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The Actin-Bundling Protein L-Plastin Dissociates CCR7 Proximal Signaling from CCR7-Induced Motility


Chemokines promote lymphocyte motility by triggering F-actin rearrangements and inducing cellular polarization. Chemokines can also enhance cell–cell adhesion and costimulate T cells. In this study, we establish a requirement for the actin-bundling protein L-plastin (LPL) in CCR7- and sphingosine-1-phosphate–mediated T cell chemotaxis using LPL−/− mice. Disrupted motility of mature LPL−/− thymocytes manifested in vivo as diminished thymic egress. Two-photon microscopy of LPL−/− lymphocytes revealed reduced velocity and motility in lymph nodes. Defective migration resulted from defective cellular polarization following CCR7 ligation, as CCR7 did not polarize to the leading edge in chemokine-stimulated LPL−/− mice. However, CCR7 signaling to F-actin polymerization and CCR7-mediated costimulation was intact in LPL−/− lymphocytes. The differential requirement for LPL in CCR7-induced cellular adhesion and CCR7-induced motility allowed assessment of the contribution of CCR7-mediated motility to positive selection of thymocytes and lineage commitment. Results suggest that normal motility is not required for CCR7 to function in positive selection and lineage commitment. We thus identify LPL as a molecular critical for CCR7-mediated motility but dispensable for early CCR7 signaling. The requirement for actin bundling by LPL for polarization reveals a novel mechanism of regulating actin dynamics during T cell motility. The Journal of Immunology, 2010, 184: 3628–3638.
in intranodal motility strikingly similar to those of CCR7−/− T cells. Despite normal expression of the receptors CCR7 and S1P1, LPL−/− thymocytes exhibited diminished chemotaxis in vitro toward the ligands CCL19 and S1P, and mature LPL−/− T cells did not move efficiently toward CCL19. Defective migration resulted from defective cellular polarization following CCR7 ligation. LPL is thus required for the establishment or maintenance of cellular polarity that enables directed chemotaxis, without being required for initial adhesion that is required for cell motility.

The differential requirement for LPL in CCR7-induced cellular adhesion and CCR7-induced motility allowed assessment of the contribution of CCR7-mediated motility to positive selection and lineage commitment. Results suggest that normal motility is not required for CCR7 to function in positive selection and lineage commitment. We thus identify LPL as a molecule critical for CCR7-mediated motility but dispensable for early CCR7 signaling. Furthermore, the requirement for the actin-bundling protein LPL for polarization following CCR7 ligation reveals a novel mechanism of regulation of actin dynamics, at a step downstream from actin polymerization and the cytoskeletal changes required for adhesion, but nonetheless necessary for T cell motility.

Materials and Methods

Mice

The generation of mice deficient for LPL has been previously described (26). Mice were backcrossed to generate LPL−/− mice on a B6 background. n3.L2 transgenic mice were bred as described (31). Directly conjugated Abs were commercially available: CD8-FITC, CD45.2-allophycocyanin, CD4-allophycocyanin/Alexa Fluor750, CD45.1-PE/Cy7 (eBioscience, San Diego, CA), Vα2-FITC, CD24-PE, CD62L-PE, CD69-PE, CCR7-PE, CD4-allophycocyanin, (Biolegend, San Diego, CA), β7-integrin-PE, CD8-PerCP (BD Biosciences, San Jose, CA). Staining for CCR7 was performed either at phycocyanin (Biolegend, San Diego, CA), or without Hb(64–76) peptide, with or without CCL21 (100 ng/ml). Cell numbers were counted by dividing the number of migrated cells, gated as indicated, by the total number of equivalently gated input cells.

Flow cytometry

The Ab against S1P1 was generously provided by J. Cyster (University of California at San Francisco, San Francisco, CA) (30). The clonotypic Ab for the n3.12 TC (26) was originaly obtained from M. Nussenzweig (The Rockefeller University, New York, NY) (29). All animals were maintained in specific pathogen-free housing, and all experiments were performed in accordance with protocols approved by the Washington University Animal Studies Committee.

IntrathyMIC FITC injection

Intrathymic FITC injections were performed as described (32), with injection of 10 μl FITC (1 mg/ml in sterile PBS) into one lobe of the directly visualized thymus of anesthetized mice age 6–10 wk. Thymus, lymph nodes, spleens, and 100 μl blood were harvested 48 h postinjection. Cell numbers were counted by hemocytometer and subpopulations and FITC labeling determined by flow cytometry. To control for the variability intrinsic to intrathymic injection, data are presented as percentages normalized to the total number of FITC+ thymocytes and peripheral cells isolated. Percentages were calculated by dividing the number of cells per organ, gated as indicated, by the sum total number of equivalently gated cells from the thymus, lymph nodes, spleen, and blood from each mouse. Each experiment was performed on a single pair of age-matched mice with the indicated number of replicate experiments.

Transwell assays

Transmigration assays were performed as described (8) using 5 μm transwell filters (Corning Costar, Lowell, MA) with CCL19 (R&D Systems, Minneapolis, MN) or S1P (Sigma-Aldrich, St. Louis, MO) at the indicated concentrations. Postincubation for 3 h at 37°C, cells were recovered from the lower chamber and counted using a hemocytometer with subpopulations determined by flow cytometry. Percentage of migrated cells was determined by dividing the number of migrated cells, gated as indicated, by the total number of equivalently gated input cells.

Two-photon microscopy

CD8+ T cells were purified from the lymph nodes of OT-1 wild-type (WT) and OT-1 LPL−/− mice by MACS-based negative selection (Miltenyi Biotec, Auburn, CA), then labeled for 30 min at 37°C with 20–50 μM CMAC or 10 μM CMTPX (Invitrogen, Carlsbad, CA). In the experiment depicted in Fig. 3H, OT-1 WT mice were labeled blue (CMAC) and OT-1 LPL−/− T cells were labeled red (CMTPX). In a replicate experiment, OT-1 WT T cells were labeled red (CMTPX) and OT-1 LPL−/− T cells were labeled blue (CMAC) without any alteration in results. T cells (3 × 10^4–20 × 10^5) were resuspended in 200 μl PBS, adoptively transferred by tail vein injection into C11c−YFP mice (29), and allowed to home for 2 h. Exploated lymph node cells were secured to coverslips with a thin film of VetBond (3M, St. Paul, MN) and placed in a flow chamber and maintained at 37°C by perfusion with warm, high-glucose DMEM bubbled with a mixture of 95% O2, and 5% CO2. Time-lapse imaging was performed with a custom-built two-photon microscope, fitted with two Chameleon Ti:Sapphire lasers (Coherent Radiation, Santa Clara, CA) and an Olympus XLUMPlanFI 20× objective (Olympus, Melville, NY; water immersed; numerical aperture, 0.95) and controlled and acquired with ImageWarp (A&B Software, New London, CT). For imaging of YFP, excitation wavelength was 980–915 nm; for CMTPX and CMAC, 780–800 nm was used. Signals from fluorescent dyes and YFP were separated by dichroic mirrors (490 nm, 515 nm, and 560 nm). To create time-lapse sequences, we typically scanned with 2 Z-steps of 2.5 μm each at 45–55 s intervals for up to 60 min. For data analysis, cells were detected based on fluorescence intensity and cell tracks obtained with Velocity (PerkinElmer, Waltham, MA) or Imaris (Bitplane, Zurich, Switzerland) software. Only cells that could be tracked for at least eight time points were included in the analysis. The median of instantaneous velocities in each cell track was reported as the velocity for that cell. Motility coefficients (mm²/min) were calculated for individual tracks by linear regression of displacement versus time plots with T Cell Analysis (John Dempster, University of Strathclyde, Glasgow, Scotland, U.K.).

Generation of bone marrow chimeras

Bone marrow was harvested from WT and LPL−/− mice, mixed in a 1:1 ratio, and injected retro-orbitally into sublethally irradiated (500 rad) RAG−/− mice. After 5 to 6 wk, mice were sacrificed, and thymocytes, PBMCs, lymph node cells, and splenocytes were assessed for expression of CD4, CD8, CD45.1, CD45.2, CD69, and CD24 by flow cytometry.

Rac assays

CD4+ T cells isolated from WT and LPL−/− mice were rested overnight in reduced serum medium (Opti-MEM, Invitrogen), then stimulated with CCL19 and CCL21 (100 ng/ml; R&D Systems) for 15 s. Cell lysates were generated and assayed for GFP-Rac using the Rac1,2,3 Activation Assay G-LISA kit (Cytoskeleton, Denver, CO).

F-actin content

CD4+ T cells were isolated from lymph nodes with magnetic beads (Miltenyi Biotec) and were stimulated in suspension with CCL19 (100 ng/ml). Thymocytes were stained with CD8-PerCP and CD4-allophycocyanin and sorted (FACS/Aria, BD Biosciences) to isolate CD4SP cells, which were then stimulated in suspension with CCL19 and CCL21 (100 ng/ml each) for 20 min. Cells were fixed in 3.6% paraformaldehyde, then permeabilized with 0.1% Triton-X 100. F-actin content was determined by incubation with Alexa Fluor 488-phalloidin (Invitrogen) followed by flow cytometry (33).

Cell conjugation assays

Cell conjugation assays were performed as described (13) with minor modifications. CH27 B cells (H-2b, I-Eb) were used as APCs (34). MACS bead-purified CD4+ T cells from n3.L2 WT and n3.L2 LPL−/− mice were labeled with CellTrace Far Red DDAO (Invitrogen), and CH27 cells were labeled with CFSE (Invitrogen). Cells were mixed and incubated with or without Hb(64–76) peptide, with or without CCL21 (100 ng/ml), and with or without blocking anti–LEA-1 Ab (anti-C11a clone M174; Biolegend; 10 μg/ml) for 30 min at 37°C. Cells were fixed with 2% paraformaldehyde and percentage of T cells that formed conjugates determined by flow cytometry.
**CD69 upregulation**

The agonist Hb(64–76) peptide, sequence GKKVITAFNEGLK, was synthesized, purified, and analyzed as previously described (31). MACS bead-purified CD4+ T cells isolated from n3.L2 WT and n3.L2 LPL−/− lymph nodes were incubated overnight with congenic splenocytes with or without the indicated concentration of Hb peptide and with or without the chemokines CCL21 or CCL19 (100 ng/ml). Upregulation of CD69 on n3.L2+ CD4+ T cells was assessed by flow cytometry.

**Confocal microscopy**

Coverslips were coated with 10 μg/ml recombinant mouse ICAM-1/Fc chimera (R&D Systems). Cells were incubated on coated coverslips with or without CCL19 as indicated, then fixed with 4% paraformaldehyde for 20 min at RT. Cells were stained with anti-CD43 (unconjugated; BD Pharmingen) or anti-CCR7 [either unconjugated (eBioscience) or biotinylated (Biolegend)] for a minimum of 30 min at RT prior to permeabilization. Either goat anti-rat Ig Alexa Fluor 546 (Invitrogen) or streptavidin-Alexa Fluor 546 was used as a secondary. Cells were permeabilized with 0.5% Triton X-100 for 4 to 5 min at RT. Actin was stained with Alexa Fluor 488-phalloidin (Invitrogen). LPL was stained with mAb 12A2 (16.5 μg/ml). The mAb 12A2 was generated by immunizing LPL−/− mice with recombinant human LPL and screened for binding to recombinant murine LPL by ELISA. Specificity of binding was confirmed by immunoblot and immunofluorescence, with WT and LPL−/− cells as positive and negative controls (data not shown). The secondary Ab for anti-LPL was goat anti-mouse–Alexa Fluor 546. Confocal and differential interference contrast images were acquired using the Zeiss LSM 510 microscope (Zeiss, Oberkochen, Germany) fitted with a 1.3-narrow aperture ×40 Fluor objective. For quantitation, images of each cell in at least two randomly selected fields of each sample were acquired. Cells that appeared dead or were in contact with other cells were excluded from analysis. Images were randomized, and polarization of each cell was determined by an independent, blinded observer.

**Statistics**

For normally distributed data, either paired or unpaired Student t test was used to determine statistical significance, with p < 0.05 considered significant. If data were not normally distributed, then the Mann-Whitney or Wilcoxon ranked sum test was used. All statistical analyses were performed using GraphPad Prism version 4 software (GraphPad, La Jolla, CA).

**Results**

Mature thymocytes accumulated in n3.L2 LPL−/− mice due to diminished thymocyte egress

LPL is upregulated as thymocytes successfully undergo positive selection along with CCR7 and other molecules associated with motility, such as gelsolin (23). Neutrophil migration was not affected in LPL−/− mice (26), but no role for LPL in T cell motility and development has yet been explored. To determine how LPL might be required for thymocyte maturation and subsequent lymphocyte motility, LPL−/− mice transgenic for the n3.L2 TCR were generated.

**FIGURE 1.** Mature n3.L2 CD4SP thymocytes accumulate in n3.L2 LPL−/− mice. Flow cytometry of CD4 and CD8 (A) and n3.L2 TCR (β) expression on thymocytes from n3.L2 WT (gray histogram) and n3.L2 LPL−/− (solid line) mice. C. Number of total and n3.L2+ CD4SP thymocytes from n3.L2 WT (gray; n = 10) and n3.L2 LPL−/− (filled; n = 11) mice analyzed in 10 independent experiments. Each symbol represents the value for an individual mouse. Mean indicated by bar; p value determined with Wilcoxon signed rank test. D. Expression of the maturation markers CD69, CD24, CD62L, and β7-integrin on n3.L2 CD4SP thymocytes from n3.L2 WT (gray histogram) and n3.L2 LPL−/− (solid line) mice. E. CD4 and CD8 expression of lymphocytes from peripheral lymph nodes of n3.L2 WT and n3.L2 LPL−/− mice. F. Number of total and CD4+ cells isolated from four lymph nodes of n3.L2 WT (gray; n = 10) and n3.L2 LPL−/− (filled; n = 10) mice, analyzed in 10 independent experiments. Each symbol represents the value for an individual mouse. Mean indicated by bar; p values determined using Wilcoxon signed rank test. A, B, D, and E. Flow cytometry representative of at least seven pairs of mice analyzed in seven independent experiments.
Use of a transgenic TCR model enabled a more detailed analysis of the maturation of thymocytes with a defined TCR specificity. The n3.L2 TCR recognizes Hb(64–76)/I-Ek, and thymocyte development in n3.L2 mice has been well characterized (28, 31).

The number and percentage of CD4SP, TCR-high thymocytes was dramatically increased in n3.L2 LPL−/− mice compared with n3.L2 WT mice, though the total number of thymocytes was not affected (Fig. IA–C). The accumulated n3.L2 CD4SP LPL−/− thymocytes were more phenotypically mature than n3.L2 CD4SP WT thymocytes, as they were CD69+RBC, CD24low, CD62Lhigh, and β7-integrinhigh (Fig. 1D). The accumulation of phenotypically mature, TCR-high, CD4SP thymocytes in n3.L2 LPL−/− mice suggested a defect in thymocyte egress, as the same phenotype has been observed in other systems in which thymic egress is diminished (7, 8, 33, 35). The finding of smaller thymic egress, as the same phenotype has been observed in other systems in which thymic egress is diminished (7, 8, 33, 35). The finding of smaller thymic egress, as the same phenotype has been observed in other systems in which thymic egress is diminished (7, 8, 33, 35). The finding of smaller thymic egress, as the same phenotype has been observed in other systems in which thymic egress is diminished (7, 8, 33, 35). The finding of smaller thymic egress, as the same phenotype has been observed in other systems in which thymic egress is diminished (7, 8, 33, 35).

Intrathymic injection of FITC confirmed that LPL deficiency resulted in diminished thymic egress. The percentage and number of FITC-labeled n3.L2 CD4+ T cells recovered from peripheral blood, lymph nodes, and spleens of intrathymically injected n3.L2 LPL−/− mice were reduced (Fig. 2A, 2B, and data not shown). Reduction of the number of FITC-labeled cells from peripheral blood suggests that the accumulation of n3.L2 CD4SP thymocytes and relative paucity of CD4+ T cells in n3.L2 LPL−/− mice was due to decreased thymocyte egress and not to reduced entry into peripheral lymphoid organs. B cells recovered from the spleens of mice injected intrathymically with FITC were not labeled with FITC (data not shown), indicating that FITC-positive T cells from the peripheral lymphoid organs were recent thymic emigrants and not nonspecifically labeled during injection. Thus, intrathymic FITC injection confirmed diminished thymocyte egress in n3.L2 LPL−/− mice.

**Diminished in vitro motility of n3.L2 LPL−/− thymocytes**

CCR7 and the chemokine ligand CCL19 regulate thymic egress in newborn mice (4), and S1P1 and its ligand S1P are absolutely required for thymocyte egress (7, 8). Thymocytes from n3.L2 LPL−/− mice expressed normal levels of the receptors S1P1 and CCR7 (Fig. 2C). We hypothesized that the failure of thymic egress in n3.L2 LPL−/− mice was due to a failure to migrate toward S1P1 and CCL19 and therefore assessed in vitro motility of mature CD4SP thymocytes using transwell chemotaxis assays. Mature (CD62L<sup>bhigh</sup>) CD4SP thymocytes from n3.L2 WT mice migrated toward CCL19 and S1P as expected, based on previously published reports (8, 36). In contrast, mature CD4SP thymocytes from n3.L2 LPL−/− mice demonstrated a severe defect in migration toward both chemotactants (Fig. 2D). Defective migration toward chemotactants explains the observed defect in thymic egress.

**Diminished motility of mature OT-1 LPL−/− T cells**

CCR7 and its ligands have been demonstrated to regulate the motility of mature T cells in lymph nodes, as both the velocity and motility coefficient of T cells was reduced in the absence of the receptor CCR7 or its ligands CCL19 and CCL21 (5, 6). If LPL is required for naive T cell motility in response to CCR7 ligands, then LPL−/− lymphocytes should demonstrate intranodal motility defects similar to those found in CCR7−/− lymphocytes. To test this hypothesis, we generated OT-1 LPL−/− mice. The OT-1 receptor is restricted to the H-2<sup>d</sup> background, which enabled the use of the CD11c-YFP H-2<sup>d</sup> mice and allowed simultaneous visualization of lymphocytes and dendritic cells.

We first determined that the OT-1 LPL−/− mouse exhibited a similar phenotype to the n3.L2 LPL−/− mouse. There was a relative increase in mature (CD62L<sup>bhigh</sup>) TCR-high, CD8SP cells in the OT-1 LPL−/− thymus (Fig. 3A–C). TCR<sup>high</sup> CD8SP thymocytes from OT-1 LPL−/− mice were phenotypically more mature (CD69<sup>pos</sup>, CD24<sup>low</sup>) than those isolated from the OT-1 WT mouse (Fig. 3D). Fewer mature CD8+ T cells were isolated from the periphery of OT-1 LPL−/− mice (Fig. 3E). Intrathymic FITC injection demonstrated diminished thymic egress in OT-1 LPL−/− mice (Fig. 3F). CD8SP thymocytes from OT-1 LPL−/− mice exhibited defective in vitro motility toward CCL19 and S1P, despite comparable levels of expression of the

**FIGURE 2.** Diminished thymic egress and diminished in vitro motility of n3.L2 CD4SP LPL−/− thymocytes. A, Labeling of n3.L2 CD4+ T cells recovered from peripheral blood, lymph nodes, and spleens of intrathymically injected n3.L2 LPL−/− mice (48 h after intrathymic injection of FITC (representative of four independent experiments). B, Normalized percentage of FITC<sup>−</sup> n3.L2 CD4+ T cells recovered from the blood and lymph nodes of mice injected with FITC intrathymically. Each symbol represents the value from one mouse. Mean represented by bar; p value determined using Mann-Whitney U test. Data from four independent experiments each with a pair of age-matched mice. C, Equivalent expression of S1P1 and CCR7 on n3.L2 CD4SP thymocytes from n3.L2 WT and n3.L2 LPL−/− mice. Expression of S1P1 and CCR7 on DP thymocytes from n3.L2 WT mice included as negative control. Representative of at least two independent experiments. D, Transwell migration of mature (CD62L<sup>bhigh</sup>) CD4SP thymocytes from n3.L2 WT (gray bars) and n3.L2 LPL−/− (filled bars) mice toward the chemotactants CCL19 and S1P. Data shown are mean ± SEM of duplicate or triplicate samples within a single experiment; representative of at least 3 independent experiments. Value of p determined by unpaired t test.
FIGURE 3. OT-1 LPL−/− mice exhibit similar phenotypic defects as n3.L2 LPL−/− lymphocytes exhibit diminished intranodal motility. Expression of CD4 and CD8 (A) and Vα2 (B) on thymocytes from OT-1 WT (gray) and OT-1 LPL−/− mice (solid line). C, Number of total, CD62Llow Vα2high CD8SP, and CD62Lhigh Vα2high CD8SP thymocytes from OT-1 WT (gray; n = 11 or 10) and OT-1 LPL−/− (filled; n = 12 or 11) mice. Each symbol represents the value from an individual mouse; p value from unpaired t test. D, Expression of CD69 and CD24 on Vα2high CD8SP thymocytes from OT-1 WT (gray) and OT-1 LPL−/− (solid line) mice. E, Numbers of total and CD8+ T cells from lymph nodes of OT-1 WT (gray; n = 8) and OT-1 LPL−/− (filled; n = 8) mice 48 h after intrathymic FITC injection. Each symbol represents the value from individual mouse; data from four independent experiments; p value determined using Mann-Whitney test. F, Normalized percentage of FITC-labeled Vα2 CD8+ cells recovered from peripheral blood and lymph nodes of OT-1 WT and OT-1 LPL−/− mice. Data shown are mean ± SEM of duplicate or triplicate samples with p values determined by unpaired t test; representative of two independent experiments. G, Vα2 CD8SP thymocytes from OT-1 LPL−/− mice did not migrate efficiently in transwell assays toward CCL19 or S1P. Data shown are mean ± SEM of duplicate or triplicate samples with p values determined by unpaired t test; representative of two independent experiments. H, Two-photon time-lapse image sequences of CD8+ OT-1 WT (blue) and CD8+ OT-1 LPL−/− (red) cells in naive lymph nodes 2 h post-injection into CD11c-YFP mice. Dendritic cells appear green. Representative cell tracks are shown (white lines). Scale bar, 20 μm. I, Velocity (mean OT-1 WT, 7.8 μm/min; mean OT-1 LPL−/−, 5.3 μm/min), motility (mean OT-1 WT, 104 μm²/min; mean OT-1 LPL−/−, 61 μm²/min), and meandering index of CD8+ cells from OT-1 WT (gray circles; n = 56) and OT-1 LPL−/− (filled circles; n = 55) mice. Each point represents a single cell tracked for a minimum of eight frames. Mean of each population is indicated, with p values determined by Mann-Whitney U test. Data pooled from two independent experiments.
receptors CCR7 and S1P (Fig. 3G and data not shown). LPL is thus required for motility of both CD4 and CD8 T lymphocytes.

Intranodal motility of OT-1 LPL−/− T cells and OT-1 WT T cells was then compared using two-photon microscopy (Fig. 3H, Supplemental Video). As predicted, CD8+ T cells isolated from OT-1 LPL−/− mice demonstrated reduced velocity and motility. The degree to which the velocity and motility of LPL−/− lymphocytes was reduced was comparable to published findings with CCR7−/− mice.

FIGURE 4. Nontransgenic LPL−/− cells demonstrate in vitro motility defects, though the in vivo phenotype of nontransgenic LPL−/− mice varies from that of transgenic LPL−/− mice. A, Peripheral T cells isolated from LN of nontransgenic LPL−/− mice do not migrate efficiently in transwell assays toward CCL19 (100 ng/ml). Data shown are the mean ± SEM of triplicate samples in an individual experiment; representative of three independent experiments. *p values determined by unpaired t test. B, Number of total, CD4SP, and CD8SP thymocytes recovered from nontransgenic WT (gray circles; n = 6) and LPL−/− (filled circles; n = 6) mice. Each symbol represents the value from one mouse; data from six independent analyses; mean represented by bar. C, Expression of CD4 and CD8 on thymocytes from nontransgenic WT and LPL−/− mice. D, Expression of the maturation markers CD69, CD24, and CD62L on CD4SP and CD8SP thymocytes from nontransgenic LPL−/− (solid line) and WT (gray histogram) mice. E, Number of total, CD4+, and CD8+ cells recovered from four lymph nodes of nontransgenic WT (gray circles; n = 7) and LPL−/− (filled circles; n = 7) mice. Each symbol represents the value from one mouse; mean represented by bar; data from seven independent analyses. Flow cytometry in C and D represents at least four pairs of mice.

FIGURE 5. LPL−/− SP thymocytes are at a competitive disadvantage during thymic egress. A, CD4 and CD8 expression of thymocytes isolated from sublethally irradiated RAG1−/− mouse reconstituted with bone marrow from WT (CD45.1+) and LPL−/− (CD45.2+) mice mixed in a 1:1 ratio. B, Percentage of thymocytes derived from either WT (CD45.1+) or LPL−/− (CD45.2+) donors that were CD4SP or CD8SP. C, CD69 expression on CD4SP or CD8SP thymocytes derived from either WT (CD45.1+) or LPL−/− (CD45.2+) donors. D, Percentage of CD4SP or CD8SP thymocytes that were CD69+ derived from either WT (CD45.1+) or LPL−/− (CD45.2+) donors. E, Ratio of CD4SP:CD8SP thymocytes derived from WT (CD45.1+) or LPL−/− (CD45.2+) donors. F, Percentage of DP thymocytes that were CD69+ derived from either WT (CD45.1+) or LPL−/− (CD45.2+) donors. A and C, Representative data from one of nine chimeric mice shown. C–F, Each symbol represents data from one chimeric mouse; mean represented by bar; data from four independent experiments. Values of *p determined using Wilcoxon signed rank test.
lymphocytes (6). CCR7−/− lymphocytes demonstrated a minor but statistically significant reduction in the meandering index, which measures the degree to which a cell’s movement varies from a straight line. We observed a very slight decrease in the meandering index of LPL−/− lymphocytes, though the difference was not significant. LPL was thus required for normal mature CD8+ T cell motility in lymph nodes, and the reduction in LPL−/− lymphocyte motility was consistent with loss of CCR7-mediated motility.

Nontransgenic T cells are defective in CCR7-mediated motility

The motility of nontransgenic lymphocytes was also examined (Fig. 4). Both CD4+ and CD8+ mature T cells from nontransgenic LPL−/− mice demonstrated diminished motility toward CCL19 in chemotaxis transwell assays (Fig. 4A), indicating that defective motility in LPL−/− cells was not due to the transgenic system. However, the in vivo phenotype of nontransgenic LPL−/− mice differed slightly from that of the transgenic mice, in that the total number of thymocytes recovered from LPL−/− mice was reduced (Fig. 4B) and the percentage of CD4SP and CD8SP thymocytes was not increased (Fig. 4C). Similar to LPL−/− mice expressing transgenic TCRs, the SP thymocytes recovered from nontransgenic LPL−/− mice were CD69low, CD24low, and CD62Lhigh and thus phenotypically more mature (Fig. 4D). This phenotype was observed with temporary inhibition of S1P1. Treatment with the S1P1-selective agonist SEW2871 increased the proportion of SP thymocytes exhibiting a mature phenotype without changing the total percentage of SP thymocytes (10). Furthermore, there was a paucity of mature T cells isolated from the lymph nodes of nontransgenic LPL−/− mice (Fig. 4E). Although the in vivo phenotype differed in some respects in the nontransgenic LPL−/− mouse, the core findings of the increase of phenotypically mature thymocytes, smaller lymph nodes, and defective in vitro CCR7-mediated motility remained constant across n3.L2, OT-1, and nontransgenic LPL−/− mice.

The participation of CCR7 in positive selection and subsequent maturation of thymocytes has been described (1, 2). Overexpression of CCR7 in thymocytes increased the commitment of positively selected thymocytes to the CD8SP lineage. Increased motility generated by increased CCR7 expression was hypothesized to contribute to the increase in commitment to the CD8SP lineage (2). To reveal stages of thymocyte development and maturation that might be affected by diminished motility toward CCR7, we

![Figure 6](http://www.jimmunol.org/)

***FIGURE 6.*** Early CCR7 signaling and CCR7-mediated TCR costimulation are not dependent upon LPL. A, Levels of GTP-Rac in CD4+ T cells from WT (gray circles) and LPL−/− (filled circles) mice incubated with or without CCR7 ligands for 15 s. Duplicate samples indicated with symbols and mean indicated by bars; representative of two independent experiments. B, F-actin content of CD4+ cells isolated from peripheral LN of WT (gray circles) and LPL−/− (filled circles) mice stimulated with CCL19 (100 ng/ml) and of CD4SP thymocytes sorted by FACS from WT (gray circles) and LPL−/− (filled circles) mice stimulated with CCL19 and CCL21 (100 ng/ml each). Data normalized to unstimulated cells. Each symbol represents results from one of two independent experiments with mean indicated by bars. C, Conjugate formation of CFSE-labeled CD4+ T cells and DDAO-labeled CH27 cells incubated ± cognate peptide and ± CCL19 plus CCL21. Data shown as percentage of T cells that formed conjugates. D, Percent of cells positive for CD69 following overnight stimulation with APCs ± cognate peptide and ± CCL21. C and D, Symbols represent replicate samples within one experiment, and bars represent means. Representative of at least two independent experiments.
Ligation of G protein-coupled receptors by chemoattractants results in a rapid burst of actin polymerization. This initial burst of polymerization is dependent upon signaling through the small GTPase CCR7, which is required for thymocyte motility. However, CCR7-dependent motility is only partially reconstituted in the context of mixed bone marrow chimeras, indicating that other factors may also contribute to thymocyte migration. These findings suggest that CCR7 plays a critical role in thymocyte egress and that additional factors may be involved in determining the final pattern of CD4–CD8 T cell ratios in the periphery.
Rac. Cells unable to activate Rac in response to chemoattractant ligation demonstrate deficient initiation of F-actin polymerization and defective motility (33, 37, 38). Rapid activation of Rac following CCR7 stimulation with CCL19 was not inhibited in LPL−/−T cells (Fig. 6A). Furthermore, the rapid burst of F-actin polymerization following CCL19 ligation was observed in LPL−/−CD4+ T cells and CD4SP thymocytes (Fig. 6B). LPL deficiency did not disrupt signaling immediately proximal to CCR7 ligation, nor did it prevent the initiation of F-actin polymerization.

Costimulation of TCR signaling is intact in LPL−/− T cells

Ligation of CCR7 has been demonstrated to costimulate TCR signaling (13, 39). CCR7 costimulation depends upon CCR7-mediated increases in LFA-1 affinity. Increased affinity of LFA-1 increases the likelihood and/or duration of T cell–DC contacts that then enable TCR engagement and signaling (13). In both WT and LPL−/− n3.L2 T cells, CCR7 ligation enhanced the formation of peptide-specific T cell–APC conjugates (Fig. 6C) and increased the upregulation of CD69 following TCR engagement (Fig. 6D). The function of CCR7 as a costimulatory molecule did not depend upon LPL.

Failure of CCR7 polarization in LPL−/− T cells following CCL19 stimulation

Acquisition of a motile phenotype correlates with polarization of the T cell. Chemokine receptors, activated integrins, and a concentration of F-actin can be found at the lamellipod of the polarized T cell, whereas markers such as CD43 and CD44 are found in the uropod (16, 40–43). Cells unable to polarize in response to chemokine stimulation exhibit motility defects (16, 43). Following chemokine stimulation, LPL colocalized with F-actin and appeared to be concentrated with F-actin in the lamellipod of CCL19-stimulated T cells (Fig. 7A). LPL was also present in the uropod and was therefore not itself polarized. However, in the absence of LPL, many fewer T cells exhibited polarized F-actin (Fig. 7B) upon CCL19 stimulation. Colocalization of polarized CCR7 and F-actin was determined in WT and LPL−/− cells stimulated with CCL19 (Fig. 7C). Fewer LPL−/− stimulated cells demonstrated polarized, colocalized CCR7 and F-actin (Fig. 7D). Cellular polarization was further assessed in LPL−/− T cells by staining for F-actin and the uropod marker CD43 (Fig. 7E). Although some polarization of LPL−/− T cells occurred in response to CCL19 stimulation, the percentage was reduced (Fig. 7F). Without affecting Rac activation or F-actin polymerization, LPL deficiency resulted in diminished CCR7-mediated polarization of T cells (Fig. 8). Failure to polarize in response to chemokine stimulation would explain the defective motility observed in LPL−/− T cells.

Discussion

The actin-bundling protein LPL has no previously identified function beyond stabilization of higher-order structures of F-actin microfilaments. In this study, we demonstrate that LPL is required for CCR7-mediated polarization and T cell motility, but not early CCR7 signaling and CCR7-mediated increased intracellular adhesion (Fig. 8). The function of LPL as an actin-bundling protein distinguishes LPL from other actin-binding proteins required for lymphocyte motility that primarily regulate early chemokine receptor signaling (17). For instance, knockdown of cortactin resulted in diminished ERK signaling and diminished motility upon CXCR4 ligation (44). Mice deficient for the formin mDia1, which promotes F-actin polymerization, exhibited defective thymic egress and impaired lymphocyte trafficking (20). A mutation in coronin1A, a protein that regulates the Arp2/3 complex and thus actin polymerization, has recently been described as the molecular defect underlying the phenotype of delayed thymic egress in the cataract Shionogi mouse strain (19, 22).

LPL differed from these actin-binding proteins, as LPL was not required for proximal CCR7 signaling to the initiation of F-actin polymerization. Instead, LPL deficiency resulted in a failure to either generate or maintain T cell polarization. Phalloidin staining of total cellular F-actin has been sufficiently sensitive to detect the defects in actin polymerization associated with impaired chemokine receptor proximal signaling (33). However, subtle defects in actin dynamics in LPL−/− cells have not been excluded. It is possible that actin bundling is required for local maintenance or regulation of actin polymerization that is not apparent at the whole-cell level. The disruption of higher-order microfilament structures in the presence of intact Rac activation and initiation of F-actin polymerization in LPL−/− lymphocytes illustrates a previously unrecognized level of regulation of actin dynamics.

Defective motility of LPL−/− lymphocytes manifested in vivo as diminished thymic egress and reduced intranalonal velocity and motility. Thymic egress is regulated by at least two chemoattractant receptors, S1P1 and CCR7. Consistent with a defect in thymocyte egress, nontransgenic, n3.L2, and OT-1 LPL−/− mice all demonstrated a shift of SP thymocytes to a more mature phenotype (CD69low, CD24low, and CD62Lhigh) (7, 10). The most dramatic phenotype was noted in n3.L2 LPL−/− mice, with a tripling of the number of n3.L2CD4SP thymocytes recovered from n3.L2 LPL−/− mice. There was no increase in percentage or number of CD4SP or CD8SP thymocytes in nontransgenic LPL−/− mice. Selective agonism of S1P1 with SEW2871 also resulted in a shift of SP thymocytes to a more mature phenotype without an overall increase in the percentage or number of SP thymocytes (10). Mixed bone marrow chimeras revealed a competitive disadvantage during egress for mature SP thymocytes derived from nontransgenic LPL−/− donors, as the percentage of LPL−/−/derived mature SP thymocytes was increased compared with WT-derived mature SP thymocytes. Defective thymocyte egress was confirmed by intrathymic FITC injection in both n3.L2 and OT-1 LPL−/− mice. Use of two distinct and well-characterized transgenic models makes it unlikely that the observed defect in thymocyte egress in LPL−/− mice is unique to a single transgenic system or restricted to the CD4 or CD8 lineage. Cells deficient for LPL from both transgenic and nontransgenic mice failed to migrate normally in transwell assays.
to the chemoattractants CCL19 and S1P, so it is unlikely that expression of transgenic TCRs altered cell-intrinsic actin dynamics. More likely, altered actin dynamics manifested differently during thymic maturation in the presence of transgenic TCRs.

Two-photon microscopy of LPL−/− mature T lymphocytes revealed motility defects that phenocopied the defects in velocity and motility of CCR7−/− T lymphocytes in lymph nodes. Velocity and motility of LPL−/− T cells were reduced to a similar degree as CCR7−/− T cells (6). We did not find as great a decrease in meandering index in LPL−/− T cells as was seen in CCR7−/− T cells. This difference in results may be dependent on different experimental conditions or possibly that LPL is required for the cytoskeletal apparatus that functions in speed or forward motion mediated by CCR7, but is not required for the apparatus required for changes in direction. In support of this latter possibility is the observation that proximal CCR7 signaling was not impaired in LPL−/− T cells. LPL was also dispensable for CCR7-mediated promotion of cell–cell interactions and costimulation of TCR signaling. LPL was required, however, for efficient polarization of T cells downstream of CCR7 ligation. Thus, LPL is essential for normal in vivo motility of T cells and for CCR7-induced cellular polarization required for directed motility.

In addition to regulating thymocyte egress in some systems and intranodal motility of mature T cells, CCR7 and its ligands CCL19 and CCL21 have been implicated in other processes critical to thymocyte development, such as positive selection, corticomedullary migration of positively selected thymocytes, lineage commitment, and negative selection (1, 2, 45–48). CCR7 is upregulated during positive selection and appears to promote thymocyte–DC contacts during this process (1). Overexpression of CCR7 promoted commitment to the CD8SP lineage, and increased motility due to CCR7 overexpression was hypothesized to drive this lineage choice (2). We used the ability of LPL to dissociate CCR7-mediated cell–cell interactions from CCR7-induced motility (Fig. 8) to probe which downstream functions of CCR7 were critical to positive selection and lineage commitment. In competitive bone marrow chimeric mice, we found no inhibition of positive selection or alteration of lineage commitment of LPL−/− thymocytes. These results indicate that cell–cell interactions promoted by CCR7 ligation contribute to the processes of positive selection and lineage commitment. Intact positive selection in LPL−/− thymocytes confirms results in which CCR7 overexpression increased thymocyte–DC interaction during positive selection but did not increase the motility of positively selected thymocytes (1).

Although many of the molecules required for chemokine signaling and T cell polarization have been identified, the mechanisms by which proximal signaling is linked to cytoskeletal rearrangements and later stages of T cell polarization remain unclear (17, 21). Initial signaling pathways dependent upon small GTPases such as Rap1, Rac, and cdc42 have been described. Rac activation results in the formation of lamellipodia. Polarization occurs after the stabilization of one of the lamellipodia and establishment of an anterior–posterior axis (17). The localization of LPL to the lamellipod suggests that a possible function for LPL is the stabilization of the lamellipod that then allows the generation of polarity. In the absence of LPL, lamellipodia cannot be adequately stabilized and polarity cannot be established or maintained. Alternatively, LPL may be required to stabilize the actomyosin cytoskeleton that supports retrograde flow toward the uropod (49). Further work will be required to investigate these possibilities. Interestingly, some polarization occurs in the absence of LPL. This partial reduction of polarization is reflected in the phenotype of LPL−/− mice, as thymic egress is reduced, as is seen in mice treated with FTY720 or SEW2871, but not completely ablated, as is seen in either KLF2−/− or S1P1−/− mice (7, 8, 10, 35). Either other actin-binding proteins can partially compensate for the loss of LPL or polarization is less stable in LPL−/− cells.

Whether LPL has a general role in T cell polarization and is therefore required for other functions dependent upon polarization, such as cytokine secretion and delivery of cytotoxic granules, is also under study. In mature human T cells, phosphorylation of LPL enabled the upregulation of activation markers following TCR stimulation (50). We did not see a requirement for LPL in the upregulation of CD69 following TCR ligation, as might have been predicted (50). The difference in observed results is likely due to different experimental systems, as the previous work was performed by overexpression of a nonphosphorylatable LPL mutant in human PBLs, and we are examining murine cells genetically deficient for LPL.

Although T lymphocytes deficient for LPL demonstrated both in vivo and in vitro motility defects, neutrophil deficiency for LPL did not (26). In fact, LPL was not required for adhesion, spreading, or in vivo or in vitro migration of neutrophils. LPL is thus similar to several other proteins involved in chemokine signaling or motility, such as ERK or dedicator of cytokinesis 2, that are used differently by neutrophils than by lymphocytes (17, 51). Nonetheless, it is intriguing to speculate that the role for LPL in cell polarization demonstrated in this study underlies the integrin signaling defect previously demonstrated in LPL−/− neutrophils.

In summary, LPL is required for maximal polarization of the chemokine receptor CCR7 in T lymphocytes in response to stimulation with ligand CCL19. In the absence of CCR7 polarization, T cells fail to fully polarize and fail to migrate toward chemokine. Failure to respond to the chemoattractants CCL19 and S1P in vitro correlated with in vivo motility defects; LPL−/− T cells demonstrated reduced intranodal motility, and thymic emigration of mature LPL−/− thymocytes was diminished. The requirement for the actin-bundling protein LPL in normal T cell motility reveals a mechanism beyond actin polymerization for regulating actin dynamics at the level of higher-order actin structures.

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Disclosures
The authors have no financial conflicts of interest.

References


Legend for Supplemental Material for Morley et al., "The actin-bundling protein L-plastin dissociates CCR7 proximal signaling from CCR7-induced motility"

**Supplemental movie.** OT-1 LPL\(^{-/-}\) lymphocytes exhibited decreased velocity and motility in explanted lymph nodes. CD8\(^{+}\) T cells isolated from OT-1 WT (blue) and OT-1 LPL\(^{-/-}\) (red) mice were injected into CD11c-YFP mice and allowed to home for two hours. Explanted lymph nodes were imaged as described in the Methods. The movie was generated by combining time lapse images taken approximately every 56 seconds. Time stamp is in lower left corner. CD11c\(^{+}\) cells appear green. A representative 22 minute-clip of a 56 minute video is shown.