary report by Lambolez et al. (103) was published. In this paper, they proposed a previously unknown pathway for T-cell

development that has implications for the thymus every bit as great as for the gut. Reexamining assumptions germane to the basic biology of T-cell development in the thymus, they noticed that neonatal thymus grafts leak specific subsets of early T-committed progenitor cells that will repopulate both the host thymus, in which they rapidly complete normal T-cell development, and the host gut, where they survive for many weeks and give rise via defined intermediates to type b IELs (103). This finding conflicted with previous assumptions that the thymus is a source solely of mature T cells and that it does not have the potential for long-term reconstitution. Furthermore, this report emphasizes that the leaked cells are early CD4⁻CD8⁻CD3⁻ triple-negative stage 2 (TN2) and TN3 progenitors that are T-cell committed but that have not yet rearranged their TCR genes. However, CD8 α IELs could not be reconstituted even after adoptive transfer of other more mature DP or single-positive thymocyte populations (Fig. 3).

Using an indirect assay, the authors (103) also showed that the leaked TN cells are much more efficient at completing T-cell differentiation in the gut than are cells derived directly from the bone marrow (Fig. 3). These results strongly suggest that stroma cells in the thymus and/or other thymocytes could help 'persuade' progenitors to differentiate into unconventional CD8 α IELs. The thymic imprinting might include gene products that favor survival and/or are biased toward unconventional IEL development.

It should be underlined that recent studies using H-Y TCR transgenic mice seem to exclude the possibility that DP thymocytes could repopulate the intestine (104) because H-Y TCR β and TCR α transgenes in this model are expressed abnormally in DN thymocytes and the mice have no TN2–TN4 cells. Those phenotypes were shown to be artifacts of the transgenic systems. In conditions in which the TCR α chain is physiologically expressed in DP thymocytes of the H-Y mice, the male mice have many DP thymocytes expressing the self-reactive $\alpha\beta$ T cells, but no increase in *Smyc* peptide-specific CD8 α IELs is found in the gut. Thus, the accumulation of self-reactive TCR transgenic cells in the gut of conventional H-Y transgenic mice seems to be because of the anomalous expression of the TCR α transgene in early TN thymocytes.

It is not clear whether leaked TN2–TN3 cells home exclusively to the gut or to several mucosal sites. CPs may be special features of the gut that attract immature cells. In this regard, Onai et al. (105) reported that CD11c⁺ DCs in CP aggregates express CCL25 and that the signaling via the chemokine receptor CCR9 that is present on c-kit⁺ CP cells might play a central role in CP and CD8 α IEL development. When the expression of CCR9 on the surface of c-kit⁺ bone

marrow cells was blocked by a viral construct expressing an intracellular form of CCL25, bone-marrow-recipient mice largely failed to develop CP aggregates and CD8 α IELs. By contrast, the cellularity and composition of the thymus and spleen of these mice were well developed. These findings led Onai et al. (105) to hypothesize that CCR9–CCL25 interaction is required for CP maturation in the generation of TCR⁻ and TCR⁺ IEL subsets.

The gut itself then seems to retain immature thymic emigrants and to nurture their development. As shown before, this nurturing includes local production of IL-7, which promotes local TCR gene rearrangement (97). A capacity for gene rearrangement might seem unexpected, given a report that a reporter gene expressed from the *Rag-2* promoter is not detected outside the thymus (98). Nonetheless, it has been shown that *Rag-1* is expressed in the gut, albeit in low numbers, from an alternative 5' exon that may be conserved in the mouse (106). The reporter construct that has been used previously (95) might not register expression by an unorthodox *Rag* exon (98). In summary, the gut is a viable site for T-cell differentiation because it expresses *Rag*, locally produces IL-7, and can accommodate immature T-cell-committed progenitors.

Cytokine requirements for IEL development

IELs require cytokines for their development. As mentioned above, γ_c -deficient mice show impaired development of IELs expressing either $\alpha\beta$ TCRs or $\gamma\delta$ TCRs (107, 108). Several cytokines (IL-2, IL-4, IL-7, IL-9, IL-15, and IL-21) share γ_c as a part of their receptors. Consistent with impaired CP development, IL-7- or IL-7 receptor (IL-7R)-deficient mice did not have $\gamma\delta$ IELs but possessed substantial numbers of $\alpha\beta$ IELs (109, 110). Local transgene expression of IL-7 by the ECs of IL-7^{-/-} mice restored the development of $\gamma\delta$ IELs associated with CP development (97). Although both $\alpha\beta$ TCR- and $\gamma\delta$ TCR-positive CD8 α IELs belong to type b IELs, the developmental pathway of each is distinct, at least in terms of IL-7 requirements. $\gamma\delta$ IELs could be preferentially impaired at least in part because IL-7-mediated signaling is essential for the rearrangement of TCR γ genes (111). Additionally, the presence of substantial numbers of $\alpha\beta$ IELs in IL-7- or IL-7R-deficient mice implies that other cytokines using γ_c play a pivotal role in the regulation of IEL development, particularly of $\alpha\beta$ IELs.

Much attention is currently focused on IL-15, a cytokine that is produced by several kinds of cells including DCs and ECs and that includes an IL-2 receptor β chain (IL-2R β) as a part of its receptor. As one might expect from the elimination of $\alpha\beta$ and $\gamma\delta$ IELs from IL-2R β -deficient mice (112), disruption of the IL-15- or IL-15R α -encoding gene also resulted in a significant

reduction in IELs expressing either $\alpha\beta$ TCRs or $\gamma\delta$ TCRs (113, 114). Reciprocally, IL-15 stimulates IEL proliferation and restricts the TCR repertoire of IELs by selectively regulating local $V\gamma$ gene chromatin modification (115, 116). In regard to IL-15-induced IEL development, a recent finding has indicated that MyD88 (myeloid differentiation factor 88), an adapter molecule associated with Toll-like receptors, mediates IL-15 production by ECs. As expected, the number of IELs significantly decreased in MyD88-deficient mice (117), suggesting that a MyD88 regulation pathway may help maintain the intestinal homeostasis against commensal microorganisms via the IEL and EC cross-talk system.

IEL trafficking pathway

Determination of IEL tropism by chemokine

It is generally accepted that chemokines play an important role in the regulation of lymphocyte trafficking (118, 119). For the regulation of IEL trafficking into the small intestine, CCR9 has been considered to be a key chemokine (120, 121) (Fig. 1). Its ligand, thymus-expressed chemokine (CCL25), is constitutively expressed on the ECs in the small intestine (122, 123) (Fig. 1). CCL25 has been shown to mediate the localization of type a IELs, especially CD103⁺-naive cells and, more recently, to activate CD69⁺ cells in the small intestine (124). CCL25 also regulates precursor differentiation during the generation of type b IELs, but it is less involved in steps that involve IEL retention (125). Mice with defective responses against CCL25 exhibited a significant reduction in CPs and IELs in the small intestine. In contrast to these implications, however, disruption of the CCR9 gene resulted in only a modest decrease in IELs, indicating that other chemokines might be involved in the IEL migration into the small intestine (126, 127). Furthermore, although there are CCR9⁺ cells in the large intestine, the CCR9/CCL25-mediated pathway is not involved in IEL recruitment into the large intestine because its ligand (CCL25) is poorly expressed on large intestinal ECs (120, 122, 123). It would stand to reason then that chemokines other than CCR9⁺ are involved in the regulation of IEL trafficking in the large intestine. Indeed, CCL28, which acts as a ligand for CCR10⁺-IgA-secreting cells (128, 129), is the predominant chemokine expressed on ECs in the large but not the small intestine (130). Although memory-type T cells in the intestine did not express CCR10 (128), it is possible that a CCL28/CCR10-mediated pathway might contribute to the regulation of large intestinal IEL trafficking, especially naive-type IELs preferentially residing in the large intestine. In addition, the various chemokine receptors expressed on IELs, such as CCR3, CCR4, CCR5, and

CXCR3 (131–133), may play a pivotal role in the alternative pathway of IEL trafficking. Although CCR6 is not expressed on mature IELs, CCR6-deficient mice showed impaired development of CD4 CD8 $\alpha\alpha$ $\alpha\beta$ IELs and CD8 $\alpha\alpha$ $\alpha\beta$ IELs but not CD8 $\alpha\alpha$ $\gamma\delta$ IELs because of the dysregulated development of CPs (134). These findings suggest that CD8 $\alpha\alpha$ $\alpha\beta$ IEL and CD8 $\alpha\alpha$ $\gamma\delta$ IEL developments have different chemokine requirements.

In addition to chemokine-mediated pathways, we recently found that sphingosine-1-phosphate (S1P) plays a role in IEL trafficking, especially in the large intestine (Jun Kunisawa, unpublished data). S1P, a lipid mediator originating from sphingomyelin, has been linked to the regulation of lymphocyte emigration from the secondary lymphoid organs and thymus by an accumulating body of evidence (118). Our new finding suggests that some type a IELs also use S1P in their exit from the GALT (Jun Kunisawa, unpublished data). Because previous studies proposed a mutual interaction between S1P- and chemokine-mediated pathways in lymphocyte trafficking (135, 136), the cooperative pathway mediated by both S1P and chemokines may determine the selective trafficking of IELs into the small and large intestine.

Integrin-mediated IEL trafficking and distribution

In addition to chemokines, adhesion molecules also contribute to the regulation of IEL trafficking (Fig. 1). Among them, β 7 integrin has been considered to be an important molecule in determining the localization of intestinal IELs (Fig. 1). Thus, β 7-integrin-deficient mice showed critically reduced numbers of IELs in the intestine (137, 138). IEL numbers were equally reduced in the small and large intestine, indicating that a chemokine-mediated rather than an adhesion-molecule-mediated pathway determines whether IELs migrate to the small or large intestine.

Several lines of evidence have demonstrated that DCs derived from the GALT (PPs and MLNs) but not those derived from the spleen are capable of determining the gut tropism of intestinal T cells by the induction of high levels of α 4 β 7 integrin and CCR9 expression (139–141). A recent study showed that retinoic acid, which induces the α 4 β 7 integrin and CCR9 expression on T cells (142), is dominantly expressed by GALT DCs (Fig. 1). Thus, mucosal T cells activated by orally administered antigens presented by GALT DCs tend to migrate into distant intestinal effector sites, including the intestinal epithelium, by obtaining mucosal trafficking molecules (e.g. α 4 β 7 integrin and CCR9). Regarding the integrated integrin- and chemokine-mediated induction pathway, CCL25, a ligand for CCR9, is expressed in the crypt region of the small intestine, which is close to the mucosal addressin cell adhesion molecule 1 (MAdCAM-1)⁺ vessels (123, 127). These findings further indicate that α 4 β 7

integrin–MAdCAM-1 and CCL25–CCR9 interactions synergistically recruit gut-tropic IELs into the small intestine.

The unique migration pathway of RTE into the small intestine

Because the migration of RTE into the small intestinal epithelium was detected in the lymphotoxin- α -deficient mice lacking organized secondary lymphoid organs such as PPs and MLNs, their migration must have been independent of the GALT structure (32). In contrast, the migration of RTEs was abolished in CCR9-deficient mice or mice treated with anti-CCL25 antibody or anti- $\alpha 4\beta 7$ integrin antibody, suggesting that the migration was dependent on a CCL25/CCR9-mediated and $\alpha 4\beta 7$ -integrin-mediated pathway (32) (Fig. 1). As one might expect given the preferential expression of CCR9 on RTEs, RTEs homed to the small intestinal epithelium more efficiently than did non-RTE-naïve T cells (32). Although RTEs share with type a IELs a similar dependency on CCL25/CCR9 and $\alpha 4\beta 7$ integrin, RTEs do not require antigen-mediated activation. Thus, much unlike type a IELs, RTEs isolated from OT-I⁺Rag1^{-/-} mice can migrate into the small intestinal epithelium without antigen stimulation (32, 143).

Retention molecules on IELs

The intimate integrin-mediated interaction between ECs and IELs plays a pivotal role not just in the infiltration of the IELs into the intestine but also in the retention of IELs at the epithelium. Many attempts have been made at identifying the molecules that tether IELs and ECs together in the small intestine. CD103 (αE integrin), which interacts with the E-cadherin expressed on ECs, is expressed on IELs (144) (Fig. 1). It was previously reported that transforming growth factor- β (TGF- β) induced downregulation of $\alpha 4\beta 7$ integrin and simultaneously upregulated CD103 (145). Thus, $\alpha 4\beta 7$ integrin expression was reduced following IEL entry into the small intestinal epithelium, a reduction that coincided with an increase in CD103 expression (146). In addition to the TGF- β -mediated pathway, a recent study (146) proposed that CCR9-mediated signaling promoted the induction of CD103. In CCR9-deficient CD8⁺ T cells, upregulation of CD103 after migration into the small intestinal epithelium was markedly delayed (146). Additionally, the interaction between CD103 and E-cadherin was involved in the destruction of the intestinal epithelium during graft-versus-host disease (147), indicating that CD103 is a pivotal molecule regulating lymphocyte trafficking not only under natural conditions but also during pathological situations. Further, although CD103 was not required for the entry into the intestine (138), CD103-deficient mice exhibited

a decreased number of IELs (148). However, the reduction was modest, suggesting that other molecules contribute to the retention of IELs at the epithelium. In this regard, several lines of evidence have demonstrated that IEL retention may be mediated by their expression of some integrins (e.g. $\alpha 1\beta 1$ integrin, $\alpha 4\beta 1$ integrin, and $\beta 2$ integrin) (149–151). As in CD103-deficient mice, no or only a partial reduction in IELs was seen following the disruption of each of these integrins, suggesting that other integrins and/or molecules may compensate for their lack (151, 152). Additionally, we previously reported that IELs expressed the EC adhesion molecule (Ep-CAM) (153) (Fig. 1). Because Ep-CAM is also expressed on ECs and mediates homophilic cell-to-cell adhesion (154), interaction between IELs and ECs via Ep-CAM may play a pivotal role in IEL retention. Further, a recent study (155) indicates that IELs express tight-junction molecules such as occludins and zonula occludens-1 (Fig. 1). Taken together, the intimate biological interactions between ECs and IELs provide physiological and immunological barriers that act as a first line of defense at the intestine.

Physiological and biological functions of IELs

CD8 $\alpha\alpha$ $\alpha\beta$ IELs in immune protection and mucosal homeostasis

Despite their high frequency, potentially autoreactive type b CD8 $\alpha\alpha$ $\alpha\beta$ IELs are not self-destructive under normal conditions. Rather, their presence in the gut epithelium correlates with immune regulation and immune quiescence. Because endogenous, natural, specific self-antigens for these CD8 $\alpha\alpha$ $\alpha\beta$ IELs have not yet been identified, studies analyzing their functional ability are limited to the use of TCR transgenic mice, which express a cognate antigen for this TCR either naturally or through expression of a second transgene.

These transgenic mice, which have few conventional $\alpha\beta$ T cells expressing the transgenic TCR in the thymus, spleen, and lymph nodes, generate enhanced numbers of transgenic CD8 $\alpha\alpha$ $\alpha\beta$ IELs specific for the self-antigen (63) that are not anergic (83, 86, 156). Upon a systemic LCMV infection in double-transgenic mice expressing both an LCMV epitope (GP33-41)-specific TCR transgene and the cognate antigen transgene (GP33-41) driven by an MHC class I promoter, CD8 $\alpha\alpha$ $\alpha\beta$ IELs showed signs of virus-induced activation (86). This activation was antigen specific as the CD8 $\alpha\alpha$ $\alpha\beta$ IELs from double-transgenic mice expressing an OVA-peptide-specific TCR and the OVA antigen did not show activation following LCMV infection. Unlike conventional (GP33-41-specific) CD8 $\alpha\beta$ T cells, which lose self-tolerance upon LCMV infection, the activation of these self-specific CD8 $\alpha\alpha$ $\alpha\beta$ IELs did not result in

a loss of tolerance. The activated CD8 $\alpha\alpha$ $\alpha\beta$ IELs did not show antigen-specific cytotoxicity nor did they promote an inflammatory response. Instead, the responding CD8 $\alpha\alpha$ $\alpha\beta$ IELs secreted enhanced levels of TGF- β . The constitutive expression of measurable levels of mRNA for immunoregulatory cytokines, including IFN- γ , TGF- β , and IL-10, even without specific antigenic stimulation, further underscores that these type b IELs might exert regulatory functions rather than affording immune protection against specific pathogens.

The regulatory ability of these self-reactive T cells was first showed by analysis of gene-targeted animals, including IL-2/IL-2R-deficient mice, which have specific defects in this mucosal T-cell subset. These mice, in which the CD8 $\alpha\alpha$ $\alpha\beta$ IEL population is reduced, are highly susceptible to gut inflammatory diseases (157, 158).

Direct evidence that CD8 $\alpha\alpha$ $\alpha\beta$ IELs could regulate immune responses by conventional mucosal T cells was provided by sophisticated adoptive transfer experiments of CD8 $\alpha\alpha$ $\alpha\beta$ IELs into immune-deficient Rag^{-/-} recipient mice that also received pathogenic CD4⁺CD45RB^{high} splenocytes (156). In the absence of CD8 $\alpha\alpha$ $\alpha\beta$ IELs, transferred pathogenic CD4⁺ T cells migrate to the intestine and induce an uncontrolled inflammatory immune response, whereas mice that had previously received CD8 $\alpha\alpha$ $\alpha\beta$ IELs were protected against this CD4⁺ T-cell-mediated pathology. Surprisingly, CD4⁺CD45RB^{high} T-cell-induced colitis was prevented by transfer of CD8 $\alpha\alpha$ $\alpha\beta$ IELs isolated from H-Y TCR transgenic male donor mice (156). This protection, however, was observed only in H-2D^b male recipient mice and not in female mice, indicating that TCR must recognize self-antigen if it is to mediate regulatory function.

CD8 $\alpha\alpha$ $\gamma\delta$ IELs and immune protection

The antigen specificity of CD8 $\alpha\alpha$ $\gamma\delta$ IELs is poorly defined, and cell-transfer studies have indicated that the CD8 $\alpha\alpha$ $\gamma\delta$ IELs have only minimal pathogen-specific activity (159, 160). Nevertheless, in the case of recipient mice infected with *Toxoplasma gondii*, it was shown that successful protection by CD8 $\alpha\alpha$ $\alpha\beta$ IELs was, in part, dependent on the presence of CD8 $\alpha\alpha$ $\gamma\delta$ IELs (160). Although these CD8 $\alpha\alpha$ $\gamma\delta$ IELs show only minimal pathogen-specific activity, their ability to respond rapidly to unprocessed self-antigens expressed by stressed or transformed ECs and to control the activity of the non-self-reactive CD8 $\alpha\beta$ $\alpha\beta$ IELs has led to the idea that these cells might provide the first line of defense against invading pathogens (54).

Alternatively, because activated $\gamma\delta$ IELs can produce keratinocyte growth factor (KGF), which is important for epithelial growth and the repair of damaged tissue, some $\gamma\delta$ IELs could be involved in the repair of tissue damage elicited during

inflammatory immune responses (161). The synthesis of KGF by $\gamma\delta$ IELs has led to the hypothesis that these cells are important for the integrity and healing of the epithelium. Consistent with this hypothesis, the epithelium of the intestine in TCR δ -chain-deficient mice had reduced numbers of crypts (162). By contrast, the epithelium had a higher mitotic index in the presence of $\gamma\delta$ TCR transgene (163). In addition, TCR δ -chain-deficient mice show increased susceptibility to epithelial damage caused by 2,4,6-trinitrobenzene sulfonic-acid-induced colitis; indeed, the transfer of $\gamma\delta$ IELs to TCR δ -chain-deficient mice ameliorated the hapten-induced colitis, which correlated with decreased IFN- γ and TNF- α production and increased TGF- β production by IELs (164).

It remains to be determined if the effect of $\gamma\delta$ IELs on ECs is mediated solely by KGF or if other factors are involved. $\gamma\delta$ IELs synthesize mRNA for TGF- β 1 (18) and TGF- β 3 (165), and TGF- β has been reported to aid in the healing of epithelial damage (166). Furthermore, a lack of $\gamma\delta$ T cells often results in severe pathological conditions mediated by autoantibodies and/or in destructive cell-mediated immune responses by $\alpha\beta$ T cells (167). TCR δ -deficient mice that have been orally infected with *Eimeria vermiformis*, an enterocyte-specific parasite, can efficiently control the infection, but they also show severe bleeding in the small intestine because of uncontrolled immune responses by infiltrating CD4⁺ $\alpha\beta$ T cells (168). Transfer of normal IELs to these TCR δ -deficient mice could prevent mucosal injury. These results suggest that $\gamma\delta$ IELs exert significant control over the immune responses mediated by mucosal $\alpha\beta$ T cells.

Not all immune responses mediated by $\gamma\delta$ IELs are beneficial. For example, in the case of celiac disease, it was shown that dysregulated IL-15 secretion by wheat gliadin elicited over-expression of MICA/MICB on ECs and IELs, possibly including $\gamma\delta$ T cells. When isolated from patients with active celiac disease, $\gamma\delta$ T cells exhibited NKG2D-mediated cytotoxicity of the EC line HT29 possessing MICA (30).

$\gamma\delta$ IELs for mucosal IgA responses

$\gamma\delta$ IELs appear to be necessary for the induction and maintenance of humoral immune responses in the intestine. Interestingly, in TCR α -chain-deficient mice, an increase in the numbers and overall proportion of $\gamma\delta$ IELs is seen after environmental microbial challenge (169), which is accompanied by B-cell maturation and production of high levels of Igs (170–172). Collaboration between B and non- $\alpha\beta$ T cells, including $\gamma\delta$ T cells as well as $\beta\beta$ TCR⁺ T cells ($\beta\beta$ T cells), sustains the production of germinal centers, lymphoid follicles that ordinarily are the anatomical signature of $\alpha\beta$ T-cell and B-cell interactions.

Murine $\gamma\delta$ IELs, if administered in the periphery, can enhance antigen-specific IgA responses to orally ingested antigens and thus might be capable of abrogating oral tolerance (173, 174). However, because it is unlikely that $\gamma\delta$ IELs recirculate from the intestinal mucosa, their potential role in inducing or suppressing systemic oral tolerance is questionable. Although antibody responses against some T-cell-dependent antigens are impaired in mice lacking $\alpha\beta$ T cells, mucosal $\gamma\delta$ T cells may enhance antigen-specific as well as polyclonal S-IgA responses in the lamina propria of the small intestine (175).

Conclusions and future perspectives

In the harsh environment of the gut, the mucosal immune system must prevent the invasion of pathogenic micro-

organisms from the lumen while remaining quiescent against commensal microorganisms and food antigens, if disordered immune responses are to be prevented. Remarkably, IELs play a pivotal role in both immune surveillance and immune regulation, helping to maintain intestinal homeostasis by adjusting their immunological function to the different circumstances that prevail in the small and large intestines. However, the origin and development of these key players remain a subject of debate, and so it is crucial that we work to define their precise function and delineate their developmental pathway. Further investigation of IELs in both the small and large intestine will undoubtedly be key to the development of novel mucosal vaccines against infectious diseases and to immune therapy for inflammatory bowel diseases, food allergies, and celiac diseases.

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Role of Peyer's patches in the induction of *Helicobacter pylori*-induced gastritis

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Edited by Jeffrey I. Gordon, Washington University School of Medicine, St. Louis, MO, and approved April 12, 2007 (received for review October 12, 2006)

***Helicobacter pylori* is a Gram-negative spiral bacterium that causes gastritis and peptic ulcer and has been implicated in the pathogenesis of gastric adenocarcinoma and mucosa-associated lymphoid tissue lymphoma. Although Th1 immunity is involved in gastritis and the accumulation of *H. pylori*-specific CD4⁺ T cells in the *H. pylori*-infected gastric mucosa in human patients, how T cells are primed with *H. pylori* antigens is unknown because no apparent lymphoid tissues are present in the stomach. We demonstrate here that Peyer's patches (PPs) in the small intestine play critical roles in *H. pylori*-induced gastritis; no gastritis is induced in *H. pylori*-infected mice lacking PPs. We also observed that the coccoid form of *H. pylori* is phagocytosed by dendritic cells in PPs. We propose that *H. pylori* converts to the coccoid form in the anaerobic small intestine and stimulates the host immune system through PPs.**

CD4 T cells | coccoid form | dendritic cells | gastric epithelial cells | inflammation

H*elicobacter pylori* is a Gram-negative microaerophilic bacterium that infects human gastric epithelial cell (gEC) surfaces and the overlying gastric mucin. More than 50% of the world's population is infected by *H. pylori*, although most patients have no remarkable symptoms (1). However, in some of patients, *H. pylori* infection leads to active chronic gastritis or peptic ulcer (2). In addition, *H. pylori* has also been implicated in the pathogenesis of gastric adenocarcinoma and mucosa-associated lymphoid tissue lymphoma (3). When *H. pylori* colonizes gastric mucosa, effector molecules are injected into gastric epithelial cells or the submucosal area through the type IV secretion system (1, 4). For example, the CagA effector is phosphorylated in the target cells and activates a signaling pathway to elicit growth factor-like responses. Another effector molecule, VacA, causes the massive vacuolar degradation of epithelial cells, thus disrupting the gastric epithelial barrier. VacA also interferes with the activation and proliferation of T lymphocytes within the gastric lamina propria (gLP) (5).

It was originally proposed that effector molecules, including CagA, trigger the secretion of chemokines such as IL-8 and RANTES from gECs, which attract neutrophils and mononuclear cells into the gLP (4). However, it was later shown that *H. pylori* did not induce gastritis in lymphopenic SCID mice, although gastritis was induced after adoptive transfer of naive CD4⁺ T cells (6). The importance of CD4⁺ T cells was underscored by the fact that *H. pylori* is not eliminated from gastric mucosa in MHC class II-deficient mice (7).

Gastritis is more severe in Th1-prone mice than Th2-prone mice on infection with the mouse-adapted *H. pylori* strain SS1 (8). Furthermore, the accumulation of *H. pylori*-specific CD4⁺ T cells in the *H. pylori*-infected gastric mucosa in human patients (9) suggests that CD4⁺ T cell-mediated Th1 immune responses play a critical role in *H. pylori*-induced gastritis. However, how CD4⁺ T cells are primed by *H. pylori* antigens in the stomach where no apparent lymphoid tissues are present and how the

H. pylori-induced chronic inflammation is maintained by T cells is unknown.

Although *H. pylori* forms an actively dividing, spiral-shaped morphology in the stomach, it is able to convert to a nonculturable, but viable, coccoid form under unfavorable conditions such as an anaerobic environment, increased oxygen tension, and long-term culture (10, 11). The coccoid form is thought to be important for transmission to new hosts by an oral-oral or oral-feces route because this form is more resistant to environmental stresses. Although the coccoid form is not culturable *in vitro*, transcription and translation actively take place in the coccoid form (12, 13). However, it is unknown whether the coccoid form is involved in the pathogenesis of *H. pylori*-induced gastritis.

In this study, we demonstrate that *H. pylori* antigen-specific CD4⁺ T cells are necessary and sufficient for the induction of gastritis by *H. pylori*. We also demonstrate that CD4⁺ T cells are likely primed with *H. pylori* antigens captured in the small intestine, where the coccoid form of *H. pylori* is taken up by dendritic cells (DCs) in Peyer's patches (PPs).

Results

Adoptive Transfer of Naive CD4⁺ T Cells Induces Gastritis in *H. pylori*-Infected Rag2^{-/-} Mice. The *H. pylori* SS1 strain induces more severe gastritis in Th1-prone C57BL/6 than Th2-prone BALB/c mice as demonstrated by the infiltration of neutrophils and lymphocytes into the gLP and the submucosal area (Fig. 1*a* and data not shown). In contrast, when C57BL/6-Rag2^{-/-} mice lacking T and B cells were infected with *H. pylori*, no gastritis was observed (Fig. 1*b*), as previously shown with SCID mice (6). The clearance of bacteria in Rag2^{-/-} mice was impaired because >10⁷ cfu/g tissues of *H. pylori* colonized the gastric mucosa (Table 1), and the colonization of *H. pylori* was readily detected by anti-*H. pylori* antibody staining (Fig. 1*c*). However, adoptive transfer of naive splenic CD4⁺ T cells into Rag2^{-/-} mice 2 months after *H. pylori* infection induced severe gastritis, with massive infiltration of neutrophils and lymphocytes into the gLP and the submucosal area (Fig. 1*d*). This massive infiltration resulted in the exclusion of colonized *H. pylori* from gastric mucosa (Table 1).

Author contributions: S.N., H.K., C.S., and S.K. designed research; S.N., H.M., T.Y., Y.B., K.M., T.N., and T.S. performed research; and S.N., C.S., and S.K. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

Abbreviations: DKO, double knockout; LP, lamina propria; gLP, gastric LP; gEC, gastric epithelial cell; DC, dendritic cell; PPs, Peyer's patches; OVA, ovalbumin; NK, natural killer; APC, antigen-presenting cell; β -Rag DKO, IL-2 receptor β chain (IL-2R β)^{-/-} Rag2^{-/-} DKO; γ -Rag DKO, cytokine receptor common γ chain (γ)^{-/-} Rag2^{-/-} DKO; GALT, gut-associated lymphoid tissue; ILF, isolated lymphoid follicle; BMDC, bone marrow-derived cell; SED, subepithelial dome; mLN, mesenteric lymph node.

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This article contains supporting information online at www.pnas.org/cgi/content/full/0609014104/DC1.

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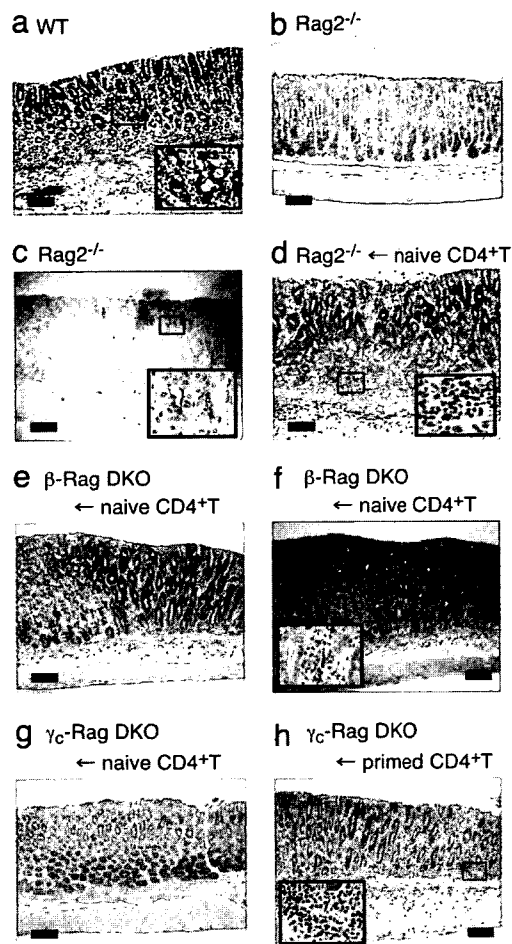


Fig. 1. Naive CD4⁺ T cells did not induce gastritis in *H. pylori*-infected γ_c -Rag double knockout (DKO) mice. (a–c) Wild-type (a) or *Rag2*^{-/-} (b and c) mice were infected with *H. pylori*. Two months after the infection, gastric specimens were prepared. (d–h) *Rag2*^{-/-} (d), β -Rag DKO (e and f), and γ_c -Rag DKO (g and h) mice were infected with *H. pylori*. Two months after the infection, naive (d–g) or primed (h) splenic CD4⁺ T cells were transferred. Two months after the cell transfer, gastric specimens were prepared. Specimens were stained with H&E (a, b, d, e, g, and h), anti-*H. pylori* antisera (brown) (c), or chloroacetate esterase (red) for infiltrated neutrophils and mast cells (f). (Scale bars: 200 μ m.)

***H. pylori* Antigen-Specific CD4⁺ T Cells Are Indispensable for Induction of Gastritis.** Primary gECs secrete MIP-2, a functional homolog of IL-8, on *H. pylori* infection *in vitro*, and the amount produced by *Rag2*^{-/-} gECs was comparable to that produced by wild-type gECs [supporting information (SI) Fig. 5a]. CD4⁺ T cells isolated from the gLP of *H. pylori*-infected mice were able to produce larger amounts of MIP-2 than splenic CD4⁺ T cells from the same mice in response to *H. pylori* antigens (SI Fig. 5b). Moreover, the amounts of MIP-2 produced by gLP CD4⁺ T cells were much larger than those produced by gECs (compare SI Fig. 5a and b). The importance of CD4⁺ T cells for neutrophil infiltration on *H. pylori* infection was further confirmed by the depletion of CD4⁺ T cells from wild-type mice that had already developed gastritis by *H. pylori* infection. After depleting CD4⁺ T cells by the i.v. injection of anti-CD4 mAb, the gastritis became milder (Table 1 and SI Fig. 6a and b), and the number of bacteria in the gastric mucosa increased (Table 1). These results indicate the critical role of CD4⁺ T cells for both triggering and maintaining gastritis. When CD4⁺ T cells from OT-II transgenic mice

on a *Rag2*^{-/-} background (OT-II-Rag mice), specific for an ovalbumin (OVA_{323–339} peptide on an MHC class II molecule I-A^b), were transferred into *H. pylori*-infected *Rag2*^{-/-} mice, no gastritis was induced (SI Fig. 6c). Similarly, when OT-II-Rag mice were infected with *H. pylori*, no gastritis was induced despite the presence of CD4⁺ T cells (Table 1 and SI Fig. 6d). Furthermore, when OVA protein or OVA_{323–339} peptide was administered into *H. pylori*-infected OT-II-Rag mice, no inflammation was observed, although CD4⁺ T cells were activated in these mice (Table 1, SI Fig. 6e, and data not shown). These results collectively indicate the importance of *H. pylori* antigen recognition by CD4⁺ T cells in the induction of gastritis.

CD4⁺ T Cells Are Not Primed with *H. pylori* Antigen in γ_c -Rag DKO Mice. IFN γ , a key cytokine for Th1 immune responses, is important for the pathogenesis of *H. pylori*-induced gastritis (14). Natural killer (NK) cells and antigen-presenting cells (APCs) including DCs are able to produce IFN γ to prevent bacterial infection (15). The interaction between DCs and NK cells enhances the production of IFN γ during *H. pylori* infection (16, 17). To test the importance of DC–NK interaction in the *H. pylori*-induced inflammatory response, we transferred splenic CD4⁺ T cells into *H. pylori*-infected IL-2 receptor β chain (IL-2R β)^{-/-}*Rag2*^{-/-} (β -Rag DKO) mice and cytokine receptor common γ chain (γ_c)^{-/-}*Rag2*^{-/-} DKO (γ_c -Rag DKO) mice. These mice lack NK cells because of impaired IL-15 signaling, which is critical for NK cell development (18, 19). In addition, the production of IL-12 and IFN γ by APCs from these mice is impaired (20). As shown in Fig. 1e and f, gastritis was induced in *H. pylori*-infected β -Rag DKO mice when naive CD4⁺ T cells were transferred. Clearance of bacteria was also achieved by the naive CD4⁺ T cell transfer (Table 1), indicating that NK cells and NK–DC interaction are dispensable for the induction of gastritis by *H. pylori* infection.

Surprisingly, there was no gastritis induced in γ_c -Rag DKO mice even after the transfer of naive CD4⁺ T cells (Fig. 1g), NK cells (SI Fig. 7a), or NK cells with naive CD4⁺ T cells (SI Fig. 7b), suggesting that γ_c -Rag DKO mice have additional defects compared with β -Rag DKO mice. Interestingly, when splenic CD4⁺ T cells isolated from *H. pylori*-infected wild-type mice were transferred, gastritis was induced in *H. pylori*-infected γ_c -Rag DKO mice (Fig. 1h), and the clearance of bacteria was evident (Table 1). These results suggest that CD4⁺ T cells were not primed in γ_c -Rag DKO mice. In fact, splenocytes from these mice did not respond to DCs preincubated with *H. pylori* lysate, whereas splenocytes from wild-type mice infected with *H. pylori* strongly responded and produced IFN γ in response to the same DC preparation (data not shown). It should be noted that there were no apparent defects in DCs from γ_c -Rag DKO mice compared with those from wild-type mice with regard to their ability to induce T cell activation and present antigen as examined by the induction of CD69 expression and IFN γ production by splenic CD4⁺ T cells (SI Fig. 8).

PPs Are a Critical Tissue for Priming CD4⁺ T Cells with *H. pylori* Antigen.

One difference between β -Rag DKO and γ_c -Rag DKO mice is that the latter lack gut-associated lymphoid tissues (GALT) such as PPs and isolated lymphoid follicles (ILFs) due to impaired IL-7 signaling (21) (Fig. 2a–d). Thus, we hypothesized that CD4⁺ T cells are primed in GALT such as PPs or ILFs. To test this possibility, we generated PP-null mice by administration of anti-IL-7R α mAb *in utero* (22) (Fig. 2e and f). As observed in γ_c -Rag DKO mice, no gastritis was induced in PP-null mice 2 months after *H. pylori* infection, and a large number of *H. pylori* was detected in the gastric mucosa (Fig. 2h and i and Table 1). We also generated PP-null mice on a *Rag2*^{-/-} background (PP-null-*Rag2*^{-/-} mice) (Fig. 2g). The adoptive transfer of CD4⁺ T cells from *H. pylori*-infected wild-type mice, but not naive CD4⁺ T cells, induced strong inflammation in PP-null-*Rag2*^{-/-}

Table 1. PP-dependent bacterial clearance in *H. pylori* infection

Mouse*	n	Cells transferred†	Bacterial colonization,‡ cfu/g tissue × 10 ⁻⁶	Neutrophils, average (range)	Active inflammation, average (range)	GAIS,§ average (range)
Wild type	7	None	2.2 ± 1.3	1.6 (0-3)	1.4 (0-3)	13.6 (0-34)
Wild type	4	CD4 ⁺ T cell-depleted	25 ± 7	0 (0)	1.0 (0-2)	0 (0)
Rag2 ^{-/-}	4	None	14 ± 4	0 (0)	0 (0)	0 (0)
Rag2 ^{-/-}	5	Naive CD4 ⁺ T	0.15 ± 0.17	2.0 (2)	2.0 (2)	5.0 (2-8)
Rag2 ^{-/-}	5	OT-II-Rag CD4 ⁺ T	34 ± 11	0 (0)	0 (0)	0 (0)
β-Rag DKO	4	Naive CD4 ⁺ T	0.67 ± 0.39	1.5 (1-2)	1.3 (1-2)	9.2 (3-18)
γ _c -Rag DKO	6	Naive CD4 ⁺ T	18 ± 11	0 (0)	0 (0)	0 (0)
γ _c -Rag DKO	6	Primed CD4 ⁺ T	0.22 ± 0.39	0.66 (0-1)	1.5 (0-2)	1.6 (0-5)
γ _c -Rag DKO	3	Primed CD4 ⁺ T from PPs	<0.01	1.0 (1)	2.0 (2)	1.0 (0-2)
γ _c -Rag DKO	3	Primed CD4 ⁺ T from mLN	0.88 ± 0.78	0.5 (0-1)	1.5 (1-2)	4.0 (3-5)
γ _c -Rag DKO	3	Primed CD4 ⁺ T by coccoid form	1.8 ± 1.6	0.33 (0-1)	0.66 (0-1)	2.0 (0-6)
PP-null-wild type	8	None	16 ± 9	0 (0)	0.37 (0-1)	0 (0)
PP-null-Rag2 ^{-/-}	3	Naive CD4 ⁺ T	5.3 ± 4.0	1.0 (0-2)	1.0 (0-2)	4 (0-7)
PP-null-Rag2 ^{-/-}	3	Primed CD4 ⁺ T	0.53 ± 0.28	2.6 (2-3)	2.6 (2-3)	21 (13-33)

*All mice were on a C57BL/6 background. Although not shown, the degrees of bacterial colonization in β-Rag DKO, γ_c-Rag DKO, and PP-null-Rag2^{-/-} mice without CD4⁺ T cell transfer were similar to those of Rag2^{-/-} mice.

†Splenocytes were used for cell transfer unless otherwise stated. Five million cells were transferred except for the transfer of PP-derived cells, where 5 × 10⁵ cells were used.

‡Mean ± SD.

§Gland active inflammatory score. See *Materials and Methods*.

mice just as in the γ_c-Rag DKO mice (Fig. 2 *j* and *k* and Table 1). These results strongly suggest that PPs are critical for priming CD4⁺ T cells in *H. pylori* infection, but dispensable for the effector phase.

The Coccoid, but Not Helical, Form of *H. pylori* Is Phagocytosed by DCs in PP. Although *H. pylori* is helical in the stomach, it transforms to the coccoid form under anaerobic conditions, such as in the small intestine (23). Interestingly, the coccoid form of *H. pylori* induced higher IL-12 production from bone marrow-derived cells (BMDCs) than the helical form (SI Fig. 9). There is a possibility that *H. pylori* transforms to the coccoid form in the intestine and is then captured by DCs present in PPs to induce a Th1 response. To test this possibility, the coccoid and helical forms of *H. pylori* were inoculated into ligated small intestinal loops. As shown in Fig. 3*a*, immunofluorescence staining detected *H. pylori* in the subepithelial dome (SED) region of PPs in a time-dependent manner, and the number of bacteria in the SED region inoculated with the coccoid form was larger than that inoculated with the helical form of *H. pylori*. In addition, double immunofluorescence staining with anti-*H. pylori* antibody and anti-CD11c mAb demonstrated that the bacteria were captured by CD11c⁺ DCs in the SED region (Fig. 3*b*). Although the helical form of *H. pylori* kept the rod shape 1.5 h after inoculation (SI Fig. 10 *a* and *b*), the bacteria phagocytosed by CD11c⁺ DCs in PPs were round (SI Fig. 10 *c* and *d*). These results suggest that the coccoid, but not the helical, form of *H. pylori* is captured by DCs in PPs and activates immune responses by generating *H. pylori*-specific pathogenic CD4⁺ T cells. Consistent with this observation, CD4⁺ T cells from the PPs as well as mesenteric lymph node (mLN) of *H. pylori*-infected wild-type mice were also able to eliminate the bacteria in γ_c-Rag DKO mice infected with *H. pylori* (Table 1).

When wild-type mice were infected with the coccoid form of *H. pylori*, gastritis was not induced because the coccoid form of *H. pylori* was unable to colonize in the stomach (Fig. 4*a* and data not shown). However, CD4⁺ T cells from these mice induced gastritis in γ_c-Rag DKO mice infected with the helical form of *H. pylori* (Fig. 4 *b* and *c* and Table 1). These results indicate that CD4⁺ T cells primed with the coccoid form of *H. pylori* in the

intestine are sufficient to induce inflammation in the stomach infected with the helical form of *H. pylori*.

Discussion

We showed here that *H. pylori* antigen-specific CD4⁺ T cells are required to induce and maintain gastritis on infection with *H. pylori*. Because *H. pylori* interacts with and injects pathological molecules into gECs, it is generally thought that neutrophils infiltrating the gLP are attracted by chemokines produced by gECs. However, neutrophil infiltration was not observed in Rag2^{-/-} mice, although gECs of Rag2^{-/-} mice were able to secrete MIP-2 on *H. pylori* infection. Thus, the secretion of chemokines by gECs seems insufficient for the induction of gastritis. In addition, adoptive transfer of CD4⁺ T cells recognizing *H. pylori*-independent antigens did not induce gastritis, suggesting that *H. pylori*-specific CD4⁺ T cells directly or indirectly regulate production of chemokines that attract neutrophils. In fact, a large amount of MIP-2 was produced by activated CD4⁺ T cells derived from the gLP of *H. pylori*-infected mice. In addition, infiltrated neutrophils were located around CD4⁺ T cells in the gLP of *H. pylori*-infected mice (data not shown). It is known that another keratinocyte-derived chemokine is able to recruit neutrophils. However, the amounts of keratinocyte-derived chemokine produced by both gECs and CD4 T cells were much lower than those of MIP-2 (data not shown).

Oral or intra-PP immunization with *H. pylori* antigens was effective in enhancing *H. pylori*-specific CD4⁺ T cell responses and reducing *H. pylori* colonization in the stomach (24, 25). These reports are consistent with our current observation that PPs play critical roles in priming CD4⁺ T cells, and *H. pylori* is indeed captured by DCs in PPs. *H. pylori* antigen-specific CD4⁺ T cells would be primed by DCs in PPs or mLNs where DCs migrate after capturing antigens. Interestingly, CD4⁺ T cells cannot be primed by DCs in the gLP or gEC, both of which are capable of expressing class II MHC and presenting antigens. The lack of antigen presentation is partly due to the fact that the helical form of *H. pylori* is resistant to phagocytosis in a type IV secretion system-dependent manner, although the molecular mechanisms of antiphagocytic activity remain to be determined (26). It should be noted that the transformation of the helical to the coccoid form is accompanied by changes in the composition

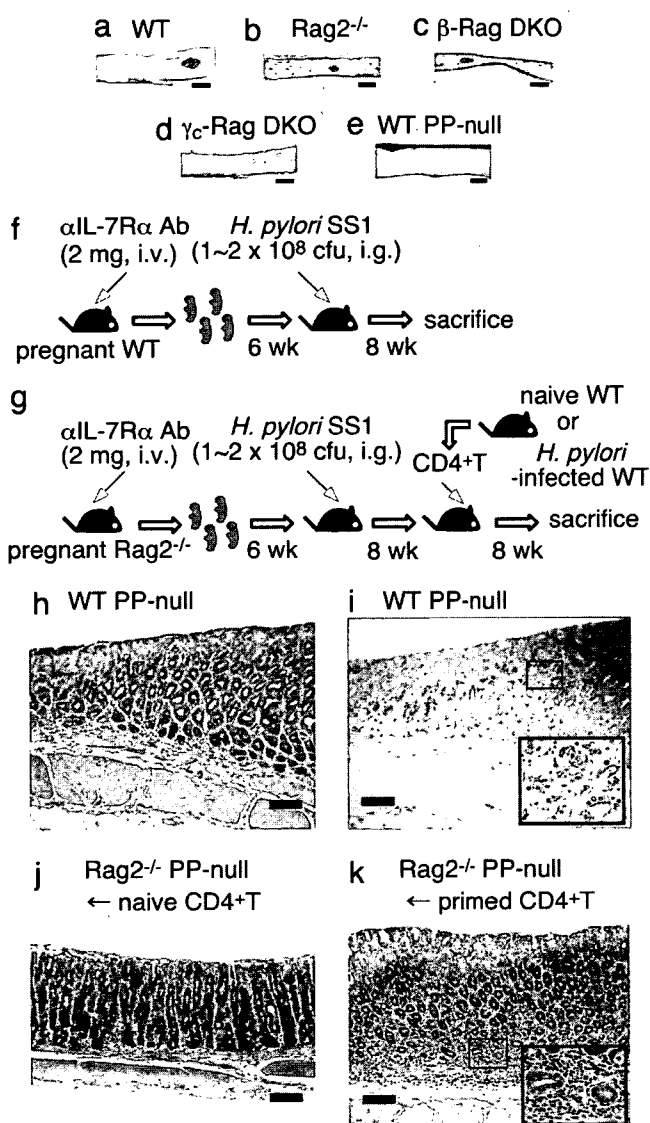


Fig. 2. PPs are critical for the priming of CD4⁺ T cells in *H. pylori* infection. (a–e) Small intestines from wild-type (WT) (a) and PP-null (e) mice were stained with anti-B220 mAb, and small intestines from Rag2^{-/-} (b), β -Rag DKO (c), and γ -Rag DKO (d) mice were stained with anti-CD45 mAb. (f and g) Schemes of the generation of PP-null WT (f) or PP-null-Rag2^{-/-} (g) mice. (h and i) PP-null WT mice were infected with *H. pylori*. Two months after the infection, gastric specimens were prepared. (j and k) PP-null-Rag2^{-/-} mice were infected with *H. pylori*. Two months after the infection, naive (j) or primed (k) splenic CD4⁺ T cells were transferred. Two months after the cell transfer, gastric specimens were prepared. Specimens were stained with H&E (h, j, and k) or anti-*H. pylori* antisera (brown) (l). (Scale bars: 200 μ m.)

of surface proteins and/or carbohydrates, which may make the bacteria susceptible to phagocytosis (27, 28), a subject worthy of further studies. It has been shown that mast cells are able to present *H. pylori* antigens to *H. pylori*-specific CD4⁺ T cells, which in turn activate mast cells to degranulate (29). When we infected W/W^v and SI/SI^d mice lacking mast cells with *H. pylori*, gastritis was readily induced in both strains of mice on infection (S.N., T.Y., Y.B., and S.K., unpublished data), indicating that mast cells are not essential in priming CD4⁺ T cells. In an *in vitro* experiment, BMDCs infected with the coccoid form of *H. pylori* produced larger amounts of IL-12 than those infected with the helical form of *H. pylori*, suggesting that the coccoid form of

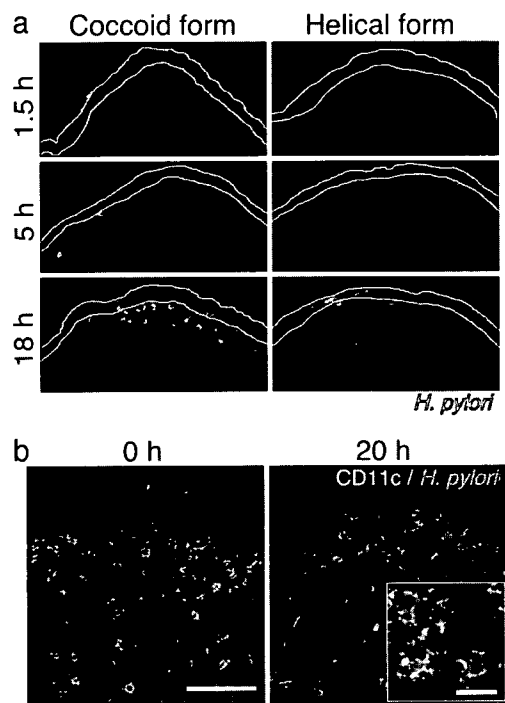


Fig. 3. The coccoid form of *H. pylori* is captured by DCs in PPs. The coccoid or helical form of *H. pylori* was inoculated into the ligated small intestines of wild-type mice. (a) After the indicated incubation times, PPs were stained with anti-*H. pylori* antibody. (b) Twenty hours after inoculation of the coccoid form, PPs were stained with anti-CD11c mAb (green) and anti-*H. pylori* antibody (red). (Scale bars: 0 h, 100 μ m; 20 h, 20 μ m.)

H. pylori easily induces Th1 immune responses on *H. pylori* infection.

Importantly, CD4⁺ T cells primed with the coccoid form of *H. pylori* were able to induce gastritis in *H. pylori*-infected GALT-null γ -Rag DKO mice where CD4⁺ T cells are not

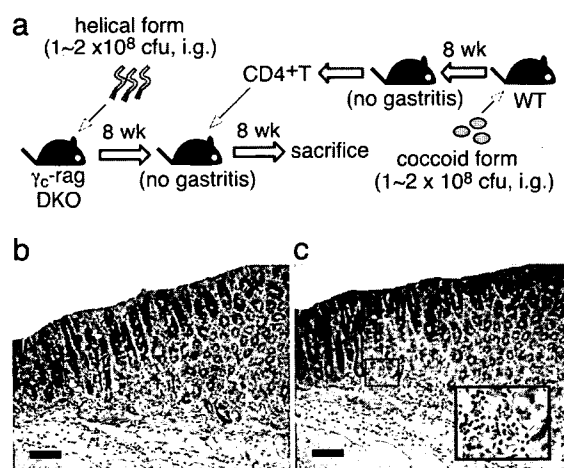


Fig. 4. Gastritis is induced by CD4⁺ T cells primed by the coccoid form of *H. pylori*. Two months after infection of γ -Rag DKO mice with the helical form of *H. pylori*, splenic CD4⁺ T cells from wild-type (WT) mice orally infected by the coccoid form of *H. pylori* were transferred to the infected γ -Rag DKO mice (a). Two months after the cell transfer, gastric specimens were prepared. Specimens were stained with H&E (b) or chloroacetate esterase (red) for infiltrated neutrophils and mast cells (c). (Scale bars: 200 μ m.)

primed with *H. pylori* antigen at all. Thus, the following scenario emerges from our results: *H. pylori* transforms to the coccoid form when entering the intestinal tract and is captured by DCs in PPs. *H. pylori* antigens presented by DCs are recognized by CD4⁺ T cells in PPs or mLN, and activated T cells migrate to the gastric mucosa to induce and maintain inflammatory responses. We noted that PP-null-Rag2^{-/-} mice exhibited modest inflammation in the stomach after naive CD4⁺ T cell transfer, compared with γ_c -Rag DKO mice, which showed no sign of inflammation. Because treatment by anti-IL-7R α mAb *in utero* suppresses PP development, but leaves ILFs and recently discovered villous M cells (30) intact, ILFs and/or villous M cells may also participate in the capture of *H. pylori* in the intestine. Although luminal antigens can be taken up by M cells located over the follicular epithelia of ILF, the tissue is predominantly occupied by B cells (31). Furthermore, in a separate study, we have shown that no apparent organized lymphoid structure is developed under villous M cells (data not shown).

H. pylori is also implicated in the cause of other diseases such as idiopathic thrombocytopenic purpura (32) and Sjögren syndrome (33). Indeed, it has been shown that T cells migrate from the intestine to the salivary gland in Sjögren syndrome patients (34). It will be of interest to examine the functional relationship between these diseases and the coccoid form of *H. pylori* captured via PPs.

Materials and Methods

Mice. All mice used in this study were on a C57BL/6 background and were maintained at Taconic (Germantown, NY) or in our animal facility under specific pathogen-free conditions. Wild-type C57BL/6 mice were purchased from Sankyo Labo Service (Shizuoka, Japan) and CLEA Japan (Tokyo, Japan). Rag2^{-/-} mice, γ_c -Rag DKO mice, and OT-II-Rag mice were obtained from Taconic. IL-2R β ^{-/-} mice (35) were generously provided by T. W. Mak (Ontario Cancer Institute, Toronto, ON, Canada). β -Rag DKO mice were obtained by crossing IL-2R β ^{-/-} with Rag2^{-/-} mice (20). All experiments were approved by the Animal Care and Use Committee of the Keio University School of Medicine and were performed in accordance with institutional guidelines.

Antibodies. Fluorescein-conjugated antibodies for flow-cytometric analysis and biotin anti-mouse CD2_c (HL3) were purchased from BD Bioscience (San Jose, NJ). Anti-*H. pylori* antibodies were purchased from Biomeda (Foster City, CA) or DAKO (Glostrup, Denmark).

Bacteria. *H. pylori* strain SS1, a mouse-adapted human isolate, was used for all experiments. To prepare the helical form of *H. pylori*, SS1 was grown on 5% sheep blood agar plates for 2 days. Before inoculating into mice, bacteria were grown in *Brucella* broth with 5% FCS overnight at 37°C under microaerobic conditions with gentle agitation. To prepare the coccoid form, SS1 was grown on 5% sheep blood agar plates under microaerobic conditions for 3 days at 37°C and then cultured under anaerobic conditions for 7 days at 37°C.

In Vivo Infection of Mice. Bacteria were prepared from logarithmic phase cultures. Mice were intragastrically infected with 1–2 \times 10⁸ cfu *H. pylori* in 0.15 ml of broth. After the indicated time period, mice were killed and the stomach was aseptically removed. The stomach was then bisected along the greater and lesser curvatures. Half of the stomach was homogenized for the determination of bacterial colonization by a plate-dilution method. The rest of the stomach was sectioned transversely into two strips for frozen and paraffin-embedded sections.

In Situ Infection of Intestinal Loop. Wild-type mice (6-week-old females) were anesthetized by an i.m. injection of 2 mg of

ketamine hydrochloride (Sankyo, Tokyo, Japan) and 0.1 mg of Xylazine per mouse. An \approx 4-cm-long piece of the small intestine containing one or two PPs was ligated at both ends with surgical thread. *H. pylori* (1 \times 10⁹) suspended in 0.2 ml of saline was inoculated into the loop. After the indicated time periods, PPs were removed and extensively washed with PBS. After fixation in 4% paraformaldehyde in PBS, specimens were processed for histopathological examination.

Generation of PP-Null Mice. PPs were depleted from a small intestine as previously described (22). Briefly, 14.5 days postcoitum pregnant wild-type or Rag2^{-/-} mice were i.v. injected with 2 mg of anti-IL-7R α mAb (A7R34; kindly provided by S.-I. Nishikawa, RIKEN CDB, Kobe, Japan). To confirm the depletion of PPs, a dissected small intestine from one of the offspring was fixed with acetone and stained with anti-B220 or anti-pan CD45 mAb (BD Biosciences).

Adoptive Transfer of CD4⁺ T Lymphocytes. Naive or *H. pylori* antigen-primed CD4⁺ T cells were purified from splenocytes, mLN, and PPs by using anti-mouse CD4-microbeads and AutoMACS (Miltenyi Biotech, Sunnyvale, CA) according to the manufacturer's instruction. The purity of isolated cells was >95%. Isolated cells (5 \times 10⁶ per mouse for splenocytes and mLN, 5 \times 10⁵ per mouse for PP-derived cells) were injected i.v. into recipient mice infected with *H. pylori* for 8 weeks. Eight weeks after the transfer, mice were killed for the indicated analyses.

Histological Analysis. An excised stomach was fixed in a neutral-buffered 10% formalin solution and cut into four strips. Samples were processed by standard methods, embedded in paraffin, and sectioned at 4 to 5 μ m. Specimens were stained with H&E or used for cytochemical and immunohistochemical studies. The Leder method was used to assess naphthol-AS-d-chloroacetate esterase detection (36). Immunohistochemical analysis was performed with formalin-fixed and paraffin-embedded tissue sections by using heat-induced epitope retrieval and the ABC (Vectastain ABC kit; Vector Laboratories, Burlingame, CA) method. Anti-*H. pylori* serum from DAKO was used for *H. pylori* staining. In some cases, frozen sections (7 μ m) were prepared, fixed with 4% paraformaldehyde in PBS, and blocked with 2% BSA-PBS, and immunofluorescence was performed using the tyramide amplification method (TSA-Plus Fluorescein System; PerkinElmer Life and Analytical Sciences, Boston, MA) and then incubated with anti-*H. pylori* antibody from Biomeda, followed by Cy5 (GE Healthcare Bioscience AB, Uppsala, Sweden) or TRITC-linked rabbit IgG (Sigma-Aldrich, St. Louis, MO). The specimens were mounted with Vectashield (Vector Laboratories) and examined with a confocal laser-scanning microscope LSM510 by using version 3.2 software (Carl Zeiss, Thornwood, NY). The zymogenic zone of middle corpus \approx 3 mm from the FS/Z transition zone was examined in each sample.

Histological Score. For assessment of gastric histopathology, blinded sections stained with H&E were examined by light microscopy. Neutrophil infiltration was assessed by the presence of neutrophils in the gastric mucosa. Active inflammation was assessed by the degree and area of damages of mucosal tissue and muscular layers because of infiltrations of neutrophils, lymphocytes, and/or macrophages. The scoring was graded as 0 (no), 1 (mild), 2 (moderate), or 3 (severe). The total number of glands with neutrophil infiltration in the crypt and lumen was also counted to produce a gland active inflammatory score.

Whole-Mount Immunohistochemistry. Small intestines were removed and stained with antibodies as described previously (37). Briefly, small intestines were washed, incubated twice in HBSS containing 5 mM EDTA at 37°C for 20 min, and fixed with ice-cold formalin