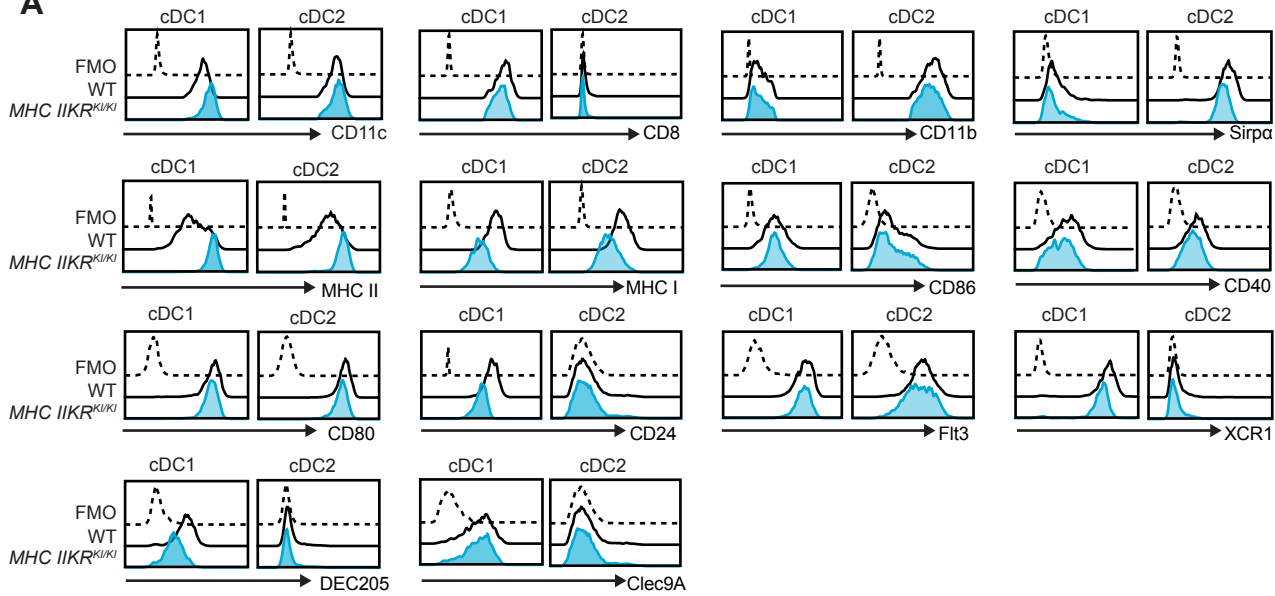
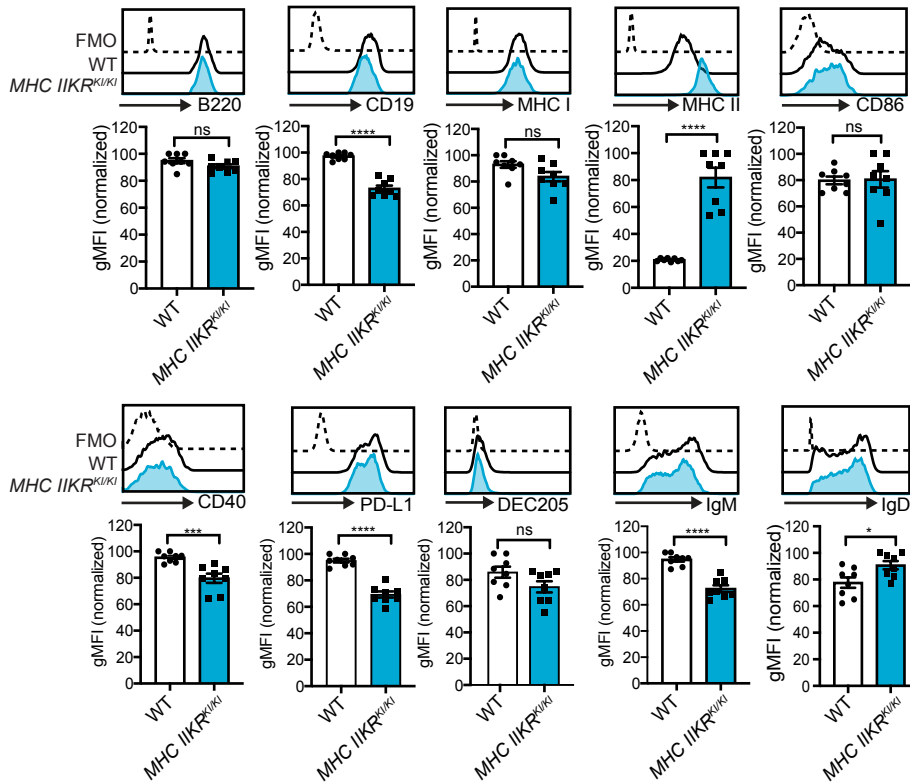


## Supplementary Figure 1

**A**

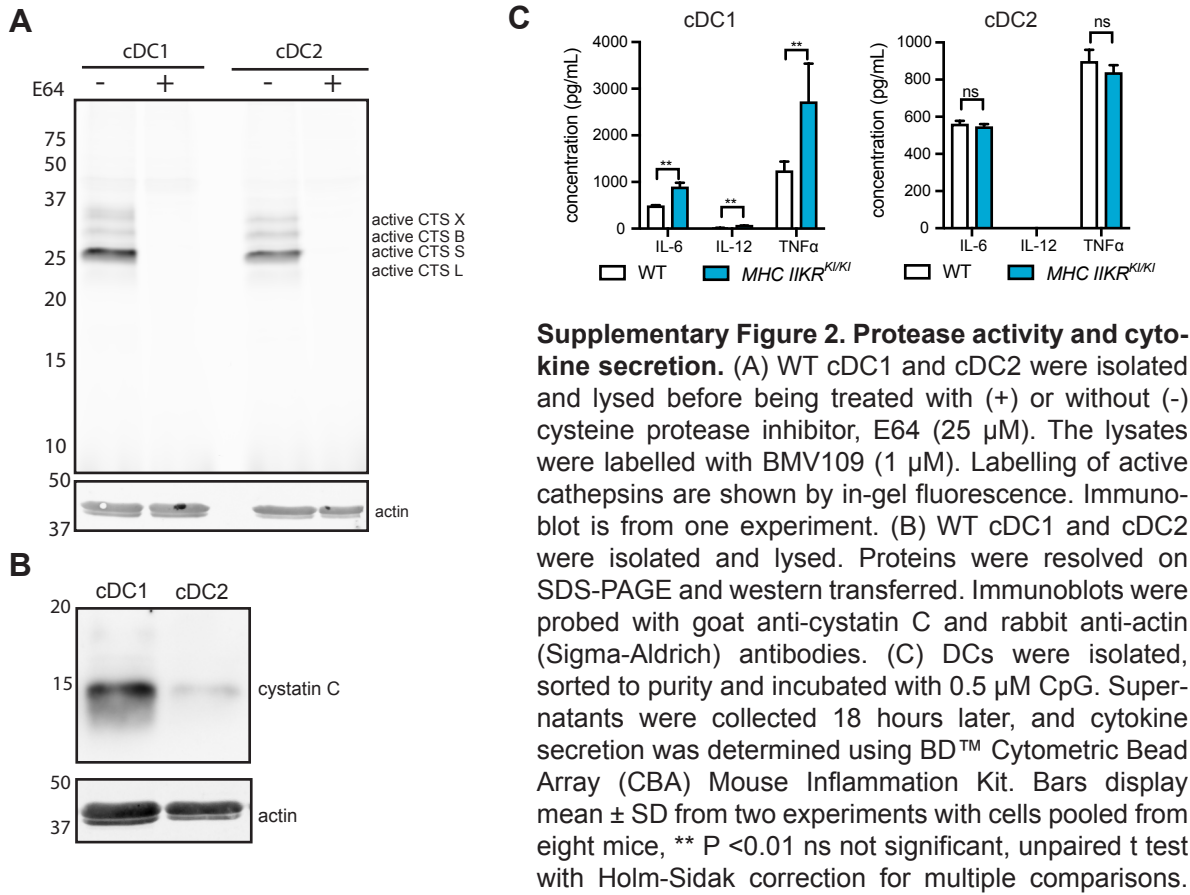


**B**



**Supplementary Figure 1. Lack of MHC II ubiquitination alters splenic DC and B cell cell-surface phenotype.** WT and *MHC IIKR<sup>K/IKI</sup>* splenic (A) DC or (B) B cell surface marker expression determined by flow cytometry. Representative histograms are shown for fluorescence minus one (FMO, dashed line), WT (black line) and *MHC IIKR<sup>K/IKI</sup>* (blue filled). gMFI for B cell markers has been quantified by normalizing the gMFI signal to the maximum gMFI signal. Symbols represent individual mice, with data pooled from a minimum of 2 independent experiments. \*\*\*\* P < 0.0001 \*\*\* P < 0.001\* P < 0.05, ns not significant, unpaired t test.

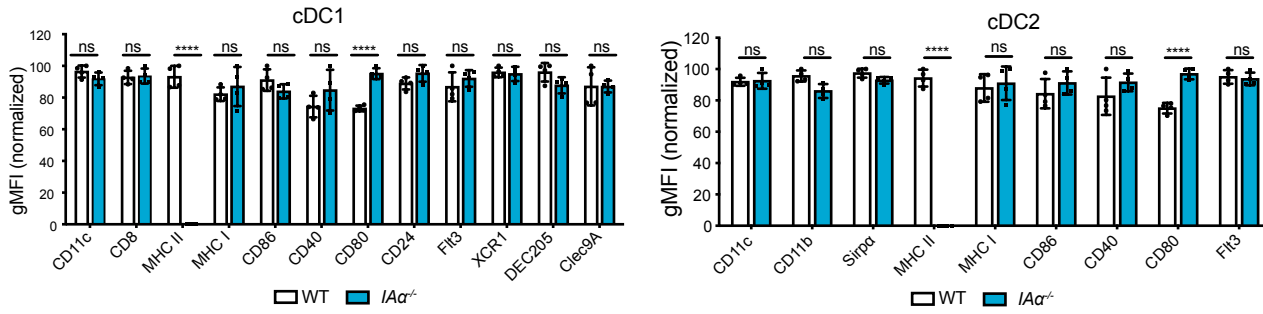
## Supplementary Figure 2



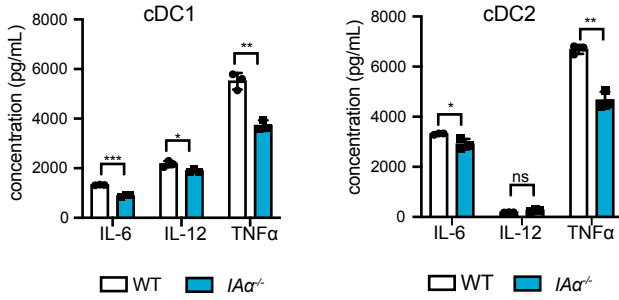
**Supplementary Figure 2. Protease activity and cytokine secretion.** (A) WT cDC1 and cDC2 were isolated and lysed before being treated with (+) or without (-) cysteine protease inhibitor, E64 (25  $\mu$ M). The lysates were labelled with BMV109 (1  $\mu$ M). Labelling of active cathepsins are shown by in-gel fluorescence. Immunoblot is from one experiment. (B) WT cDC1 and cDC2 were isolated and lysed. Proteins were resolved on SDS-PAGE and western transferred. Immunoblots were probed with goat anti-cystatin C and rabbit anti-actin (Sigma-Aldrich) antibodies. (C) DCs were isolated, sorted to purity and incubated with 0.5  $\mu$ M CpG. Supernatants were collected 18 hours later, and cytokine secretion was determined using BD™ Cytometric Bead Array (CBA) Mouse Inflammation Kit. Bars display mean  $\pm$  SD from two experiments with cells pooled from eight mice, \*\* P < 0.01 ns not significant, unpaired t test with Holm-Sidak correction for multiple comparisons.

# Supplementary Figure 3

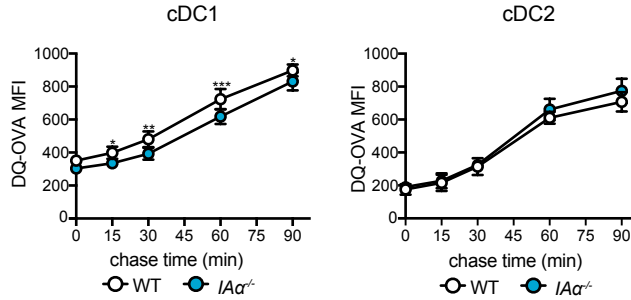
**A**



**B**

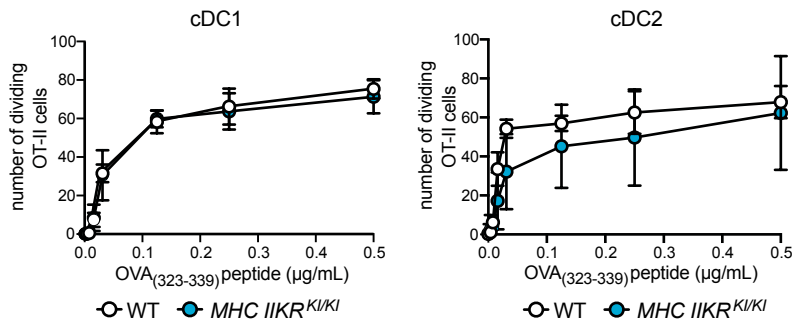


**C**



**Supplementary Figure 3. DC lacking MHC II have altered function.** (A) WT and *IAα*<sup>-/-</sup> splenic cDC1 and cDC2 cell-surface marker expression determined by flow cytometry. gMFI has been normalized to the maximum gMFI signal. Symbols represent individual mice. Data are from one experiment. \*\*\*\* P < 0.0001, ns not significant, unpaired t test. (B) WT and *IAα*<sup>-/-</sup> splenic cDC1 and cDC2 cytokine secretion. DC were sorted to purity and incubated with 0.5 μM CpG, 50 ng/mL IFNγ and 20 ng/mL GM-CSF. Supernatants were collected 18 hours later, and cytokine secretion was determined using BD™ Cytometric Bead Array (CBA) Mouse Inflammation Kit. Bars display mean ± SD from two experiments with cells pooled from eight mice, \*\*\* P < 0.001 \*\* P < 0.01 \* P < 0.05, ns not significant, unpaired t test. (C) *IAα*<sup>-/-</sup> and WT cDC1 and cDC2 were isolated, sorted to purity and pulsed with 20 μg/mL DQ-OVA for 15 min. Cells were washed twice and DQ-OVA signal was measured by flow cytometry at different chase time points. \*\*\*\* P < 0.0001\*\*\* P < 0.001\* P < 0.05, two-way ANOVA with Bonferroni's test for multiple comparisons.

## Supplementary figure 4



**Supplementary figure 4. MHC II ubiquitination does not impact in vitro presentation of OVA<sub>(323-339)</sub> peptide.**  $2.5 \times 10^4$  sorted WT and MHC IIKR<sup>KI/KI</sup> cDC1 and cDC2 were incubated with increasing concentration of OVA<sub>(323-339)</sub> peptide and  $2.5 \times 10^4$  OT-II cells. 68 hours later, the number of divided OT cells was determined as described in Materials and Methods. Data is pooled from two independent experiments.