

## Supplementary Information

### Supplementary Tables

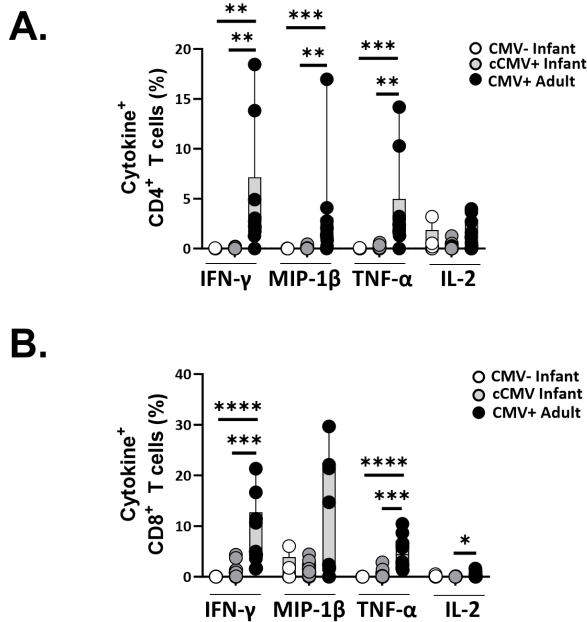
**Table S1. Summary of blood samples collected according to age.**

Age (months)	ALL	0-2	3-8	9+
CMV (n)	30	21	18	9
Control (n)	15	7	3	5

**Table S2. Flow cytometry panels and fluorochrome-conjugated antibodies.**

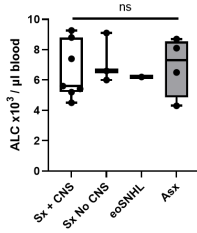
Laser	Marker	Fluorophore	Clone ID	Company	Catalog #
Surface Repertoire Panel					
405	Anti-CD8a	BV510	RPA-T8	Biolegend	300460
405	Anti-CD3	BV605	UCHT1	Biolegend	303528
405	Anti-CD57	BV785	QA17A04	Biolegend	302906
488	Anti-CD28	FITC	CD28.2	Biolegend	621614
488	Anti-PD-1	PerCP/Cy5.5	A17188B	Biolegend	359412
561	Anti-CD45RO	PE/Dazzle 594	UCHL1	Biolegend	317414
561	Anti-CD4	PE/Cy7	OKT4	Biolegend	302610
640	Anti-CD16	APC-Cy7	3G8	BD Bioscience	557758
640	Fixable Viability Stain 780	N/A	N/A	BD Bioscience	565388
Intracellular Cytokine Staining (ICS) Panel					
405	Anti-IFN- $\gamma$	BV421	4S.B3	Biolegend	301048
405	Anti-CD8a	BV510	RPA-T8	Biolegend	300460
405	Anti-CD3	BV605	UCHT1	Biolegend	328640
405	Anti-CD107a (LAMP-1)	BV711	H4A3	Biolegend	353230
405	Anti-CD197 (CCR7)	BV785	G043H7	Biolegend	304106
488	Anti-CD45RA	FITC	HI100	Biolegend	621614
488	Anti-CD279(PD-1)	PerCP/Cy5.5	A17188B	Biolegend	550078
561	Anti-MIP1 $\beta$ (CCL4)	PE	D21-1351	BD Bioscience	304248
561	Anti-CD45RO	PE/Dazzle 594	UCHL1	Biolegend	317414
561	Anti-CD4	PE/Cy7	OKT4	Biolegend	500310
640	Anti-IL-2	APC	MQ1-17H12	Biolegend	502928
640	Anti-TNF- $\alpha$	AF700	MAb11	Biolegend	565388
640	Fixable Viability Stain 780	N/A	N/A	BD Bioscience	565388

Supplemental Figures



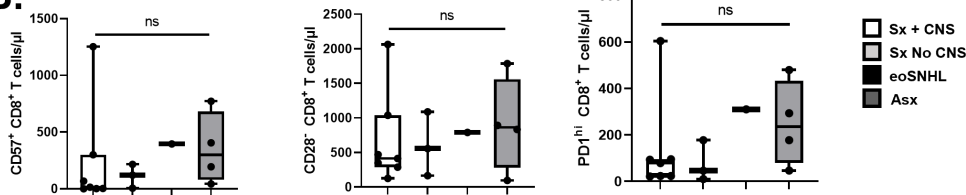
**Figure S1. T cell responses to SEB in cCMV infected neonates, uninfected infants, and CMV seropositive adults.** Peripheral blood mononuclear cells (PBMCs) of cCMV infected infants (n = 21) or uninfected infants (n = 5) at age  $\leq$  60 days and CMV seropositive adults (n = 10) were unstimulated or stimulated with staphylococcal enterotoxin B (SEB) analyzed by intracellular cytokine staining and flow cytometry. Frequencies of cytokine secreting cells in unstimulated conditions (“background”) were subtracted from those after SEB stimulation. The background-subtracted frequencies of cytokine-secreting cells after SEB stimulation are shown as frequencies of CD4<sup>+</sup> (**A**) and CD8<sup>+</sup> (**B**) T cells expressing IFN- $\gamma$ , MIP-1 $\beta$ , TNF- $\alpha$ , and IL-2 for CMV uninfected infants (CMV- infant), cCMV infected infants (CMV+ infant), and CMV seropositive adults (CMV+ adult). Groups were compared using the Kruskal-Wallis test with Dunn’s multiple comparisons for all panels. \* p < 0.05; \*\* p < 0.01; ns, not significant (p > 0.05).

**A.**

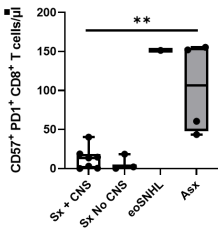


	Asx	eoSNHL	Sx No CNS	Sx With CNS
N	4	1	3	7
ALC x 10 <sup>3</sup> /μl, median [IQR]	6.5 [4.3-8.7]	6.2 [n/a]	6.6 [6.0-9.1]	5.6 [5.2-8.8]

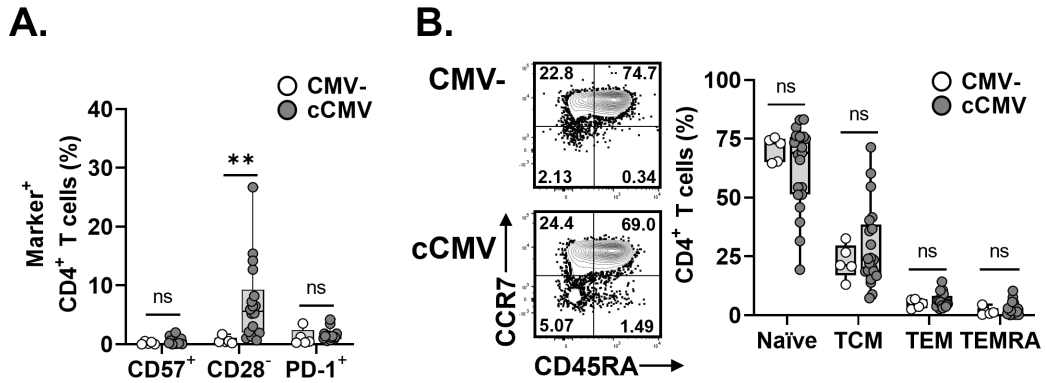
**B.**



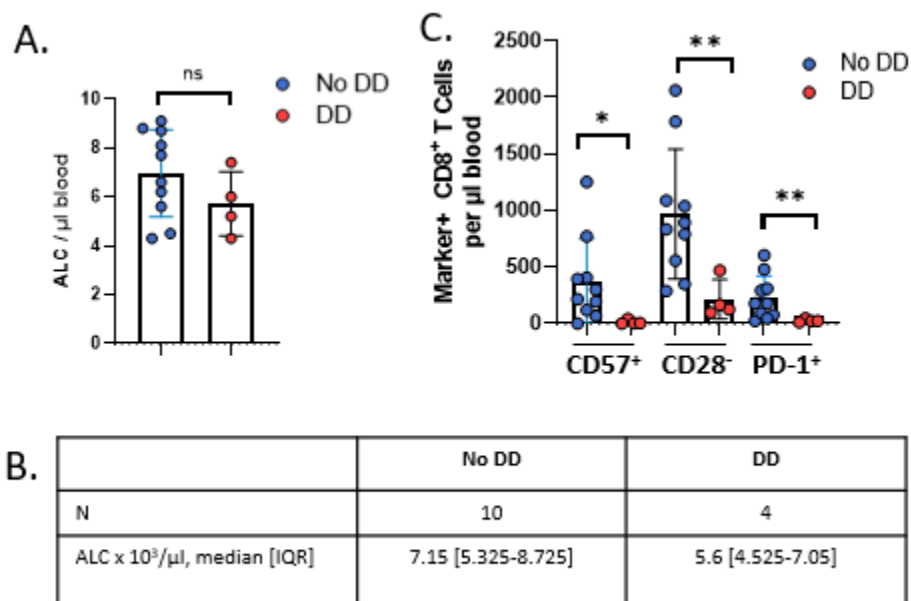
**C.**



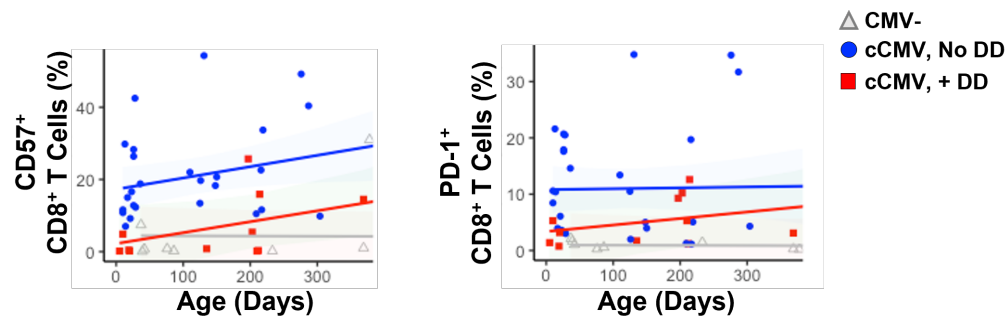
**Figure S2. Cell quantities of CD8<sup>+</sup> T cells expressing markers of differentiation in cCMV infected infants according to clinical symptoms at birth.** The cCMV infected infants were classified according to clinical symptoms as Symptomatic with CNS findings (Sx + CNS, n=7), Symptomatic without CNS findings (Sx No CNS, n=3); isolated early onset SNHL (eoSNHL, n=1), or asymptomatic (Asx, n=4) at birth. PBMCs of at age ≤ 60 days were analyzed by flow cytometry for frequencies of global populations of CD8<sup>+</sup> T cells expressing CD57, CD28, PD-1. (A) Absolute lymphocyte counts (x 10<sup>3</sup>/μl blood) were derived from the complete blood counts and compared between groups. (B) Absolute quantities of C57<sup>+</sup>, CD28<sup>+</sup>, or PD-1<sup>+</sup> CD8<sup>+</sup> T cells per μl blood were calculated using flow cytometry frequencies of CD3<sup>+</sup> CD8<sup>+</sup> T cells expressing the marker of interest and compared between clinical symptom groups. (C) Absolute quantities of C57<sup>+</sup> PD-1<sup>+</sup> CD8<sup>+</sup> T cells were calculated and compared between groups. Groups were compared using the Kruskal-Wallis test with Dunn's multiple comparisons for all panels. \* p < 0.05; \*\* p < 0.01; ns, not significant (p > 0.05).



**Figure S3. Global CD4<sup>+</sup> T cell differentiation and memory in cCMV infected infants.** PBMCs from infants at age  $\leq 60$  days were analyzed by flow cytometry for **(A)** frequencies of CD57<sup>+</sup>, CD28<sup>+</sup>, or PD-1<sup>+</sup> CD4<sup>+</sup> T cells (cCMV infected, n = 18; uninfected, n = 4), or **(B)** CCR7/CD45RA memory subsets (cCMV infected, n = 21; uninfected, n = 5). Groups were compared using the Mann-Whitney U test for all panels. \* p<0.05; \*\*\* p<0.001; ns, not significant (p > 0.05).



**Figure S4. Cell quantities of CD8<sup>+</sup> T cells expressing markers of differentiation in cCMV infected infants according to developmental outcome.** Of the cCMV infected infants with or without developmental delay who had CBCs performed (DD, n=4; No DD, n=10), groups were compared for (A-B) absolute lymphocyte counts (x 10<sup>3</sup>/μl blood), and (C) absolute quantities of C57<sup>+</sup>, CD28<sup>-</sup>, or PD-1<sup>+</sup> CD8<sup>+</sup> T cells per μl blood. Groups were compared using the Mann-Whitney U test for all panels. \* p < 0.05; \*\* p < 0.01; ns, not significant (p > 0.05).

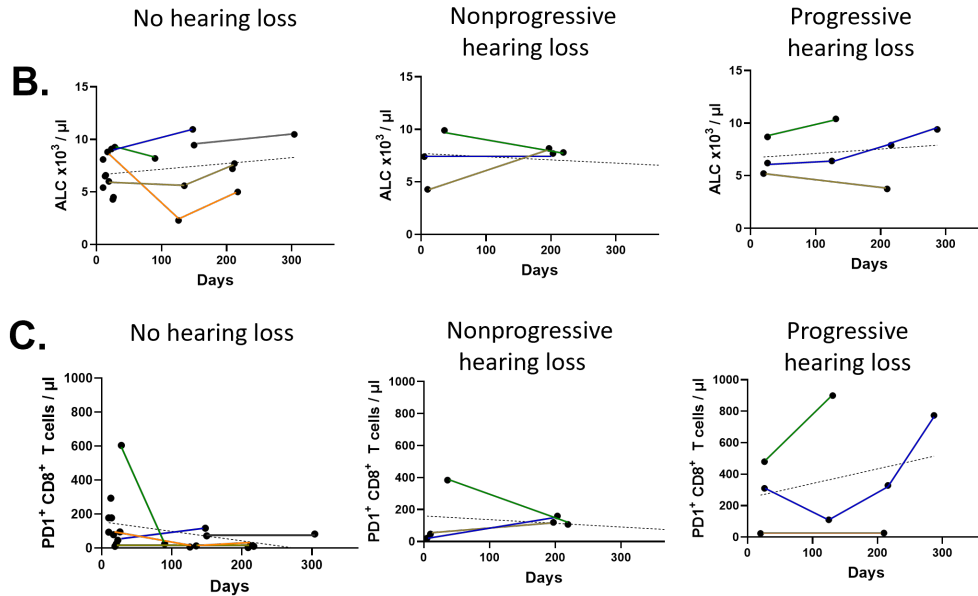


**Figure S5. Longitudinal comparison of CD8<sup>+</sup> T cells according to developmental outcome.**

For all infants with longitudinal samples available for analysis, frequencies of CD57<sup>+</sup> or PD-1<sup>+</sup> CD8<sup>+</sup> T cells are shown over the first 365 days of age for those with no DD (n = 14), or DD (n = 4) with 95% confidence intervals shown as shaded areas using linear mixed models.

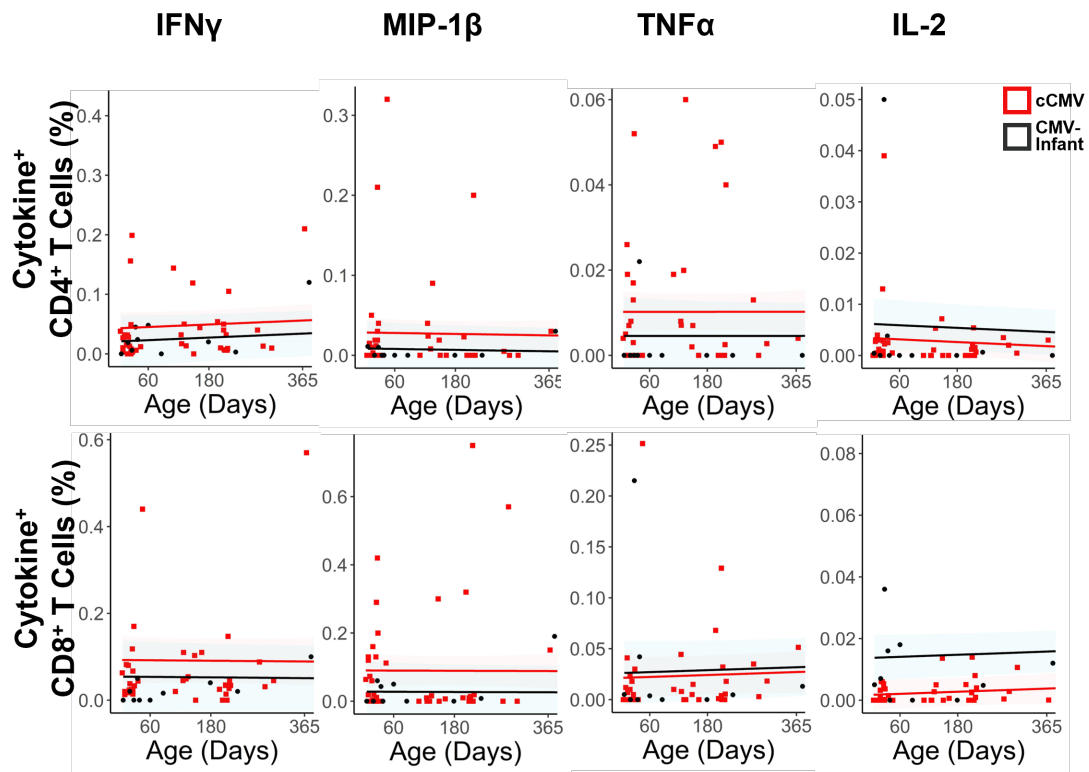
**A.**

	No Hearing Loss	Nonprogressive hearing loss	Nonprogressive hearing loss
N	12	5	3
ALC x 10 <sup>3</sup> /μl, median [IQR] (min-max)	7.2 [5.4-9.1] (2.3-10.96)	7.55 [5.1-8.1] (3.9-9.9)	7.2 [5.5-9.2] (3.7-10.4)

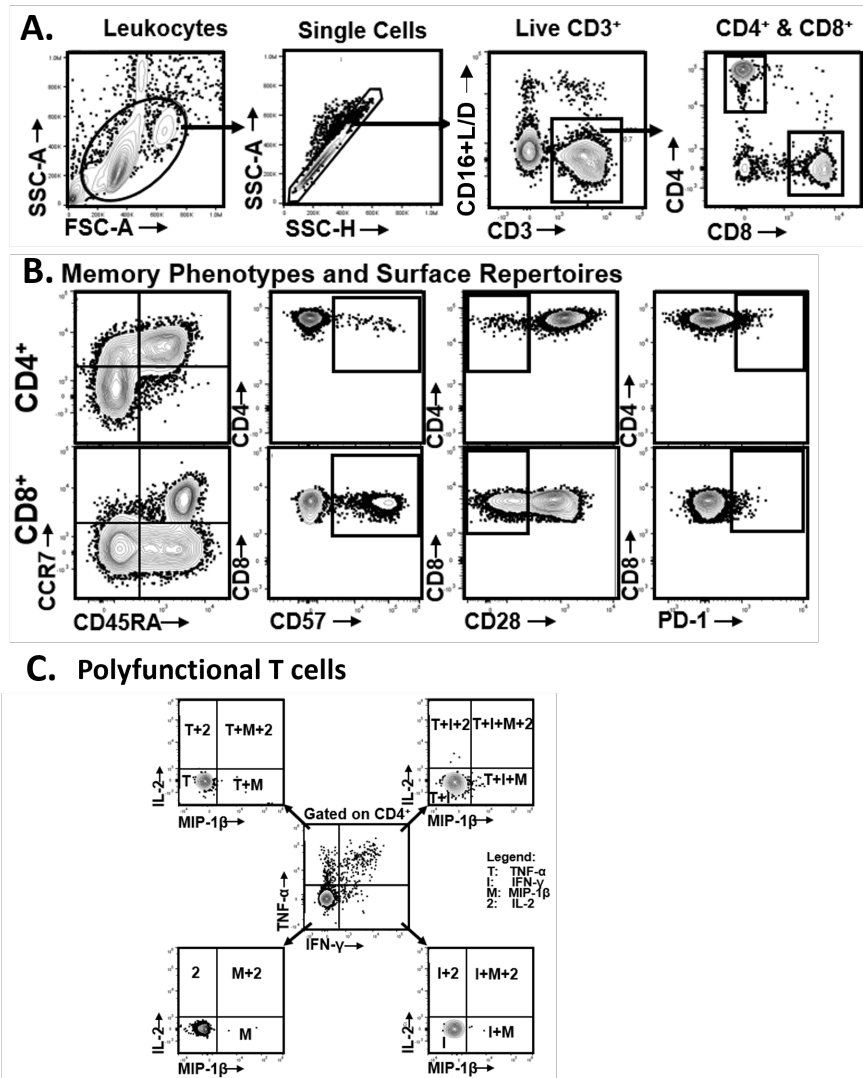


**Figure S6. Longitudinal Absolute lymphocyte counts and PD-1<sup>+</sup> CD8<sup>+</sup> T cells according to hearing outcome.** For cCMV infected infants with available CBCs, those with no hearing loss (n=12) were compared to those with nonprogressive (n=5) or progressive SNHL (n=3). **(A-B)** Absolute lymphocyte counts (x 10<sup>3</sup>/μl blood) are shown for infants with no hearing loss, nonprogressive SNHL, and progressive SNHL. In panel B, repeated observations over time are linked with colored lines representing individual subjects. **(C)** Absolute quantities of PD-1<sup>+</sup> CD8<sup>+</sup> T cells per μl blood were calculated based on flow cytometry frequencies of CD3<sup>+</sup> CD8<sup>+</sup> T cells expressing PD-1. Repeated observations over time are connected with colored lines corresponding to the same subjects shown within each group in panel B. Linear regression shown as dotted lines.





**Figure S7. Frequencies of CMV pp65-specific cytokine expressing CD4<sup>+</sup> and CD8<sup>+</sup> T cells over the first year of age.** Frequencies of CD4<sup>+</sup> and CD8<sup>+</sup> T cells expressing IFN- $\gamma$ , MIP-1 $\beta$ , TNF- $\alpha$ , or IL-2 after pp65 stimulation are shown for CMV uninfected infants (black, n=30) or cCMV infected infants (red, n=13) over 365 days. 95% confidence intervals shown as shaded areas using linear mixed models.



**Figure S8. Gating Strategy**

Debris and doublets were excluded using forward and side scatter area vs. height. Singlets were gated for viable CD3<sup>+</sup> T cells, followed by gating for CD4<sup>+</sup> and CD8<sup>+</sup> cells. Within the CD4<sup>+</sup> or CD8<sup>+</sup> population, CD45RA and CCR7 were used to define naïve (CD45RA<sup>+</sup>CCR7<sup>+</sup>), central memory (T<sub>CM</sub>; CD45RA<sup>-</sup>CCR7<sup>+</sup>), effector memory (T<sub>EM</sub>; CCR7<sup>-</sup>CD45RA<sup>-</sup>) and TEMRA (CCR7<sup>-</sup>CD45RA<sup>+</sup>) populations. Independently, CD4<sup>+</sup> or CD8<sup>+</sup> T cells were gated for CD57<sup>+</sup>, CD28<sup>+</sup>, or PD-1<sup>+</sup> cells. To determine polyfunctionality, CD4<sup>+</sup> T- cells were first gated for TNF- $\alpha$  and/or IFN- $\gamma$  expression. Each quadrant of TNF- $\alpha$ /IFN- $\gamma$  gating were further gated for IL-2 and/or MIP-1 $\beta$ . The frequencies of single vs. multiple cytokine expressing cells were expressed as percentages of total CD4<sup>+</sup> T cells. For SPICE analysis, frequencies of single and multiple cytokine expressing CD4<sup>+</sup> T cells were independently obtained using Boolean gating within FlowJo<sup>TM</sup> software.