

# Vav1/2/3-null Mice Define an Essential Role for Vav Family Proteins in Lymphocyte Development and Activation but a Differential Requirement in MAPK Signaling in T and B Cells

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## Abstract

The Vav family of Rho guanine nucleotide exchange factors is thought to orchestrate signaling events downstream of lymphocyte antigen receptors. Elucidation of Vav function has been obscured thus far by the expression of three highly related family members. We generated mice lacking all Vav family proteins and show that Vav-null mice produce no functional T or B cells and completely fail to mount both T-dependent and T-independent humoral responses. Whereas T cell development is blocked at an early stage in the thymus, immature B lineage cells accumulate in the periphery but arrest at a late "transitional" stage. Mechanistically, we show that the Vav family is crucial for both TCR and B cell receptor (BCR)-induced Ca<sup>2+</sup> signaling and, surprisingly, is only required for mitogen-activated protein kinase (MAPK) activation in developing and mature T cells but not in B cells. Thus, the abundance of immature B cells generated in Vav-null mice may be due to intact Ras/MAPK signaling in this lineage. Although the expression of Vav1 alone is sufficient for normal lymphocyte development, our data also reveal lineage-specific roles for Vav2 and Vav3, with the first demonstration that Vav3 plays a critical compensatory function in T cells. Together, we define an essential role for the entire Vav protein family in lymphocyte development and activation and establish the limits of functional redundancy both within this family and between Vav and other Rho-guanine nucleotide exchange factors.

Key words: thymocyte • antigen receptor • signal transduction • Ca<sup>++</sup> • mitogen-activated protein kinase

## Introduction

T and B lymphocytes differentiate along an ordered program controlled by crucial signals emanating from surface receptors (1). In the T cell lineage, the expression of the pre-TCR induces both differentiation and proliferative expansion of

the pool of thymic progenitors, presumably effected by Ras/mitogen activated protein kinase (MAPK) signaling (2–6). In a parallel pathway, the pre-B cell receptor (pre-BCR) induces differentiation and expansion of the pool of

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*Abbreviations used in this paper:* BCR, B cell receptor; DAG, diacylglycerol; DN, double negative thymocyte; DP, double positive thymocyte; GEF, guanine nucleotide exchange factor; FO, follicular B cell; IP3, inositol triphosphate; MAPK, mitogen-activated protein kinase; MZ, marginal zone B cell; NF, newly formed B cell; PI3K, phosphoinositide 3-OH kinase; PLC, phospholipase C; SP, single positive thymocyte; TD, T-dependent; TI-2, T-independent type 2; TNP, trinitrophenyl.

pre-B cells (7–9). Since antigen receptors of both immature and mature lymphocytes lack any intrinsic signaling capability, the pre-TCR or pre-BCR, and the TCR or BCR, transmit their signals via associated transmembrane chains (CD3 and TCR $\zeta$  in T cells or Ig $\alpha$  and Ig $\beta$  in B cells) (10–13) that become tyrosine phosphorylated and recruit adaptor proteins (14, 15) and Vav guanine nucleotide exchange factors (GEFs) (16, 17). Subsequent activation of phosphoinositide 3-OH kinase and phospholipase C (PLC) $\gamma$ 1/2 generates second messengers diacylglycerol (DAG) and inositol triphosphate that induce Ras/MAPK or Ca<sup>2+</sup> signaling, respectively (10–13).

The Vav family of three highly conserved Rho-GEFs was initially implicated in signaling downstream of both the TCR and BCR by virtue of inducible tyrosine phosphorylation and the presence of structural domains characteristic of proteins involved in signal transduction (16, 17). Receptor-induced tyrosine phosphorylation and the presence of SH2 and SH3 domains distinguish Vav proteins from the plethora of over 50 known mammalian Rho-GEFs (18). Thus, Vav proteins can interact directly with several tyrosine kinases (through SH2 domain interactions with tyrosine phosphorylation sites) and indirectly with activated tyrosine kinases by binding to adaptor proteins that contain SH2 and/or SH3 binding motifs. Like Vav1 (19, 20), Vav2 and Vav3 are inducibly phosphorylated after stimulation of antigen receptors in immortalized T and B cell lines (21, 22); however, overexpression studies indicate functional differences in the activities of the three family members in coupling to distinct downstream pathways (16, 17). As exchange factors for Rho GTPases, Vav proteins have the potential to regulate antigen receptor-induced actin cytoskeletal rearrangements and other Rho-GTPase-regulated cellular pathways (16, 17). Indeed, loss of Vav1 prevents efficient capping of the TCR through effects on actin polymerization (23, 24). Although the exact mode of Vav regulation in vivo is currently unknown, it is thought that the exchange activity of the DH domain, responsible for regulation of Rho-GTPases, may be controlled by tyrosine phosphorylation of an NH<sub>2</sub>-terminal autoinhibitory extension (25); however, Vav may have both DH-dependent and DH-independent functions in T cells (26).

The importance of Vav1 in lymphocytes was first demonstrated in mice lacking Vav1 (Vav1<sup>ko</sup>) which showed activation defects in T and B lymphocytes (27–30) and in NK (31) and  $\gamma\delta$  T cells (32). However, virtually all of those defects were partial, and Vav1<sup>ko</sup> mice still developed significant numbers of T cells, whereas development of B cells was nearly intact (27–30). Moreover, Vav1<sup>ko</sup> lymphocytes retained significant functional ability, presumably due to redundancy between Vav family members. Indeed, analyses of mice deficient in both Vav1 and Vav2 revealed a compensatory function for Vav2 in B lineage cell development and function (33, 34). Although its role in T cell activation has not yet been defined, Vav2 did not appear to have any major function in T cell development (33, 34). Little is known about the function of Vav3 in primary lymphocytes, but analyses of Jurkat T cells (21, 22) and a Vav3-

deficient chicken DT40 B cell line (35) suggest that it is involved in both TCR and BCR signaling. Since mice lacking all three Vav proteins have not yet been reported, efforts to define the role of the entire Vav family in lymphocytes have been complicated by the functional redundancy among individual family members.

In this report, we generate mice congenitally lacking all three Vav isoforms and present analyses of lymphocyte development and function. We provide genetic and biochemical evidence that the Vav family is critical and non-redundant in T and B lymphocyte development and function and is essential in the formation of the adaptive immune system. Although the involvement of Vav1 and Vav2 in Ca<sup>2+</sup> signaling is well documented (23, 24, 29, 33, 34), our analyses establish that the entire Vav family is indispensable for both TCR- and BCR-induced Ca<sup>2+</sup> fluxes. Notably, we observed a dissociation in Vav function at the level of MAPK signaling in which Vav was required in T cells but not in B cells.

## Materials and Methods

**RNAse Protection Assays.** To generate antisense riboprobes, a 198-bp cDNA fragment of murine Vav1 (nucleotides 664–862), a 200-bp cDNA fragment of murine Vav2 (nucleotides 652–852), and a 197-bp cDNA fragment of murine Vav3 (nucleotides 676–873) were generated by RT-PCR using either thymus (Vav1) or spleen RNA (Vav2, Vav3). Each fragment was sequenced and subcloned into pBluescript II SK in antisense orientation with respect to the T7 promoter. Riboprobes were synthesized with T7 RNA polymerase, labeled with  $\alpha$ -<sup>32</sup>P UTP, and PAGE purified. RNase protection assay was performed essentially as described (43) with minor modifications. Briefly, 2.5  $\mu$ g of total RNA from each sample was hybridized with antisense RNA probes (10<sup>5</sup> cpm) in 30  $\mu$ l of 80% formamide, 40 mM PIPES, pH 6.5, 1 mM EDTA, and 0.4 M NaCl at 45°C overnight. The hybrids were digested in 400  $\mu$ l of 10 mM Tris-HCl, pH 7.5, 5 mM EDTA, 0.3 M NaCl, and 5  $\mu$ g/ml RNase A at 30°C for 30 min. After proteinase K treatment (125  $\mu$ g/ml, 37°C, 30 min) and phenol/chloroform extraction, the protected RNA probes were analyzed by 6% PAGE containing 8 M urea. Signal was detected using x-ray film or PhosphorImager scanning with the same exposure time for all samples, and data was analyzed with ImageQuant software (Molecular Dynamics). RNA was isolated using Trizol reagent (Invitrogen).

**Generation of Vav3-deficient Mice.** Murine *vav3* genomic clones were obtained by screening a 129 strain phage genomic library (Stratagene) with a 438-bp fragment (probe K) corresponding to nucleotides 1,140–1,578 of murine *vav3* cDNA, which was generated as described previously (22). This yielded several overlapping *vav3* genomic fragments. One of the clones extending toward the 5' end was used to generate a new probe, which included the 5' end of DH domain-encoding sequences (probe H). Using both probes, four overlapping clones were characterized that included exons 3–17. A targeting construct was made to replace exon 10, encoding part of the DH domain, with the neomycin-resistance gene in the reverse transcriptional orientation. A thymidine kinase (tk) gene was used for negative selection of cells with randomly integrated constructs. The targeting construct was linearized and transfected into TC-1 ES cells by electroporation. Positive and negative selection of transfectants was performed in

media containing G418 (0.4 mg/ml) and gancyclovir (1  $\mu$ M). Genomic DNA from individual double resistant (G418 and gancyclovir) clones was digested with EcoRI and probed with a 5' flanking probe on the Southern blot. Several targeted clones were identified by the presence of a 6.8-kb EcoRI band upon hybridization with the 5' probe, and KpnI digestions were performed to ensure proper homologous recombination at the 3' end of the targeted region, as identified by hybridization of a 22.0-kb KpnI band with the Neo probe. To rule out any undesired integration events, EcoRI-digested DNA was also hybridized with the Neo probe. Two independently derived heterozygous Vav3 mutant ES cells were subcloned and injected into C57BL/6 blastocysts to generate somatic chimera mice, which were bred with C57BL/6 females to transmit the targeted allele to the germline. F1 heterozygotes were then intercrossed to generate homozygous mutant mice.

**Generation of Vav3 Antibodies.** To generate polyclonal Vav3 antibodies, we immunized rabbits with a fusion protein comprising glutathione S-transferase and a Vav3 fragment, including the DH and PH domains (aa 357–525) expressed in *Escherichia coli*, using standard methods. The specificity of these antibodies for Vav3 and the lack of cross-reactivity with other family members was confirmed by Western blotting of cell lysates of 293T cells transiently transfected with either an empty vector or expression constructs encoding GFP fusions of Vav1, Vav2, or Vav3. mAbs against murine Vav3 were raised by producing a recombinant glutathione S-transferase–Vav3 fusion protein (encompassing 230 aa including the PH and the NH<sub>2</sub>-terminal SH3 domains) expressed in *E. coli* and purified over a glutathione-Sepharose column (Amersham Biosciences), which was used for immunization of Vav3-deficient mice. Splenocytes of the immunized mouse were fused with P3U1 mouse myeloma cells, and hybridoma clones were selected and screened for binding to a maltose binding protein–Vav-3 fusion protein by ELISA. One positive hybridoma clone, 6D42, IgG1 isotype, was established and used for production of mAbs, which showed no cross-reactivity with Vav1 or Vav2.

**Mice, Cell Suspensions, Antibodies, and Flow Cytometry.** Germ-line Vav1<sup>-/-</sup> and Vav2<sup>-/-</sup> mice were described previously (29, 34) and were gifts from Dr. Tybulewicz (National Institute for Medical Research, London, UK) and Dr. Turner (The Babraham Institute, Cambridge, MA). These mice were bred with Vav3-deficient mice in the mixed B6/129 background and maintained in the SPF facility of Washington University School of Medicine. All mice analyzed in this report were littermates derived from mating heterozygous animals. Cell suspensions were prepared, counted, and stained with antibodies following standard procedures. The following antibody conjugates were used (BD Biosciences): phycoerythrin (PE)–H129.19 (anti-CD4), fluorescein (FITC)–, PE–, cytochrome C (CyC)–, or biotin–53–6.7 (anti-CD8 $\alpha$ ), PE–3C7, or FITC–7D4 (anti-IL2R $\alpha$ ), FITC–145–2C11 (anti-CD3 $\epsilon$ ), CyC–RA3–6B2 (anti-B220), PE–B3B4 (anti-CD23), and FITC–7G6 (anti-CD21/35). Biotinylated antibodies were detected with streptavidin (SAV)–PE or SAV–CyC (BD Biosciences). Subsets of double negative (DN) thymocytes were analyzed based on expression of CD44 and CD25 after gating out all cells staining with a cocktail of biotinylated antibodies to CD4, CD8, B220, Mac-1, and Gr-1, followed by Streptavidin Cy-Chrome (BD Biosciences). For intracellular staining of TCR $\beta$ , cells were first stained with PE–CD4 and Cy-Chrome–CD8 $\alpha$ . Subsequently, cells were fixed, permeabilized in 1% saponin, and stained with FITC-labeled anti-C $\beta$ -specific antibodies (H57 clone) and analyzed on FACSCalibur flow cytometer (Becton

Dickinson) with CellQuest or FlowJo software. Data are displayed as histograms or dot plots with logarithmic scale. Each plot represents analysis of  $\geq 2 \times 10^5$  events collected as listmode files.

**T and B Cell Stimulation and Proliferation Assays.** T cells were purified from LN cell suspensions by magnetic sorting and removal of B cells with anti-Ig-coated Dynabeads (Dyna) to avoid receptor cross-linking during the purification process using standard procedures. The purity of resulting T cells exceeded 90% as confirmed by FACS<sup>®</sup> analysis. B cells were purified from spleen cell suspensions by magnetic removal of cells stained with biotinylated antibodies anti-Thy1.2, CD4, CD8, CD3, Mac-1, and Gr-1 using streptavidin-coated Dynabeads (Dyna). Purity of B cells was confirmed in each preparation by FACS<sup>®</sup> analysis and exceeded 85%. Lymphocytes were cultured in DME medium supplemented with 10% FCS, pen/strep, L-glutamine, Na-pyruvate, nonessential amino acids, and 2-mercaptoethanol. T cells were stimulated at  $5 \times 10^6$  cells/ml with either soluble or plate-bound anti-CD3 $\epsilon$  antibodies (2C11; a gift from Dr. Kanagawa, Washington University, St. Louis, MO) (0.01–10  $\mu$ g/ml as indicated) and/or anti-CD28 (2  $\mu$ g/ml) (NA/LE formula; BD Biosciences) as indicated, and B cells were stimulated with soluble anti-IgM Fab<sub>2</sub> fragments (10  $\mu$ g/ml) (Jackson ImmunoResearch Laboratories) for the indicated times at 37°C. For proliferation assays, purified T and B lymphocytes were cultured at a concentration of  $5 \times 10^4$  cells/100  $\mu$ l in 96-well flat-bottom tissue culture plates as indicated. Cells were pulsed with 1  $\mu$ Ci <sup>3</sup>H-thymidine at 48 h for an additional 12 to 16 h and then collected and scintillation counted. The data are displayed as raw cpm values. All assays were conducted in triplicates.

**Immunizations and Antigen-specific Ig ELISA.** To determine the T-independent type 2 (TI-2) immune response, 6–8-wk-old mice were immunized intraperitoneally with 100  $\mu$ g trinitrophenyl (TNP)-Ficoll (Biosearch Technologies) in alum (Pierce Chemical Co.). T-dependent (TD) immune response was determined by injection of 100  $\mu$ g TNP-KLH (Biosearch Technologies) in alum. Mice were bled before immunization (day 0) and at days 7 and 14. After boost with 100  $\mu$ g antigen at day 14, mice were bled at days 21 and 28. Concentrations of TNP-specific serum antibody isotypes were determined by ELISA. ELISA plates (Nunc Maxisorp) were coated at 100  $\mu$ l/well with 5 mg/ml TNP-BSA (Biosearch Technologies) or with PBS only, as a control, and incubated overnight at 4°C. After overnight incubation, plates were washed three times with wash buffer (Tris–Cl, pH 7.6, plus 2% Triton X-100) and blocked in 1% FCS in PBS (blocking buffer) for 2 h at 4°C. Serum samples were serially diluted in blocking buffer and applied to the ELISA plates in triplicate. After a 2-h incubation at room temperature with gentle rocking, ELISA plates were washed three times with wash buffer. TNP-bound serum antibodies were detected using 100  $\mu$ g/well HRP-conjugated isotype-specific antibodies (Southern Biotechnology Associates, Inc.) diluted 1:5,000 in blocking buffer. Plates were incubated at room temperature for 1 h with gentle rocking, washed, and developed using ABTS (Sci-Mart) containing H<sub>2</sub>O<sub>2</sub> as a substrate. Optical densities (OD<sub>450</sub>) were read at 405 nm (Titertek). For each isotype, a standard curve was generated based on OD<sub>450</sub> values using sera from a series of immunized WT mice. Subsequently, the experimental samples were serially diluted such that at least two values fell into the linear range of the standard curve. Concentrations of TNP-specific antibodies are shown as relative units compared with the standard curve determined for each specific isotype.

**Cell Culture, Transfections, and Immunoblotting.** Cells were cultured in DMEM with 10% FCS. For immunocytochemistry, cells

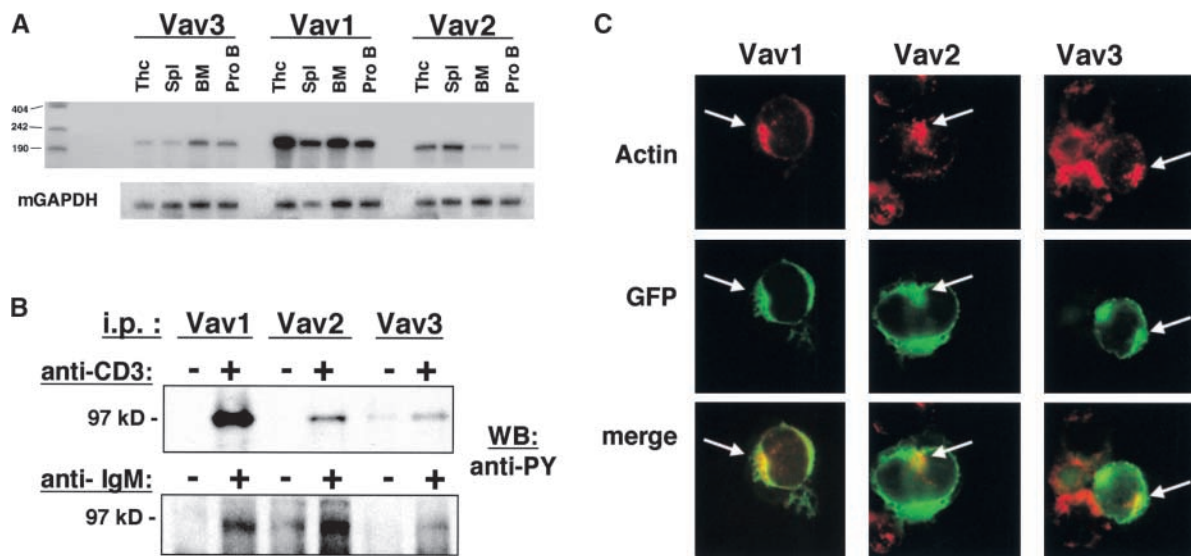
were plated in coverslips at  $5 \times 10^4$  cells per well in 24-well plates. For immunoprecipitation and immunoblotting, cells were plated on 100-mm dish at the density of  $1.8 \times 10^6$  cells per dish, and cells were transfected using SuperFect (QIAGEN) according to manufacturers instructions. At 36 h after transfection, cells were washed with ice cold PBS and then lysed in cold NP-40 lysis buffer (50 mM Tris-Cl, pH 7.6, 150 mM NaCl, 10 mM NaF, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 10% glycerol, 1% NP-40, 1 mM phenylmethylsulfonyl fluoride, 1  $\mu$ g/ml each of leupeptin, aprotinin, and pepstatin) for 10 min at 4°C. Plates were then scraped and crude lysates cleared by centrifugation at 14,000 *g* for 10 min at 4°C. For immunoprecipitation, total lysates were incubated with 2  $\mu$ g of antibodies and a 50:50 slurry of protein A-conjugated Sepharose beads (Amersham Biosciences) with rotation for 3 h at 4°C. Immunoprecipitates were washed four times with cold NP-40 lysis buffer, resuspended in 2 $\times$  SDS sample buffer, and analyzed by Western blotting following standard procedures. Primary antibodies were developed with HRP-conjugated secondary antibodies (Bio-Rad Laboratories, Inc.) and visualized by chemiluminescence (Amersham Biosciences). For analyses of tyrosine phosphorylation events, stimulated cells were lysed in RIPA buffer (PBS, 1% NP40, 0.5% deoxycholate, 0.1% SDS, 10 mM EDTA) supplemented with a protease and phosphatase inhibitor cocktail (Boehringer). Postnuclear lysates were prepared, resolved by SDS-PAGE, and analyzed with antibodies to phosphotyrosine (4G10; Upstate Biotechnology). Immune complexes were detected by enhanced chemiluminescence (Amersham Biosciences).

**Ca<sup>2+</sup> Fluxes.** Ca<sup>2+</sup> signaling was measured by preloading purified B or T cells with either Indo-1-AM or Fluo-4-AM (Mo-

lecular Probes) at the concentration of 3–5  $\mu$ g/ml,  $10^6$  cells/ml at 37°C for 45 min, with occasional vortexing. In some experiments, cells were stained with anti-CD4-APC conjugates after loading to allow “gating” on CD4<sup>+</sup> T cells during analyses. Cells were washed, resuspended in complete media, and analyzed by flow cytometry by running unstimulated samples for 20 s to establish the baseline, and then cells were stimulated by addition of either anti-IgM antibodies (Jackson ImmunoResearch Laboratories) or anti-CD3 immediately followed by cross-linking with anti-hamster antibodies (BD Biosciences) at the indicated concentrations. Ca<sup>2+</sup> fluxes were measured for 8 min followed by the addition of ionomycin for additional 1 min at 0.5  $\mu$ g/ml.

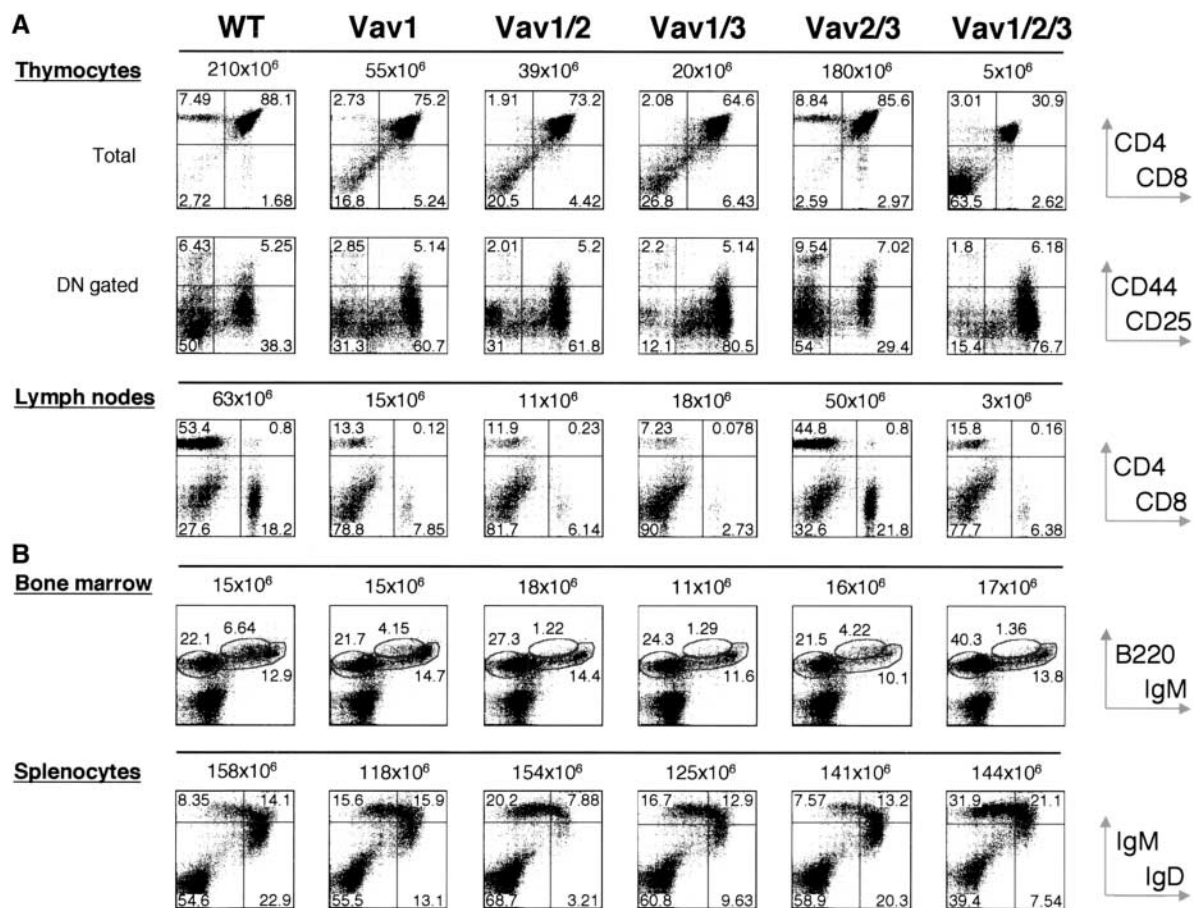
**Intracellular Staining for Phospho-p44/42 MAPK (pErk).** To determine Erk activation in T cells, intracellular staining was done. Thymocyte cell suspensions were prepared directly in pre-warmed (37°C) 4% paraformaldehyde (2% final) to fix cells, immediately followed by permeabilization in 100% ice cold methanol for 30 min. As a positive control, fresh cells were stimulated with PMA (20 ng/ml) in PBS for 5 min. Cells were stained with phospho-p44/42 MAPK (pErk) (Thr202/Tyr204, E10 clone; Cell Signaling Technology) in the presence of Fc-block at room temperature for 15–30 min. Anti-mouse IgG1-PE and CD4 CyC were added to each sample and incubated at RT for 15–30 min. Cells were analyzed by flow cytometry and gated electronically based on expression of surface markers as indicated.

**Online Supplemental Material.** Fig. S1 depicts analyses of lymphocyte development in Vav2<sup>ko</sup> and Vav3<sup>ko</sup> mice. Flow cytometry analyses of thymocytes, splenocytes, LN, and BM cells from WT mice (WT), Vav2<sup>ko</sup> (Vav2), or Vav3<sup>ko</sup> (Vav3) mice are shown. Fig. S2 shows the strategy for generation of Vav3-defi-



**Figure 1.** Involvement of Vav proteins in antigen receptor signaling in T and B lymphocytes. (A) Expression of the Vav1, Vav2, and Vav3 in lymphocytes. Cells were harvested, and equal amounts of total RNA (5  $\mu$ g) were analyzed by RNase protection assay with specific riboprobes as indicated. As a loading control, the same preparations of RNAs were hybridized with antisense probe for GAPDH (bottom). This experiment was performed several times using different sets of specific probes with similar results. (B) TCR- and BCR-induced tyrosine phosphorylation of Vav1, Vav2, and Vav3. Freshly isolated LN T cells (top) or splenic B cells (bottom) were either left untreated (–) or treated with stimulatory antigen receptor antibodies (+) for 2 h as indicated. Vav proteins were immunoprecipitated (i.p.) from cell lysates with specific antibodies, and the level of phosphotyrosine was analyzed by immunoblotting with antiphosphotyrosine (anti-PY) antibodies (4G10; Upstate Biotechnology). The mobilities of Vav proteins are indicated. The data are representative of three independent experiments; the relative levels of phosphorylation of individual Vav proteins varied somewhat between experiments. (C) Jurkat T cells were transfected with GFP-tagged full-length Vav1, Vav2, or Vav3 cDNA constructs. Cells were incubated for 16–24 h and then either left untreated or treated with stimulatory anti-CD3 antibodies for 5 h, cytospun, and processed for immunofluorescence microscopy. Staining with rhodamine-phalloidine (red, top) visualized accumulations of F-actin (patches) indicated by white arrows. GFP fluorescence is shown in green (middle). Two colors were merged (bottom) to visualize staining overlap.





**Figure 2.** Defective lymphocyte development in Vav-deficient mice. (A) Impaired development of Vav-deficient T cells. Single cell suspensions of thymocytes or LN cells from WT mice, Vav1<sup>ko</sup> (Vav1), Vav1/2<sup>ko</sup> (Vav1/2), Vav1/3<sup>ko</sup> (Vav1/3), Vav2/3<sup>ko</sup> (Vav2/3), or Vav1/2/3<sup>ko</sup> (Vav1/2/3) mice were counted and stained with fluorescent antibody conjugates and analyzed by flow cytometry as indicated. DN thymocytes were gated as live cells that did not stain with CD4, CD8, B220,  $\gamma\delta$  TCR, Gr1, and Mac1, as described in Materials and Methods. Total numbers and percentages of gated lymphocyte populations are shown for representative samples ( $n > 5$ ). (B) Impaired development of Vav-deficient B cells. Cell suspensions of BM or spleen samples were prepared as above and stained and analyzed by flow cytometry as indicated. Total numbers and percentages of gated lymphocyte populations are shown for representative samples ( $n > 5$ ). Data are displayed as dot plots with logarithmic scale.

cient mice. Southern blotting analyses of targeted ES cells used to generate chimeric mice and their offspring and Western blotting analyses of Vav3<sup>ko</sup> splenocyte lysates with Vav3-specific antibodies showing no full-length or truncated Vav3 protein present are also shown. Fig. S3 presents analyses of humoral TD responses in Vav3-deficient mice. Concentrations of TNP-specific IgM, IgG1, IgG2b, and IgG3 antibodies analyzed at various time points post-primary or secondary immunization with TNP-conjugated keyhole hemagglutinin protein (TNP-KLH) are shown. Fig. S4 presents analyses of cytokine production by Vav-deficient T cells showing defective IL-2 and IFN $\gamma$  production. Figs. S1–4 are available online at <http://www.jem.org/cgi/content/full/jem.20030874/DC1>.

## Results

*All Three Vav Proteins Are Tyrosine Phosphorylated and Recruited by Antigen Receptors in Primary T and B Lymphocytes.* Unlike Vav1, whose expression is predominantly restricted to hematopoietic lineage cells, Vav2 and Vav3 are expressed more broadly (16, 17). To examine their expression

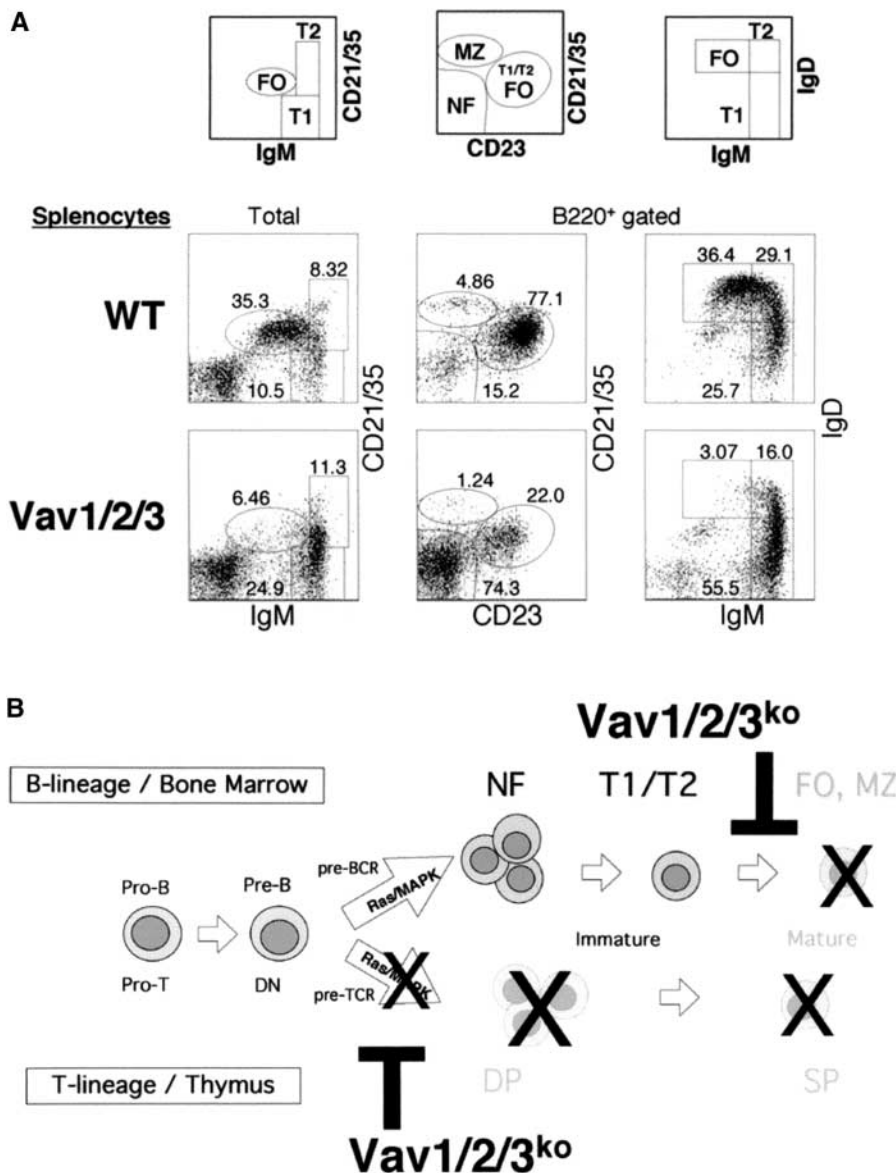
in lymphocyte subsets, we used an RNase protection assay and found that Vav1, Vav2, and Vav3 were all expressed in immature and mature populations of T and B cells (Fig. 1 A; unpublished data). Although tyrosine phosphorylation of Vav1 in response to antigen receptor stimulation in lymphocytes is well documented (19, 20), less is known about phosphorylation of Vav2 or Vav3 in T and B lymphocytes (21, 22, 35, 36). To this end, we stimulated freshly isolated LN T cells or splenic B cells with either anti-CD3 or with anti-IgM antibodies, respectively, and found that all three Vav proteins were inducibly tyrosine phosphorylated (Fig. 1 B). Moreover, since Vav1 is implicated in actin cytoskeleton changes that accompany formation of the immunological synapse (23, 24, 37), we examined the localization of all three Vav proteins fused to a GFP tag in anti-CD3-stimulated Jurkat T cells (Fig. 1 C). We found that upon treatment with anti-CD3 antibodies, each one of the Vav fusion proteins colocalized with antigen receptor formed caps and F-actin patches (Fig. 1 C). Together, these results indicate that all three Vav family proteins become tyrosine

phosphorylated and colocalize with actin at the TCR caps in lymphocytes.

*Mice with Combined Vav Deficiencies Reveal a Major Compensatory Role for Vav3 in T Cell Development.* Previous studies of *Vav1*<sup>ko</sup> mice demonstrated partial defects in T cell development and activation (27–30); however, *Vav1*<sup>ko</sup> T cells still retained substantial developmental and functional capability. To determine if this was due to the expression of *Vav2* in these cells, we examined mice lacking both *Vav1* and *Vav2* (*Vav1/2*<sup>ko</sup>). As expected based on previous studies (27–30, 33, 34), *Vav1*<sup>ko</sup> mice showed diminished numbers of thymocytes and peripheral T cells (approximately two- to threefold; Fig. 2 A), whereas *Vav2*<sup>ko</sup> mice had normal T cell development (Fig. S1 A, available at <http://www.jem.org/cgi/content/full/jem.20030874/DC1>). *Vav1/2*<sup>ko</sup> mice had populations of CD4<sup>+</sup>8<sup>-</sup> (DN), CD4<sup>+</sup>8<sup>+</sup> (double positive [DP]), and CD4<sup>+</sup>8<sup>-</sup> or CD4<sup>-</sup>8<sup>+</sup> (single positive [SP]) thymocytes and

peripheral T cells comparable to those in *Vav1*<sup>ko</sup> mice (Fig. 2 A). Moreover, subsets of DN thymocytes distinguished based upon expression of CD44 and CD25 (DN1, CD44<sup>+</sup>25<sup>-</sup>; DN2, CD44<sup>+</sup>25<sup>+</sup>; DN3, CD44<sup>-</sup>25<sup>+</sup>; and DN4, CD44<sup>-</sup>25<sup>-</sup>) were similar in *Vav1*<sup>ko</sup> and *Vav1/2*<sup>ko</sup> mice (Fig. 2 A). These experiments suggest that *Vav2* has no major function in the development of T lymphocytes in the absence of *Vav1*.

To determine if *Vav3* compensates for the loss of *Vav1* in *Vav1*<sup>ko</sup> T cells, we generated *Vav3*-deficient (*Vav3*<sup>ko</sup>) mice using standard gene targeting approaches (Fig. S2, available at <http://www.jem.org/cgi/content/full/jem.20030874/DC1>). Although *Vav3*<sup>ko</sup> mice showed normal T cell development (Fig. S1 A), the lack of *Vav3* significantly exacerbated the T cell phenotype of *Vav1*<sup>ko</sup> mice (Fig. 2 A). Thus, populations of DN4, DP and SP thymocytes, and peripheral T cells were all significantly reduced in *Vav1/3*<sup>ko</sup> mice compared with *Vav1*<sup>ko</sup> mice (Fig.



**Figure 3.** Late developmental block in *Vav*-null B cells. (A) Late developmental block at the T1/T2 stage and accumulation of NF cells in *Vav*-null mice. Splenocyte suspensions from WT mice or *Vav1/2/3*<sup>ko</sup> (*Vav1/2/3*) were stained and analyzed by flow cytometry as indicated. Percentages of gated lymphocyte populations are shown for representative samples ( $n = 3$ ). Total numbers of corresponding populations are shown in Table I. Data are displayed as dot plots with logarithmic scale. (B) Differential requirement for *Vav* family proteins in T and B lymphocyte development and MAP kinase activation. Schematic representation of developmental defects of lymphoid cells observed in *Vav1/2/3*<sup>ko</sup> (*Vav1/2/3*) mice. See Results and Discussion for details.

**Table 1.** *B Lymphocyte Populations in Vav-null Mice*

	Total	Newly formed	T1	T2	Follicular	MZ
Wild type	94.5 ± 38.3	4.5 ± 1.4	5.3 ± 2.9	6.3 ± 2.0	12.9 ± 8.6	1.9 ± 1.3
Vav null	88.7 ± 7.6	20.6 ± 3.9	15.6 ± 3.5	2.5 ± 1.1	0.7 ± 0.3	0.5 ± 0.1

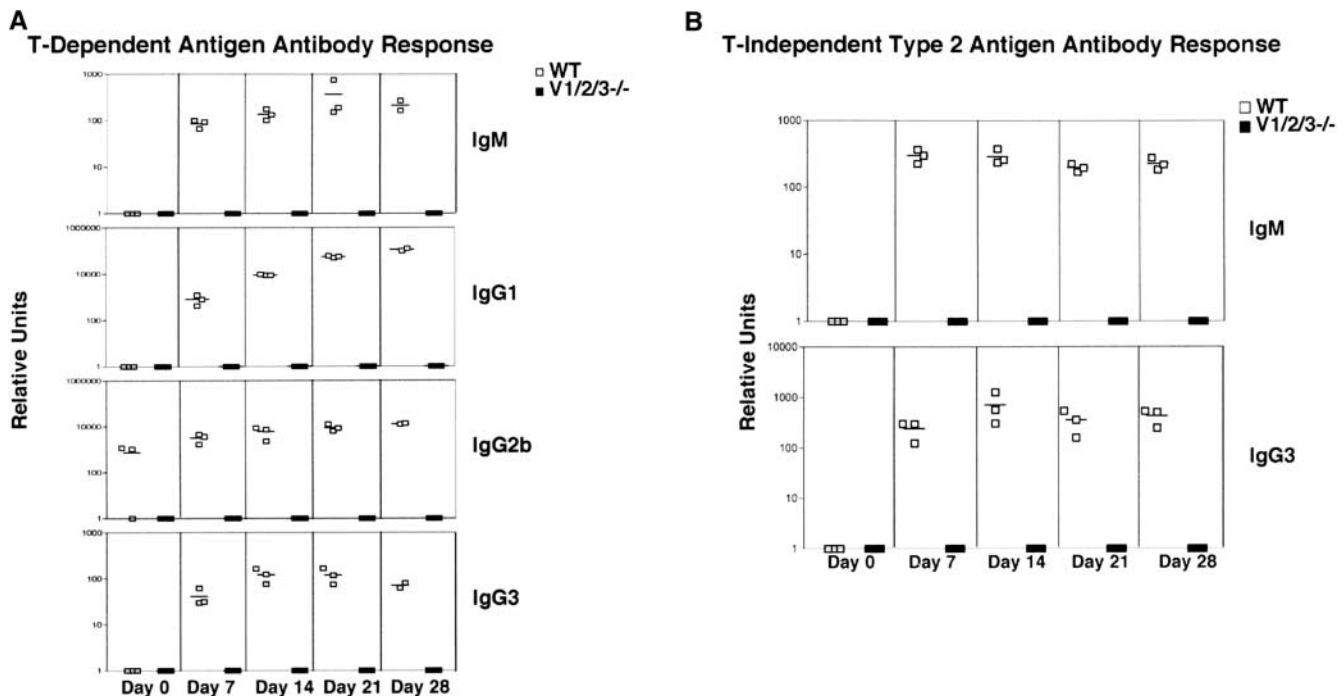
Cells from spleens of WT and Vav1/2/3<sup>ko</sup> (Vav-null) mice were counted and stained for cell surface expression of B220, IgM, IgD, CD23, and CD21/35, as shown in Fig. 3 A, and analyzed by flow cytometry. Numbers of indicated populations are shown for newly formed (NF or B220+CD21<sup>int</sup>CD23<sup>-</sup>), transitional 1 (T1 or B220+CD21<sup>lo</sup>IgM<sup>hi</sup>IgD<sup>lo</sup>), transitional 2 (T2 or B220+CD21<sup>hi</sup>IgM<sup>hi</sup>IgD<sup>hi</sup>), follicular (FO or B220+CD21<sup>int</sup>CD23<sup>hi</sup>IgM<sup>lo</sup>IgD<sup>hi</sup>), and marginal zone (MZ or B220+CD21<sup>hi</sup>CD23<sup>lo</sup>) cells. Data are presented as means ± SDs and represent multiples of 10<sup>6</sup> cells (*n* = 3). Total cells were counted by gating live lymphoid cells.

2 A), indicating that Vav3 is critical for development of T cells lacking Vav1.

**Vav-null T Cells Are Developmentally Blocked at the DN3 Stage.** To establish the role of the entire family of Vav proteins in lymphocytes, we generated mice lacking Vav1, Vav2, and Vav3 (Vav1/2/3<sup>ko</sup>). These mice were viable and showed no gross abnormalities of internal organs upon histological examination at 4–6 wk of age (unpublished data). Vav1/2/3<sup>ko</sup> mice had drastically reduced thymocyte numbers (~50- to 100-fold compared with WT) with few DP and SP cells or peripheral T cells (which were reduced by ~100-fold compared with WT) (Fig. 2 A). The block in thymocyte development occurred at the DN3 to DN4 tran-

sition, since Vav1/2/3<sup>ko</sup> mice showed predominantly DN3 cells and only few DN4 cells (Fig. 2 A). In addition, large DN4 cells were essentially absent in Vav1/2/3<sup>ko</sup> mice (<2% compared with 15–20% in WT; unpublished data). These results clearly demonstrate that Vav proteins are indispensable in T cell development, although we could still detect few DP cells in the thymus and few T cells in the peripheral lymphoid organs of Vav-null mice suggesting some “leakiness” of the developmental block (Fig. 2 A and Fig. 3 B).

Notably, mice lacking Vav2 and Vav3 (Vav2/3<sup>ko</sup>) showed no obvious T cell abnormalities (Fig. 2 A), indicating that expression of Vav1 alone is sufficient to support T cell development. Thus, these results suggest a hierarchy of



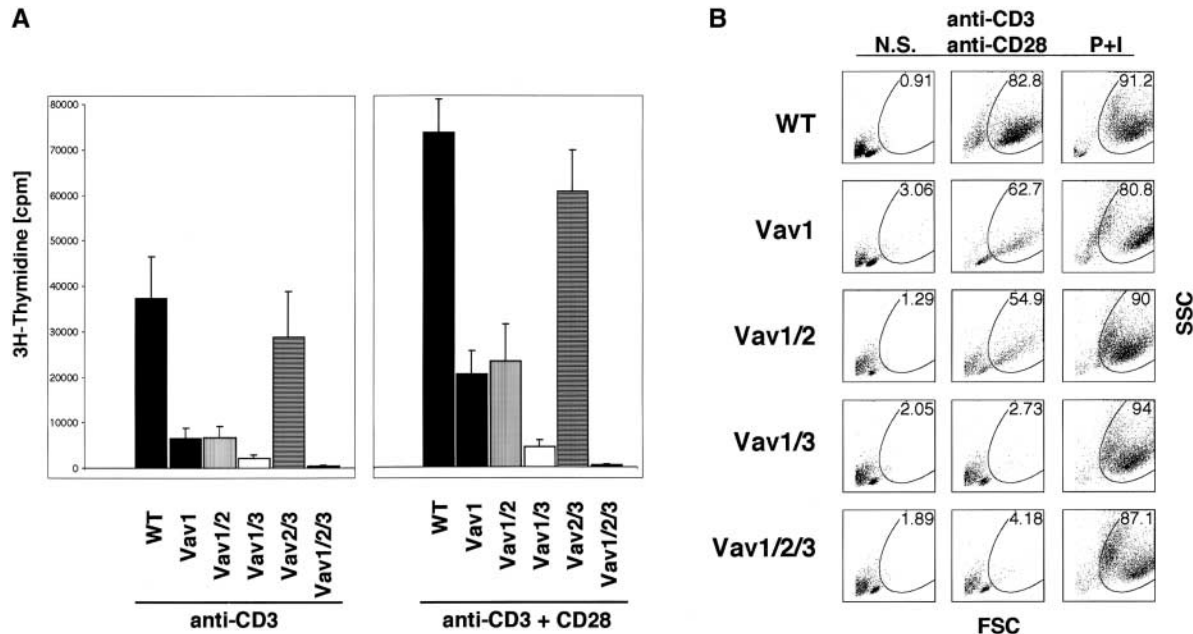
**Figure 4.** Failure to mount antigen-specific humoral responses in Vav-null mice. (A) TD responses. WT or Vav1/2/3<sup>ko</sup> mice were immunized with trinitrophenyl (TNP)-conjugated keyhole haemagglutinin protein (TNP-KLH), and the concentrations of TNP-specific IgM, IgG1, IgG2b, and IgG3 antibodies were analyzed at various time points post-primary or secondary immunization by antigen-specific ELISA as described in Materials and Methods. Each symbol represents an individual mouse, and mean values are indicated by horizontal bars. WT mice (□) and Vav1/2/3<sup>ko</sup> mice (■). (B) T-independent type 2 (TI-2) responses. WT or Vav1/2/3<sup>ko</sup> mice were immunized with a TI-2 antigen, TNP-Ficoll, and the concentrations of TNP-specific IgM and IgG3 antibodies were analyzed as in A. Each symbol represents an individual mouse and mean values are indicated by horizontal bars. WT mice (□) and Vav1/2/3<sup>ko</sup> mice (■).

Vav proteins in T cell development with Vav1 playing a major role that can be compensated for, at least in part, by Vav3. On the other hand, a compensatory function of Vav2 is only apparent in developing T cells lacking both Vav1 and Vav3.

**Vav-null B Cells Show a “Leaky” Developmental Block at the T1/T2 Stage.** In contrast to T cells, immature populations of B cells in BM including progenitors (B220<sup>low</sup>CD43<sup>hi</sup>IgM<sup>-</sup>), precursors (B220<sup>+</sup>CD43<sup>low</sup>IgM<sup>-/low</sup>), and immature (B220<sup>+</sup>CD43<sup>low</sup>IgM<sup>+</sup>) B cells appeared to develop normally in mice with single or combined Vav deficiencies (Fig. 2 B; Fig. S1 B; unpublished data). Notably, Vav1/2<sup>ko</sup>, Vav1/3<sup>ko</sup>, and Vav1/2/3<sup>ko</sup> mice showed reduced populations of “recirculating” BM B220<sup>hi</sup>IgM<sup>lo</sup> B cells (Fig. 2 B). Mature IgM<sup>lo</sup>IgD<sup>hi</sup> B cells appeared normal in Vav2<sup>ko</sup>, Vav3<sup>ko</sup>, and Vav2/3<sup>ko</sup> mice and marginally decreased in Vav1<sup>ko</sup> and Vav1/3<sup>ko</sup> mice (Fig. 2 B and Fig. S1 B) but were more significantly decreased in Vav1/2<sup>ko</sup> and Vav1/2/3<sup>ko</sup> mice (approximate two- to threefold reduction compared with WT; Fig. 2 B). To more precisely determine the developmental defect in B cells lacking all Vav proteins, we used anti-CD21/35 and anti-CD23 antibodies to subdivide B cells into populations of newly formed (NF or B220<sup>+</sup>CD21<sup>-</sup>CD23<sup>-</sup>), transitional 1 (T1 or CD21<sup>lo</sup>IgM<sup>hi</sup>IgD<sup>lo</sup>), transitional 2 (T2 or CD21<sup>hi</sup>IgM<sup>hi</sup>IgD<sup>hi</sup>), follicular (FO or CD21<sup>int</sup>CD23<sup>hi</sup>IgM<sup>lo</sup>IgD<sup>hi</sup>), and marginal zone (MZ or CD21<sup>hi</sup>CD23<sup>lo</sup>) cells. Strikingly, these analyses revealed

drastically reduced subsets of FO and MZ B cells in Vav1/2/3<sup>ko</sup> mice compared with WT (Fig. 3 A and Table I). Moreover, T2 cells appeared slightly reduced (approximately twofold) in Vav1/2/3<sup>ko</sup> mice, whereas the numbers of T1 and NF cells were increased, indicating a “leaky” developmental block at T1/T2 stage (Fig. 3, A and B; Table I).

**Vav-null Mice Fail To Mount Humoral Responses to TD and TI Antigens.** To determine if lymphocytes generated in Vav-null mice are functional in vivo, we immunized Vav1/2/3<sup>ko</sup> or WT mice with the TD antigen, trinitrophenyl (TNP)-conjugated keyhole hemagglutinin (TNP-KLH), and measured the concentrations of TNP-specific IgM, IgG1, IgG2b, and IgG3 antibodies at various time points post-primary and secondary immunization (Fig. 4 A). In this regard, although Vav1<sup>ko</sup> and Vav2<sup>ko</sup> mice were shown previously to have impaired production of switched antibodies against TD antigens, their IgM responses were completely normal (33, 38). In contrast, Vav3<sup>ko</sup> mice showed robust humoral responses and the ability to isotype switch (Fig. S3, available at <http://www.jem.org/cgi/content/full/jem.20030874/DC1>). Strikingly, Vav1/2/3<sup>ko</sup> mice completely failed to produce any detectable anti-TNP-specific IgM antibodies at any time point after immunization (Fig. 4 A). Moreover, Vav1/2/3<sup>ko</sup> mice generated no detectable IgG1, IgG2b, or IgG3 anti-TNP antibodies (Fig. 4 A). These data demonstrate that the en-



**Figure 5.** Defective proliferative responses and blast formation by Vav-deficient T cells. (A) Impaired proliferation of antigen receptor-stimulated T cells lacking Vav proteins. Purified LN T cells from WT (black bars), Vav1<sup>ko</sup> (Vav1, white dotted bars), Vav1/2<sup>ko</sup> (Vav1/2, black dotted bars), Vav1/3<sup>ko</sup> (Vav1/3, white bars), Vav2/3<sup>ko</sup> (Vav2/3, horizontal striped bars), or Vav1/2/3<sup>ko</sup> (Vav1/2/3, gray bars) mice were cultured in either media alone or in the presence of stimulatory anti-CD3 and/or anti-CD28 antibodies as indicated. Proliferation was determined at 48 h by <sup>3</sup>H-thymidine incorporation and scintillation counting. The data are displayed as raw cpm values. The results are representative of three independent experiments. All assays were conducted in triplicates, and SDs are shown as vertical lines. (B) Defective blast formation of antigen receptor-stimulated Vav-deficient T cells. LN T cells from indicated mice were cultured in either media alone or in the presence of stimulatory anti-CD3 and anti-CD28 antibodies, or with phorbol myristyl acetate (PMA) and ionomycin (I) for 36 h as described in Materials and Methods. Cells were then harvested and analyzed by flow cytometry. Forward scatter dot plots of CD4<sup>+</sup> gated cells in representative samples are shown ( $n > 5$ ) and numbers represent percentages of blasting cells.



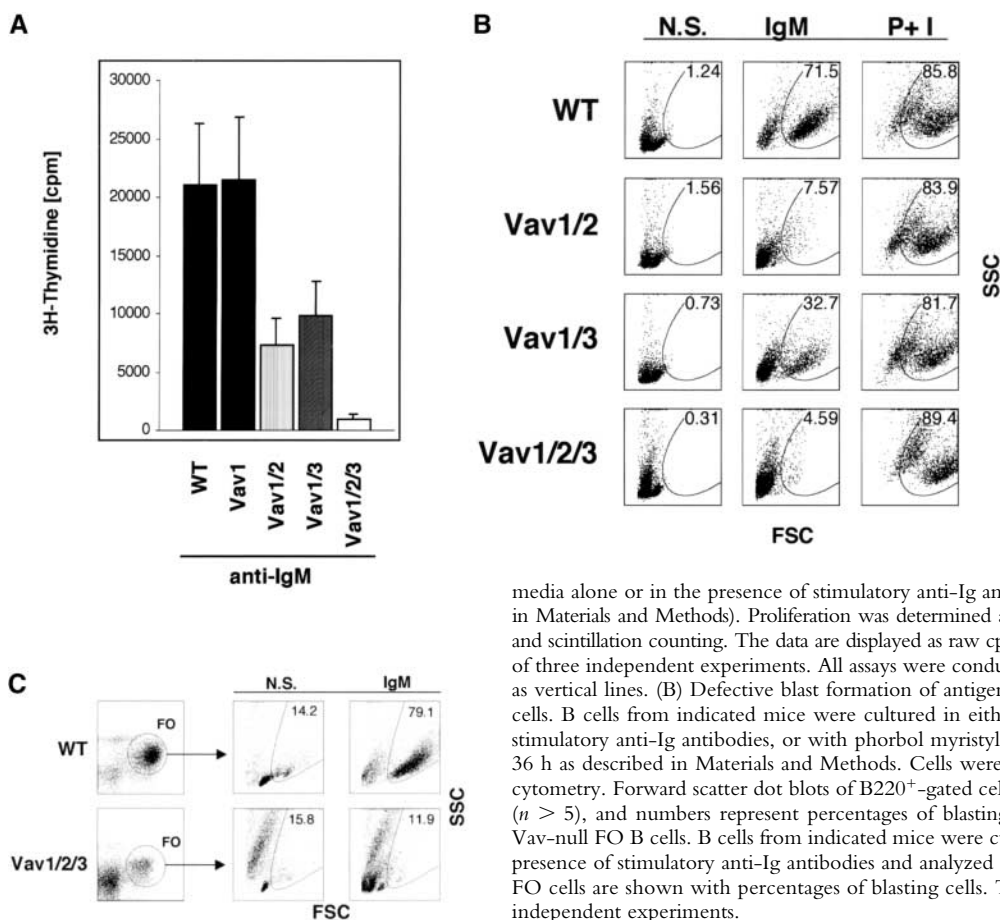
tire Vav family is absolutely required for generation of antigen-specific TD humoral responses.

To determine if Vav-null mice could mount TI-2 responses, we immunized WT or Vav1/2/3<sup>ko</sup> mice with TNP-conjugated Ficoll (TNP-Ficoll) and determined concentrations of TNP-specific IgM and IgG3 antibodies after primary or secondary immunization. In this regard, Vav1<sup>ko</sup> mice were shown previously to generate normal responses to TI-2 antigens (38), although Vav2<sup>ko</sup> and Vav1/2<sup>ko</sup> mice had defective TI-2 responses (33, 34). Strikingly, Vav1/2/3<sup>ko</sup> mice completely failed to produce any detectable anti-TNP antibodies in response to TI-2 antigen at any time point after immunizations (Fig. 4 B). Together, these data show a strict requirement for Vav proteins in mounting both TD and TI humoral responses.

**Vav3 Functionally Compensates for the Loss of Vav1 in T Cell Proliferative Responses.** Next, we examined in vitro proliferative responses of T cells lacking Vav proteins. Consistent with previous reports (27–30), Vav1<sup>ko</sup> T cells showed a partial defect in proliferative responses to stimulation with anti-CD3 with or without anti-CD28 antibodies as measured by <sup>3</sup>H-thymidine incorporation (Fig. 5 A). Notably, Vav1/2<sup>ko</sup> T cells proliferated equally to Vav1<sup>ko</sup> T cells, indicating that Vav2 does not substantially compensate for the loss of Vav1 (Fig. 5 A). Strikingly, however, T cells from Vav1/3<sup>ko</sup> or Vav1/2/3<sup>ko</sup> mice vir-

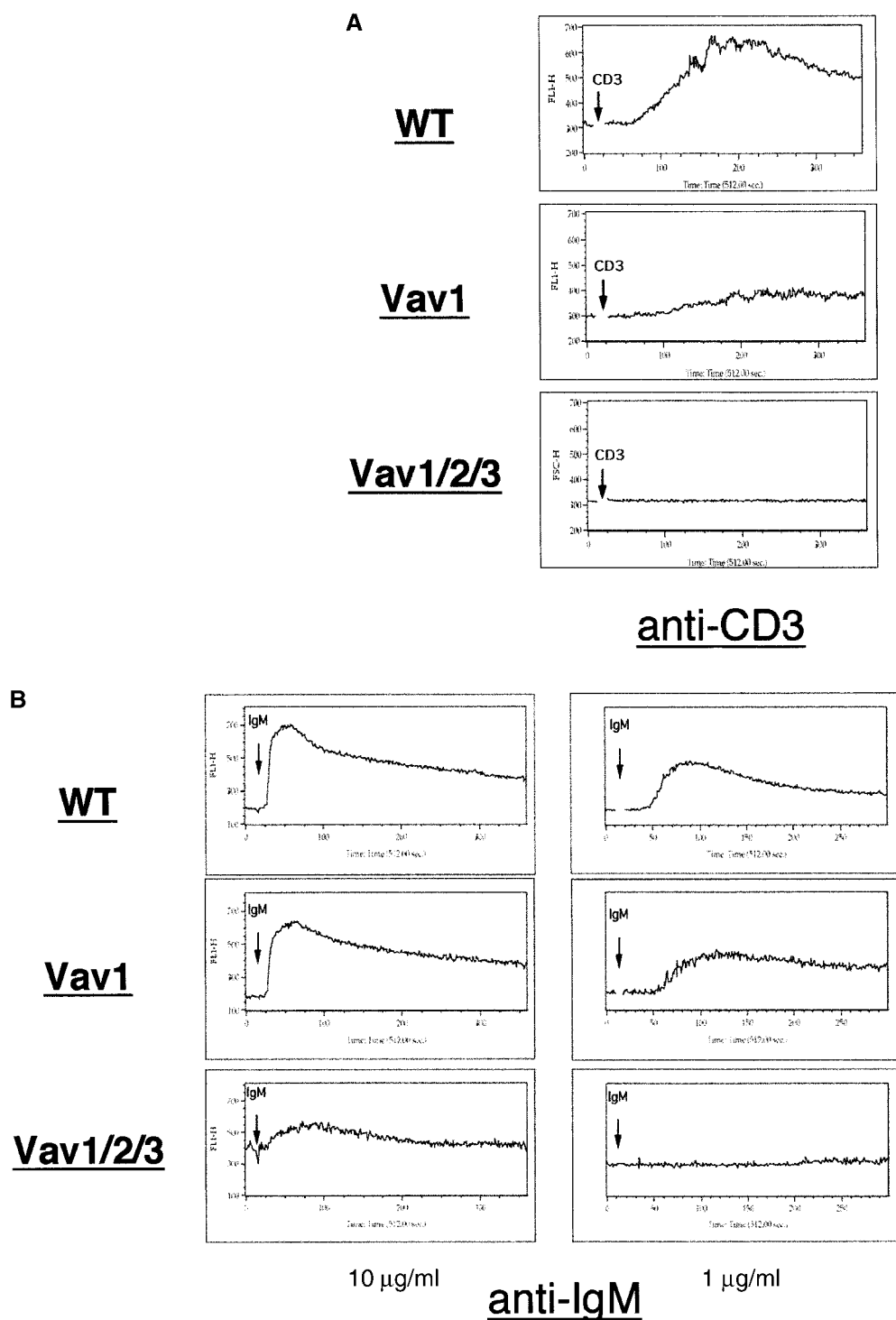
tually failed to incorporate <sup>3</sup>H-thymidine in response to stimulation with anti-CD3 and anti-CD28 antibodies compared with WT (Fig. 5 A). These data demonstrate that, unlike Vav2, Vav3 appears critical for function of T cells that do not express Vav1. T cells from Vav2<sup>ko</sup> or Vav3<sup>ko</sup> mice showed normal proliferative responses, indicating that neither of these proteins is essential in T cells expressing Vav1 (unpublished data).

As an alternative measure of T cell proliferation, we also assayed for the production of large (blasting) cells. In this assay, nonstimulated fresh LN T cells remained small at various time points in culture (indicated by their low forward and side scatter; Fig. 5 B, left). On the other hand, cultures of WT T cells stimulated with anti-CD3 and anti-CD28 antibodies contained predominantly blasting cells (Fig. 5 B, middle). In contrast, the ability of Vav1<sup>ko</sup> or Vav1/2<sup>ko</sup> T cells to generate blasts was reduced by approximately three- to fourfold (Fig. 5 B), whereas T cells from Vav1/3<sup>ko</sup> or Vav1/2/3<sup>ko</sup> mice virtually failed to produce blasts in response to anti-CD3 and anti-CD28 antibody stimulation. Thus, unlike Vav1<sup>ko</sup> T cells, which retain a significant proliferative capacity, T cells lacking both Vav1 and Vav3 have severely disrupted proliferative responses. All T cells responded normally to PMA and ionomycin (Fig. 5 B; unpublished data), indicating no intrinsic cell cycle defects in Vav-deficient T cells.



**Figure 6.** Defective proliferative responses and blast formation by Vav-deficient B cells. (A) Impaired proliferation of antigen receptor-stimulated B cells lacking Vav proteins. Purified spleen B cells from WT (black bars), Vav1<sup>ko</sup> (Vav1, white dotted bars), Vav1/2<sup>ko</sup> (Vav1/2, black dotted bars), Vav1/3<sup>ko</sup> (Vav1/3, diagonal striped bars), or Vav1/2/3<sup>ko</sup> (Vav1/2/3, white bars) mice were cultured in either

media alone or in the presence of stimulatory anti-Ig antibodies as indicated (and as described in Materials and Methods). Proliferation was determined at 48 h by <sup>3</sup>H-thymidine incorporation and scintillation counting. The data are displayed as raw cpm values. The results are representative of three independent experiments. All assays were conducted in triplicates, and SDs are shown as vertical lines. (B) Defective blast formation of antigen receptor-stimulated Vav-deficient B cells. B cells from indicated mice were cultured in either media alone, or in the presence of stimulatory anti-Ig antibodies, or with phorbol myristyl acetate (PMA) and ionomycin (I) for 36 h as described in Materials and Methods. Cells were then harvested and analyzed by flow cytometry. Forward scatter dot blots of B220<sup>+</sup>-gated cells in representative samples are shown ( $n > 5$ ), and numbers represent percentages of blasting cells. (C) Defective proliferation of Vav-null FO B cells. B cells from indicated mice were cultured in either media alone or in the presence of stimulatory anti-Ig antibodies and analyzed as in B. Forward scatter blots of gated FO cells are shown with percentages of blasting cells. The results are representative of three independent experiments.



**Figure 7.** Disruption of  $Ca^{2+}$  signaling in Vav-null lymphocytes. (A) LN T cell and spleen B cells (B) were loaded with Fluo4-AM as described in Materials and Methods. T cell suspensions were also surface stained with anti-CD4 antibodies conjugated to APC. Cell suspensions were analyzed by flow cytometry. Samples were run at  $\sim 10^3$  cells/s for 15 s to determine background, and then for an additional 7 min after stimulation. T cells were stimulated by the addition of anti-CD3 antibodies (2C11) immediately followed by anti-hamster antibodies; B cells were stimulated by the addition of anti-IgM antibodies at indicated concentrations. Data was analyzed using FlowJo software and displayed as histograms of relative fluorescence intensity over time (in seconds). Representative analyses are shown ( $n > 3$ ).

In addition, we examined cytokine production by T cells in response to TCR stimulation and found that the production of IL-2 and IFN $\gamma$  appeared equally diminished in Vav1<sup>ko</sup> and Vav1/2<sup>ko</sup> T cells compared with WT but was virtually undetectable in Vav1/3<sup>ko</sup> and Vav1/2/3<sup>ko</sup> T cells (Fig. S4, available at <http://www.jem.org/cgi/content/full/jem.20030874/DC1>). Together, these results demonstrate that in contrast to mice lacking individual Vav

proteins no functional T cells can be generated in mice lacking all Vav family members. Moreover, individual Vav proteins contribute unevenly in this lineage, since Vav3 but not Vav2 has a major compensatory function.

*Vav-null B Cells Fail To Proliferate in Response to Ig Receptor Stimulation.* To determine if B cells developing in Vav-deficient mice still retain their ability to proliferate in vitro in response to stimulation with anti-Ig antibodies, we

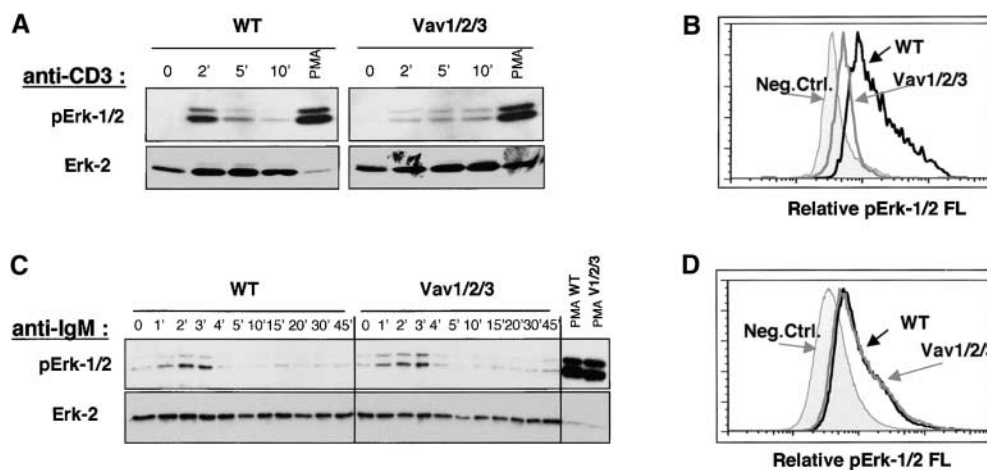
used  $^3\text{H}$ -thymidine incorporation assay. Although responses of  $\text{Vav1}^{\text{ko}}$  B cells to Ig cross-linking were comparable to WT, proliferation of  $\text{Vav1/2}^{\text{ko}}$  B cells was severely reduced (Fig. 6 A). Thus, consistent with previous reports (33, 34),  $\text{Vav1}$  and  $\text{Vav2}$  appear to overlap functionally in B cells.  $\text{Vav1/3}^{\text{ko}}$  B cells also showed diminished proliferation, whereas  $\text{Vav1/2/3}^{\text{ko}}$  B cells neither incorporated  $^3\text{H}$ -thymidine nor produced blasting cells in response to anti-Ig stimulation (Fig. 6, A and B). Notably, all B cells proliferated vigorously to stimulation with PMA and ionomycin (Fig. 6 B).

In normal mice, only mature B cells respond to Ig receptor cross-linking with proliferation, whereas the majority of immature cells undergo cell death. Because  $\text{Vav1/2/3}^{\text{ko}}$  B cells show a reduction of mature (FO) B cells and a relative increase in the proportion of immature (NF) cells, it is possible that the failure of such cells to proliferate and to incorporate  $^3\text{H}$ -thymidine is a reflection of these developmental abnormalities. To directly test the ability of mature (FO)  $\text{Vav1/2/3}^{\text{ko}}$  B cells to proliferate, we used flow cytometry and showed that the residual FO B cells present in  $\text{Vav1/2/3}^{\text{ko}}$  mice failed to respond to stimulation with anti-Ig antibodies (Fig. 6 C), demonstrating that the  $\text{Vav}$  family proteins are essential for *in vitro* B cell proliferative responses to anti-Ig stimulation.

**The  $\text{Vav}$  Family Proteins Are Indispensable for  $\text{Ca}^{2+}$  Signaling in Both T Cells and B Cells.** To determine if intracellular signaling was impaired by the loss of  $\text{Vav}$  function, we examined the ability of  $\text{Vav}$ -deficient T and B cells to mobilize  $\text{Ca}^{2+}$  ions.  $\text{Ca}^{2+}$  fluxes induced by anti-CD3 antibody cross-linking were diminished in  $\text{Vav1}^{\text{ko}}$  T cells (Fig.

7 A), consistent with previously published data (23, 24, 29). However,  $\text{Ca}^{2+}$  fluxes were totally disrupted in  $\text{Vav1/2/3}^{\text{ko}}$  T cells even under maximal stimulation conditions (Fig. 7 A). In contrast,  $\text{Vav1}^{\text{ko}}$ ,  $\text{Vav1/2}^{\text{ko}}$ ,  $\text{Vav1/3}^{\text{ko}}$ ,  $\text{Vav2}^{\text{ko}}$ ,  $\text{Vav3}^{\text{ko}}$ , and  $\text{Vav2/3}^{\text{ko}}$  B cells had  $\text{Ca}^{2+}$  fluxes comparable to WT when stimulated with maximal doses of anti-Ig antibodies (Fig. 7 B; unpublished data), whereas  $\text{Vav1}^{\text{ko}}$ ,  $\text{Vav1/2}^{\text{ko}}$ , and  $\text{Vav1/3}^{\text{ko}}$  B cell  $\text{Ca}^{2+}$  fluxes were only minimally reduced at lower doses of stimulatory antibodies (Fig. 7 B; unpublished data). Notably, anti-Ig induced  $\text{Ca}^{2+}$  responses were severely disrupted in  $\text{Vav1/2/3}^{\text{ko}}$  B cells even under maximal stimulation conditions (Fig. 7 B). Together, these data indicate that the  $\text{Vav}$  family is indispensable for  $\text{Ca}^{2+}$  signaling downstream of both the TCR and BCR.

**$\text{Vav}$ -null Mice Reveal a Differential Requirement for  $\text{Vav}$  Proteins in  $\text{Erk}$  Activation in T Cells and B Cells.** Activation of Ras and MAP kinases is crucial for the induction of lymphocyte proliferation and cytokine production. To examine if the loss of  $\text{Vav}$  perturbed the ability of T cells to activate the Ras/MAPK pathway, we probed immunoblots with phospho-specific antibodies that recognize active forms of the p44 and p42 MAP kinases ( $\text{Erk-1}$  and  $\text{Erk-2}$ ). In this regard,  $\text{Vav1}^{\text{ko}}$  T cells showed attenuated  $\text{Erk-1/2}$  activation in response to anti-CD3 antibodies (39 and unpublished data), whereas  $\text{Vav1/2/3}^{\text{ko}}$  T cells virtually failed to activate  $\text{Erk-1/2}$  in response to this treatment (Fig. 8 A). Importantly,  $\text{Erk-1/2}$  could still be induced in  $\text{Vav1/2/3}^{\text{ko}}$  T cells by PMA treatment, indicating a defect upstream of DAG production (Fig. 8 A). Surprisingly, anti-Ig cross-linking resulted in normal  $\text{Erk-1/2}$  activation in  $\text{Vav1/2/3}^{\text{ko}}$



**Figure 8.** Disruption of  $\text{Erk-1/2}$  activation in  $\text{Vav1/2/3}^{\text{ko}}$  T cells but not B cells. (A) Impaired TCR induced  $\text{Erk-1/2}$  phosphorylation in WT and  $\text{Vav1/2/3}^{\text{ko}}$  T cells. Freshly isolated LN T cells were either left untreated (0') or treated with anti-CD3 antibodies immediately followed by anti-hamster antibodies for the indicated period of time. PMA was used as a positive control at 100 ng/ml for 5 min. Cell lysates were prepared by direct lysis with SDS buffer followed by boiling and sonication and then analyzed by immunoblotting with anti-phospho-p42/p44  $\text{Erk-1/2}$  antibodies (as described in Materials and

Methods). Protein loading was verified by stripping and reprobing blots with anti- $\text{Erk-2}$  antibodies. The data are representative of three independent experiments. (B) Disruption of  $\text{Erk-1/2}$  activation in thymocytes undergoing "β selection." Thymocytes from WT or  $\text{Vav1/2/3}^{\text{ko}}$  mice were surface stained with CyC-labeled anti-CD4 and anti-CD8 and FITC-labeled anti-CD25 antibodies, and were permeabilized and stained with PE-labeled anti-phospho-p42/p44  $\text{Erk-1/2}$  antibodies as described in Materials and Methods. Histograms representing phospho- $\text{Erk-1/2}$  staining intensity are shown for gated populations of  $\text{CD4}^{\text{lo}}\text{CD8}^{\text{lo}}\text{CD25}^{\text{hi}}$  "large" cells from WT or  $\text{Vav1/2/3}^{\text{ko}}$  mice as indicated. (C) Normal BCR induced  $\text{Erk-1/2}$  activation in  $\text{Vav1/2/3}^{\text{ko}}$  B cells. Freshly isolated spleen B cells were either left untreated (0') or treated with anti-Ig antibodies or PMA and analyzed as in A. (D) Normal  $\text{Erk-1/2}$  activation in large pro-B cells. BM cells from WT or  $\text{Vav1/2/3}^{\text{ko}}$  mice were surface stained with CyC-labeled anti-B220 and FITC-labeled anti-CD43 antibodies and then were permeabilized and stained with PE-labeled anti-phospho-p42/p44  $\text{Erk-1/2}$  antibodies as described in Materials and Methods. Histograms representing phospho- $\text{Erk-1/2}$  staining intensity are shown for gated populations of  $\text{B220}^{\text{lo}}\text{CD43}^{\text{hi}}$  large cells from WT or  $\text{Vav1/2/3}^{\text{ko}}$  mice as indicated.

$3^{\text{ko}}$  B cells and in B cells lacking any other combination of Vav proteins (Fig. 8 C; unpublished data). Thus, it appears that in contrast to T cells, Vav proteins are not required for Erk-1/2 activation in B cells, indicating a fundamentally distinct mechanism of antigen-receptor signal transduction in cells of the T and B lineages.

Since Ras/MAPK signaling is implicated in the DN to DP transition in the thymus and pre-B cell differentiation and BM egress (2–9), we investigated if Erk-1/2 activity could be detected in large DN3 cells and in large pro-B cells in WT and Vav1/2/3<sup>ko</sup> mice. In this regard, intracytoplasmic staining of ex vivo BM cells revealed increased levels of activated Erk-1/2 in large pro-B cells in both WT and Vav1/2/3<sup>ko</sup> mice, whereas large DN3 cells in Vav1/2/3<sup>ko</sup> mice had significantly reduced levels of activated Erk-1/2 compared with WT (Fig. 8 B). Phospho-Erk-1/2 staining was equivalent in all cells treated with PMA (unpublished data). Collectively, these data indicate that the loss of Vav proteins leads to a disruption of Ras/MAP kinase signaling in both immature and mature T cells but not in developing or mature B cells.

## Discussion

In this report, we examined the roles of Vav Rho-guanine nucleotide exchange factors in lymphocytes. We describe the generation and analyses of mice in which all known members of the Vav family, Vav1, Vav2, and Vav3, have been inactivated via gene targeting. We show that in the lymphoid system this combined Vav deficiency prevents the development of functionally mature T or B cells and disrupts signaling downstream of both T and B cell antigen receptors.

In primary lymphocytes, all three Vav proteins are functionally coupled to signaling events downstream of the TCR and the BCR as indicated by their inducible tyrosine phosphorylation and recruitment to structures associated with the immunological synapse. Despite their common ancestry, conserved structural features, and the ability of each Vav protein to couple with lymphocyte antigen receptors, individual Vav proteins have both common and specialized functions in lymphocytes *in vivo*. For example, our analyses of Vav1/2<sup>ko</sup> mice indicate that Vav2 does not functionally compensate for Vav1 in T cells as it does in B cells (33, 34). Thus, Vav2 may be specifically involved in signaling downstream of the BCR but not the TCR. Nevertheless, Vav2 is able to substitute for some functions of Vav3, since there is a more severe developmental block of Vav1/2/3<sup>ko</sup> thymocytes compared with Vav1/3<sup>ko</sup>. On the other hand, analyses of lymphocytes in Vav2/3<sup>ko</sup> mice indicate that Vav1 itself is sufficient to support apparently normal development and function of T and B cells, suggesting mainly auxiliary role(s) for Vav2 and Vav3. Thus, although we have not yet identified any strictly nonredundant functions for any of the individual Vav proteins, genetic evidence presented in this report clearly demonstrates that of the ~50 Rho-GEFs that exist in mammalian cells (18) the

Vav family Rho-GEFs play a role that is both specialized and nonredundant in T and B lymphocytes (Fig. 3 B).

Developing T and B cells receive differentiation and expansion signals from distinct classes of surface receptors, such as cytokine receptors and the pre-TCR or pre-BCR, all of which could conceivably involve Vav proteins. For example, Vav has been implicated in signaling by receptors sharing the common  $\gamma$ -chain, which have crucial functions in lymphocyte ontogeny (1). In this regard, normal development of DN1, DN2, and DN3 thymocytes as well as pro-B and pre-B cells in Vav1/2/3<sup>ko</sup> mice indicates that Vav proteins are not required in signaling by receptors such as the IL-7R. In fact, both WT and Vav1/2/3<sup>ko</sup> BM cells proliferate in response to IL-7 treatment *in vitro* (unpublished data). Moreover, the accumulation of immature populations of B cells, including NF and T1 (Table I), in Vav1/2/3<sup>ko</sup> mice indicates that Vav proteins are not required for signaling downstream of the pre-BCR, yet they appear essential for the transition to more mature developmental stages that may require stronger antigen receptor signals. Clearly, our data indicate that in B lineage cells Vav proteins are superfluous in Ras/MAPK activation through the BCR (Fig. 8, C and D) and thus may allow for development of all immature populations of B cells in Vav1/2/3<sup>ko</sup> mice despite the uncoupling of BCR signaling from Ca<sup>2+</sup> fluxes. In this regard, our data are consistent with previously published reports by us and other groups that the activation of Ras/MAPK pathway alone is sufficient to promote development of B lineage cells beyond the pre-B cell stage (7–9).

On the other hand, our observations indicate a critical role for Vav in signaling by the pre-TCR (Fig. 8 B). Specifically, we show that Vav deficiency results in developmental arrest at the DN3 to DN4 transition. This block is not due to defects in assembly and/or expression of TCR $\beta$  chain genes, since a significant fraction of Vav1/2/3<sup>ko</sup> DN3 cells expressed cytoplasmic TCR $\beta$  chains (unpublished data). Rather, drastically reduced numbers of DP thymocytes and large (blasting) DN3/DN4 thymocytes in Vav1/2/3<sup>ko</sup> mice compared with WT suggest a pivotal role for Vav proteins in pre-TCR signaling.

Several studies have implicated the activation of the Ras/MAPK pathway as a crucial event in the induction of differentiation and expansion of both T and B cell precursors (2–9). Given the crucial role for Ras/MAPK signaling in the development of both T and B lineage cells (2–5, 7–9) and the strikingly different requirement for Vav in these cells, it is possible that Vav may not be required for regulation of MAPK, or alternatively, T and B cells may utilize diverse mechanisms of MAPK activation. Indeed, two lines of experimental evidence presented in this report support the latter possibility. First, analyses of MAP kinase (Erk-1/2) activity in antigen receptor-stimulated T cells and B cells indicated a complete disruption of Erk-1/2 signaling in Vav1/2/3<sup>ko</sup> T cells, which is in contrast to normal activation of Erk-1/2 in Vav1/2/3<sup>ko</sup> B cells. Second, whereas WT “large” DN3 thymocytes showed significantly in-



creased Erk-1/2 activity (presumably resulting from pre-TCR signaling), Erk-1/2 activity was drastically reduced in the equivalent population of Vav1/2/3<sup>ko</sup> pro-B cells. Yet, both WT and Vav1/2/3<sup>ko</sup> “large” pro-B cells showed similarly increased Erk-1/2 activity. Together, these results support a view in which the regulation of Ras/MAPK signaling in T and B lineage cells is fundamentally different. In this context, recent evidence suggests that DAG-dependent Ras guanine nucleotide release protein (Ras-GRP1) may be specifically required in the activation of Erk-1/2 in T cells, but not in B cells (40). In fact, the defects in Erk-1/2 activation in Vav1/2/3<sup>ko</sup> T cells resemble those observed in RasGRP1<sup>ko</sup> T cells (Fig. 8) (40), suggesting that TCR induction of Erk-1/2 maybe regulated by Vav via PLC $\gamma$ 1 and RasGRP1. Thus, it is plausible that the TCR complex may require DAG-dependent pathway for Erk-1/2 activation, whereas the BCR could, conceivably, use a DAG-independent mechanism, such as the one involving Grb2/SOS. Indeed, Grb2 haplo-insufficiency in T cells resulted in decreased activity of JNK and p38 but not Erk-1/2 (41), consistent with such a view. Our data also indicate that the induction of Ras/MAPK in B cells does not depend on Ca<sup>2+</sup> signaling and, presumably, PLC $\gamma$  activity.

In this paper, we also extended our previous studies of Vav function in Ca<sup>2+</sup> signaling (23). We found that unlike Vav1<sup>ko</sup> lymphocytes, which show only partial Ca<sup>2+</sup> defects, Vav1/2/3<sup>ko</sup> T and B cells have disrupted Ca<sup>2+</sup> signaling downstream of their antigen receptors. These data are consistent with observations that Vav may be involved in the activation of PLC $\gamma$ 1 (42), although the exact mechanism is still unknown. Nevertheless, our observations establish that Vav family proteins are essential to determine the signaling output of antigen receptors in both T cells and B cells, although they appear to differentially link the TCR and BCR with Ras/MAPK induction.

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