

Supplemental Materials

	Gene	Forward (5' – 3')	Reverse (5' – 3')
Genomic PCR	V γ 2-J γ 1	TGGACATGGGAAGTTGGAG	CAGAGGGAATTACTATGAGC
	V γ 3-J γ 1	GATCAGCTCTCCTTTACCC	
	V γ 4-J γ 1	CTGGGGTCATATGTCATCAA	
	V γ 5-J γ 1	GCTAACCTACCATTCTCTGT	
	V γ 1.1-J γ 4	CTCCATATTTCTCCAACACAGC	ACTACGAGCTTTGTCCCTTGG
	V γ 1.2-J γ 2		ACTATGAGCTTTGTTCCTTCTGCAA
	V δ 4	CCGCTTCTCTGTGAACTTCC	CAGTCACTTGGGTTCCCTTGTC
	V δ 5	CAGATCCTTCCAGTTCATCC	
	Gapdh	AGGTCGGTGTGAACGGATTG	TGTAGACCATGTAGTTGAGGTCA
C δ	TTGAGAGAGAGAACCCGTAT	CTCTCCTGTGTGACTATTGG	
Copy number assay	C δ (primer)	CTGGAATGGTACCGCACAGTAG	AAGAGGGAGCGGTGGTATCA
	C δ (probe)	CACCAGCCTTCCCA	
RT-PCR	V γ 2-C γ	GGAGTACAAGAAAATGGAGG	CTTATGGAGATTTGTTTCAGC
	V γ 3-C γ	CATCGGATGAAGCCACGTA	
	V γ 4-C γ	AGTGACAGAAGAGGACACG	
	V γ 5-C γ	CGATTCTGCTCTGTACTACT	
	V γ 1.1-C γ	AACTTCTACCTCAACCTTGA	
	V γ 1.2-C γ	AAGTTCTACCTCAACCTTGG	
	Gapdh	AGGTCGGTGTGAACGGATTG	TGTAGACCATGTAGTTGAGGTCA

Table S1. List of primer/probe sequences

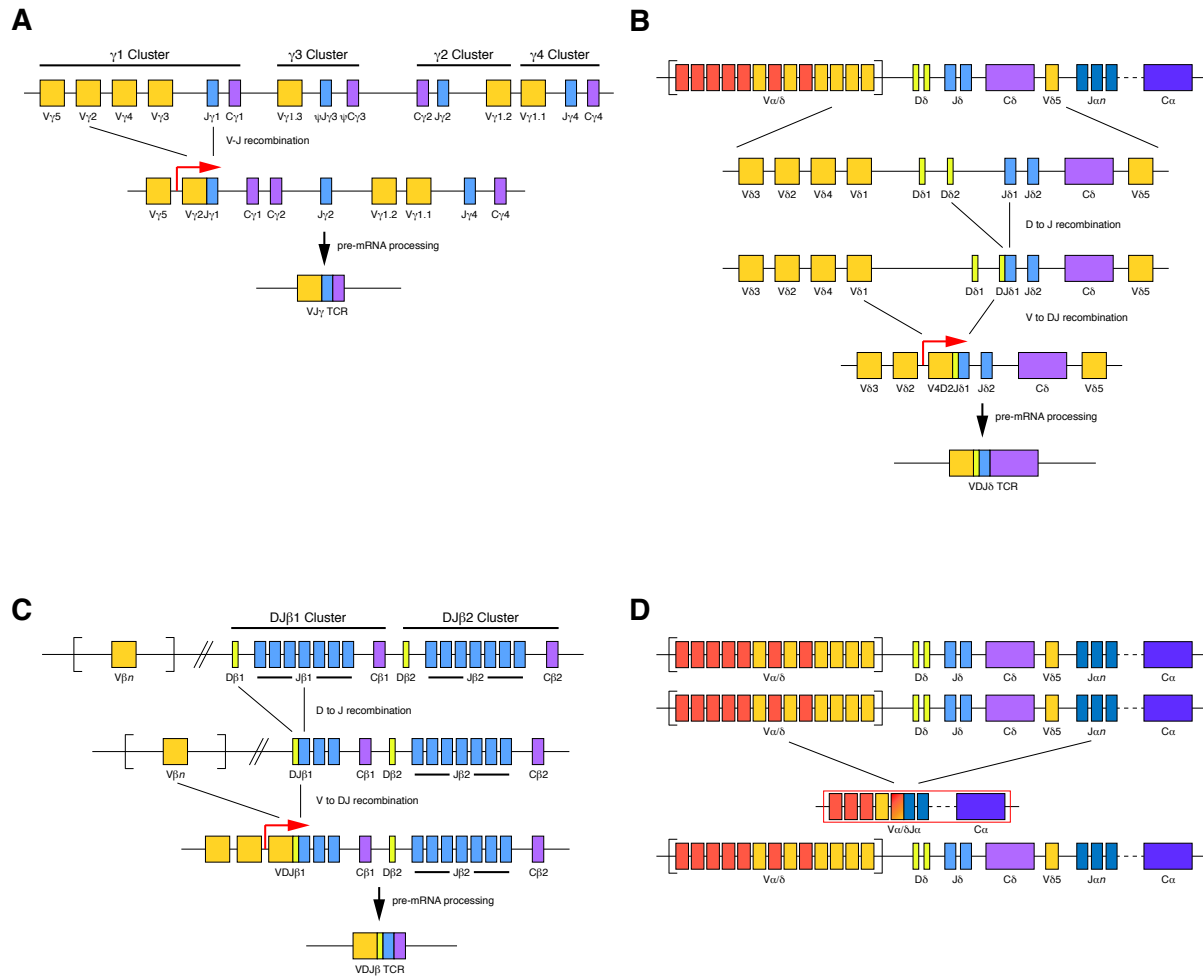


Figure S1. Comprehensive schematic of TCR γ (A), TCR δ (B), TCR β (C) and TCR α (D), and their rearrangement mechanism. Transcription is marked by a red arrow.

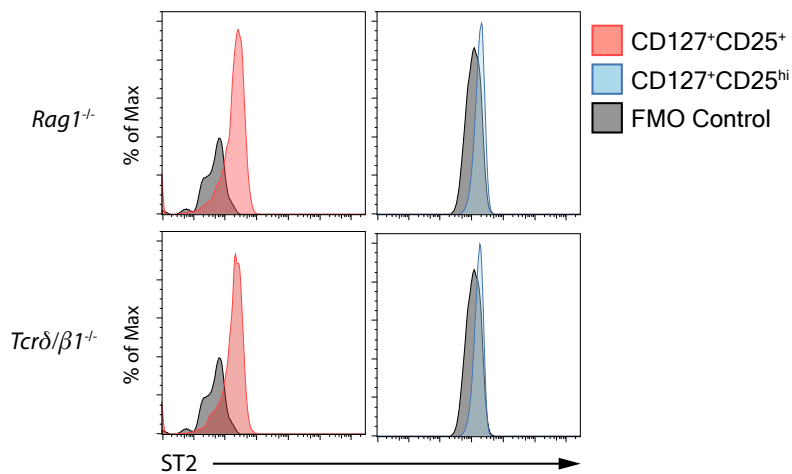


Figure S2. Expression of ST2 in *Rag1*^{-/-} and *Tcr δ/β* ^{-/-} thymic ILC2s. Thymic CD45.2⁺Lin⁻Thy1.2⁺ cells in both *Rag1*^{-/-} and *Tcr δ/β* ^{-/-} mice were further characterized into CD127⁺CD25⁺

and CD127⁺CD25^{hi} populations and the expressions of ST2 were analyzed. CD127⁺CD25⁺ (red) and CD127⁺CD25^{hi} (green) subsets were compared against ST2 null (grey).