# Supplemental Table 1. Vκ-Jκ usage

Family	R4A	R4A x FcγRIIB <sup>-/-</sup>
<b>Vκ1</b> BB1-Jκ1 <sup>a</sup> BB1-Jκ 2 BB1-Jκ4 <sup>a</sup> BB1-Jκ5 <sup>b</sup> <b>Total</b>	2 4 3 3 <b>12</b>	16 6 10 0 <b>32</b>
VK4/5 AH4-JK5 KB4-JK2 KB4-JK4 KF4-JK5 KK4-JK5 Total	5 2 2 3 5 <b>19</b>	2 1 0 1 1 <b>5</b>
Vk9/10 Bv9-Jk5 CE9-Jk4 CE9-Jk5 Total	3 0 0 <b>3</b>	3 1 3 7
Vk12 12-44-Jk2 12-44-Jk4 Total	0 0 <b>0</b>	2 1 <b>3</b>
Vk21 21-2-Jk1 21-4-Jk1 21-2-Jk2 21-2-Jk4 21-5-Jk2 21-12-Jk5 Total	4 0 0 0 0 0 4	0 3 2 2 2 2 4 <b>13</b>
Vk23 23-48-Jk4 23-48-Jk5 Total	0 0 <b>0</b>	4 3 <b>7</b>
<b>Vk24/Vk25</b> HF24-Jk4 HF24-Jk5	2 2 <b>4</b>	000
Total sequenced	42	67

VK families were identified using the IgBLAST algorithm. <sup>a</sup>Germline genes that generate high affinity anti-DNA antibodies when paired with the R4A H chain.

<sup>b</sup>Germline gene that generate a low affinity anti-DNA antibodies when paired with the R4A H chain.

## Supplemental Table II. Phenotype of FcγRIIB-/- DCs.

	% CD86 <sup>+</sup>	%CD83 <sup>+</sup>
WT	34.8 (± 16.3)	20.1 (± 4.7)
FcγRIIB <sup>-/-</sup>	52.7% (± 3.1) <sup>a</sup>	35.4 (± 6.8) <sup>b</sup>

The percentage of CD86<sup>+</sup> and CD83<sup>+</sup> DCs was determined by flow cytometry. Data are presented as the mean  $\pm$  SD.  $^ap$  = 0.057;  $^bp$  < 0.029 determined by Mann-Whitney.

## Supplemental Table III. Cytokine profile of medium and LPS-treated DCs

	WT		FcγRIIB <sup>-/-</sup>	
	medium	LPS	medium	LPS
TNFα	1.6 (± 0.3)	6.7 (± 4.0)	1.2 (± 0.2)	5.6 (± 2.3)
IL-6	1.7(± 0.4)	554.0 (± 127.3)	3.1 (± 1.6)	819.0 (± 245.8)
IL-10	6.6 (± 4.2)	3052 (± 1813)	6.2 (± 4.8)	3319 (± 1298)
IL-12	6.4 (± 2.2) <sup>a</sup>	78.4 (± 17.2)	2.4 (± 1.0)	94.6 (± 42.6)

Cytokine responses to medium and LPS were measured by qPCR. Values are expressed as the mean  $\pm$  SD of the relative expression levels normalized to *polr2a*. There were no statistically significant differences in the cytokine responses to medium or LPS between WT and Fc $\gamma$ RIIB<sup>-/-</sup> DCs, with the exception of IL-12<sup>a</sup>, which was lower in medium-treated Fc $\gamma$ RIIB<sup>-/-</sup> DCs (p < 0.03 as determined by Mann-Whitney).

## Supplemental Table IV. Cytokine profile of sera-treated DCs

		WT mDCs		FcγRIIB <sup>-/-</sup> mDCs	
sera	R4A	R4A x FcγRIIB-/-	R4A	R4A x FcγRIIB-/-	
TNFα	0.9 (± 0.1)	1.4 (± 0.4)	0.9 (± 0.1)	1.6 (± 0.7)	
IL-6 <sup>a</sup>	1.6 (± 0.2)	3.2 (± 0.8)	1.8 (± 0.2)	4.0 (± 2.0)	
IL-10 <sup>a</sup>	6.7 (± 1.2)	45.2 (± 30.2)	5.0 (± 2.1)	12.2 (± 18.7)	
IL-12	3.9 (± 1.7)	4.2 (± 1.0)	2.2 (± 0.8)	2.9 (± 1.2)	

Cytokine responses of DCs to sera treatment were measured by qPCR. Values are expressed as the mean  $\pm$  SD of the relative expression levels normalized to *polr2a*.