## **Requirement for AHNAK1-mediated calcium signaling during T lymphocyte cytolysis**

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Contributed by Richard A. Flavell, April 23, 2009 (sent for review January 20, 2009)

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**Cytolytic CD8 T cells (CTLs) kill virally infected cells, tumor cells, or other potentially autoreactive T cells in a calcium-dependent manner. To date, the molecular mechanism that leads to calcium intake during CTL differentiation and function has remained unresolved. We demonstrate that desmoyokin (AHNAK1) is expressed in mature CTLs, but not in naive CD8 T cells, and is critical for calcium entry required for their proper function during immune response. We show that mature AHNAK1-deficient CTLs exhibit reduced** Ca<sub>v</sub>1.1 α1 subunit expression (also referred to as L-type calcium channels or  $\alpha$ 1S pore-forming subunits), which recently were suggested to play a role in calcium entry into CD4<sup>+</sup> T cells. **AHNAK1-deficient CTLs show marked reduction in granzyme-B** production, cytolytic activity, and IFN- $\gamma$  secretion after T cell **receptor stimulation. Our results demonstrate an AHNAK1-dependent mechanism controlling calcium entry during CTL effector function.**

Calcium plays critical roles in T-cell differentiation and func-<br>tion, such as activation, proliferation, and cytokine production  $(1, 2)$ . Cytotoxic CD8<sup>+</sup> effector T cells (CTLs) primarily use the perforin/granule exocytosis pathway to kill virus-infected and tumor cells (3). The role of calcium in CTL-mediated cytolysis has been studied extensively, mainly by using calcium antagonists (e.g., see ref. 4). The exact requirements for calcium and its molecular mode of entry during CTL function still are largely uncertain (4, 5).

The calcium release-activated calcium channel (CRAC) pathway is the most studied plasma membrane store operated calcium (SOC) channel through which calcium enters after T-cell stimulation. Surprisingly, ORAI1 (also known as ''CRACM1'' or ''TMEM142A'')-deficient CTLs indeed show reduced calcium entry but only partial IFN- $\gamma$  production and normal granzyme-B expression, suggesting that other pore subunits, possibly ORAI2 or ORAI3, or different channels altogether, are involved in this process (6).

In addition to CRAC channels (7), we and others (8, 9), found that L-type calcium channel (Ca<sub>v</sub>1) subunits  $\alpha$ 1,  $\alpha$ 2,  $\beta$ ,  $\gamma$ , and  $\delta$ , which constitute the major route of calcium entry in excitable cells and take up calcium in response to membrane depolarization, also are expressed by T cells (10). Previously, we have shown that CD4<sup>+</sup> T cells express  $\alpha$ 1 subunits of the Ca<sub>v</sub>1 family and that functional Ca<sub>v</sub>  $\beta$ 4 and  $\beta$ 3 regulatory subunits are necessary for normal TCR-triggered calcium response, nuclear factor of activated T cells (NFAT) nuclear translocation, and cytokine production (8, 11).

The AHNAK family of scaffold PDZ proteins consists of 2 giant proteins (700 kDa), AHNAK1 (desmoyokin) and AHNAK2 (12–14). AHNAK1 is involved with calcium signaling through protein–protein interactions (15–18). Furthermore, in cardiomyocytes, AHNAK1 associates with the  $\beta$ -subunit of cardiac  $Ca<sub>v</sub>$  channels at the plasma membrane and is phosphorylated by protein kinase A (PKA) in response to  $\beta$ adrenoreceptor stimulation (19).

Using AHNAK1-deficient mice (14), we recently described a novel mechanism for the regulation of calcium signaling through

Ca<sub>v</sub>1.1  $\alpha$ 1 subunits mediated by AHNAK1 in peripheral CD4<sup>+</sup> T cells. AHNAK1 is associated with the regulatory  $\beta$ 2 subunit of Cav1 channels and is required for normal expression of the  $Ca<sub>v</sub>1.1$   $\alpha$ 1 subunit and intact calcium influx following TCR cross-linking (11). Here, we demonstrate that CTLs employ AHNAK1 to mediate calcium entry required for cytolytic activity late in primary TCR stimulation through the regulation of Cav1.1 channels.

## **Results**

**AHNAK1 Is Expressed in Mature CTLs.** Previously, we have shown that AHNAK1 is required for differentiation of naive  $CD4<sup>+</sup>$  T cells (11). In  $CD8<sup>+</sup>$  T cells, we found that AHNAK1 was not expressed in naive  $CDS^+$  T cells [\(Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1)A) but was highly expressed in mature CTLs 5 days after primary stimulation in vitro [\(Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1)*A*). Further examination of AHNAK1 expression showed that it was highly expressed late after primary differentiation in vitro, starting on day 4, throughout day 5, and 24 h after restimulation, as shown by densitometry of immunoblot analysis [\(Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1)*B*).

*Ahnak1*-/- mice showed normal thymocyte development and normal lymphoid composition in their thymus and spleen (11) as well as normal  $CD8<sup>+</sup>$  T cell numbers in the spleen [\(Fig. S2\)](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF2). In addition, using CD44, CD62L, and CD69 as markers, we found that  $CD8<sup>+</sup>$  T cells exhibited normal numbers of naive and memory populations as well as normal activation in culture. [\(Fig.](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3) [S3](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3)*A* shows an analysis of unstimulated cells and purified CD8 T cells 48 h after primary stimulation.)

Naive  $Ahnak1^{-7-}$  CD8<sup>+</sup> T cells proliferated normally, as determined by the 5 (and 6)-carboxyfluorescein diacetate succinimidyl ester (CFSE) assay 72 h after primary TCR stimulation [\(Fig. S3](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3)*B*), and secreted normal amounts of IL-2 and IFN- $\gamma$ , as measured 24 h (not shown) and 48 h after activation [\(Fig. S3](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3) *C* [and](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3) *D*). Collectively, our observations indicate that AHNAK1 is neither expressed nor required for early stages of in vitro CD8 T-cell differentiation.

Ahnak1<sup>-/-</sup> CTLs Display Reduced Ca<sub>v</sub>1.1  $\alpha$ 1 Subunit Expression and **Calcium Influx.** AHNAK1 has been shown previously to be required for calcium influx after TCR activation of naive CD4 T cells  $(8, 11)$ . AHNAK1 is highly expressed in naive CD4<sup>+</sup> T cells and seems to be required for the targeting and/or stabilization of  $Ca<sub>v</sub>1.1 \alpha1$  subunits at the plasma membrane. Reduced

Author contributions: D.M., V.T.M., and R.A.F. designed research; D.M., A.B., M.K.J., T.W., A.A., and S.S. performed research; T.W. and K.K. contributed new reagents/analytic tools; and D.M. and R.A.F. wrote the paper.

The authors declare no conflict of interest.

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Fig. 1. Reduced Ca<sub>v</sub>1.1 a1 subunit expression and calcium entry in response to TCR stimulation in *Ahnak1<sup>-1-</sup>* CTLs. (A) The expression of Ca<sub>v</sub>1.1 and Ca<sub>v</sub>1.2 was examined in purified naive CD8<sup>+</sup> T cells and in mature CTLs 5 days after primary stimulation using plate-bound anti-CD3 and anti-CD28 (amount of antibodies is described in [Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1) A) in wild-type and A*hnak1<sup>-I –</sup>* cells. (B) Kinetics of Ca<sub>v</sub>1.1 and Ca<sub>v</sub>1.2 expression by immunoblot analysis [using anti-a1S (Ca<sub>v</sub>1.1) and anti-a1C (Ca<sub>v</sub>1.2)] was as described in [Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1)*B*. Expression of Ca<sub>v</sub>1.1 and Ca<sub>v</sub>1.2  $\alpha$ 1 subunit was normalized to *β*-actin for each time point. This final densitometry plot of  $Ca<sub>v</sub>1.1$  and  $Ca<sub>v</sub>1.2$   $\alpha$ 1 subunit expression is an average of 3 independent experiments. There is a statistically significant difference in  $Ca<sub>v</sub>1.1$  expression, but not in Ca<sub>v</sub>1.2, «1 subunit expression, between wild-type and Ahnak1<sup>-/-</sup> cells (\*P < 0.05) at day 5 after primary stimulation. Experiment *B* was performed in addition to and was independent of experiment *A*. (C) CTLs from wild-type and *Ahnak1<sup>-/-</sup>* mice, obtained after 5 days of primary stimulation as in [Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1) *A*, were incubated with anti-CD3 (10  $\mu$ g/mL) for 30 min on ice and subsequently were cross-linked by goat anti-hamster at the indicated time (TCR stimulation). Then [Ca<sup>2+</sup>]i was measured by a ratiometric method using Fura-2 as a probe. An average of 3 independent experiments is shown (The individual experiments are shown in [Fig.](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF4) [S4\)](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF4). There is statistically significant difference between wild-type and  $Ahnak1^{-/-}$  cells ( $P < 0.05$ ).

expression of  $Ca<sub>v</sub>1.1$  in CD4<sup>+</sup> T cells from both *Ahnak1<sup>-/-</sup>* and  $\beta \bar{4}^{-/-}$  mice resulted in diminished calcium entry and consequent functional abnormalities after TCR stimulation (8, 11). We therefore examined whether  $Ca<sub>v</sub>1.1 \alpha1$  also was expressed in CTLs (Fig. 1 *A* and *B*). Interestingly, unlike expression in CD4 T cells,  $Ca<sub>v</sub>1.1$  and  $Ca<sub>v</sub>1.2 \alpha1$  subunits were not expressed in naive wild-type CD8<sup>+</sup> T cells (Fig. 1*A*); instead, their expression was detected only in CTLs after primary stimulation (Fig. 1*A*).

Fig. 1*B* shows the expression, measured by immunoblot assay and densitometry, of  $\hat{C}a_v1.1$  and  $Ca_v1.2$  at various time points during primary stimulation, which lasted for 5 days, followed by restimulation for 24 h. Ca<sub>v</sub>1.1 but not Ca<sub>v</sub>1.2  $\alpha$ 1 subunit expression, which correlated remarkably with AHNAK1 protein expression, was up-regulated by day 4 after primary stimulation, and remained highly expressed after restimulation (cf. Fig. 1*B* and [Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1)*B*). Although expression of a second pore-forming subunit,  $Ca<sub>v</sub>1.2$ , was up-regulated late during primary stimulation, its expression was drastically down-regulated after restimulation (Fig. 1*B Lower*). AHNAK1- deficient CTLs showed reduced induction of  $Ca<sub>v</sub>1.1$  but not of  $Ca<sub>v</sub>1.2$ ,  $\alpha$ 1 subunit expression (Fig. 1*B*), possibly because peak amounts of AHNAK1 were required. These data suggest that AHNAK1 is likely to regulate Ca<sub>v</sub>1.1  $\alpha$ 1 protein expression in CTLs in a way similar to its role in  $CD4^+$  T cells.

Reduced Ca<sub>v</sub>1.1  $\alpha$ 1 subunit expression correlated with reduced calcium entry into  $CD4+T$  cells after TCR cross-linking (11). Because Ca<sub>v</sub>1.1 expression also was reduced in  $Ahnak1^{-/-}$ CTLs after primary stimulation, we examined calcium entry after TCR stimulation in these cells. Intracellular calcium concentration in response to TCR stimulation was measured in wild-type and *Ahnak1<sup>-/-</sup>* CTLs by a ratiometric method using Fura-2-acetoxymethyl ester (Fura-2) as a probe. In 3 independent experiments, averaged in Fig. 1*C* and shown individually in [Fig. S4,](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF4) we found significant reduction ( $P < 0.05$ ) in *Ahnak1<sup>-/-</sup>* 

CTL calcium response to TCR cross-linking compared with wild-type T cells.

**Ahnak1/ CTLs Show Reduced Killing Activity.** Because CTL require calcium entry for the killing activity, we next examined CTL activity in the absence of AHNAK1 during late stages after primary stimulation in vitro. Therefore, we collected CTLs from wild-type and *Ahnak1*-/- mice late after primary stimulation (on day 5, when AHNAK1 is highly expressed) and tested their lytic activity in the presence or absence of anti-CD3 using A20 cells as targets in a redirected lysis assay in vitro. Ahnak1<sup>-/-</sup> CTLs showed marked impairment in their cytolytic activity in a standard 4-h assay (Fig. 2*A*).

We next tested *Ahnak1<sup>-/-</sup>* CTL function in vivo using a previously described model system (20–22). For this purpose, wild-type dendritic cells pulsed with H-2Kb-binding SIINFEKL peptide (OVA257–264) were transferred into wild-type and *Ahnak1<sup>-/-</sup>* mice. Under these conditions, after 7 days, we observed normal expansion of the SIINFEKL peptide (OVA<sub>257-264</sub>)-specific CTLs in vivo in both wild-type and  $Ahnak1^{-/-}$  mice as detected by staining with an H-2K<sup>b</sup>/OVA<sub>257-264</sub> peptide tetramer. (Fig. 2*B* shows representative mice, and Fig. 2*C* demonstrates average SIINFEKL peptide (OVA257–264)-specific cell numbers from 3 individual mice.) As controls we used OT-I TCR transgenic mice, which express a TCR specific for the SIINFEKL  $(OVA_{257-264})$ peptide, or naive, unimmunized wild-type mice (Fig. 2*B*) (23). This result corroborated our in vitro proliferation results [\(Fig.](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3) [S3\)](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3) and further suggests that in vivo differentiation and expansion of CTLs does not require AHNAK1.

We examined the cytotoxicity of SIINFEKL peptide (OVA257–264)-specific CTLs to transferred CFSE-labeled (0.5  $\mu$ M, CFSE-low) SIINFEKL peptide (OVA<sub>257–264</sub>)-loaded target cells 7 days after immunization of wild-type or  $A$ hnak $1^{-/-}$  mice with peptide-pulsed dendritic cells. As control, we co-transferred



**Fig. 2.** *Ahnak1*-/- CTLs show deficient cytolytic activity in vitro and in vivo. (*A*) Calcein-AM-loaded A20 were used as targets in the calcein-AM retention CTL assay, with wild-type or *Ahnak1<sup>- ו -</sup>* CTLs at the noted effector/target ratios and redirected with anti-CD3. Results shown are mean (SD) for 3 individual mice tested per group. We obtained similar results from 2 similar independent experiments. There is a statistically significant difference between wild-type and Ahnak1<sup>-/-</sup> CTL killing (\*, *P* < 0.05). (*B* and C) Wild-type dendritic cells, pulsed with H-2K<sup>b</sup>-binding SIINFEKL peptide (OVA), were transferred into wild-type and *Ahnak1<sup>-/-</sup>* mice. Splenocytes were analyzed for CTL expansion in vivo by H-2k<sup>b</sup>/OVA<sub>257-264</sub> (SIINFEKL) peptide tetramer staining 7 days later. A representative mouse is shown in *B*. An OT-I mouse is shown as positive control for the tetramer staining, and a naive wild-type mouse is shown as negative control. (*C*) An average absolute cell number was calculated for 3 individual mice. (*D* and *E*) Wild-type and *Ahnak1*-/- mice were immunized as in *B*. After 7 days, we co-transferred SIINFEKL peptide (OVA)-loaded CFSE-low-labeled (0.5  $\mu$ M) mixed with unloaded CFSE-high (5  $\mu$ M) target wild-type splenocytes. Splenocytes were harvested 3 h later and were analyzed by flow cytometry. Representative naive unimmunized wild-type control mice, immunized wild-type mice, and Ahnak1<sup>-/-</sup> mice are shown in D. The left and the right peaks shown in the histogram represent the peptide-loaded and unloaded target cells, respectively. Specific lysis was calculated from the ratio between peptide-loaded and unloaded target cells in unimmunized wild-type control and immunized wild-type or *Ahnak1<sup>–/–</sup>* mice. An average of 7 mice from each group is shown in *E*. Significant differences were found between wild-type and *Ahnak1*-/- mice (*P* 0.05).

CFSE-labeled (5  $\mu$ M, CFSE-high) naive, unloaded target cells. We observed a typical 40% antigen-specific lysis in wild-type mice (Fig. 2*E*), measured as a ratio between unloaded CFSEhigh and peptide-loaded CFSE-low target cells in each individual mouse (Fig. 2D). In contrast to wild-type animals, Ahnak1<sup>-/-</sup> mice showed significantly reduced antigen-specific cytotoxicity, consistent with the in vitro results in Fig. 2*A*.

Altogether, we observed a consistent and significant reduction in cytotoxicity (roughly 30%–40%) by *Ahnak1<sup>-/-</sup>* CTLs in vitro and in vivo.

**Reduced Expression of Granzyme B by AHNAK1-Deficient CTLs.** To examine why *Ahnak1<sup>-/-</sup>* CTLs display deficient cytotoxic activity, we tested the production of granzyme B, the main granzyme produced by CTLs during primary stimulation (24, 25). We used protein extracts from purified naive  $CD8<sup>+</sup>$  T cells as well as from wild-type and *Ahnak1<sup>-/-</sup>* CTLs to examine the expression of granzyme B by immunoblot. We detected very low granzyme-B expression in *Ahnak1<sup>-/-</sup>* CTLs obtained by primary TCR stimulation in vitro for 5 days (Fig. 3*A*). Fig. 3*B* shows the expression of granzyme B at various time points during primary stimulation (which lasted for 5 days) followed by restimulation for 24 h measured by immunoblot assay followed by densitometry. We found that granzyme-B expression correlated with AHNAK1 expression (Fig. 3B). Although *Ahnak1*<sup>-/-</sup> effector CD8<sup>+</sup> T cells up-regulated granzyme-B expression normally during the first 4 days after stimulation (Fig. 3*B*), they displayed significant ( $P <$ 0.05) reduction in granzyme-B expression on the fifth day after primary stimulation. Granzyme B was neither produced (Fig. 3 *A* and *B*) nor secreted by *Ahnak1*<sup>-/-</sup> CTLs on the fifth day after primary stimulation, as determined by ELISPOT assay (Fig. 3*C*).

**Low Levels of IFN- Are Produced by AHNAK1-Deficient CTLs.** Because Ahnak1<sup>-/-</sup> CTLs showed severely reduced cytolytic activity, we further examined their ability to secrete IFN- $\gamma$  (26). As shown in [Fig. S3,](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF3) AHNAK1 was not required for IFN- $\gamma$  production during the early stages of primary stimulation. Interestingly, we found that *Ahnak1*-/- CTLs (obtained after 5-day stimulation in vitro)



**Fig. 3.** Reduced granzyme-B production in the absence of AHNAK1. (*A*) The expression of granzyme B was examined in wild-type and Ahnak1<sup>-/-</sup>-naive CD8<sup>+</sup> T cells versus CTLs. (*B*) Kinetics of granzyme-B expression were obtained as in Fig.  $S1B$ .  $\beta$ -actin was used as the loading control. We performed 3 independent experiments. There is a statistically significant difference in granzyme-B expression in wild-type and *Ahnak1<sup>-/-</sup>* cells on the fifth day after primary stimulation (\*,  $P$ < 0.05). Experiment *B* was performed in addition to and is independent of experiment *A*. (*C*) The granzyme-B ELISPOT assay was performed using wild-type or Ahnak1<sup>-/-</sup> CTLs, stimulated for 5 days as described in [Fig. S1.](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1) We performed 3 independent experiments. There is a statistically significant difference in granzyme-B secretion in wild-type and Ahnak1<sup>-/-</sup> cells  $(P < 0.05)$ .

were defective in IFN- $\gamma$  secretion following in vitro TCR restimulation for 24 h (Fig. 4*A*). In addition, we found that  $Ahnak1^{-/-}$  CTLs produced normal IFN- $\gamma$  mRNA levels; therefore AHNAK1 probably is required for posttranscriptional events [\(Fig. S5\)](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF5). AHNAK1-deficient CTLs also exhibited normal expression of T-Bet and Eomesodermin, which regulate IFN- $\gamma$ mRNA production [\(Fig. S5\)](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF5) (27–29).

To test IFN- $\gamma$  production by CD8<sup>+</sup> T cells in the absence of AHNAK1 in vivo, we used the model system described earlier (see in vivo CTL assay, Fig. 2). Thus, wild-type dendritic cells primed with H-2K<sup>b</sup>-binding SIINFEKL peptide  $(OVA<sub>257-264</sub>)$ were transferred to wild-type and *Ahnak1<sup>-/-</sup>* mice. We examined the ability of antigen-specific CTLs to produce IFN- $\gamma$  7 days after transfer. As shown in Fig. 4 *B* and *C*, the wild-type response in the spleen resulted in 2–3% of antigen-specific IFN- $\gamma$ producing CTLs. In contrast, *Ahnak1<sup>-/-</sup>* CTLs showed a significant deficiency in IFN- $\gamma$  production, consistent with the in vitro results shown in Fig. 4*A*. In addition, this finding suggests that the deficiency in IFN- $\gamma$  production in *Ahnak1<sup>-/-</sup>* CTLs is cell autonomous, because wild-type dendritic cells were used for the transfers in this assay.

## **Discussion**

For decades, numerous studies have shown that calcium is required for CTL-mediated killing. Most of these studies were performed using calcium antagonists, which significantly inhibited the killing by CTLs (e.g., see ref. 4). However, the mechanism responsible for calcium entry required for the killing process is largely unknown.

The functional presence of  $Ca<sub>v</sub>1$  channels in T lymphocytes has been suggested previously (30–35). Ca<sub>v</sub>1.1  $\alpha$ 1 subunits are critical for TCR-induced calcium entry into  $CD4^+$  T cells, as suggested by our recent studies of  $Ca<sub>v</sub>1 \beta4^{-/-}$  and  $Ahnak1^{-/-}$ mice (8, 11). In all the cases reported thus far, reduced  $Ca<sub>v</sub>1.1 \alpha1$ subunit expression results in deficient calcium entry through the plasma membrane and does not affect calcium release from intracellular stores (11).

In the present study, we demonstrate that AHNAK1 is required for CTL function, probably by regulating Ca<sub>v</sub>1.1  $\alpha$ 1 subunit expression. In T cells, AHNAK1 interacts with  $Ca<sub>v</sub>1.1 \alpha1$  subunits and regulates their membrane expression through its interaction with the regulatory  $\beta$ 2 subunit (11). AHNAK1 and Ca<sub>v</sub>1.1  $\alpha$ 1 subunits are expressed only in CTLs, but not in naive CD8<sup>+</sup> T cells, days after primary stimulation. This expression pattern explains our finding that, unlike CD4<sup>+</sup> T cells, *Ahnak1<sup>-/-</sup>*-naive CD8<sup>+</sup> T cells do not differ from their wild-type counterparts in activation and proliferation following primary TCR stimulation both in vitro and in vivo. Our observations suggest that AHNAK1 is required after  $CD8<sup>+</sup>$  T cells differentiate to become CTLs. When the requirement for AHNAK1 by CTLs arises, a phenotype is revealed in  $A$ *hnak*1<sup>-/-</sup> CTLs, and they display severe abnormality in Ca<sub>v</sub>1.1  $\alpha$ 1 subunit expression levels and deficient calcium entry after TCR restimulation, resulting in deficient CTL activity and IFN- $\gamma$ and granzyme-B production.

Collectively,  $CD4^+$  and  $CD8^+$  T cells display differential requirements for AHNAK1 and  $Ca<sub>v</sub>1.1 \alpha1$  subunits during the immune response. We therefore conclude that T cells probably require multiple calcium pathways, including CRAC, AHNAK1,  $Ca<sub>v</sub>$ , or others, to mediate calcium entry during various differentiation stages in the course of an immune response. Further studies will be required to identify those pathways and to clarify their functional requirement during the immune response. Here we describe a critical function for AHNAK1 during CTL response, which normally is required to fight cancer or viral infections.

## **Materials and Methods**

Reagents and Antibodies. For reagents and antibodies we used anti- $\beta$ -actin (sc-1616, Santa Cruz Biotechnology), anti-CD3 (145–2C11), anti-CD28 (37.1), anti-AHNAK1-C2 (a kind gift from the Hasse-H Max Delbrück Center for Molecular Medicine), Anti-Granzyme-B (R&D, AF1865), anti-Ca<sub>v</sub>1.1 (Santa Cruz), anti-Cav1.2 (Alomone), anti-CD44 CyChrome, anti-CD62L FITC, anti-CD69 FITC, anti-TCR<sub>B</sub> (H57) PE, and anti-CD8 PE CyChrome (all PharMingen), and H-2K<sup>b</sup>/OVA<sub>257-264</sub> peptide tetramer (kindly provided by Tim Willinger).

Mice. Ahnak1<sup>-/-</sup> mice were previously described (14). Wild-type littermates were used as control. Mice within experiments were age and sex matched. Mice were cared for in accordance with protocols approved by the Institutional Animal Care and Use Committee at the Yale University School of Medicine Animal Facility.



Fig. 4. In vivo and in vitro deficient IFN-<sub>Y</sub> production in the absence of AHNAK1. (A) Wild-type and Ahnak1<sup>-/-</sup> CTLs were obtained as in [Fig. S1](http://www.pnas.org/cgi/data/0902844106/DCSupplemental/Supplemental_PDF#nameddest=SF1)A, were washed, and were restimulated using plate-bound anti-CD3 only (2  $\mu$ g/mL) for 24 h. Supernatants from in vitro-stimulated CD8<sup>+</sup> T cells were collected, and IFN- $\gamma$ production was measured by ELISA. Results shown are mean (SD) for 3 mice per group. There is a statistically significant difference between wild-type and Ah*nak1<sup>-/-</sup> cells (P* < 0.05). We obtained similar results with 2 other similar experiments. (*B*) Wild-type dendritic cells pulsed with H-2K<sup>b</sup>-binding SIINFEKL peptide (OVA) were transferred into wild-type and Ahnak1<sup>-/-</sup> mice. Splenocytes were purified and restimulated in vitro with OVA for 6 h, and 7 days later CD8 and IFN-<sub>Y</sub> expression was analyzed by flow cytometry. The normalized percentages of IFN- $\gamma$ -producing CD8<sup>+</sup> T cells are shown. Results shown are mean (SD) for 6 mice per group. We performed 2 independent experiments. There is a statistically significant difference between wild-type and *Ahnak1*-/- cells (\**P* 0.05). (*C*) Results from one representative wild-type and Ahnak1<sup>-/-</sup> are shown.

**Immunoblotting.** Cell lysis and immunoblotting were performed as previously described (11).

**ELISPOT Assay.** The ELISPOT assay was performed using the granzyme-B ELISPOT assay kit (R&D, EL1865)

**In Vitro T-Cell Activation/Differentiation Experiments.** Splenic CD8<sup>+</sup> T cells were isolated from 6- to 8-week-old mice by MACS sorting using anti-CD8-coupled beads and columns (Miltenyi Biotec). Cells were cultured in Bruff's medium (10% FCS, penicillin, streptomycin, and L-glutamine). T cells were stimulated using coated plates as described previously (36). Plates were coated for 2 h at 37 °C with anti-CD3 and anti-CD28 (2  $\mu$ g/mL of each) in PBS. Typically, 2 million T cells/well were plated in a 24-well plate. All effector CD8<sup>+</sup> T cells used in this study were incubated as described earlier in this article for a 5-day period.

**In Vitro Cytotoxicity Assay.** Cytotoxicity analysis was performed by using calcein-AM retention assays (37). Briefly, A20 target cells plated in Ubottomed microtiter plates at a concentration of  $2 \times 10^4$  cells per well were washed once with PBS and labeled with 0.02  $\mu$ g/mL calcein-AM (Molecular Probes) in serum- and phenol red-free medium (GIBCO BRL) for 30 min at 37 °C. Labeling efficiency was assessed initially by using a fluorescence plate reader. Effector cells were added at different effector/target ratios in quadruplicate. Phenol red-free medium was added to a 6-well set of target cells for estimation of retention of calcein-AM in medium alone. Anti-CD3 was added at 10  $\mu$ g/mL, and a control without anti-CD3 was included. Maximal lysis was determined by solubilizing 6 wells of target cells in lysis buffer (50 mM sodium botate 0.1% Triton X-100, pH 9.0). After 4 h of incubation at 37 °C, the assays were terminated by washing the plates twice, and the remaining fluorescence was read. The percentage of specific cytotoxicity was calculated as follows: % cytotoxicity = [(retention experimental well  $-$  retention maximal lysis)/(retention in medium – retention maximal lysis)]  $\times$  100. Retention values were calculated by normalizing measured fluorescence with initial labeling of the same well.

**ELISA.** ELISA protocols were described previously (8, 11).

**Analysis of Intracellular Calcium Concentration.** Calcium concentration was measured using Fura2/AM (Molecular Probes) as described previously (8, 11).

**5 (and 6)-Carboxyfluorescein Diacetate Succinimidyl Ester (CFSE) Labeling.** CFSE (Molecular Probes) was added to purified  $CDB<sup>+</sup> T$  cells to a final concentration of 3  $\mu$ M, and cells were incubated for 25 min at 37 °C. At the end of the incubation period, the cells were washed immediately 3 times in PBS containing 10% FCS. Cells then were stimulated for the 72-h period, and CFSE staining was measured by flow cytometry.

**Real-Time PCR.** Real-time PCR protocols were described previously (8, 11).

**In Vivo Priming of CD8<sup>+</sup> T Cells, CTL Assay, and IFN-** $\gamma$  **Production.** The method for generating bone marrow dendritic cells was adapted from Lutz et al (21). Briefly, bone marrow cells were harvested from wild-type mice and cultured for 7 days in petri dishes in complete Bruff's medium containing GM-CSF. On day 7, cells were centrifuged at 300  $\times$  g for 5 min and were resuspended in complete Bruff's medium containing GM-CSF. Cells then were incubated overnight with H-2K<sup>b</sup>-binding SIINFEKL peptide (OVA<sub>257-264</sub>) (10  $\mu$ g/mL) and LPS (200 ng/mL). On the next day, cells were washed 3 times with sterile PBS and finally were resuspended in PBS for injection.

**In Vivo IFN-** $\gamma$  **Production.** We injected 1 million cells in 200  $\mu$ L PBS i.p. into wild-type and Ahnak1<sup>-/-</sup> mice. After 7 days mice were killed, spleens were harvested, and splenocytes were stimulated in vitro with SIINFEKL peptide (10  $\mu$ g/mL) or with Phorbol Myristate Acetate (PMA) (100 ng/mL) plus ionomycin (1  $\mu$ M) or were left unstimulated in complete Bruff's medium for 6 h in the presence of Golgi stop (BD Biosciences, Cat # 554715). Cells were stained with CD8 and then were fixed and permeabilized for intracellular staining of IFN-y. Stained cells were analyzed by flow cytometry on a FACSCalibur and analyzed using CellQuest software from BD Biosciences or Flowjo software from Treestar.

In Vivo CTL Assay. Wild-type and *Ahnak1<sup>-/-</sup>* mice were immunized as described previously for 7 days, followed by co-transfer of SIINFEKL peptide (OVA)-loaded CFSE-low-labeled (0.5  $\mu$ M) (Molecular Probes) or unloaded CFSE-high (5  $\mu$ M) target wild-type splenocytes. Splenocytes were harvested 3 h later and analyzed by flow cytometry. We injected 5 million mixed target cells per mouse.

The percentage of specific lysis was calculated as { $1 -$  [(ratio in a naive mouse: unpulsed CFSE-high/pulsed CFSE-low)/(ratio in an immunized mouse: unpulsed CFSE high/pulsed CFSE low)]}  $\times$  100, as described previously (21).

**Statistical Analysis.** Results are shown as average (SD). Statistical differences were determined by an analysis of 2-tailed Student's *t* test. Values of  $P < 0.05$ were considered statistically significant.

**ACKNOWLEDGMENTS.** The authors thank F. Manzo for manuscript preparation. D.M. and R.A.F. thank Prof. Gillian M. Griffiths for helpful discussions related to this work. R.A.F. is an Investigator of the Howard Hughes Medical Institute and a recipient of grants from the National Institutes of Health (NIH). D.M. and S.S. were supported by Cancer Research Institute postdoctoral fellowships. D.M. is currently supported by a fellowship from the Israeli

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Ministry of Immigrant Adsorption. S.S. also was supported by NIH Grants CA121974 and DK051665. A.B. was supported by the Arthritis National Research Foundation. M.K.J. is supported by an Arthritis Foundation postdoctoral fellowship. T.W. is supported by a James Hudson Brown-Alexander Brown Coxe postdoctoral fellowship. A.A. is supported by a Richard K. Gershon fellowship.

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