

(a) Representative plots displaying pre-sort and post-sort purity of splenic NK cells from WT and SMAD4-deficient mice. (b) Quantification of surface expression of the indicated markers by splenic NK cells from WT and SMAD4-deficient mice. (c) Representative histograms showing expression of the indicated proteins by splenic NK cells from WT and SMAD4-deficient mice. (d) Representative plots and quantification of siLP ILC3 populations in WT and SMAD4-deficient mice. (e) Representative plots and quantification of siLP ILC3 after stimulation by IL-23. (f) Representative histograms of cell surface expression of CD49a in NKp46<sup>-</sup> or NKp46<sup>+</sup> NK cells from the bone marrow of WT and Smad4<sup>t/f</sup> x II7r<sup>Cre</sup> mice. \*\*P<0.01, \*\*\*P < 0.001; (unpaired Student's t-test). Data are pooled from four (b) or two (d,e,f) independent experiments with 1-2 mice of each genotype for each independent experiment (error bars represent mean  $\pm$  s.d. in **b**,d,e).



(a) Representative plots of Ly49H<sup>+</sup> and Ly49H<sup>-</sup> splenic NK cells (gated on CD3<sup>-</sup>CD19<sup>-</sup>) from naïve WT and *Smad4<sup>t/f</sup> x Ncr1<sup>iCre</sup>* mice. (b) Ly49H<sup>+</sup> splenic NK cells from WT and *Smad4<sup>t/f</sup> x Ncr1<sup>iCre</sup>* mice positive for Annexin V at days 3 and 6 of MCMV infection. (c) Splenic NK cells from WT and *Smad4<sup>t/f</sup> x Ncr1<sup>iCre</sup>* mice positive for Annexin V ex vivo or after 24 and 48 h of culture with IL-2. Data are pooled (b, c) from two independent experiments with at least 1-2 mice of each genotype for each independent experiment (error bars represent mean ± s.d. in b, c).



Representative histograms and quantification of CD49a, CD73, and TRAIL surface expression by (**a**) WT and (**b**) SMAD4-deficient NK cells after 48 h of culture with TGF- $\beta$ 1 + IL-2, BMP4 + IL-2, activin A + IL-2, or IL-2 alone. \*\**P*<0.01, \*\*\**P* < 0.001 (unpaired Student's *t*-test). Data are pooled from three independent experiments with 1-2 mice of each genotype for each independent experiment (error bars represent mean ± s.d.).



Impact of SMAD4, DKO, TGFβR2, and BMPR2 deficiency on conventional NK cells and SG ILC1.

(a) Representative histograms of TGF $\beta$ R2 expression by splenic NK cells from WT, *Smad4<sup>t/t</sup> x Ncr1<sup>iCre</sup>*, DKO, and *Tgfbr2<sup>t/t</sup> x Ncr1<sup>iCre</sup>* mice *ex vivo*. (b) Representative histograms intracellular phospho-SMAD2/3 and phospho-SMAD1/8 in splenic NK cells after 60 min of culture with TGF- $\beta$ 1 and IL-2. (c) Representative histograms and quantification of CD49a expression by SG ILC1 from WT, *Smad4<sup>t/t</sup> x Ncr1<sup>iCre</sup>*, DKO, and *Tgfbr2<sup>t/t</sup> x Ncr1<sup>iCre</sup>* mice. (d) Relative gene expression of type II receptors by splenic NK cells of WT mice (Immgen.org). (e) Representative histograms of cell surface expression of CD49a, CD73, and TRAIL (after 48 h of culture with IL-2) by splenic NK cells from WT, *Smad4<sup>t/t</sup> x Ncr1<sup>iCre</sup>*, *Tgfbr2<sup>t/t</sup> x Ncr1<sup>iCre</sup>*, and *Bmpr2<sup>t/t</sup> x Ncr1<sup>iCre</sup>* mice. (f) Quantification of Annexin V<sup>+</sup> NK cells from WT, *Smad4<sup>t/t</sup> x Ncr1<sup>iCre</sup>*, DKO, and *Tgfbr2<sup>t/t</sup> x Ncr1<sup>iCre</sup>*, mice after 48 h of culture with IL-2 or TGF- $\beta$ 1 plus IL-2 and DMSO or SB431542. \*\*\**P* < 0.001 (unpaired Student's *t*-test). Data are pooled from at least two independent experiments with 1-2 mice of each genotype for each independent experiment (error bars represent mean ± s.d.).