



(A) Representative FACS plots to identify CD11b<sup>+</sup> and CD8<sup>+</sup> subsets in S1<sup>AF647</sup>-captured migratory or resident DCs as gated in Fig 4D. Representative plots of two independent experiments are shown.

(B)  $S1^{AF647+}$  cell frequencies in the CD11b<sup>+</sup> and CD8<sup>+</sup> subsets in migratory or resident DCs in draining lymph nodes on day 2 after DT+S1<sup>AF647</sup>-injection. Representative of two independent experiments. The data are presented as the mean  $\pm$  SEM (n = 3/group). Data were analyzed using two-way ANOVA with Tukey's multiple comparisons test. (C) Representative overlaid contour plots of CD80 and CD86 expression gated on CD11b<sup>+</sup> or CD8<sup>+</sup> DC subsets within S1<sup>AF647+</sup> resident or S1<sup>AF647+</sup> migratory DC subsets from draining lymph nodes of DT+S1<sup>AF647-</sup>injected mice are shown (WT, green; *Foxp3*<sup>DTR</sup>, red-purple). Representative of two independent experiments for FACS plots and graphics are shown. The graphic data represents as the mean  $\pm$  SEM of the MFI (n=3/ group). Data were analyzed using an

unpaired Student's t-test.