TRAF6 and TRAF2/3 Binding Motifs in CD40 Differentially Regulate B Cell Function in T-Dependent Antibody Responses and Dendritic Cell Function in Experimental Autoimmune Encephalomyelitis

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Expression of the costimulatory molecule CD40 on both B cells and dendritic cells (DCs) is required for induction of experimental autoimmune encephalomyelitis (EAE), and cell-autonomous CD40 expression on B cells is required for primary T-dependent (TD) Ab responses. We now ask whether the function of CD40 expressed by different cell types in these responses is mediated by the same or different cytoplasmic domains. CD40 has been reported to possess multiple cytoplasmic domains, including distinct TRAF6 and TRAF2/3 binding motifs. To elucidate the in vivo function of these motifs in B cells and DCs involved in EAE and TD germinal center responses, we have generated knock-in mice containing distinct CD40 cytoplasmic domain TRAF-binding site mutations and have used these animals, together with bone marrow chimeric mice, to assess the roles that these motifs play in CD40 function. We found that both TRAF2/3 and TRAF6 motifs of CD40 are critically involved in EAE induction and demonstrated that this is mediated by a role of both motifs for priming of pathogenic T cells by DCs. In contrast, the TRAF2/3 binding motif, but not the TRAF6 binding motif, is required for B cell CD40 function in TD high-affinity Ab responses. These data demonstrate that the requirements for expression of specific TRAF-binding CD40 motifs differ for B cells or DCs that function in specific immune responses and thus identify targets for intervention to modulate these responses. *The Journal of Immunology*, 2023, 211: 1814–1822.

The costimulatory molecule CD40 is widely expressed on multiple APC types and has been shown to play an important role in diverse immune responses, including induction/maintenance of autoimmune diseases and primary T-dependent (TD) Ab responses (1). These observations have led us to analyze in our previous studies the identity of the cell populations in which CD40 functions to mediate the autoimmune disease model experimental autoimmune encephalomyelitis (EAE) (2) and the TD germinal center (GC) response that generates class-switched high-affinity Ab production (3, 4).

Multiple sclerosis (MS) is a disease of the CNS that has an autoimmune component and that leads to progressive neurologic deficits and disability (5). Analysis of the immune mechanisms underlying MS has been informed by the murine EAE model, which approximates the pathological features of MS. EAE can be induced by immunization with CNS Ags, including the myelin oligodendrocyte (MOG) protein and peptide (6, 7). In EAE, Ag-specific T cells, including differentiated IFN- γ -producing Th1 and IL-17-producing Th17 cells, are primed by APCs in peripheral lymphoid organs. Th17 cells can also produce IFN- γ (IFN- γ and IL-17 double-positive cells) mediated by T-bet and Runx1 or Runx3, and this developmental flexibility has been linked to the pathogenicity of Th17 cells in multiple autoimmune

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diseases, including EAE (8). Pathogenic T cells cross the bloodbrain barrier, are restimulated by local APCs in the CNS, and recruit additional immune cells that mediate tissue damage (9). In addition to the role of pathogenic Th cells, B cells are also required for EAE induced by recombinant human myelin oligodendrocytes glycoprotein (rhMOG) (10, 11), paralleling the observation that B cell infiltration, Abs to myelin components, and activated complement are routinely detected in patients with MS (12, 13). Our recent study showed that CD40 expression on both B cells and dendritic cells (DCs) is required for induction of EAE by rhMOG and that CD40 on these two cell types functions through distinct and complementary mechanisms, with CD40 on DCs functioning through priming of pathogenic Th17 cells, whereas the role of CD40 on B cells is mediated by production of MOG-specific IgG (2).

In addition to a critical function in EAE, CD40 also plays an essential role in TD Ab production, which is a critical aspect of the adaptive immune response to pathogens and vaccine development. To generate class-switched, high-affinity, Ag-specific Abs, Ag-activated B cells enter the GC, where cognate interaction with T follicular helper cells (Tfhs) results in class-switch recombination, somatic mutation of BCR genes, and clonal expansion with selection of B cells expressing high-affinity Ag-specific BCRs (14, 15). GC

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Abbreviations used in this article: BM, bone marrow; DC, dendritic cell; DLN, draining lymph node; EAE, experimental autoimmune encephalomyelitis; GC, germinal center; KLH, keyhole limpet hemocyanin; MOG, myelin oligodendrocyte; MS, multiple sclerosis; NP, 4-hydroxy-3-nitrophenylacetyl; TD, T dependent; Tfh, T follicular helper cell; WT, wild type.

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B cell responses and TD Ab responses are dependent on costimulatory interactions with T cells, including a requirement for CD40, as demonstrated by the inhibitory effects of blocking anti-CD40 or CD154 Abs and studies of CD40- or CD154-deficient mice (16–19). Using conditional knockouts and bone marrow (BM) chimeras, we have demonstrated that signals provided by CD40 expressed on B cells, but not on DCs, are required for primary GC responses (3, 4).

It has been reported that the CD40 cytoplasmic tail contains multiple functional domains, including two distinct intracellular motifs that regulate the binding to either TRAF6 or TRAF2/3, adapter proteins that are involved in the CD40 signaling cascade (20–22). It was therefore of interest to determine whether the same or distinct CD40 TRAF binding motifs are important for CD40 function of B cells and DCs in immune responses. To address this question, we have generated mice expressing CD40 TRAF binding motif mutations by CRISPR/Cas9 technology. Using these mice, together with BM chimeras, we have identified distinct requirements for TRAF6 and TRAF2/3 binding motifs of CD40 on DC and B cells during both primary GC responses and the autoimmune pathogenesis of EAE.

Materials and Methods

Mice

CD40 cytoplasmic domain mutant mice were generated using CRISPR/Cas9 technology (Genome Modification Core, National Cancer Institute, National Institutes of Health). C57BL/6 (wild-type [WT]) mice were purchased from Charles River Laboratories. CD40^{-/-} (B6.129P2-Cd40tm1Kik/J) and μ MT (B6.129S2-Ighmtm1Cgn/J) mice were purchased from The Jackson Laboratory. CD40^{-//-} mice were provided by Dr. David Wagner (University of Colorado Denver, Anschutz Medical Campus, Aurora, CO) as described previously (2). Mixed BM chimera mice were generated by reconstitution of 6×10^6 total T cell-depleted BM cells from μ MT and CD40 WT/mutant

FIGURE 1. Generation of endogenous CD40 cytoplasmic domain mutant mice. (**A**) Schematic representation of WT CD40 and mutants. CD40 Δ T6 was generated by deletion of the sequence ²³⁴ RQDPQE²³⁹. CD40 Δ T2 was made by deletion of the sequence ²⁵¹ PVQET²⁵⁵. CD40 Δ T2* was made by truncation of a sequence distal to amino acid 245 by inserting two stop codons (TAATAG). (**B**) Representative FACS analysis of CD40 surface expression on the same mouse spleen DCs (PI⁻CD11c⁺B220⁻) and B cells (PI⁻ CD11c⁻B220⁺) that were in 4°C overnight (gray) or incubated with CpG overnight (red). (**C**). CD40 was immunoprecipitated from cell lysates of the indicated mice, and we analyzed bound proteins by immunoblotting using TRAF2 Ab. The lower two panels show expression of TRAF2 and CD40 in whole-cell lysates. The arrow indicates the position of TRAF2, and *ns indicates the position of a nonspecific band present even in CD40^{-/-} mice. donor mice at a 1:1 ratio into irradiated (950 rad) WT host mice by i.v. injection and were used after 6 wk. Mice were maintained in accordance with National Institutes of Health guidelines. All animal experiments were approved by the National Cancer Institute Animal Care and Use Committee.

Immunoprecipitation and Western blotting

A combined stimulation/immunoprecipitation protocol was used to isolate CD40 as previously described (23). Briefly, protein G magnetic beads (Dynal) were coated with goat anti-mouse IgG (5 μ g/10 μ l beads) (Jackson ImmunoResearch) followed by anti-mouse CD40 mAb 3/23 (10 μ g/10 μ l beads). After stimulation of splenocytes with CpG ODN1826 (0.2 μ g/ml) overnight, the cells were washed and incubated with anti-CD40–coated magnetic beads for 20 min at 37°C. Following incubation, beads and cells were pelleted by centrifugation (2 min at 300 × g), and the medium was discarded. Cells were lysed and beads were washed as described (24). Bead-bound proteins were eluted in SDS-PAGE sample loading buffer by boiling and analyzed by SDS-PAGE and immunoblotting using anti-TRAF2 (MBL M112-3) or anti-CD40 mAb (Abcam ab65853).

Immunizations and adoptive transfer studies

Mice were immunized i.p. with 100 μ g 4-hydroxy-3-nitrophenylacetyl (NP)keyhole limpet hemocyanin (KLH) or NP-OVA (Biosearch Technologies) mixed 1:1 with Imject Alum (Thermo) in a total volume of 200 μ l. For adoptive transfer experiments, CD45.1 OT-II Tg CD4⁺ T cells were purified using a MACS CD4⁺ T cell isolation kit (Miltenyi Biotec), and 5 × 10⁵ cells were transferred by i.v. injection into host mice 1 d before immunization as described previously (4).

EAE induction and scoring

EAE induction was performed by mixing 150 µg rhMOG (Protein Expression Laboratory, Leidos Biomedical Research, Frederick National Laboratory for Cancer Research, Frederick, MD) or 200 µg of MOG_{35-55} peptide (MEVG-WYRSPFSRVVHLYRNGK) (GenScript) in CFA containing *Mycobacterium tuberculosis* H37Ra (Difco Laboratories). Pertussis toxin (120 ng) (List Biological Laboratories) diluted in PBS was administered by i.p. injection on days 0 and 2 after immunization. The clinical severity of EAE was scored by an observer who was unaware of mouse genotypes using a grading scale of 0–5 as previously described (2, 7).



Isolation of mononuclear cells from peripheral lymph nodes and CNS parenchyma

Inguinal and axillary draining lymph nodes (DLNs) were mixed and digested with 2.5 mg/ml collagenase D (Roche) and 1 mg/ml DNase I (Roche) for 30 min at 37° C. The samples were then filtered through 70-µm mesh strainers to generate single-cell suspensions. For isolation of CNS-infiltrating cells, brain and spinal cord tissues were minced and digested with collagenase D and DNase I as above. The samples were then filtered through 70-µm mesh strainers and centrifuged through a discontinuous Percoll (Sigma) density gradient (38% and 70%). Mononuclear cells at the interphase were removed, washed, and resuspended in culture medium for analysis by flow cytometry.

Flow cytometry

Cells were washed with FACS buffer (HBSS containing 0.2% BSA and 0.05% sodium azide), treated with anti-FcR (24G2), and then stained using specific Abs. Anti-mouse CD4, CD8, B220, CD19, CD11c, MHC class II, and CD40 Abs were purchased from BioLegend. For intracellular cytokine staining, mononuclear cells from DLNs and CNS were stimulated with PMA/ ionomycin for 2 h, at which time GolgiStop (BD Biosciences) was added for an additional 2 h. The cells were fixed and permeabilized with the BD Fix/Perm kit (BD Biosciences) according to the manufacturer's instructions and stained with anti–IFN- γ , anti–IL-17A, and isotype control Ab (BD Biosciences) for 30 min. Data were collected with a FACS LSR II or FACS Fortessa flow cytometer (BD Biosciences) and analyzed using FlowJo software.

Ab ELISA

NP-specific and MOG-specific IgG was measured by ELISA. In brief, NP30-BSA (Biosearch Technologies) or rhMOG (10 µg/ml) was coated on ELISA plates (Immulon 4HBX; Thermo Fisher) at 4°C overnight. The high-affinity IgG Ab titer was determined by coating plates with NP2-BSA overnight. The plates were washed with ELISA wash buffer (0.5% Tween in PBS), serially diluted sera were applied to the plates, and plates were incubated for 2 h at room temperature. After washing, anti-mouse IgG HRP (SouthernBiotech) was used to detect bound serum IgG. After a wash step, 2,2'-azino-di-(3-ethylbenzthiazoline-6-sulfonate) substrate (KPL) was added to the wells, and the enzyme reaction was stopped after 20 min using 2,2'-azino-di-(3-ethylbenzthiazoline-6-sulfonate) HRP Stop Solution (KPL). The OD at 405 nm was measured using a SpectraMax iD3 plate reader.

Statistical analyses

The Student *t* test with two-tailed distributions was performed for statistical analyses with single comparisons. For multiple comparisons, statistical analysis was performed with one-way ANOVA followed by Dunnett multiple comparison test. A *p* value <0.05 was considered statistically significant.

Results

Generation of CD40 cytoplasmic domain mutant knock-in mice

In vitro studies have suggested that there are multiple functional motifs in the CD40 cytoplasmic domain that are involved in CD40 signaling. Among these are two cytoplasmic motifs that are thought to be important for the recruitment of the adapter molecules TRAF6 or TRAF2/3 to CD40 (25, 26). We generated mice expressing cytoplasmic domain mutations at the endogenous CD40 locus by CRISPR/Cas9 as illustrated in Fig. 1A. A proximal domain CD40 cytoplasmic tail mutant (termed CD40 Δ T6) was generated by deletion of the sequence ²³⁴RQDPQE²³⁹, which is homologous to the TRAF6 binding motif in human CD40 (27). A distal domain CD40 cytoplasmic tail mutant (termed CD40 Δ T2) was made by deletion of the sequence ²⁵¹PVQET²⁵⁵, which corresponds to the TRAF2 binding motif ²⁵¹PxQxT²⁵⁵sequence in human CD40 (27–29). We also established a third strain by truncation of a sequence distal to amino acid 245 by inserting two stop codons (TAATAG), which preserves the proximal TRAF6 binding domain and deletes the distal TRAF2/3 binding domain as well as CD40 cytoplasmic amino acids 246–278 (termed CD40 Δ T2*). This region of the CD40 cytoplasmic domain is identical to that in human CD40 and also contains a putative second TRAF2 binding motif (29). All CD40 cytoplasmic domain mutants express levels of CD40 on the surface of B cells and DCs that are comparable to CD40 expressed in C57BL/6 (WT) mice both at steady state and after overnight stimulation with CpG (Fig. 1B). Coimmunoprecipitation studies using CD40-stimulated mouse splenocytes confirmed that TRAF2 bound to WT CD40 and CD40 Δ T6 but was unable to bind to CD40 expressed in CD40 Δ T2 and CD40 Δ T2* mice (Fig 1C). Nearly identical results were obtained when examining the binding of epitope-tagged TRAF2 coexpressed with WT or mutant CD40 molecules in transfected HeLa cells (Supplemental Fig. 1). We were unable to reproducibly coimmuno-precipitate TRAF6 even in WT CD40 expressing mice, a finding that is consistent with other reports that failed to show direct TRAF6/CD40 interactions in vitro (27).

Both CD40 TRAF6 and TRAF2/3 binding motifs on non-B cells are required for severe EAE

CD40 expression is required for mice to develop EAE (2, 30, 31). We therefore analyzed the requirement for the distinct TRAF6 and



FIGURE 2. Both CD40 TRAF6 and TRAF2/3 binding motifs on non-B cells are required for EAE. WT or CD40 mutant mice were immunized with rhMOG, and the severity of EAE was examined over time. (**A**) Clinical score of the indicated mice immunized with rhMOG. (**B**) Radiation BM chimeras were established by mixing BM from µMT mice with BM from CD40 mutant mice. (**C**) EAE was induced in the indicated chimeric mice with rhMOG. EAE scores are combined from three independent experiments (mean ± SEM using 10–18 mice per group). Median of the clinical score during days 11–27 for WT mice was compared with each CD40 mutant using a two-tailed nonparametric Mann–Whitney *U* test. **p < 0.01, ****p < 0.0001. Comparing clinical scores of CD40^{-/-} mice with CD40ΔT6 (p < 0.001), CD40ΔT2 (p < 0.01), or CD40ΔT2* (p < 0.01) mice also revealed statistically significant differences.

TRAF2/3 binding motifs in EAE. Immunization of mice with rhMOG results in a form of EAE that requires both T and B cells, mirroring the known roles of these cell types in human MS (5, 10, 11, 32). All CD40 TRAF binding motif mutant mice showed a significant reduction of EAE severity relative to WT mice (Fig. 2A), indicating that the CD40 TRAF6 and TRAF2/3 binding motifs play important and not completely redundant roles in EAE. EAE severity in CD40^{-/-} mice was significantly lower than in any single TRAF binding motif mutant mouse, suggesting that the TRAF2/3 and TRAF6 binding motifs have partially overlapping functions and/ or that additional CD40 domains function in EAE.

We have previously shown that CD40 expression on both DCs and B cells is required for the development of EAE (2). To address whether the CD40 TRAF binding motifs on B cells or DCs (or both) play a role in EAE, we generated mixed BM chimeras in which expression of mutant CD40 was restricted to B cells, whereas all other APCs in the mouse express WT CD40 (Fig. 2B). Chimeric mice with B cells derived from CD40^{-/-} donors were completely protected from EAE, demonstrating a requirement for B cell CD40 in mediating EAE. By contrast, chimeras with B cells derived from CD40 TRAF binding motif mutants all developed severe EAE similar to the WT group (Fig. 2C). Because EAE was less severe in mice expressing CD40 TRAF binding motif mutations (in which all APC subtypes, including B cells, possess mutant CD40 proteins), these data show that the reduced EAE observed in CD40 TRAF binding motif mutant mice was a consequence of impaired CD40 function on APCs other than B cells.

EAE can also be induced in mice by immunizing mice with MOG_{35-55} (7). In contrast to rhMOG-induced EAE, MOG_{35-55} induced EAE is a B cell–independent disease (10, 11). As was observed in rhMOG-based disease, each of the CD40 TRAF binding motif mutant mice showed a significant reduction of EAE severity relative to WT mice when immunized with MOG_{35-55} (Supplemental Fig. 2). This result further supports our finding using chimeric mice and demonstrates that TRAF2/3 binding and TRAF6 binding motifs of CD40 APCs other than B cells are important in EAE.

The CD40 TRAF6 and TRAF2/3 binding motifs function in EAE through optimal priming of Th17 cells

An early event following rhMOG immunization is the priming and differentiation of inflammatory cytokine-producing CD4⁺ T cells in lymph nodes, of which IL-17A-producing Th17 cells and IL-17A and IFN- γ double-producing cells are the most pathogenic (8). Using conditional CD40^{-/-} mice, we confirmed our previous observation (2) that CD40 expression on DCs, but not on B cells, is required for priming of pathogenic Th17 cells (Fig. 3A). To determine if the TRAF binding motifs on DCs are involved in CD4 priming in DLNs, we measured the generation of cytokine-producing CD4⁺ T cells in DLNs of WT and TRAF binding motif mutant mice following rhMOG immunization (Fig. 3B). All CD40 TRAF binding motif mutant mice showed lower numbers of CD4⁺ T cells, equivalent to the defective response of $CD40^{-/-}$ mice (Fig. 3C). The frequency of IL-17A-producing Th17 cells in DLNs was markedly reduced in all CD40 TRAF binding motif mutant mice (Fig. 3D), demonstrating that the TRAF6 and TRAF2/3 binding motifs on

FIGURE 3. Both CD40 TRAF6 and TRAF2/3 binding motifs function in EAE by regulating Th17 cell priming. WT or CD40 mutant mice were immunized with rhMOG, and, 6 d after EAE induction, CD4⁺ T cells from the DLNs of the indicated mice were analyzed by FACS analvsis. (A) DLN cells from rhMOG-immunized CD40^{fl/fl} mice, CD40^{-/-} mice, or CD40^{fl/fl} mice crossed with CD19-Cre or CD11c-Cre (to delete CD40 only on B cells or DCs, respectively) were analyzed for cytokine production by FACS analysis (in the CD4⁺ B220⁻ gate), and the percentage of CD4 T cells expressing IL-17A was determined. (B) DLN CD4⁺ T cells from rhMOG-immunized WT or CD40 TRAF binding motif mutant mice were analyzed for cytokine production by FACS analysis (in the CD4⁺ B220⁻ gate). The number of CD4 T cells in the DLNs (C) and the percentage of all DLN CD4 T cells expressing either IL-17A or IFN- γ was determined (**D**). Data are combined from two independent experiments (mean ± SD using six mice per group). One-way ANOVA followed by Dunnett test against the immunized WT group was performed for multiple comparisons. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.



DCs play a nonredundant role in the induction and differentiation of pathogenic Th17 cells in the periphery of rhMOG-immunized mice.

The CD40 TRAF6 and TRAF2/3 binding motifs function in EAE by promoting the accumulation of pathogenic Th17 cells in CNS

We also monitored the presence of CD4⁺ T cells in the CNS (brain and spinal cord) of rhMOG-immunized WT and CD40 TRAF binding motif mutant mice. Immunized CD40^{-/-} mice, which did not develop disease, had very few CNS mononuclear cells and even fewer CD4⁺ T cells recoverable from brain and spinal cord (Fig. 4A). All CD40 TRAF binding motif mutant mice, which were significantly protected from EAE, had comparably reduced numbers of CD4⁺ T cells in the CNS when compared with WT mice. Substantial proportions of CD4⁺ T cells in the CNS of immunized CD40 WT mice were IL-17A⁺ and IL-17A⁺/IFN- γ^+ (Fig. 4B), whereas the small numbers of CD4⁺ T cells isolated from the CNS of CD40^{-/-} mice had significantly lower frequencies of cytokine-positive populations. All CD40 TRAF binding motif mutant mice had reduced frequencies of these pathogenic Th cells in the CNS, a finding that is consistent with reduced EAE severity and reduced frequencies of Th17 cells in the DLNs in these mice. As was observed for clinical EAE severity, the reduction in Th17 cells in CD40^{-/-} mice was even more profound than in any single TRAF binding motif mutant, suggesting that the functions of these two motifs are not completely redundant and/or that other domains of the CD40 molecule play a role in Th17 generation. Taken together with our conditional CD40 deletion data, these findings demonstrate the importance of the CD40 TRAF6 and TRAF2/3

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binding motifs on DCs in the differentiation of pathogenic Th17 cells in the periphery of rhMOG-immunized mice and in the pathogenesis of EAE.

The TRAF2/3 binding motif of CD40 expressed by B cells is required in T-dependent GC responses

We have previously reported that CD40 expression on B cells, but not on DCs, is required for TD GC responses and high-affinity Ab production (3, 4). Having established that both TRAF2/3 and TRAF6 binding motifs of CD40 are required for DC-dependent T cell priming in EAE, we assessed the role of these two TRAF binding sites on CD40 for primary TD GC responses. CD40 WT and mutant mice were immunized with NP-KLH in alum, and spleen GC B cells were analyzed 8 d after immunization by flow cytometry. In comparison with the response of mice containing WT CD40, GC B cell numbers were significantly decreased in CD40 Δ T2 and CD40 Δ T2* mutant mice to levels approximating that in complete CD40^{-/-} mice. By contrast, the response of CD40 Δ T6 mice was comparable to that of WT mice (Fig. 5A). The total B cell number in CD40^{-/-} mice was comparable to WT mice before immunization (16, 17), and the total spleen B cell numbers were also comparable among all strains after immunization (Fig. 5B, left). Like GC B cell responses, Ag-specific Tfh differentiation was significantly reduced in spleens from CD40 Δ T2 and CD40 Δ T2* mice and was unaffected in CD40 Δ T6 mice (Fig. 5C). These results indicate that whereas CD40 does not play a role in the maintenance of total spleen B cell numbers, the CD40 TRAF2 binding motif, but not the TRAF6 binding

FIGURE 4. The CD40 TRAF6 and TRAF2/3 binding motifs function in EAE by mediating the accumulation of pathogenic Th17 cells in the CNS. (A) On day 14 after EAE induction, total mononuclear cells were recovered from the CNS, and the total numbers of mononuclear cells as well as CD4⁺ T cells present in the CNS were determined by FACS analysis. (B) CNS CD45.2⁺CD11b⁻ CD4⁺ T cells were analyzed for cytokine production by FACS analysis, and the percentage of all CNS CD4 T cells expressing either IL-17A alone or both IL-17A and IFN-y was determined. Data are combined from two independent experiments (mean ± SD using six mice per group). Oneway ANOVA followed by Dunnett test comparing the immunized WT group with each CD40 mutant was performed for multiple comparisons. *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001. Comparison of the immunized CD40^{-/-} group with each CD40 mutant revealed statistically significant differences: for IL-17A⁺ CD4 T cells, p < 0.01 compared with Δ T2 and Δ T6, p < 0.05compared with $\Delta T2^*$; for IL17A⁺/IFN- γ^+ CD4 T cells, p < 0.05 compared with $\Delta T2$ and $\Delta T6$, p < 0.01 compared with $\Delta T2^*$.



TRAF6 binding motif is required for TD GC responses. WT and CD40 mutant mice were immunized with NP-KLH in alum. After 8 d, GC B cells and total B cells were analyzed in the spleen. Representative FACS analysis (A) of CD38⁻ GL7⁺ GC B cells in the CD4⁻ B220⁺ B cell gate and statistical analyses of FACS data examining the total number of spleen B cells and GC B cells in each mouse strain (B). (C) OT-II T cells (CD45.1) were transferred to the indicated recipient mice (CD45.2) and were immunized with NP-OVA/alum 1 d after cell injection. Transferred OT-II cells were analyzed at day 8 by FACS analysis. A representative flow cytometry profile of spleen OT-II cells (CD45.1⁺ CD4⁺B220⁻) analyzed for the Tfh CXCR5^{high} PD-1^{high} phenotype and statistical analyses of FACS data showing the percentage of transferred OT-II cells expressing a Tfh phenotype are shown. (D) WT and CD40 mutant mice were immunized with NP-KLH in alum. Serum was collected 3 wk after immunization, and total NP-specific Ab (anti-NP30 IgG) and high-affinity NP Ab (anti-NP2 IgG) was determined by ELISA. Data are combined from two independent experiments (mean \pm SD using six mice per group). One-way ANOVA followed by Dunnett test against the immunized WT group was performed for multiple comparisons. **p < 0.01, ***p < 0.001, ***p < 0.0001.

FIGURE 5. The CD40 TRAF2 binding motif but not the



motif, is required for Ag-specific Tfh differentiation and GC B cell generation. Despite the significantly reduced numbers of GC B cells and Tfh cells, the Ag-specific class-switched IgG response measured as binding to NP30-BSA was unaffected by mutations in any of the CD40 cytoplasmic TRAF binding motif mutants (Fig. 5D). A similar result was observed by measuring the antirhMOG IgG levels in mice induced with EAE (Supplemental Fig. 3). Notably, however, high-affinity anti-NP IgG, measured as binding to NP2-BSA, was substantially reduced in CD40 Δ T2 and CD40 Δ T2* strains relative to that observed in WT mice, although greater than that of CD40^{-/-} mice. High-affinity anti-NP2 IgG from CD40 Δ T6 mice was comparable to that found in immunized WT mice, a finding that is consistent with the observation that GC B cell numbers in immunized WT and CD40 Δ T6 mice were identical.

Although we have previously shown that CD40 on DCs is not required for TD Ab responses, we specifically examined the importance of the TRAF binding motifs on B cells during this process. We constructed radiation BM chimeras in which BM from μ MT mice was mixed with BM from CD40 cytoplasmic domain mutant mice and transferred into lethally irradiated WT recipients. The B cells in these chimeric mice possess either WT or mutant CD40, whereas all other cell types (including DCs) have intact CD40 (Fig. 6A). The total numbers of spleen B cells in each chimera were comparable (Fig. 6B); however, chimeras with CD40 Δ T2* B cells, but not CD40 Δ T6 B cells, showed significant reduction of GC B cells following immunization with NP-KLH in alum. Taken together with our studies in CD40 TRAF binding motif mutant mice, these data demonstrate that the CD40 TRAF2 binding motif, but not the TRAF6 binding motif, in B cells is required for GC development and high affinity Ab production.

Discussion

CD40 is a cell surface receptor that belongs to the TNF receptor family and serves as a costimulatory molecule expressed on multiple APC lineages. The cytoplasmic domain of CD40 has been implicated as essential in CD40-CD154 costimulatory signal transduction in a variety of in vitro systems; however, understanding of the roles played by this domain in vivo is incomplete (21, 25, 33). The data presented in the present study reveal cell type– and cytoplasmic domain–specific requirements for CD40 function in vivo. We show that both TRAF6 and TRAF2/3 binding motifs of CD40 on DCs play an essential role in rhMOG-induced EAE by regulating pathogenic Th cell priming. In contrast, the cytoplasmic membrane distal TRAF2/3 binding motif, but not the membrane proximal TRAF6 binding motif, is required for CD40 expressed on B cells for TD humoral immune responses, including optimal GC B cell production, Tfh generation, and expression of high-affinity Abs.

Previous work in our laboratory and others highlighted the critical role of CD40 in mediating autoimmune and TD GC responses. The interaction between CD40 and TRAF molecules has been intensively investigated in cell lines, which suggested that TRAFs are involved in CD40 function (22, 25, 26, 34-36). In the present study, we have generated mutations in the proposed CD40 cytoplasmic TRAF2 binding motif and have found that endogenous TRAF2 does not bind to CD40 in spleen cells in which this motif has been mutated (CD40AT2 and CD40AT2*) or when TRAF2 is coexpressed with CD40 Δ T2 or CD40 Δ T2* in HeLa cells, consistent with previous reports (27, 37). Although we were able to reliably coimmunoprecipitate CD40 with TRAF2, we were unable to detect the binding of CD40 with TRAF6 in mouse spleen cells or in transfected HeLa cells. It is important to note that this does not mean that TRAF6 does not interact with CD40 in vivo; it simply shows that under the immunoprecipitation conditions used in our study, the interaction is not preserved. This result is, however, consistent with a previous report that also failed to detect any binding between CD40 and TRAF6 in vitro (27). It is also possible that alterations in the CD40 cytoplasmic domains used in our and other studies act through CD40 interaction with as yet unidentified molecules. Despite deletion of the TRAF6 and TRAF2/3 binding motifs, both surface expression of CD40 and TLR-mediated upregulation of CD40 expression proceed normally in these mice, demonstrating that some components of the machinery driving CD40 expression and upregulation are independent of the CD40 cytoplasmic TRAF binding motifs.

Our laboratory recently demonstrated that CD40 on DCs, but not on B cells, is essential for peripheral Th1 and Th17 priming and CNS accumulation in rhMOG-induced EAE (2). In the present study, we used the same rhMOG-induced EAE model to further examine the cellular mechanism mediating the function of CD40 in this autoimmune disease. Mice expressing either TRAF6 or TRAF2/3 binding motif mutations developed less severe EAE than did mice expressing WT CD40. The use of mixed BM chimeras revealed that this was not a consequence of CD40 TRAF binding motif deletion on B cells, because EAE severity was not affected when expression of either TRAF6 or TRAF2/3 binding motif mutant was restricted to B cells. These data point to a role of these domains of CD40 in a cell other than a B cell. Because expression of CD40 on DCs, but not on B cells, impairs peripheral Th17 priming, the inability of both CD40 TRAF binding motif mutant mice to generate pathogenic Th17 cells in the periphery of rhMOG-immunized mice demonstrates that these two motifs exert distinct and nonredundant roles of DC CD40 in priming of CD4 T cells. Analysis of the T cell subsets in the CNS of EAE mice revealed that all CD40 TRAF binding motif mutant mice used in this study had impaired pathogenic Th17 CNS infiltration, a result that could be a consequence of simply reduced numbers of peripheral pathogenic Th17 cells primed by DCs or by another APC in the brain that also requires both TRAF binding motifs for full activity. It is interesting to note that a recent study reported that myeloid cells plays a role in EAE (38), leading to the possibility that the non-APCs expressing the CD40 TRAF binding motif mutations used in this study could also be due to CD40 expression on one of these cell types. The observation that clinical EAE severity as well as Th17 priming in CD40^{-/-} mice was even lower than in mice with single TRAF binding motif mutations suggests that TRAF2/3 and TRAF6 binding motifs have partially overlapping functions or that additional CD40 domains function in EAE.

Previous studies using conditional deletion of TRAFs have addressed the role of specific TRAF proteins in humoral immune responses. Conditional deletion of B cell TRAF2, but not TRAF3, suppressed TD Ab responses but not T cell-independent Ab responses (39, 40). By contrast, conditional deletion of TRAF6 in B cells did not affect GC formation or Ab affinity maturation in mice immunized with TD Ags (41), unlike the profound defect in these responses in both CD40-deficient mice or in mice lacking CD40 on B cells (3, 4). Curiously, deletion of TRAF6 on B cells suppressed Ig class switching following immunization with either TD and T cell-independent Ags (41). Because T cell-independent Ab responses do not require CD40-CD40L signaling (16), these data reveal an additional function of TRAF6 in humoral immune responses in B cells that is independent of CD40. CD40 Δ T2 or CD40 Δ T2* mutant mice had significantly reduced GC B cells and Ag-specific Tfh after TD protein Ag immunization, consistent with a role of CD40-TRAF2 interactions during the TD GC response. By generating mixed BM chimeras, we also demonstrated a B cellintrinsic effect of this process, confirming a B cell, but not DC, requirement for CD40 in TD immune responses as we have reported previously (3, 4). Interestingly, the significantly reduced GC B and Tfh cells did not result in a decrease in total Ag-specific anti-NP30 or anti-rhMOG IgG levels in the serum 3 wk after immunization, although high-affinity anti-NP Ab measured by binding to NP2-BSA was substantially reduced in CD40AT2 or CD40AT2* mutant mice. In a previous report, transgenic mice expressing TRAF2 binding motif mutant of human CD40 also showed normal serum IgG levels in mice lacking GCs after a TD Ag immunization response (42). GC-independent serum IgG was also reported in lymphotoxin- α -deficient mice, which fail to form GCs in the spleen but mount an anti-NP IgG1 response similar to WT mice after immunization with the TD Ag NP-OVA (43). Consistent with previous reports, our data suggest that normal extrafollicular plasma B cell differentiation may occur in the absence of GC response (42).

Two different reports have described CD40^{-/-} mice expressing CD40 transgenes with cytoplasmic TRAF6 and TRAF2/3 binding motif mutations with expression driven by either the I-E α promoter (28) or the IgH promoter (27). In the I-E α -driven CD40 mutants, both TRAF6 and TRAF2/3 binding motif mutations mount a normal GC response after immunization with sheep erythrocytes; however, after immunization with NP-KLH/CFA, Ag-specific IgG1 was significantly reduced in the serum from TRAF6, but not TRAF2/3, binding motif mutant mice (28). By contrast, in IgH-driven CD40 mutants, TRAF6 binding motif mutant mice mount a normal GC response and anti-KLH IgG in serum, whereas TRAF2/3 binding motif mutant mice cannot (27). Spleen B cells from our CD40 cytoplasmic TRAF6 binding motif mutant mice retain the ability to bind TRAF2 and are able to effectively mount a CD40-dependent GC response, demonstrating that this motif is not involved in TRAF2 binding and is not important for TD immune responses. Thus, our data, generated from CRISPR/Cas9 mutagenesis of endogenous CD40, and showing that the CD40 binding motif for TRAF2 is essential for CD40-mediated GC formation and high-affinity anti-NP2 IgG but not total anti-NP30 IgG production, are most consistent with the findings of Jabara et al. (27).

In conclusion, our data reveal important information regarding the functional roles of two distinct CD40 cytoplasmic domains expressed on distinct cell types in vivo. DCs require both TRAF2/3 and TRAF6 binding motifs for optimal ability to prime peripheral Th17 cells following immunization with rhMOG and in the development of EAE. In contrast, B cells require the CD40 TRAF2 binding motif, but not the TRAF6 binding motif, in the generation of TD Ab responses. These findings may have translational relevance

FIGURE 6. The CD40 TRAF2 binding motif on B cells is required for TD GC responses. (A) Radiation BM chimeras were established by mixing BM from µMT mice with BM from CD40 WT or mutant mice and transferring these BM cells into lethally irradiated B6 recipients. After 6 wk, chimeric mice express WT or mutant CD40 only on B cells, whereas all other APC cell types possess WT CD40. (B) The indicated chimeric mice were immunized with NP-KLH in alum, and, after 8 d, GC B cells and total B cells were analyzed in the spleen. Representative FACS analysis of CD38⁻GL7⁺ GC B cells in CD4⁻B220⁺ B cell gate. Statistical analyses of FACS data examining the total number of GC B cells and total spleen B cells, respectively. Data are combined from two independent experiments (mean ± SD using six mice per group). One-way ANOVA followed by Dunnett test against immunized C57BL/6 group was performed for multiple comparisons. ****p < 0.0001.

A

В

uMT 3e6

CD40 mutant

Spleen CD4-B220+

3e6

for strategies specifically targeting pathogenic versus adaptive aspects of the CD40-dependent immune response.

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Disclosures

The authors have no financial conflicts of interest.

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μMT+CD40ΔT6 μMT+CD40ΔT2 μMT+CD40ΔT2* µMT+WT 126 0.14 0.12



6 weeks

Chimera

950 rad

B6

recipient

µMT+CD40-/-

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