



Figure S1. Analysis of intracellular cytokine expression of IFNa subtype treated CD8⁺ T cells. Positively sorted Cell TraceTM Violet-labeled CD8⁺ T cells from FV-specific TCRtg mice were co-cultured with FV peptide-loaded BM-DCs in the presence or absence of different murine IFNa subtypes for 72 hours (500 units/well). Multi-parametric flow cytometry was used to determine MFI of intracellular expression of IFN γ , IL-2 and TNF α in CD8⁺ T cells. (A) Representative histograms of unstimulated, IFN α 4-stimulated CD8⁺ T cells and FMOs are shown. Intracellular expression (MFI) of (B) IFN γ , (C) IL-2 and (D) TNF α are shown. Mean values (+SEM) are indicated by bars and the IFN α subtypes were sorted in the order of their antiproliferative potency (n=15). Statistically significant differences between the IFN α -treated groups and the untreated group were tested using Kruskal-Wallis one-way or Ordinary One Way ANOVA analysis and Dunn's multiple comparison and are indicated by * for p < 0.05; ** for p < 0.01; *** for p < 0.001.



Figure S2. Influence of IFNAR expression on intracellular cytokine expression of IFNα subtype-stimulated CD8⁺ T cells. Positively enriched Cell TraceTM Violet-labeled CD8⁺ T cells from FV-specific TCRtg (WT) or IFNAR^{-/-} TCRtg (IFNAR^{-/-}) mice were co-cultured with FV peptide-loaded WT or IFNAR^{-/-} BM-DCs in the presence or absence of IFNα4, IFNα6 and IFNα9 for 72 hours (500 units/well). Multi-parametric flow cytometry was used to determine MFI of intracellular expression of (A) IFNγ, (B) IL-2 and (C) TNFα in CD8⁺ T cells. Mean values (+SEM) are indicated by bars and the IFNα subtypes were sorted in the order of their antiproliferative potency (n≥6). Statistically significant differences between the IFNα-treated groups and the untreated group were tested using Kruskal-Wallis one-way or Ordinary One Way ANOVA analysis and Dunn's multiple comparison and are indicated by * for p < 0.05; ** for p < 0.01; *** for p < 0.001.