## Cutting Edge: Innate memory CD8<sup>+</sup> T cells are distinct from homeostatic expanded CD8<sup>+</sup> T cells and rapidly respond to primary antigenic stimuli

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**Supplemental Data** 

Supplemental FIGURE 1. Changes in specific genes between HP and IMP CD8<sup>+</sup> T cells. (A) Relative *Tnfrsf1a*, *Nfkb2*, *Eomes and Bcl2111* expressional level in HP and *Itk*<sup>-/-</sup> CD8<sup>+</sup> T cells compared to WM IMP CD8<sup>+</sup> T cells as determined by microarray analysis. Quantile normalized gene expressional values were normalized to mean values of WM IMP CD8<sup>+</sup> T cells. n = 3 in each group. (B) Relative mRNA expression for *Tnfrsf1a*, *Nfkb2*, *Eomes and Bcl2111* expressional level in HP and *Itk*<sup>-/-</sup> CD8<sup>+</sup> T cells compared to WM IMP CD8<sup>+</sup> T cells, determined by qRT-PCR. Gene expressional level was first normalized to GAPDH value of each sample, than normalized to the average value of WM IMP CD8<sup>+</sup> T cells. n = 3 in each group. \*p < 0.05, "ns" = no significance by Student's *t* test.

Name	HP	ltk-/-
Tnfrsf1a	5.89	0.97
Nfkb2	3.91	1.16
Eomes	0.32	1.68
Bc/2/11	0.12	1.36

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