
Supplementary information

Disrupting Roquin-1 interaction with Regnase-1 induces autoimmunity and enhances antitumor responses

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Genotype			Roquin-1/2 DKO ^T	Regnase-1 KO ^T	Roquin-1/2 Regnase-1 TKO ^T	
<i>Cd4-Cre</i> deletion (homeostasis)	CD4⁺ T cells	activation	EM	EM/CM	EM/CM	
		T _{FH} cells (%)	++	+	+++	
		Treg cells (%)	+++	+++	++	
		glycolytic activity	++	++	+++	
		mitochondrial respiration	+	++	+++	
	GC B cells		+	++	+++	
		ANAs	unchanged	+++	++	
	CD8⁺ T cells	activation	EM	EM/CM	EM	
		glycolytic activity	++	++	+++	
		mitochondrial respiration	+	++	(++)	
		ICOS	+	++	++	
		CD38	++	+	++	
		IFN- γ + TNF+	+++	+++	+++	
		IL-2	unchanged	+++	+	
		Granzyme B	+++	+	++	
		KLRG1	+++	+	+	
		BATF	+	+	+	
		TCF-1	↓↓↓	↓	↓↓↓	
		CXCR5	unchanged	+	+	
		Tim-3	++	+	+	
		PD-1	++	+	+	
		CTLA-4	+	+	++	
		in vitro killing	+++	+	n.d.	
adoptive transfer of CD4 ⁺ T cells; acute deletion (<i>Cd4-Cre-ERT2</i>)		proliferation but no long-term persistence	proliferation and long-term persistence; induction of ANAs	proliferation but no long-term persistence		
adoptive transfer of CD8 ⁺ OTI T cells in the B16-OVA melanoma model		enhanced anti-tumor response	enhanced anti-tumor response		n.d.	

Supplementary Table 1: Summary of Roquin and Regnase knock out phenotypes in different cell types and experimental settings. n.d.: not determined, EM: effector memory, CM: central memory, GC: germinal center, ANA: anti-nuclear antibody.

Name	Sequence (5' - 3')	UPL probe
<i>Icos</i> for	AACCTTAGTGGAGGATATTGCAT	33
<i>Icos</i> rev	CTACGGGTAGCCAGAGCTTC	33
<i>Tnfrsf4</i> for	GCTTGGAGTTGACTGTGTTCC	79
<i>Tnfrsf4</i> rev	GGGTCTGCTTCCAGATAAGG	79
<i>Ywhaz</i> for	CGCTAATAATGCAGTTACTGAGAGA	2
<i>Ywhaz</i> rev	TTGGAAGGCCGGTTAATT	2
<i>Zc3h12a</i> for	GAAGCAATGTGGCCATGAG	76
<i>Zc3h12a</i> rev	CCTCGCTCCAGAAACCAG	76

Supplementary Table 2: DNA oligonucleotides and probes from the Universal Probe Library (UPL) system (Roche) used for RT-qPCR analyses.