

SUPPLEMENTAL MATERIAL

Weiss et al., <http://www.jem.org/cgi/content/full/jem.20120914/DC1>

**Table S1.** Methylation pattern of the T reg cell-specific demethylated region in different T reg cell populations

CPG-ID	CD4 <sup>+</sup> Foxp3- GFP <sup>-</sup>	Total T reg cells Foxp3-GFP <sup>+</sup>	Nrp1 <sup>+</sup> T reg cells Foxp3-GFP <sup>+</sup>	Nrp1 <sup>-</sup> T reg cells Foxp3- GFP <sup>+</sup>	OVA-iT reg cells Foxp3-GFP <sup>+</sup>
38	100	42	0	38	NA
74	100	41	0	55	37
90	100	31	0	59	32
124	97	28	0	41	22
151	94	10	0	39	20
156	97	12	0	38	21
205	95	18	0	41	32
224	94	0	0	38	40
228	100	54	0	49	42
236	100	46	0	41	42
298	94	9	0	17	28

**Table S2.** List of the PCR primers used for qPCR in Figs 1, 3, 4, 7, and 8

Gene	Forward	Reverse
Dapl1	5' -GCAGTGAAAGCTGGAGGGATGCG-3'	5' -TGTGCCGTGTGAACTGTCGCTG-3'
Foxp3	5' -GGCCCTTCTCCAGGACAGA-3'	5' -GCTGATCATGGCTGGGTGTG-3'
Helios	5' -ACACCTCAGGACCCATTCTG-3'	5' -TCCATGCTGACATTCTGGAG-3'
Igfbp4	5' -TTCATCATCCCCATTCCAAAC-3'	5' -ACCCCTGTCTTCCGATCCA-3'
Nrp1	5' -GACAAATGTGGCGGGACCATA-3'	5' -TGGATTAGCCATTCACACTTCTC-3'
Swap70	5' -TGACACAGTTATGGGAAAGG-3'	5' -ACGAACTGCTCAAAGCCATT-3'
Beta Actin	5' -GTCCACCTTCCAGCAGATGT-3'	5' -AAGGGTGTAACGCAGCTCAG-3'