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# Identification of Proliferating Dendritic Cell Precursors in Mouse Blood

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## Summary

While it has been known that dendritic cells arise from proliferating precursors in situ, it has been difficult to identify progenitors in culture. We find that aggregates of growing dendritic cells develop in cultures of mouse blood that are supplemented with granulocyte/macrophage colony-stimulating factor (GM-CSF) but not other CSFs. The dendritic cell precursor derives from the Ia-negative and nonadherent fraction. The aggregates of developing dendritic cells appear at about 1 wk of culture, with 100 or more such clusters being formed per  $10^6$  blood leukocytes. The aggregates can be dislodged and subcultured as expanding clusters that are covered with cells having the motile sheet-like processes ("veils") of dendritic cells. By about 2 wk, large numbers of single, major histocompatibility complex (MHC) class II-rich dendritic cells begin to be released into the medium. Combined immunoperoxidase and [ $^3$ H]thymidine autoradiography show that the cells that proliferate within the aggregate lack certain antigenic markers that are found on mature dendritic cells. However, in pulse-chase protocols, the [ $^3$ H]thymidine-labeled progeny exhibit many typical dendritic cell features, including abundant MHC class II and a cytoplasmic granular antigen identified by monoclonal antibody 2A1. The progeny dendritic cells are potent stimulators of the mixed leukocyte reaction and can home to the T-dependent areas of lymph node after injection into the footpads. We conclude that mouse blood contains GM-CSF-dependent, proliferating progenitors that give rise to large numbers of dendritic cells with characteristic morphology, mobility, phenotype, and strong T cell stimulatory function.

The immune system contains a system of dendritic cells that is specialized to present antigens and initiate several T-dependent immune responses. A good deal is known about the tissue distribution of dendritic cells (reviewed in reference 1). Dendritic cells are found in nonlymphoid organs either close to body surfaces, as in the skin and airways, or in interstitial regions of organs like heart and liver. Dendritic cells, possibly under the control of the cytokine GM-CSF, can undergo a maturation process that does not entail cell proliferation (2, 3). Initially, the cells process and present antigens most likely on abundant, newly synthesized MHC class II molecules, and then strong accessory and cell-cell adhesion functions are acquired (4–7). Dendritic cells can migrate via the blood and lymph to lymphoid organs (8–10). There, presumably as the "interdigitating" cells of the T area (8, 11–13), antigens can be presented to T cells in the recirculating pool (14). However, little is known about the progenitors of dendritic cells in the different compartments outlined above.

Dendritic cells in spleen (15) and afferent lymph (16, 17) are not in cell cycle but arise from a proliferating precursor. Ultimately, dendritic cells emanate from the bone marrow (15, 16, 18, 19), yet it has been difficult to generate these cells in marrow culture, except for two reports describing their formation in small numbers (20, 21). Here we describe the formation from mouse blood of proliferating cell aggregates that in turn release large numbers of mature dendritic cells. We will outline this tissue culture system, the importance of GM-CSF, and the identification of the progeny as dendritic cells on the basis of cytology, phenotype, and functional properties.

## Materials and Methods

**Mice.** We purchased BALB/c, (BALB/c  $\times$  DBA/2) $F_1$ , (BALB/c  $\times$  C57BL/6) $F_1$ , (C57BL/6  $\times$  DBA/2) $F_1$ , and C57BL/6 males and females, 6–8 wk of age, from Japan SLC Inc. (Shizuoka, Japan),

the Trudeau Institute (Saranac Lake, NY), and Charles River Wiga (Sulzberg, FRG).

**Blood.** Blood was obtained by cardiac puncture or from the carotid artery. The blood was diluted in, or allowed to drip into, RPMI 1640 with 100 U/ml heparin ( $\sim 2$  ml/mouse). Blood cells were pelleted at 1,000 rpm at 4°C, resuspended in RPMI 1640, and sedimented again. The pellet was suspended in 1 ml RPMI 1640 per mouse and mixed with an equal volume of 1.66% ammonium chloride in distilled water to lyse the red cells. After 2 min at room temperature, the suspension was spun at 1,000 rpm at 4°C. The pellet, which still contained red cells, was resuspended again in 0.5 ml RPMI and 0.5 ml  $\text{NH}_4\text{Cl}$  for 2 min, diluted in RPMI, and sedimented again. After two more washes, most platelets and red cells had been depleted.

**Blood Cultures.** The cells were cultured in 24-well dishes (25820; Costar, Cambridge, MA) in 1 ml of medium per well. The medium was RPMI 1640 supplemented with 5% FCS (JRH Biosciences, Lenexa, KA), 50  $\mu\text{M}$  2-ME, 20  $\mu\text{g}/\text{ml}$  gentamicin, and recombinant mouse (r)GM-CSF. Four preparations of rGM-CSF were evaluated with similar results, the yield of dendritic cells reaching a plateau with 30–100 U/ml. The preparations were from Dr. S. Gillis (Immunex Corp., Seattle, WA); Genetics Institute (supernatant from COS cells transfected with mGM-CSF; used at 30 U/ml or greater); and Dr. T. Sudo (supernatant from CHO cells transfected with the expression vector, pHSmGM-CSF [22]), and *Escherichia coli*-expressed material). The protocol is described in detail in Results. Cultures were fed first at day 6–7, and then every 3 d by aspirating 0.5–0.75 ml of the medium and adding back an equal volume of fresh medium with GM-CSF. To subculture the wells, we dislodged by pipetting most of the aggregates of growing dendritic cells as well as some cells in the monolayer of growing macrophages and fibroblasts. Pipetting usually disrupted the aggregate, particularly the peripheral cells that were more mature. With time in culture, e.g., at 2 wk, the aggregates of the growing dendritic cells became more stable and it was possible to dislodge the aggregates for separation by 1-g sedimentation. Typically, we applied the contents of five wells to a 6-ml column of 50% FCS-RPMI 1640 in a 15-ml conical tube (62.553.002 PS; Sarstedt, Inc., Princeton, NJ). After at least 20 min, the applied medium and top 1 ml of the column were removed. RPMI was added, the aggregates were pelleted at 1,000 rpm at 4°C for 5 min, and the cells were suspended gently for subculture in fresh medium.

**Phenotyping with mAbs.** A panel of mAbs was used to identify and characterize the cells in the GM-CSF-expanded blood cultures. The mAbs are reviewed elsewhere (23, 24) and are cited in Results. Cytospin preparations were made in a cytocentrifuge (Shandon Southern Instruments Inc., Sewickley, PA) using  $3\text{--}10 \times 10^4$  cells. The slides were stored with dessicant before fixation in acetone and staining with mAbs followed by peroxidase mouse anti-rat Ig (605-545; Boehringer Mannheim Biochemicals, Indianapolis, IN) or rabbit anti-hamster Ig (JZY-036-003; Accurate Chemical & Scientific Corp., Westbury, NY). The preparations were stained with Giemsa and mounted in Permout for bright field analysis. For cytofluorography (FACScan®; Becton Dickinson & Co., Mountain View, CA) aliquots of cells were stained with primary rat or hamster mAb followed by FITC mouse anti-rat Ig (605-540; Boehringer Mannheim Biochemicals) or biotin rabbit anti-hamster Ig (JZY-066-003; Accurate Chemical & Scientific Corp.) and FITC-avidin.

**Autoradiography.** Cultures were labeled with [ $^3\text{H}$ ]TdR to identify and phenotype the proliferating cells and their progeny. For pulse labeling, [ $^3\text{H}$ ]TdR was added to the cultures (6 Ci/mM, 1  $\mu\text{Ci}/\text{ml}$  final). 2 h later, the medium was replaced with [ $^3\text{H}$ ]TdR

free medium, and the cultures were separated into nonadherent released cells and residual adherent aggregates for examination on cytospin preparations (59900102; Shandon Southern Instruments Inc.). The cytospun cells were stained for specific antigens with mAb and immunoperoxidase as above. Also, the slides were dipped in photographic emulsion (Kodak autoradiography emulsion type NTB2 #165-4433) for exposure (5 d) before development, staining with Giemsa, and mounting in Permout. For pulse-chase experiments, a lower dose of [ $^3\text{H}$ ]TdR was used to maintain cell viability, but the cells were handled similarly otherwise. The pulse was applied at 0.1  $\mu\text{Ci}/\text{ml}$  for 2 or 16 h, the latter to provide higher initial labeling indices. The cells were washed and chased for 1–3 d before harvesting, and analysis was as above with immunoperoxidase, autoradiography, and Giemsa staining.

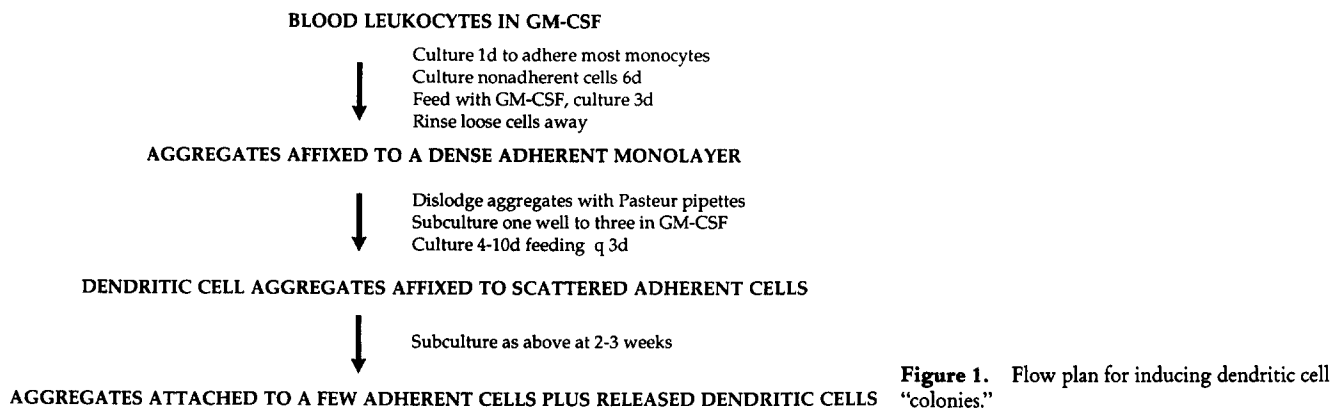
**Mixed Leukocyte Reactions.** Cells from the blood cultures were exposed to 1,500 rad ( $^{137}\text{Cs}$ ) and applied in graded doses to  $3 \times 10^5$  purified syngeneic or allogeneic T cells in 96-well, flat-bottomed microtest wells. The T cells were nylon wool–nonadherent spleen and lymph node suspensions that were treated with anti-Ia plus J11d mAbs and complement to remove residual APC. [ $^3\text{H}$ ]TdR uptake was measured at 72–86 h (6 Ci/mM, 4  $\mu\text{Ci}/\text{ml}$  final).

**Dendritic Cell Homing.** Dendritic cells or other cell types were labeled at  $2\text{--}10 \times 10^6/\text{ml}$  with carboxyfluorescein for 10 min on ice (C-1157; Molecular Probes, Eugene, OR) (30  $\mu\text{M}$  final concentration in HBSS with 5% FCS), washed in RPMI 1640, and injected in a volume of 50  $\mu\text{l}$  RPMI 1640 into the foot pads. 1 d later, the draining popliteal lymph nodes were removed, frozen in OCT medium, and sectioned (10  $\mu\text{M}$ ) in a cryostat. To sample the entire node, we took duplicate specimens at regular intervals. The sections were applied to multiwell slides (111006; Carlson Scientific, Peotone, IL), stored at  $-20^\circ\text{C}$ , dried in a dessicator 30 min before use (or left at room temperature overnight), fixed in acetone, and stained with a peroxidase-conjugated rabbit anti-FITC antibody (P404; Dakopatts, Carpinteria, CA). To verify that the dendritic cells in the lymph node were in the T-dependent areas as described (8), we added appropriate mAbs to B cells, T cells, macrophages, or dendritic cells, and visualized the latter with alkaline phosphatase-conjugated mouse anti-rat Ig (605-5357; Boehringer Mannheim Biochemicals) plus a chromogen kit (S04; Biomedica Corp., Foster City, CA). We then blocked endogenous peroxidase with “Endo Blocker” (M69; Biomedica Corp.) followed by the peroxidase anti-FITC as above.

## Results

When mouse blood leukocytes were cultured in GM-CSF at 30 U/ml, the cultures developed a large number of aggregates from which typical dendritic cells were eventually released. In the absence of GM-CSF, no colonies developed. Initially, we used cytologic criteria to detect the dendritic cells that characteristically extend large, sheet-like processes or veils (25–27). The procedure for expanding the aggregates of developing dendritic cells is described first (Fig. 1). Then, we will consider the phenotype and functional data that identify the progeny as typical mature dendritic cells.

**Aggregates of Proliferating Dendritic Cells in Blood Supplemented with GM-CSF.** Blood leukocytes, usually from (C  $\times$  D2) $\text{F}_1$  mice, were cultured in 16-mm tissue culture wells in medium supplemented with GM-CSF at 30 U/ml and at  $1.5 \times 10^6$  cells/well. After overnight culture, many monocytes adhered and the nonadherent cells were transferred to new 16-mm

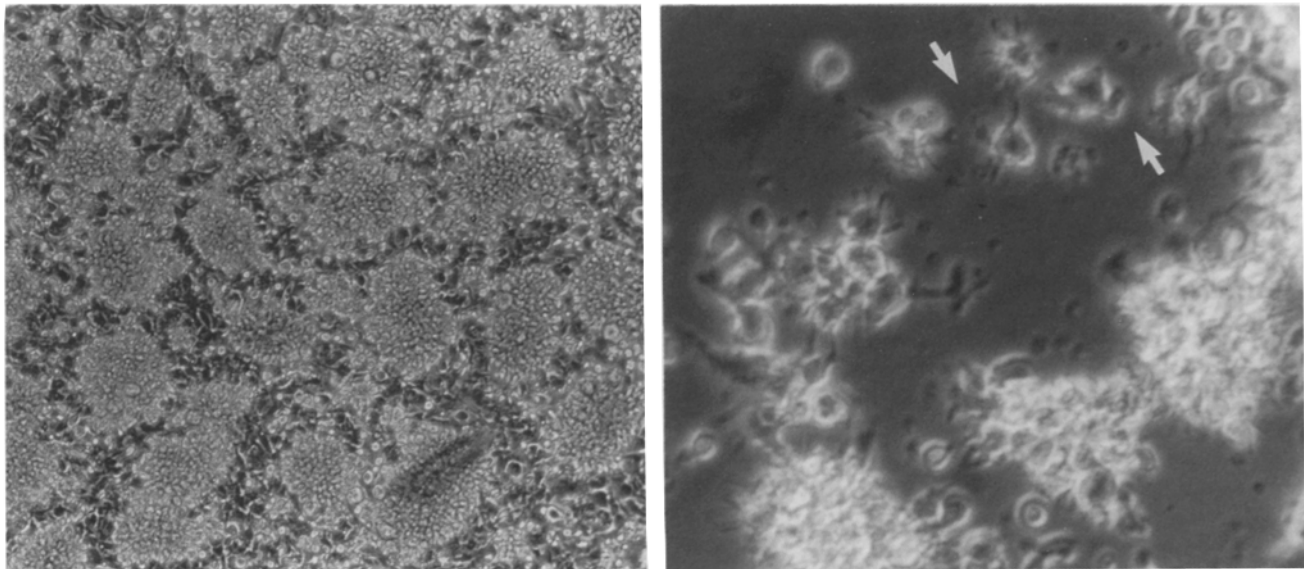


wells. The adherent cells did not develop dendritic cell colonies, but during the next week, the nonadherent populations exhibited three changes. First, most of the lymphocytes and granulocytes died or could be removed by washing. Second, the surface of the well became covered with a monolayer of tightly adherent cells that included macrophages and fibroblasts. Third, affixed to scattered sites on the monolayer, there developed small aggregates of cells. The cultures were fed with GM-CSF at day 6–7. The aggregates continued to expand in number and size. At about day 10, the cells were ready to be subcultured. Any residual loose cells could be rinsed off before dislodging the aggregates into fresh medium and GM-CSF. About  $0.8\text{--}1 \times 10^6$  dislodged cells per original well were divided into three subculture wells.

Most of the aggregates disassembled during this first subculture, while the bulk of the adherent monolayer remained attached to the original well. Upon transfer, most of the cells

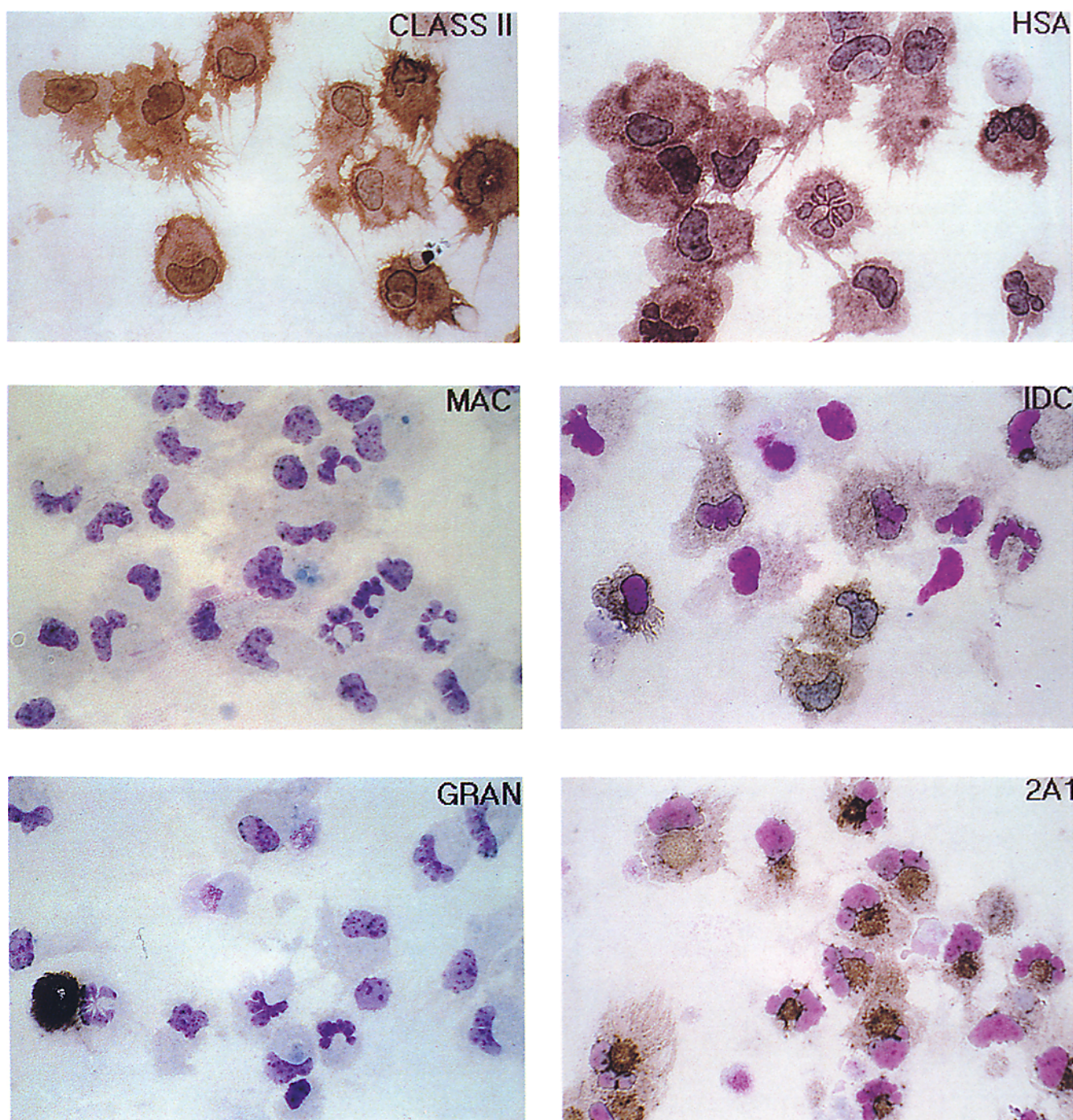
adhered as single cells to the new culture well, but over a period of 2–3 d, aggregates reappeared. The aggregates again were affixed to adherent stromal cells, but these adherent cells were much less numerous than the dense monolayer in the original culture. Over the next 4–7 d, aggregates filled the wells (Fig. 2 A). These colonies were often larger than those of the original wells and were covered with many sheet-like processes typical of dendritic cells. It was more difficult to count cells at this point, since many of the aggregates contained a core of tightly associated cells. However, the number of single cells that could be recovered per well expanded about twofold between days 10 and 17 of culture.

If the cultures were allowed to overgrow, some cells with the morphology of dendritic cells were released. More typically, we did not allow the cells to overgrow but instead dislodged the aggregates and subcultured them again at ~20 d. Before subculture, the aggregates could be purified from free



**Figure 2.** Phase contrast micrographs of dendritic cells developing in vitro. (A) Low power view of a monolayer covered with aggregates of proliferating dendritic cell progenitors (day 17 culture) ( $\times 100$ ). (B) Higher power view of typical dendritic cells (arrows) released from the aggregates ( $\times 400$ ).



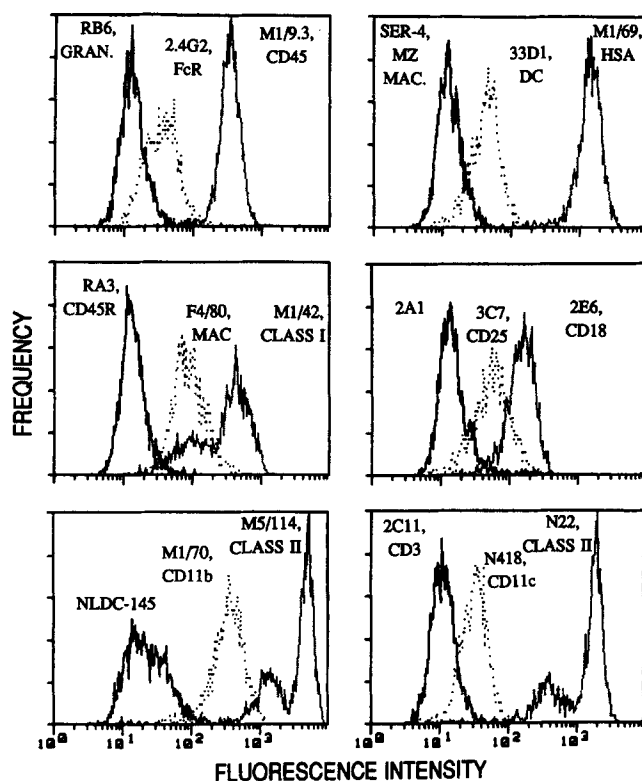


**Figure 3.** Immunoperoxidase labeling of a culture transferred after 2–3 wk to show some of the typical phenotypic features of released dendritic cell progeny. The primary antibodies are B21-2 anti-MHC class II (American Type Culture Collection [ATCC] TIB 229); M1/69 anti-heat-stable antigen (HSA, ATCC TIB 125); F4/80 antimacrophage (MAC, ATCC HB 198); NLDC 145 (13) anti-interdigitating cell (IDC, kindly provided by Dr. G. Kraal, Free University, Amsterdam); RB6 antigranulocyte (an example of a rare granulocyte is shown; GRAN); and 2A1, a mAb that reacts primarily with granules within the cytoplasm of cultured dendritic cells in vitro, and interdigitating cells in situ (Witmer-Pack et al., manuscript in preparation) ( $\times 600$ ).

cells by 1-g sedimentation. Such separations were more easily performed with longer periods of culture; i.e., it was easier to isolate intact aggregates at 3 vs. 2 vs. 1 wk of culture. With additional subculturing, the number of aggregates that were produced per well was progressively reduced. However,

colonies of growing cells, as confirmed by [ $^3\text{H}$ ]TdR labeling and autoradiography (see below), could be generated in subcultures for 1–2 mo.

After subculturing at 2–3 wk, typical single dendritic cells were now released into the medium (Fig. 2 B). By direct ob-



**Figure 4.** FACS® analyses of dendritic cells released from proliferating aggregates. Several mAbs summarized elsewhere (23, 24, 28) are shown. Except for MHC class I and II products, the phenotype of the released cells is homogeneous. Staining with no primary mAb was identical to RB6 and RA3.

servation with video recording, these released cells had the active motility of dendritic cells, continually extending and retracting large veils or sheet-like processes. In the presence of continued GM-CSF, we observed both free dendritic cells as well as expanding colonies. In the absence of GM-CSF, only free dendritic cells were released into the medium. The aggregates essentially fell apart and did not reform. The yields of free dendritic cells per subculture ranged from  $0.3$  to  $2.5 \times 10^5$ .

In summary, from a starting blood mononuclear culture of  $1.5 \times 10^6$  cells, where dendritic cells were difficult to detect, we on average obtained 5–10 subcultures each with at least  $3$ – $10 \times 10^4$  released dendritic cells at 3 wk, as well as many aggregates capable of further proliferation. Therefore, aggregates of growing cells were developing in mouse blood supplemented with GM-CSF, and these aggregates were covered with dendritic cells, many of which could be released spontaneously into the medium.

**Phenotype of the Cell Aggregates and Dendritic Cells Released Therefrom.** Cytospin preparations of 2–3-wk cultures were examined with a panel of mAbs and an immunoperoxidase method. The released cells, and many of the cells that could be dislodged from the periphery of the aggregate, were similar in their stellate shape and phenotype (Fig. 3). Most of the cells stained strongly with mAb to MHC class II (Fig. 3), the CD45 leukocyte common antigen, CR3 receptor CD11b, heat-stable antigen (HSA; Fig. 3), and CD44. Staining with

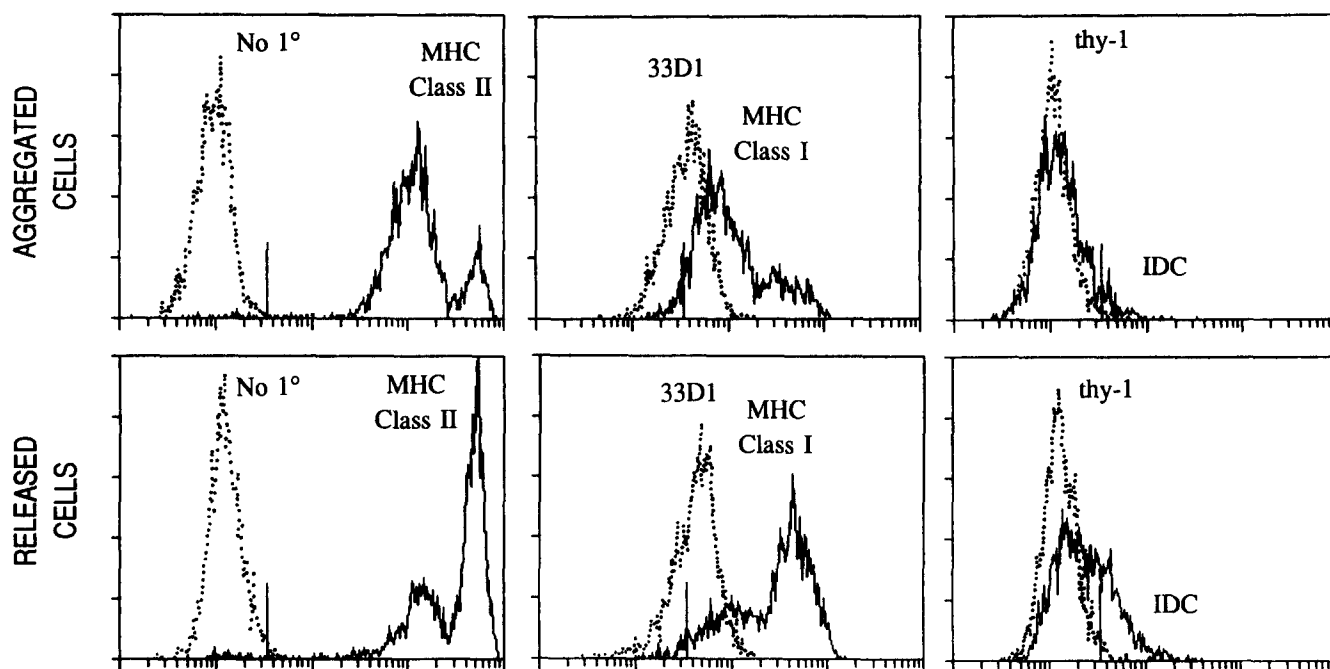
mAbs to the Fc receptor (2.4G2) and macrophage F4/80 antigen (MAC; Fig. 3) was weak or undetectable in  $>95\%$  of the cells. The cultures contained only rare B cells (B220 mAb, RA-3), T cells (Thy-1 mAb, B5-5), or granulocytes (GRAN, mAb RB6; Fig. 3). Some cells at the periphery of the aggregate, and many of the cells that were released from the aggregates, were stained with two markers that are largely restricted to dendritic cells. The interdigitating cell antigen (mAb NLDC 145 [13], IDC; Fig. 3), which also binds to thymic epithelium, stained many but not all of the dendritic profiles. Virtually all of the dendritic profiles stained with mAbs 2A1 (Fig. 3) and M342 (Witmer-Pack et al., manuscript in preparation), which stain granules in the perinuclear region of mature dendritic cells as well as interdigitating cells in sections through the T areas of lymphoid organs. Macrophages from many sites (blood monocytes; peritoneal cavity macrophages; macrophages in sections of lymph node, thymus, spleen) do not contain 2A1- or M342-reactive granules.

Cytofluorography was used to gain semiquantitative information on the expression of antigens at the cell surface. A panel of mAbs were applied to two populations: cells that could be dislodged from the aggregates by Pasteur pipetting, and cells that were released spontaneously when the aggregates were subcultured for 1 d. These “dislodged” and “released” populations were identical in their dendritic shape and in phenotype, but for some exceptions that are considered below. The phenotype of the released cells is shown in Fig. 4, and the few differences between aggregated and released cells are in Fig. 5. Virtually all the dendritic cells developing in and from the aggregates expressed high levels of the leukocyte common (CD45, mAb M1/9.3) and heat-stable (mAbs M1/69 and J11d) antigens, as well as high levels of CD44 (not shown) and CD11b (mAb M1/70). Low levels of the following antigens were detected on the cell surface: the dendritic cell antigen 33D1, the macrophage marker F4/80, the Fc $\gamma$  receptor antigen 2.4G2, the p55 IL-2 receptor CD25 antigen 3C7, and the CD11c integrin N418 (Fig. 4). These antigens were noted on all cells by FACS®, even though many of the antigens like F4/80 and 2.4G2 were weak or absent in the cytoplasm with an immunoperoxidase method (Fig. 3, see above). Several antigens were absent: RB6 granulocyte, RA3 B cell, B5-5 Thy-1, GK 1.5 CD4, and SER-4 marginal zone macrophage (Fig. 4).

Expression of class I and II MHC products by the dendritic cells in these cultures was very high but nonetheless bimodal (Figs. 4 and 5). Most of the dendritic cells that were dislodged from the aggregates had somewhat lower levels of MHC class I and II, while dendritic cells that were released from the aggregates had very high levels of MHC products. The other marker that was different in the released and loosely attached dendritic cells was NLDC 145, which was higher in the released population (Fig. 5, *top*). We conclude that the phenotype of the cells that arise from the proliferating aggregates is very much like that seen in cultured dendritic cells from skin, spleen, and thymus (24, 28), with the exception that the M1/70 CD11b marker is more abundant.

**[<sup>3</sup>H]TdR Autoradiography to Verify Growth of Dendritic Cell Precursors.** After 2 and 3 wk in liquid culture, the wells con-

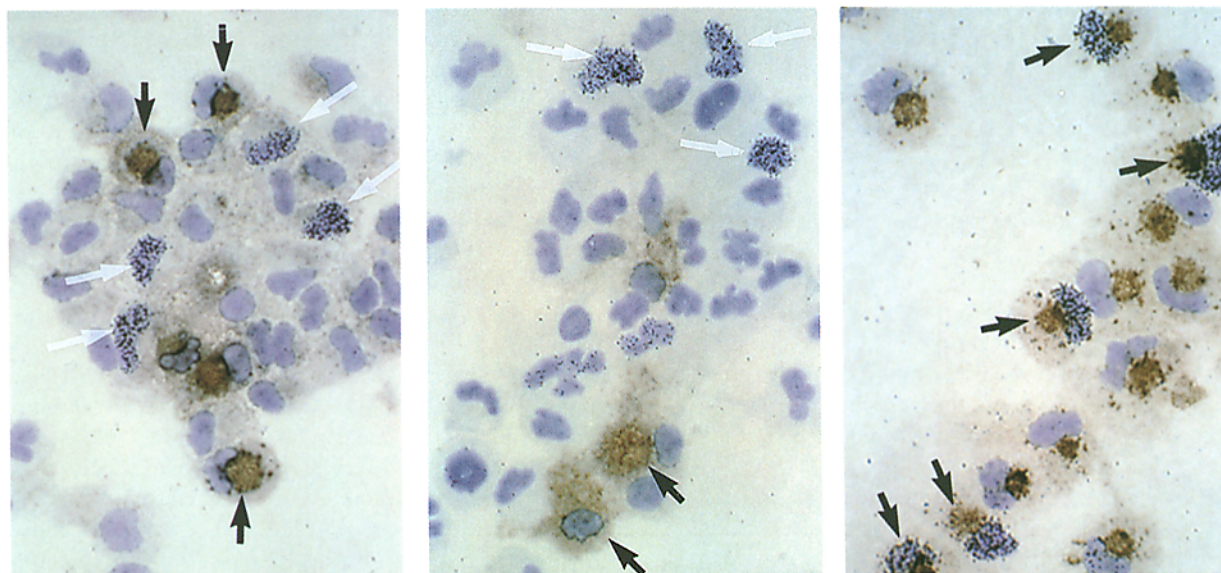




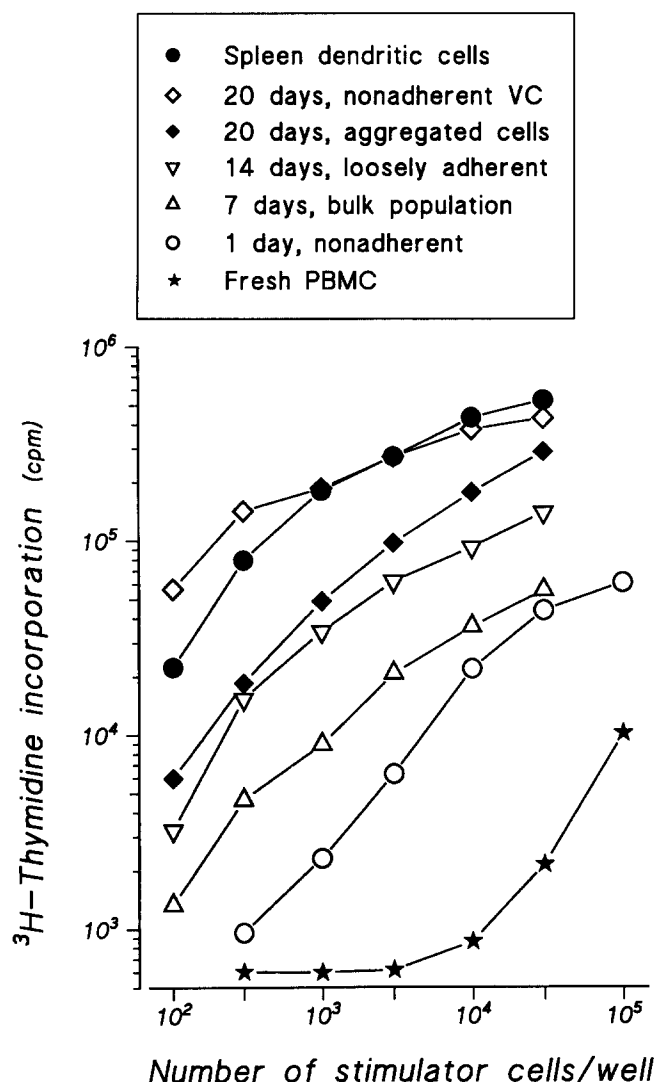
**Figure 5.** FACS® analyses of dendritic cells that could be dislodged by Pasteur pipetting of proliferating aggregates, and dendritic cells released spontaneously in culture. The mAbs are M1/42 anti-MHC class I (ATCC TIB 126); NLDC145 anti-interdigitating cell (13); M5/114 anti-MHC class II (ATCC TIB 120); 33D1 anti-dendritic cell (ATCC TIB 227); B5-5 anti-Thy-1. The staining with anti-MHC mAbs is bimodal, but the released cell fraction of dendritic cells is richest in expression of MHC class I and II.

tained numerous expanding aggregates of cells (as in Fig. 2 A), and in some cases were already releasing nonadherent dendritic cells in large numbers (Fig. 2 B). These cultures were exposed to [ $^3\text{H}$ ]TdR and examined for proliferative activity. The labeled cells were washed, spun onto slides, and

the cytopsins stained with mAbs and an immunoperoxidase method before dipping and exposure to photographic emulsion. Important markers were mAbs 2A1 and NLDC-145, which recognize intracellular granules and a cell surface antigen in mature dendritic cells, respectively (Fig. 3, see above).



**Figure 6.** [ $^3\text{H}$ ]TdR pulse and pulse-chase labeling of proliferating dendritic cell aggregates and their progeny, respectively. (Left and Middle) Pulse labeling of aggregates ( $1 \mu\text{Ci}/\text{ml}$  for 2 h) at  $\sim 2$  wk of culture. Before dipping in photographic emulsion, the cytopsins were stained with an immunoperoxidase method to identify the 2A1 anti-dendritic cell granule antigen (left) or the NLDC145 interdigitating cell antigen (middle). The cells that label for these antigens (black arrows) are not in S phase (white arrows). (Right) Pulse-chase label ( $0.2 \mu\text{Ci}/\text{ml}$  for 16 h, chase for 2 d) of aggregates that were isolated by velocity sedimentation and returned to culture to form large numbers of released progeny. In contrast to the left panel, the [ $^3\text{H}$ ]TdR-labeled cells express the 2A1 granule antigen characteristic of mature dendritic cells (black arrows).  $\times 425$ .



**Figure 7.** MLR-stimulating activity of populations isolated from the GM-CSF-stimulated mouse blood cultures (see text).

When cultures were labeled with a 2-h pulse of [ $^3\text{H}$ ]TdR, it was apparent that the labeling index in the aggregates was very high, at least 10–15% of the profiles in the aggregates being in S phase (Fig. 6, *left* and *middle*). In contrast, if [ $^3\text{H}$ ]TdR was applied to cultures that were releasing typical nonadherent dendritic cells, the released fraction contained only rare labeled profiles (not shown). If GM-CSF was removed, [ $^3\text{H}$ ]TdR labeling ceased within 1 d. Virtually all the [ $^3\text{H}$ ]TdR-labeled cells in the aggregate failed to label with mAbs to markers found on mature dendritic cells, i.e., 2A1 and NLDC145 (Fig. 6). The level of staining with anti-MHC class II mAb was less on the cells in S phase than in the released dendritic cell populations (not shown).

Pulse-chase experiments were then done to establish that labeled cells in the aggregate were giving rise to typical dendritic cells. Cultures were first exposed to a low dose of [ $^3\text{H}$ ]TdR, either for 2 or 16 h, the latter to label a larger per-

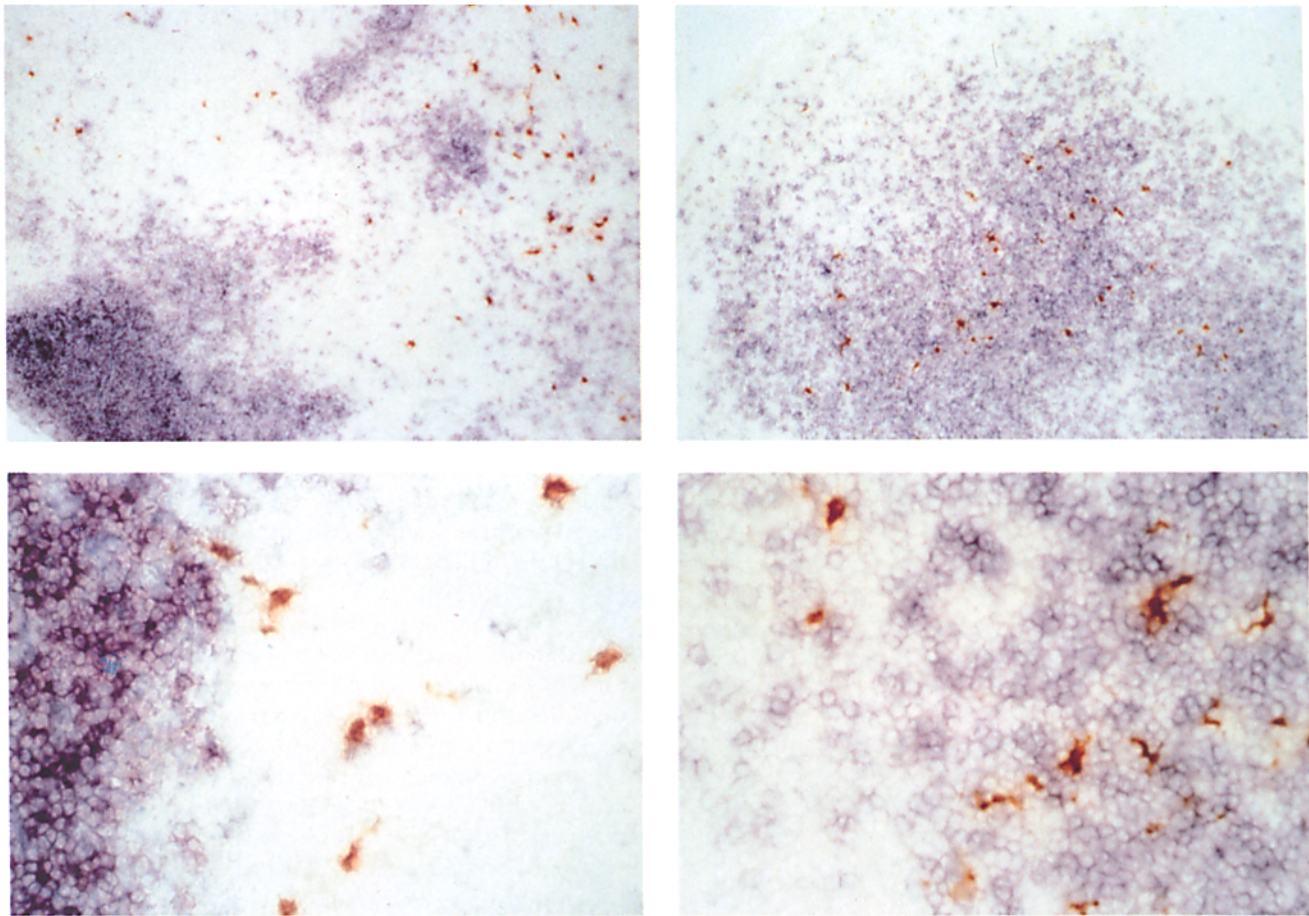
centage of the cells in the aggregates. The wells were washed free of radiolabel, and then the aggregates were dislodged and separated from free cells by 1-g sedimentation. The aggregates were transferred to fresh medium without radiolabel, and over the next 1–3 d of culture, many dendritic cells were released into the medium. When the “chased” cultures were examined, several findings were apparent. The labeling index remained high, i.e., most of the progeny of cells that were proliferating in the aggregates were not being lost from the cultures. Second, the grain counts were diluted severalfold from those apparent in the original pulse. Third, cells expressing the markers of mature dendritic cells (NLDC-145, the 2A1 granular antigen, high levels of MHC class II) were now radiolabeled (Fig. 6, *right*). Therefore, the cellular aggregates that GM-CSF was inducing in cultured mouse blood were actively proliferating and releasing nonproliferating progeny with many of the typical cytologic and antigenic features of mature dendritic cells, including the 2A1 granular antigen, the NLDC145 marker, and high levels of MHC class II.

**Accessory Cell Function for T Cell Proliferative Responses.** MLR-stimulating activity was monitored in the GM-CSF-treated blood cultures. Initially, there was little or no MLR-stimulating activity (Fig. 7, ★). Some stimulating activity was noted at day 1 of culture (Fig. 7, ○). An examination of cytopsin preparations revealed that these 1-d nonadherent blood cells had a low (<0.3%) but clear subset of Ia-rich, dendritic profiles. By day 7, when the proliferating aggregates were first evident on the monolayer, the stimulating activity of the dislodged aggregates had increased further, but was still 100 times less in specific activity than typical dendritic cells (Fig. 7, compare △ and ●), even though most of the cells at day 7 and subsequent time points were MHC class II positive. By day 14, at which time typical nonadherent dendritic cells were just beginning to be released from the aggregates, the nonadherent population had considerable MLR-stimulating activity (Fig. 7, ▽). After 3 wk, typical mature dendritic cells had become abundant, and these indeed stimulated comparably to their splenic counterparts (Fig. 7, compare ◇ and ●). Other cells in the culture, such as those dislodged from the aggregates, were ~10-fold less active than dendritic cells (Fig. 7, ◆). We conclude that the aggregates of proliferating dendritic cells have some MLR-stimulating activity but that it is the mature released cells that are fully potent, some 100–300 times more active on a per cell basis than the populations in the starting culture at 1–7 d. During days 7–20 of culture, total cell numbers also expanded at least 5–10-fold.

**Homing Activity of Dendritic Cells In Vivo** A second specialized feature of dendritic cells is their capacity to home to the T areas of peripheral lymphoid tissues (8, 10). To assess homing, we injected cells that had been labeled with carboxyfluorescein into the footpads. 1 d later, sections of the popliteal lymph nodes were stained with mAbs to B cells or T cells (secondary alkaline phosphatase anti-rat Ig conjugate) followed by peroxidase rabbit antiferretin antibody.

Blood leukocytes, even when given at a dose of  $10^6$  cells





**Figure 8.** Homing activity of dendritic cells released from proliferating cell aggregates. The dendritic cells were tagged with carboxyfluorescein and as a result could be labeled in these cryostat sections with a peroxidase-conjugated anti-FITC antibody plus diaminobenzidine (brown color). The sections were then double labeled with mAbs to B cells (*left*; RA3-6B2.1 anti-B220) or T cells (*right*; 53-6.7 anti-CD8; ATCC TIB 105). Note that the dendritic cells (brown profiles) homed to the deep cortex of the lymph node, which are the sites containing most of the recirculating T cells as well as interdigitating cells (*top*,  $\times 43$ ; *bottom*,  $\times 170$ ).

per footpad, failed to home to the lymphoid organ. When we tested dendritic cells that had been generated with GM-CSF from blood, homing to the T area was observed with injections of  $2 \times 10^5$  cells (Fig. 8). The selective localization to the T areas was confirmed by double labeling the specimens with mAbs that stain B cells or T cells (Fig. 8). Therefore, dendritic cells produced in culture have the key functional features of this lineage: homing to the T-dependent regions and strong accessory activity.

**Requirements for Generating Dendritic Cell Colonies from Blood.** The surface phenotype of the blood cell that gives rise to the dendritic cell colonies was assessed by treating the starting population with antibodies and complement. Treatment with either 33D1 anti-dendritic cell, anti-MHC class II, or anti-Thy-1 did not eliminate the colony-forming unit (not shown). Instead, removal of Thy-1<sup>+</sup> or Ia<sup>+</sup> cells enriched colony numbers severalfold. CSFs other than GM-CSF were also tested, either at the start of the 1–3-wk culture, or upon transfer of 2–3-wk-old aggregates to form veiled cells (Fig. 1). None of the CSFs tested (i.e., IL-3, M-CSF, G-CSF,

or stem cell factor) supported the formation of colonies or mature dendritic cells. Therefore, the growing dendritic colonies are very much dependent upon GM-CSF.

## Discussion

In an effort to identify proliferating precursors to the dendritic cell system, we set up cultures from several tissues that lacked mature dendritic cells and supplemented these with different growth factors, particularly the CSFs (M-CSF, G-CSF, IL-3, GM-CSF, IL-1, and SCF). Neonatal epidermis, which contains mainly Ia<sup>−</sup> Langerhans cells (29), was tried first but to no avail. Bone marrow was tried next, but it was difficult in bulk cultures to identify typical colonies or large numbers of dendritic cells. The marrow cultures became overgrown with macrophages. When we turned to blood, which has few typical dendritic cells in the mouse (30), an entirely different phenomenon was observed. Growing cell aggregates appeared after  $\sim 6$  d in culture, and these were often covered with profiles having the unusual and motile processes of den-

dritic cells. With time, typical nonadherent dendritic cells were released. The latter had the morphology and movement of dendritic cells as previously described in cultured mouse spleen, mouse skin, lymph from several species, and human blood (25–27). Therefore, to identify proliferating dendritic cells, it seems critical to begin with an appropriate starting population (in this case, blood) and to supplement the culture with GM-CSF.

We think that the initial aggregates that appeared in the cultures represented clones, since small groups of four to six cells were observed early on, e.g., day 5. We tried to prove that the aggregates were clonal by mixing blood cells from strains that were distinguished with markers to polymorphic antigens like CD44 and MHC class II. However, we could not complete the experiments since we found that mouse strains differed in the number and speed with which colonies developed. BALB/c and DBA (and F<sub>1</sub> strains derived therefrom) were the most active; B6 and B10 were several times less active; and strains like CBA/J, C3H/He, and A/J were poor sources of proliferating, dendritic cell aggregates.

The precursors to the aggregates of proliferating dendritic cells were not typical monocytes or dendritic cells, because the number of aggregates that developed could be increased substantially if one depleted monocytes by adherence or Ia-positive cells with antibody and complement. We tentatively conclude that blood contains an Ia<sup>+</sup> precursor that forms a proliferating aggregate. In the aggregate, dendritic cells mature and are released as nonproliferating progeny.

The formation of aggregates of dendritic cells required exogenous GM-CSF. If the aggregates were placed in macrophage- or granulocyte-restricted CSFs (M-CSF, G-CSF), proliferation ceased and neither macrophages nor granulocytes were formed. Because the cultures contained macrophages and some stromal cells, in addition to the dendritic cell aggregates, it was possible that other cytokines were being produced that were critical to the formation of dendritic cells. It appears, however, that the cells in the aggregates have lost responsiveness to M- and G-CSF, and that dendritic cells represent a distinct myeloid pathway of development. Perhaps the pathway originates from a common precursor in which the dendritic cell lineage is an offshoot that no longer responds to macrophage- and granulocyte-restricted CSFs.

Labeling with [<sup>3</sup>H]thymidine, using pulse and pulse-chase protocols, was important in establishing the precursor-product relationships that were taking place in these liquid cultures (Fig. 6). In a 2-h pulse, virtually every labeled cell lacked two typical markers of mature dendritic cells, i.e., the NLDC-145 interdigitating cell surface antigen (13) and the recently identified 2A1/M342 granular cytoplasmic antigens (Witmer-Pack et al., manuscript in preparation). These mAbs do not stain most macrophage populations that we have examined either as isolated cells (blood, spleen, peritoneal macrophages)

or in sections (thymus cortex, spleen red pulp, lymph node medulla). In pulse-chase protocols, large numbers of labeled progeny were released from the aggregates, and these released cells were nonadherent, motile, and strongly stimulatory in the MLR. After combined autoradiography and immunoperoxidase labeling, the labeled progeny carried the granular antigens, the NLDC-145 antigen, and very high levels of MHC class II. Each of these cytologic and antigenic markers are largely restricted to dendritic cells.

We believe that maturation to typical nonproliferating dendritic cells occurred within the aggregate. The aggregates were covered with cells with the sheet-like or veiled processes of dendritic cells. Cells with markers of mature dendritic cell markers (high MHC class II, 2A1-positive granules, NLDC antigen) were also observed at the periphery of the cell aggregates (Fig. 6, *left* and *middle*). However, it was difficult to isolate the aggregate intact, without dislodging these more mature cells. The mechanism whereby dendritic cells matured and left the aggregate was not clear. Maturation was enhanced in older cultures (>2 wk) or by removing adherent stroma cells. Both proliferation and maturation was blocked if the cultures contained too many fibroblasts.

The functional maturation that occurred in the proliferating aggregate is striking. The dendritic cells that were generated in culture were potent MLR stimulators. 10<sup>2</sup> dendritic cells induced a much stronger primary MLR than 10<sup>5</sup> blood leukocytes (Fig. 7). The increase in stimulating activity per Ia<sup>+</sup> cell was at least 2 logs between the time that the aggregates first appeared and the time that typical dendritic cells were released in large numbers. Over this time period, cell recovery increased 5–10-fold. Also, the dendritic cell progeny homed in a precise way to the T cell area of lymph node (Fig. 8), another functional property that was not detectable in blood cells (data not shown).

Given the criteria that have become evident in this paper, it will be feasible to look for proliferating progenitors of dendritic cells in other organs. It is known that proliferating precursors are giving rise to the rapidly turning over populations of dendritic cells in spleen (15) and afferent lymph (16). The proliferation of leukocytes (other than T cells) occurs in the bone marrow, but it may be that for dendritic cells, the marrow also seeds the blood and other tissues with progenitors that then proliferate extensively as shown here. By being able to prepare the otherwise trace dendritic cell in large numbers, it should be more feasible to pursue previously unexplored areas of dendritic cell function. Specifically, growing dendritic cells will facilitate molecular and clinical studies on the mechanism of action of these APCs, including their capacities to capture and retain antigens in an immunogenic form (4, 6, 14) and act as adjuvants for the generation of immunity *in vivo* (1, 14, 31).

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