

Supplementary Table	Gene	KO/WT mRNA ratio	
expression of potential miR21 target genes in Bcl6KO Tregs compared to WT Tregs	Satb1	0.44 *	
	Spry1	0.47 *	
	Krit1	0.54	
	Il12a	0.57	
	Kcna3	0.57	
	Plag1	0.57	
	Spry2	0.57	
	Tgfb1	0.57	
	Microarray expression levels of the indicated genes were normalized to average gapdh + tubb5 + RPL35 expression levels for WT and KO then gene ratios calculated	Arhgap24	0.62
		Cntfr	0.62
		Matn2	0.62
		Nfib	0.62
		Ntf3	0.62
Pitx2		0.62	
Sox2		0.62	
Xkr6		0.62	
Bcl2		0.66	
Dazl		0.66	
Pdcd4		0.66	
Reck		0.66	
Tiam1		0.66	
Aspn		0.71	
Btg2		0.71	
Pik3r1		0.71	
Psrc1		0.71	
Rasgrp1	0.71		
Rhob	0.71		
Brd1	0.76		
Fasl	0.76		
Jag1	0.76		
Pcbp2	0.76		
Tbfr2	0.76		
Yod1	0.76		
Hnrpu	0.81		
Rnf103	0.81		
Stag2	0.81		
Zcchc3	0.81		
Ski	0.87		
Spg20	0.87		
Wwp1	0.87		
Cbx4	0.93		
Mrpl9	0.93		
Pcbp1	0.93		
Ppara	0.93		
Pten	0.93		
Smad7	0.93		
Bmpr2	1		
Cdc25a	1.1		
Mtap	1.1		
Apaf1	1.2		
Peli1	1.2		
Ccl1	1.4		
Map2k3	1.5		
Casp3	1.8		
mRNA expression data from: Sawant et al (2012) J.Immunol. 189, 4759			
miR21 targets from: Hatley et al (2010) Cancer Cell 18, 282			
Lu et al (2009) J.Immunol. 182, 4994			
*asterisks show gene decreased more than 2-fold in KO Tregs			