

Supplementary Material

Molecular markers distinguishing T cell subtypes with TSDR strand-bias methylation

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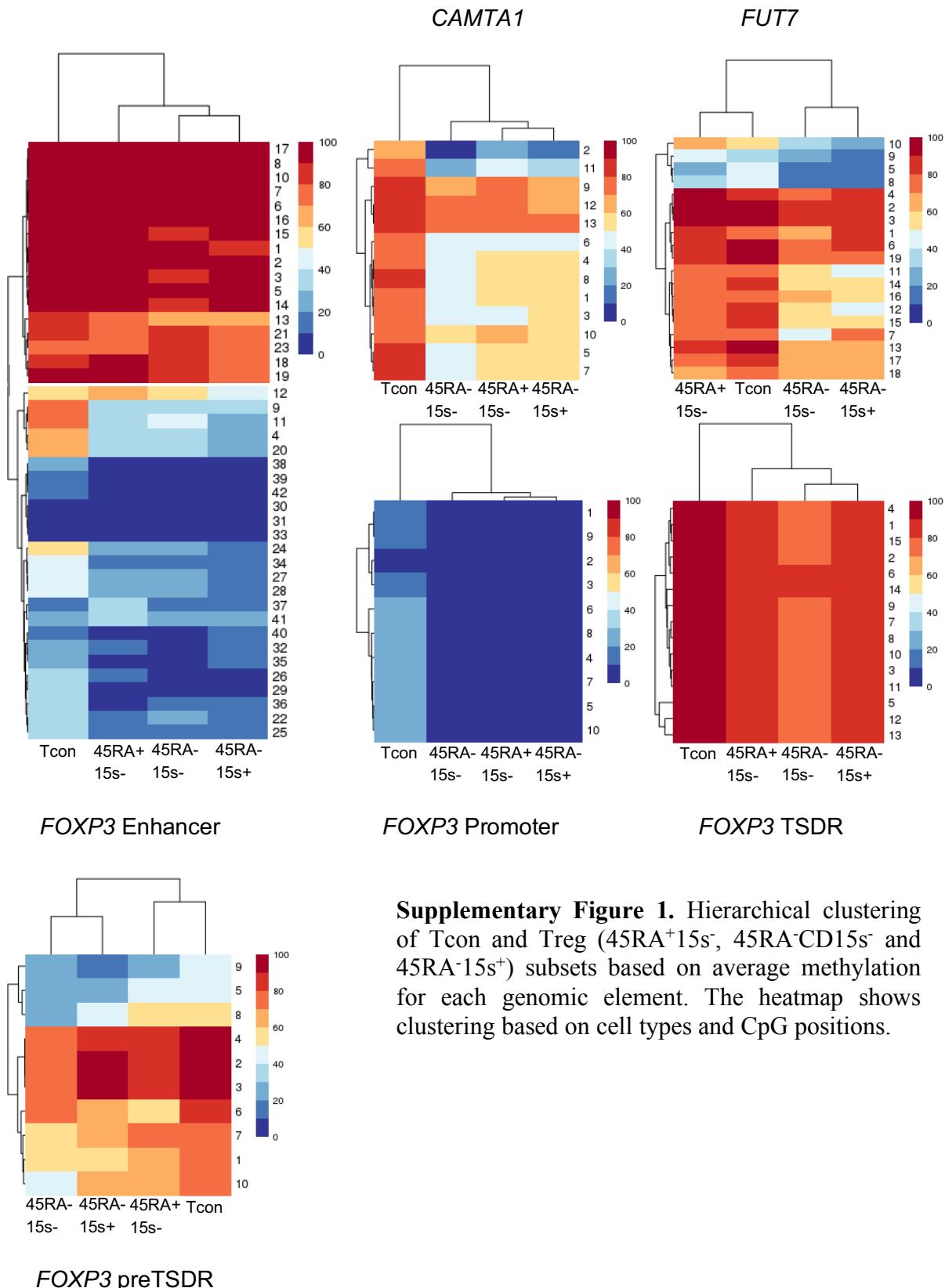
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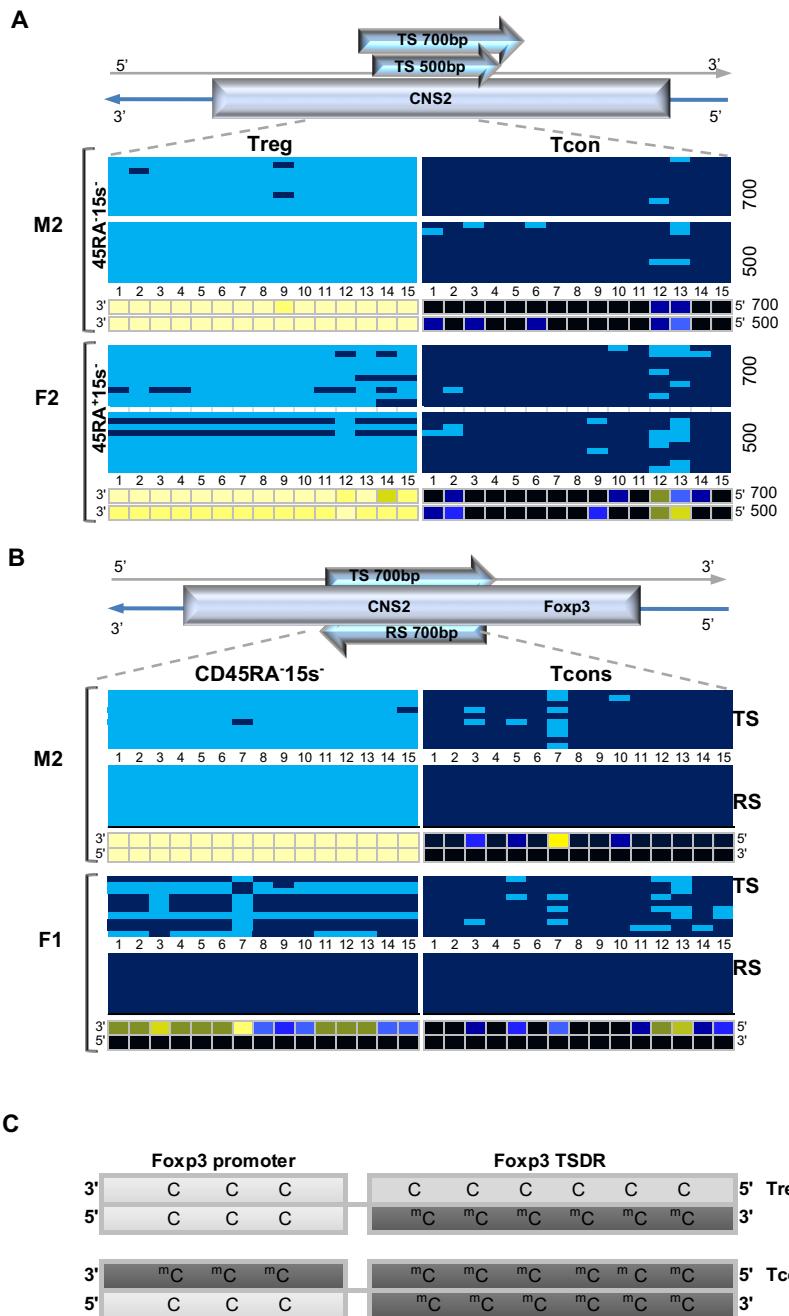
‡ These authors share senior authorship

***Correspondence:**

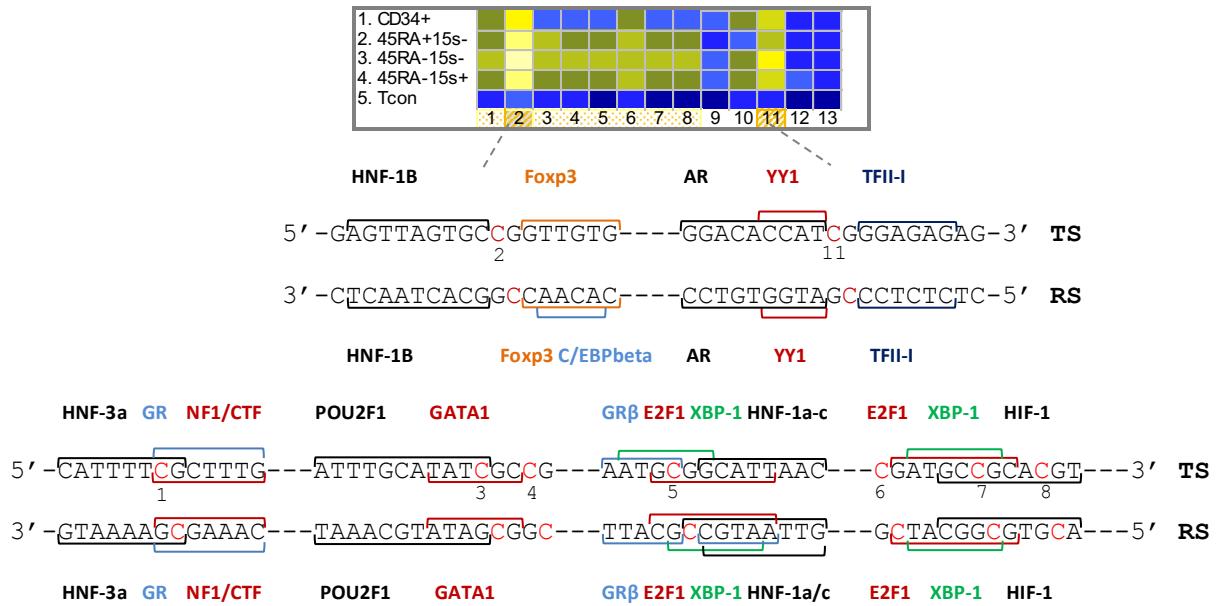
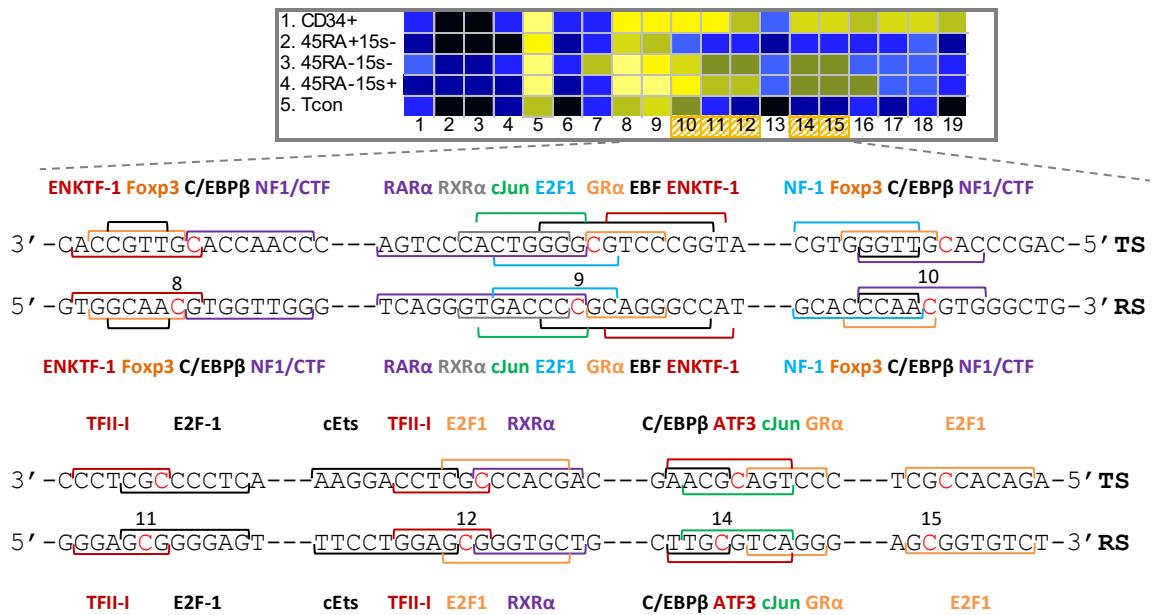
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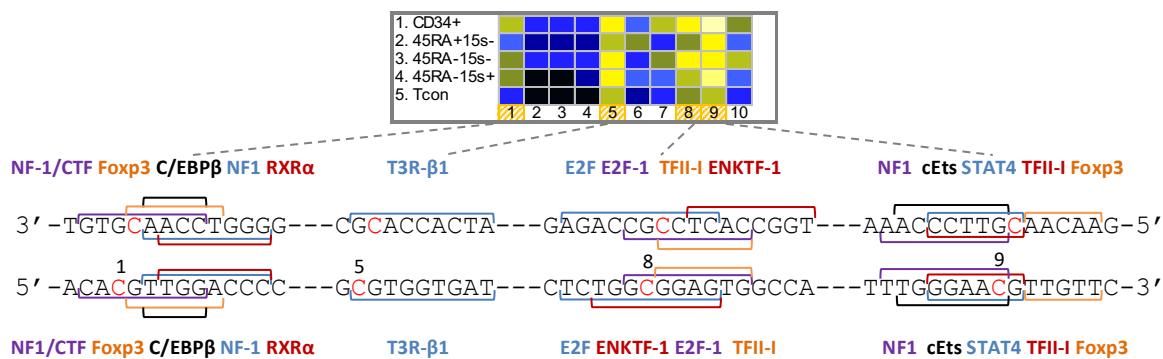
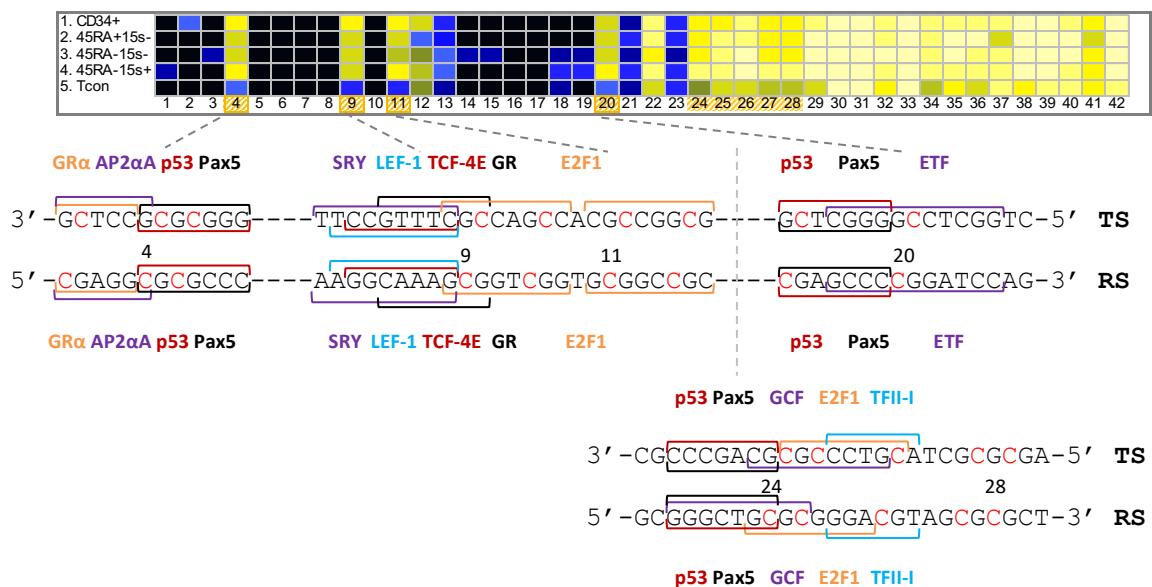
Supplementary Figure 1. Hierarchical clustering of Tcon and Treg ($45RA^+15s^-$, $45RA^-CD15s^-$ and $45RA^-15s^+$) subsets based on average methylation for each genomic element. The heatmap shows clustering based on cell types and CpG positions.



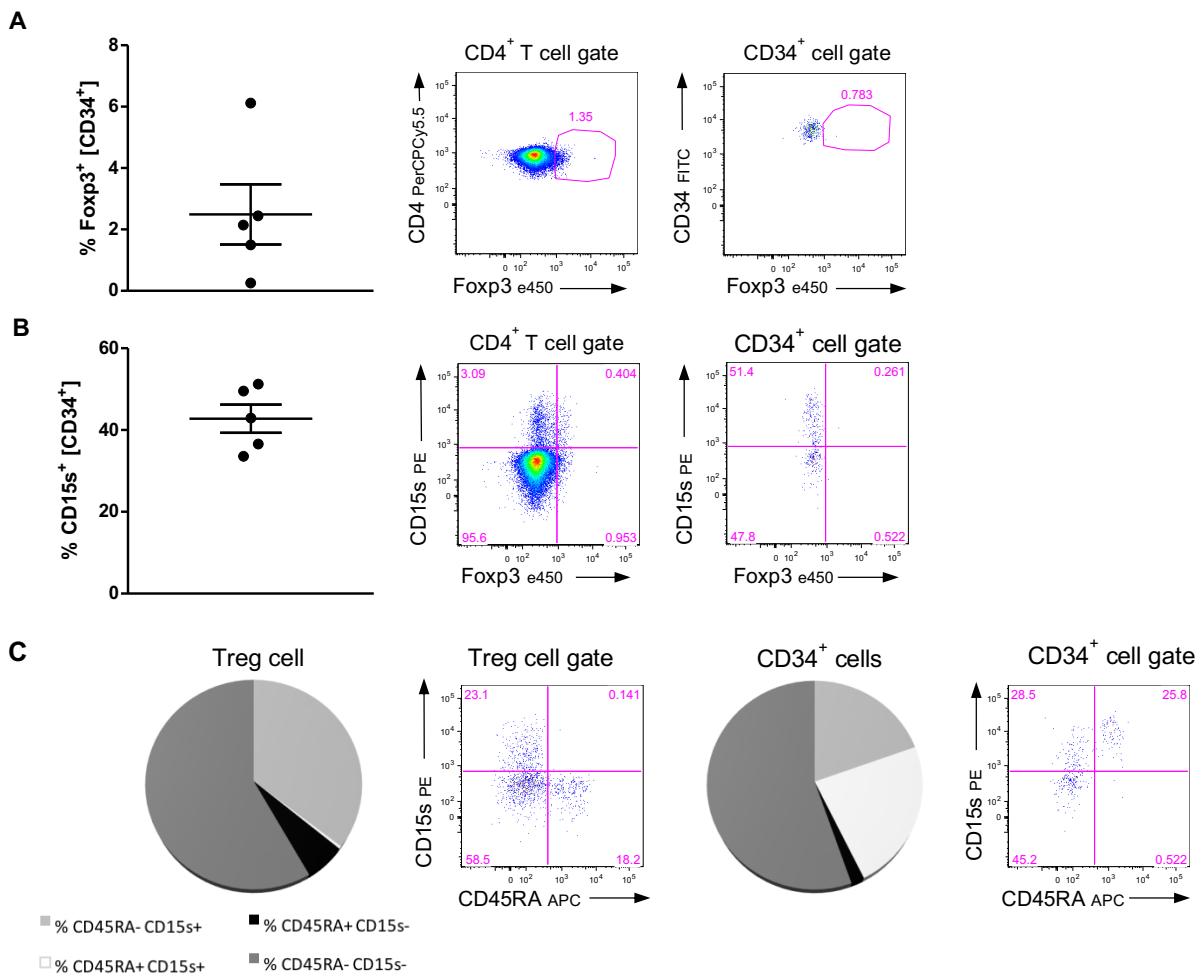
Supplementary Figure 2. Strand-specific methylation bias within the *FOXP3* TSDR region. (A). Methylation patterns of the top strand (TS) within TSDR of Treg and Tcon cells in ‘conventional’ (M2, top panel) and ‘unconventional’ (F2, lower panel) donors based on two amplicons, TS700bp and TS500bp. (B). Methylation patterns of the top strand (TS, upper panels) and reverse strand (RS, lower panels) within TSDR of CD45RA⁻CD15s⁻ and Tcon cells of donors M2 and F1. CpGs 1-15 are numbered relative to the 5'-3' direction of the coding (reverse) strand. Each horizontal line represents DNA from one cell with unmodified C in light blue and mC in dark blue. Methylation percentage for each CpG site was calculated based on the number of mCs in a total of 10 and summarised in panels below. (C). Schematic presentation of ‘unconventional’ strand-bias methylation pattern in *FOXP3* promoter and TSDR of Treg and Tcon cells.

A**CAMTA****B****FUT7**

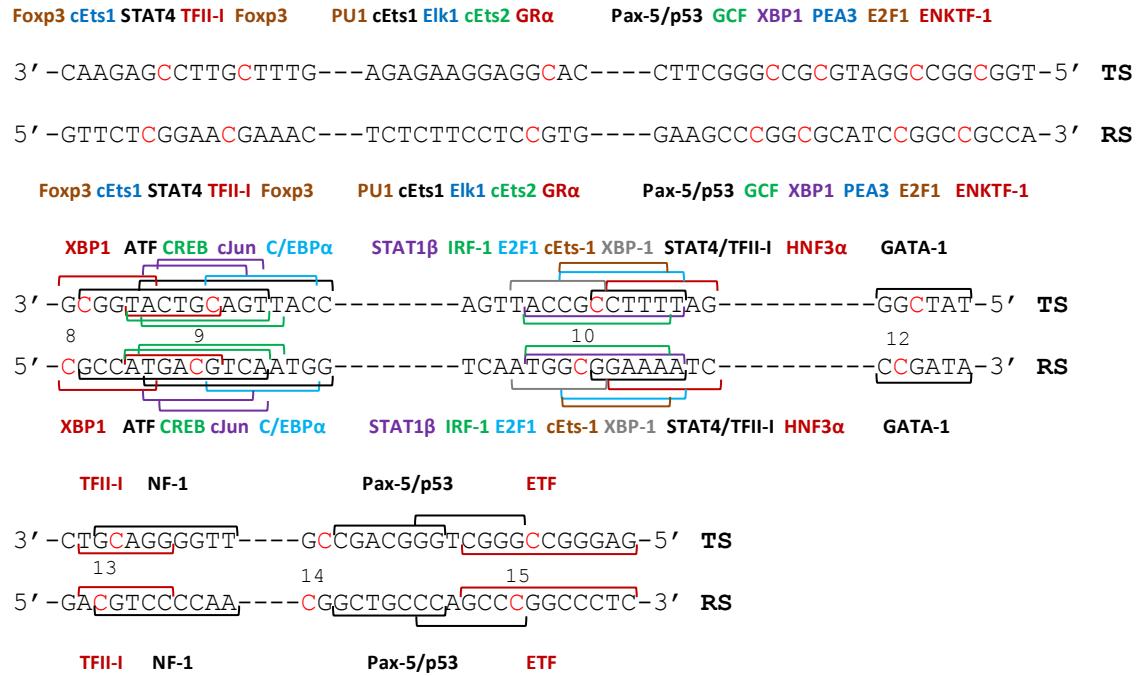
Supplementary Figure 3. Predicted TF binding sites containing or immediately adjacent to CpGs differentially methylated in *CAMTA1* and *FUT7* promoter of CD34+ cells and T lymphocyte subsets. Average methylation patterns for *CAMTA1* (A and Figure 2C) and *FUT7* (B and Figure 3C) are presented to indicate hypomethylated CpG sites. Coding strands of *CAMTA1* (TS) and *FUT7* (RS) are shown in the 5'-3' direction, Cs within 5'-CpG-3' are in red and numbered relative to the 5'-3' direction of the coding strand, TF recognition sequences are colour-coordinated with respective TFs. TS – top strand, RS – reverse strand

A**Foxp3 preTSDR****B****Foxp3 Enhancer**

Supplementary Figure 4. Predicted TF binding sites containing or immediately adjacent to CpGs differentially methylated in *FOXP3* preTSDR and enhancer regions of CD34⁺ cells and T lymphocyte subsets. Average methylation patterns for *FOXP3* preTSDR (A and Figure 5C) and enhancer (B and Figure 4C) are presented to indicate hypomethylated CpG sites. Coding (reverse) strands (RS) are shown in the 5'-3' direction, Cs within 5'-CpG-3' are in red and numbered relative to the 5'-3' direction of the coding strand, TF recognition sequences are colour-coordinated with respective TFs.



Supplementary Figure 5. Phenotypic analysis of CD34⁺ cells by flow cytometry. Whole blood samples from donors M3, M6, F1, F2 and F4 were surface stained for CD34, CD3, CD4, CD25, CD127, CD45RA and CD15s. Following fixation and permeabilization, cells were incubated with anti-Foxp3 antibody. (A). The frequency of Foxp3⁺ cells was determined within the CD34⁺ population. Dot plots from a representative donor (M6) are shown, illustrating the expression of Foxp3 within CD4 T cells and CD34⁺ cells. (B). The frequency of CD15s⁺ cells was determined within the CD34⁺ population. Dot plots from a representative donor (M6) are shown, illustrating the co-expression of CD15s and Foxp3 within CD4 T cells and CD34⁺ cells. In the scatter plots, each symbol represents an individual (n=5), mean values are shown as horizontal lines and error bars represent the standard error of the mean. (C). The frequency of CD45RA/CD15s subsets was determined within Treg and CD34⁺ cells. Pie charts show the average frequency of each subset within Treg and CD34⁺ cells (n=5). Dot plots from a representative donor (M6) are shown, illustrating the distribution of CD45RA/CD15s subsets within Treg and CD34⁺ cells.



Supplementary Figure 6. Differential binding of specific TFs to cytosines within CpG sites located on FOXP3 TSDR. While mapping to both strands, the predicted DNA binding sites of some TFs did not contain a cytosine (C) within their recognition sequences on one of the strands. Coding (reverse) strand (RS) is shown in the 5'-3' direction, Cs within 5'-CpG-3' are in red and numbered relative to 5'-3' direction of the coding strand, TF recognition sequences are colour-coordinated with respective TFs.

Primer Name	Primer Sequence, 5'-3'	Amplicon size, bp	Strand
CAMTA Fw CAMTA Rev	GCGCCATGG AATAAGTAAAGAAAATTAGTTGGTAATAAAAAGGG GCGATGCAT TATAACATCTCATCTAAATCAACCTATAACAAACACC	470	top (+) top (+)
1FUT7 Fw 1FUT7 Rev	GCGCCATGG TAATTITAATATTTGGGAGGTTGAAATTGTGG GCGATGCATT CACCTAAACCCCAACCAACCAACACC	500	reverse (+) reverse (+)
2FUT7 Fw 2FUT7 Rev	GCGCCATGG GTGGTGGGTGAAGAGGGTGTGG GCGATGCAT CCCTTAATATCTCCCACAAAAACC	454	reverse (+) reverse (+)
Foxp3 Enhancer Fw Foxp3 Enhancer Rev	GCGCCATGG GGTTGTTGGTTAGAAAGTGT GCGATGCAT AAAATTACCTAACCCCCACATT	497	reverse (+) reverse (+)
Foxp3 Promoter RS Fw Foxp3 Promoter RS Rev	GCGCCA TGG TTGTTGTTCGGGTGGTTGTGATTATTTAG GCGATGCAT ACAAAAAAATCAACCTAACTT	451	reverse (+) reverse (+)
Foxp3 Promoter TS Fw Foxp3 Promoter TS Rev	GCGCCATGG TGTTATTTTTAATGGGTTATTTGGTAGGG GCGATGCAT CAAATTCTCTTCCCCAAAACC	582	top (-) top (-)
Foxp3 preTSDR Fw Foxp3 preTSDR Rev	GCGCCATGG TTTTTAGTTAGAAGGGGAAGGGGAATTGGG GCGATGCAT AAAACCCCTAAATCTCAAATAATATAACACAAAAC	700	reverse (+) reverse (+)
Foxp3 TSDR TS700 Fw Foxp3 TSDR TS700 Rev	GCGCCA TGG TATGTGGGATGGTTGATTAGTAAAGTATA GCGATGCAT TTAACACCAAAACACTAACCTACACATAC	699	top (-) top (-)
Foxp3 TSDR TS500 Fw Foxp3 TSDR TS500 Rev	GCGCCA TGG GGATATGGAGATGATTGTTGGGGTAGAGG GCGATGCAT CTAACACTCTCAAACCTAAATCCTC	474	top (-) top (-)
Foxp3 TSDR RS Fw Foxp3 TSDR RS Rev	GCGCCA TGG TTAGGGATATTGGTTATATATTGAGATTGG GCGATGCAT CCCCAACACATATAAAAACCTAACTCAACAAAAC	700	reverse (+) reverse (+)

Supplementary Table 1. Bisulphite DNA strand-specific primers used for amplification of *CAMTA1*, *FUT7* and *FOXP3* gene regions for detection of methylation status. Coding strand is denoted as “+” (sense) and template strand is denoted as “-“ (antisense). Nucleotides comprising restriction sites NcoI (forward primers, Fw) and NsiI (reverse primers, Rev) used for directional cloning are in bold italics

Gene region	Position, 5'-3'	Primer position, 5'-3'	Amplicon size, bp	Strand
CAMTA1, Ch1				
CAMTA1 intronic region	6,965,651-6,966,120	Fw 6,965,651-6,965,689 Rev 6,966,120-6,966,083	470	+
FUT7, Ch9				
FUT7 promoter 1	137,033,577-137,033,078	Fw 137,033,577-137,033,551 Rev 137,033,078-137,033,105	500	+
FUT7 promoter 2	137,033,153-137,032,701	Fw 137,033,153-137,033,131 Rev 137,032,701-137,033,725	454	+
FOXP3, ChX				
FOXP3 Enhancer	49,270,698-49,270,202	Fw 49,270,698-49,270,676 Rev 49,270,202-49,270,228	497	+
FOXP3 Promoter RS	49,265,128-49,264,678	Fw 49,265,128-49,265,092 Rev 49,264,678-49,264,701	451	+
FOXP3 Promoter TS	49,264,607-49,265,188	Fw 49,264,607-49,264,643 Rev 49,265,188-49,265,156	582	-
FOXP3 preTSDR	49,261,770-49,261,071	Fw 49,261,770-49,261,734 Rev 49,261,071-49,261,109	700	+
FOXP3 TSDR TS700	49,260,531-49,261,229	Fw 49,260,531-49,260,563 Rev 49,261,229-49,261,197	699	-
FOXP3 TSDR TS500	49,260,579-49,261,052	Fw 49,260,579-49,260,611 Rev 49,261,052-49,261,019	474	-
FOXP3 TSDR RS	49,261,221-49,260,522	Fw 49,261,221-49,261,186 Rev 49,260,522-49,260,559	700	+

Supplementary Table 2. *CAMTA1*, *FUT7* and *FOXP3* gene regions used in the study of methylation status of the five cell populations. Fw – forward primer, Rev – reverse primer, TS-top strand, RS-reverse strand, “+” - sense (coding) strand, “-“ – antisense (template) strand.

CAMTA																						
Donor		M1					M2					M3					M4					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	N/A	25	5	100	70	N/A	50	5	10	80	N/A	35	5	10	70	25	30	25	30	70	
2	28	N/A	10	15	70	65	N/A	40	0	20	70	N/A	35	5	5	75	15	40	0	5	70	
3	57	N/A	5	20	75	90	N/A	45	0	20	85	N/A	35	10	30	80	55	45	0	30	65	
4	60	N/A	15	20	100	85	N/A	45	0	15	95	N/A	45	10	20	70	55	45	0	30	90	
5	89	N/A	25	20	100	100	N/A	45	0	25	100	N/A	50	10	30	90	60	45	5	30	100	
6	152	N/A	10	15	95	75	N/A	35	0	5	85	N/A	15	5	15	85	40	25	0	30	65	
7	158	N/A	25	25	80	100	N/A	70	5	15	100	N/A	40	5	30	95	50	35	0	30	80	
8	162	N/A	25	20	100	95	N/A	30	0	10	90	N/A	40	5	30	80	60	35	10	40	85	
9	237	N/A	70	55	100	100	N/A	85	55	90	95	N/A	70	70	95	90	85	55	25	25	95	
10	294	N/A	80	65	80	85	N/A	90	45	55	90	N/A	70	65	85	80	60	45	15	25	85	
11	317	N/A	80	60	80	85	N/A	95	45	90	95	N/A	90	55	90	90	55	65	15	10	90	
12	352	N/A	95	90	100	95	N/A	95	95	85	100	N/A	95	95	90	95	90	65	15	25	80	
13	354	N/A	85	95	100	100	N/A	100	95	100	100	N/A	100	100	90	90	65	15	25	100		
Donor		M5					M6					F1					F2					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	25	100	25	10	55	40	50	90	95	95	75	20	60	20	80	30	45	50	20	45	
2	28	15	5	0	40	0	10	5	5	70	70	10	15	0	70	25	20	25	20	35		
3	57	40	90	20	5	55	40	55	90	95	90	90	10	65	20	75	30	35	40	35		
4	60	40	100	20	5	55	40	50	90	95	90	90	10	65	10	85	30	50	35	25		
5	89	40	100	20	5	55	40	50	90	95	100	90	15	70	10	95	30	50	50	30		
6	152	40	100	20	5	55	40	50	90	95	85	85	15	65	10	60	30	50	25	35		
7	158	40	100	20	5	60	40	55	95	95	95	90	25	75	10	90	35	55	45	20		
8	162	70	100	20	5	60	40	55	95	100	90	90	20	65	10	85	35	50	40	25		
9	237	50	100	25	10	60	40	75	95	100	100	90	40	70	25	80	25	55	50	25		
10	294	45	100	20	10	60	55	65	95	95	85	90	25	55	10	75	35	45	35	25		
11	317	50	5	20	0	50	25	35	5	10	70	65	30	30	10	75	35	25	30	45		
12	352	70	100	55	5	75	80	90	100	100	90	85	35	75	30	100	35	60	30	45		
13	354	70	100	55	10	55	75	80	90	100	80	75	30	75	30	85	40	65	40	50		
Donor		F3					F4					F5					mean all donors					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	80	85	40	85	100	65	55	80	95	90	100	95	100	100	80	55	53.6	44.1	52.3	75.9	
2	28	60	45	25	40	90	5	0	10	0	70	5	10	5	0	80	24.4	20.5	10	15	66.8	
3	57	75	80	70	85	95	65	55	85	95	95	95	90	100	100	75	61.3	49.5	45.5	53.6	76.8	
4	60	80	85	65	85	90	65	55	85	95	90	95	90	100	95	85	61.9	53.6	44.5	52.3	79.5	
5	89	80	85	65	85	100	65	55	85	95	100	95	95	100	100	90	62.5	55.9	46.8	55	88.2	
6	152	80	75	65	85	90	70	55	85	95	95	95	90	100	95	65	60	47.3	42.7	50	72.3	
7	158	75	85	75	85	100	100	70	55	80	95	95	95	100	100	90	61.9	58.2	48.6	51.4	85.9	
8	162	75	85	70	85	100	70	55	80	95	95	90	95	100	100	80	66.3	53.6	45.9	54.5	81.8	
9	237	70	80	80	85	95	70	55	95	95	100	100	95	100	100	85	66.3	70.9	65.5	68.2	84.1	
10	294	35	45	25	50	90	65	55	85	100	80	100	90	95	95	90	60.6	64.5	54.5	57.3	78.2	
11	317	45	50	20	40	85	5	5	15	0	95	10	10	10	70	36.3	44.5	27.7	33.6	77.3		
12	352	75	85	100	85	100	80	55	100	100	100	100	95	100	100	95	76.3	78.2	76.8	69.5	87.3	
13	354	75	85	95	85	100	80	55	100	100	95	100	95	100	100	95	75.6	78.2	78.2	71.8	86.4	
mean CpGs 1-8		37.8	36.2	13.9	35.8	74.1	62.9	59.7	63.6	58.2	82						56.7	49	41	48	78.4	
mean all CpGs		44.4	50.7	29.4	46.8	77.2	64	60.5	64.5	57.5	82.4						59.1	56	48.5	52.6	80	
Donor		mean M1-4, F2					mean M5,6, F1,3-5															
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5											
1	1	27.5	37	18	34	67	64.2	67.5	65.8	67.5	83.3											
2	28	20	29	9	24	63	25.8	13.3	10.8	7.5	70											
3	57	42.5	33	14	38	72	67.5	63.3	71.7	66.7	80.8											
4	60	42.5	40	13	38	76	68.3	65	70.8	64.2	82.5											
5	89	45	43	17	43	86	68.3	66.7	71.7	65	90											
6	152	35	27	9	33	69	68.3	64.2	70.8	64.2	75											
7	158	42.5	45	16	35	82	68.3	69.2	75.8	65	89.2											
8	162	47.5	36	15	41	78	72.5	68.3	71.7	65.8	85											
9	237	55	67	51	67	81	70	74.2	77.5	69.2	86.7											
10	294	47.5	66	45	54	76	65	63.3	62.5	60	80											
11	317	45	71	41	60	81	33.3	22.5	16.7	11.7	74.2											
12	352	62.5	82	65	69	84	80.8	75	86.7	70	90											
13	354	65	83	69	73	88	79.2	74.2	85.8	70.8	85											

FUT7

Donor		M1					M2					M3					M4					M5					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	N/A	90	35	45	75	N/A	65	55	90	70	N/A	75	75	60	45	70	55	N/A	90	80	35	100	90	90	85	
2	65	N/A	100	80	55	90	N/A	85	90	90	95	N/A	75	100	80	80	90	85	N/A	100	100	80	100	100	90	100	
3	95	N/A	100	95	70	80	N/A	95	85	100	90	N/A	90	95	70	75	100	85	N/A	100	100	85	95	100	90	100	
4	167	N/A	100	85	45	65	N/A	75	70	90	95	N/A	95	90	70	80	70	85	N/A	100	90	55	100	100	90	95	
5	250	N/A	75	45	10	45	N/A	50	10	0	50	N/A	60	40	25	50	30	20	N/A	20	70	5	5	0	0	65	
6	266	N/A	90	80	20	90	N/A	70	80	95	95	N/A	90	85	70	90	80	60	N/A	100	90	55	95	80	90	95	
7	288	N/A	75	40	15	60	N/A	70	45	75	70	N/A	70	70	65	60	75	50	N/A	80	85	55	85	70	90	90	
8	306	N/A	90	30	10	55	N/A	70	15	10	70	N/A	75	40	25	50	25	15	N/A	20	80	15	5	0	20	50	
9	387	N/A	65	50	20	30	N/A	60	20	10	60	N/A	50	15	5	45	0	15	35	10	50	0	10	5	30	20	
10	419	N/A	70	40	10	30	N/A	75	10	0	70	N/A	60	10	10	55	0	20	25	10	60	10	90	100	70	55	
11	480	N/A	100	95	55	80	N/A	95	75	80	90	N/A	100	50	35	100	0	30	25	30	85	45	90	90	70	30	
12	524	N/A	100	85	60	80	N/A	100	85	90	95	N/A	90	75	20	95	65	25	70	30	95	70	20	5	5	10	
13	543	N/A	100	95	45	85	N/A	100	80	85	90	N/A	95	40	20	100	60	25	65	35	100	80	100	20	95	95	
14	549	N/A	95	90	35	80	N/A	100	90	80	95	N/A	100	70	30	100	45	25	35	45	95	45	95	20	55	40	
15	642	N/A	100	90	60	95	N/A	100	80	100	100	N/A	100	75	25	100	45	20	60	25	85	50	35	15	10	10	
16	667	N/A	100	80	60	80	N/A	100	60	95	90	N/A	100	55	30	75	45	15	50	25	90	45	95	100	60	50	
17	707	N/A	95	100	80	85	N/A	100	80	95	95	N/A	100	60	30	70	25	10	50	55	95	40	95	100	90	70	
18	729	N/A	85	95	85	85	N/A	65	75	85	65	N/A	80	55	10	95	25	10	45	45	80	35	100	100	100	100	
19	731	N/A	100	100	90	N/A	100	90	100	100	N/A	75	35	10	95	50	10	60	60	95	45	100	100	100	100	100	
Donor		M6					F1					F2					F3					mean all donors					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	N/A	N/A	N/A	N/A	N/A	N/A	30	N/A	80	95	100	100	95	100	95	100	85	100	95	100	95	73.8	83.6	67.1	81.4	78.8
2	65	N/A	N/A	N/A	N/A	N/A	N/A	30	N/A	95	95	100	100	100	100	100	100	95	100	100	100	100	91.3	92.1	85	87.9	95
3	95	N/A	N/A	N/A	N/A	N/A	N/A	30	N/A	85	90	100	100	100	100	100	100	95	100	100	100	100	93.8	95	85.8	90	91.3
4	167	N/A	N/A	N/A	N/A	N/A	N/A	30	N/A	90	90	100	100	95	100	100	100	85	100	95	100	100	78.8	93.6	80	84.3	88.8
5	250	N/A	N/A	N/A	N/A	N/A	N/A	0	N/A	10	0	0	15	25	5	20	0	15	10	55	13.8	30	17.9	12.9	43.8		
6	266	N/A	N/A	N/A	N/A	N/A	N/A	30	N/A	90	90	100	100	100	100	100	100	95	100	100	100	100	80	86.4	78.6	82.1	93.8
7	288	N/A	N/A	N/A	N/A	N/A	N/A	30	N/A	70	90	100	10	80	90	75	100	80	90	80	80	73.8	78.6	49.3	70.7	75.6	
8	306	N/A	N/A	N/A	N/A	N/A	N/A	10	N/A	15	5	10	15	20	10	65	0	15	10	60	27.5	37.9	17.9	16.4	48.8		
9	387	20	25	20	10	30	5	90	0	0	30	50	35	45	55	60	55	60	25	5	10	21.7	45.6	23.9	16.1	37.2	
10	419	30	35	45	15	70	5	100	0	45	55	55	25	70	70	65	75	90	45	30	80	29.2	62.8	38.3	28.9	60	
11	480	20	40	60	20	95	5	100	0	0	65	45	55	80	60	60	45	75	45	25	95	26.7	76.1	57.8	41.7	77.8	
12	524	10	45	55	10	85	0	100	25	90	90	60	85	90	80	85	65	85	45	55	90	45	72.2	59.4	48.9	80.6	
13	543	60	60	60	35	100	10	100	25	100	85	80	85	90	95	85	100	80	70	95	63.3	85	61.7	64.4	92.8		
14	549	15	45	60	20	100	0	100	30	100	80	70	85	90	65	85	65	65	40	55	100	40	78.9	58.3	53.9	86.1	
15	642	25	40	50	25	90	0	100	30	100	85	60	90	80	85	80	60	80	50	35	90	40	73.9	58.9	51.7	81.7	
16	667	35	30	75	20	100	0	100	20	45	75	55	60	80	75	70	65	85	60	80	90	40.8	76.1	64.4	54.4	80	
17	707	35	35	70	25	95	0	100	5	100	90	60	80	60	60	70	65	80	75	95	95	37.5	77.2	66.7	70	85	
18	729	55	55	65	40	90	0	100	5	100	80	40	50	75	50	25	50	80	90	40	85	34.2	69.4	67.2	61.7	78.3	
19	731	50	60	70	40	90	0	100	20	100	85	65	80	95	90	75	85	100	100	95	90	49.2	80.6	74.4	77.2	91.1	
																	mean CpGs 10-12,14,15					38.2	72.8	54.5	45	77.2	
																	mean CpGs 8-19					37.9	69.6	54.1	48.8	75	

Foxp3 Enhancer

Donor		M1				M2				M3				M4				
CpG	nt	#2	#3	#4	#5	#2	#3	#4	#5	#2	#3	#4	#5	#1	#2	#3	#4	#5
1	1	100	95	75	95	100	100	100	95	100	100	100	100	100	95	85	85	85
2	8	100	80	80	100	100	100	100	100	100	100	100	100	0	100	100	100	95
3	14	65	65	70	85	90	80	100	80	95	100	90	95	100	95	100	95	95
4	19	95	90	25	95	75	25	0	85	5	5	10	45	20	35	25	20	85
5	21	80	60	80	100	100	95	100	95	100	100	100	100	100	100	100	100	95
6	29	75	70	90	100	100	100	100	100	100	100	90	100	100	100	100	100	95
7	33	70	90	70	95	100	90	100	95	95	100	100	95	100	100	100	100	90
8	35	100	70	80	95	100	100	100	100	100	100	100	100	100	100	100	100	100
9	67	95	95	25	100	70	25	0	100	10	5	10	55	0	35	25	35	85
10	71	95	95	90	100	100	90	100	100	100	95	100	95	100	100	100	100	95
11	76	85	90	25	100	75	60	0	95	10	10	0	55	0	35	25	35	80
12	80	90	85	40	80	0	10	0	20	65	20	15	35	0	95	95	95	95
13	83	85	75	75	100	40	35	55	95	60	55	60	75	70	95	95	100	80
14	86	75	55	80	95	100	75	100	100	95	100	90	100	100	95	100	100	100
15	89	100	90	80	95	100	5	100	100	95	100	100	95	100	95	100	100	95
16	93	100	85	80	100	90	100	100	100	100	100	95	95	100	95	100	100	100
17	101	100	90	80	95	100	100	100	100	100	100	95	100	100	95	100	100	100
18	113	75	50	90	90	80	85	100	80	100	100	90	85	100	95	100	100	80
19	125	70	60	70	90	90	75	100	90	95	100	90	90	100	95	100	100	95
20	132	55	75	15	80	70	5	0	75	0	0	0	30	25	35	40	40	75
21	168	45	60	80	75	100	80	95	75	100	100	100	80	80	100	100	95	100
22	180	20	40	0	30	10	0	0	60	0	0	0	30	0	30	20	35	50
23	183	50	75	65	65	100	85	95	100	100	100	90	85	100	30	80	85	55
24	190	55	40	0	65	45	5	0	75	0	0	0	40	20	30	20	35	55
25	192	20	25	0	25	40	10	0	35	0	0	0	40	20	30	20	35	50
26	197	20	15	0	25	45	0	0	50	0	0	0	25	10	30	10	20	50
27	202	55	40	0	60	35	15	0	60	0	0	0	35	20	30	20	35	50
28	204	45	35	0	60	65	10	0	50	0	0	0	30	20	30	20	40	55
29	215	25	5	10	35	5	5	0	35	0	5	0	10	25	0	10	25	50
30	233	0	45	0	0	0	0	0	0	0	0	5	5	0	0	0	0	0
31	236	0	0	0	0	20	0	0	5	0	0	5	0	0	5	0	0	5
32	247	5	5	0	0	40	0	0	20	5	0	0	15	0	0	5	15	55
33	251	5	0	0	0	0	0	0	5	0	0	0	0	0	10	0	0	0
34	258	45	35	0	60	65	5	0	50	0	5	10	35	0	5	10	20	40
35	266	15	0	0	15	10	0	0	20	0	5	10	25	0	10	10	15	25
36	274	0	35	5	30	20	0	0	40	0	0	0	25	0	10	15	15	25
37	284	15	0	0	15	5	0	0	20	0	0	10	0	0	50	60	50	30
38	286	20	5	0	30	0	10	5	20	0	0	0	0	0	0	0	0	20
39	296	5	0	0	15	5	0	0	25	0	0	0	0	0	0	5	0	20
40	298	20	0	0	5	5	0	0	25	0	0	0	10	0	5	15	15	30
41	306	20	0	0	20	0	5	0	20	0	0	0	10	0	65	70	35	45
42	309	20	5	0	0	0	0	0	15	0	0	0	5	0	5	5	0	15

Foxp3 Enhancer

Donor		F1					F2					mean all donors				
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5
1	1	95	90	85	90	95	90	90	95	80	80	95	96.3	93.3	88.3	91.7
2	8	100	100	95	90	95	100	100	95	80	80	66.8	100	95	91.7	95
3	14	100	100	95	90	90	100	100	100	100	100	100	90.9	90	90.9	90.9
4	19	35	20	30	55	55	25	0	35	40	35	26.8	38.3	35	25	66.7
5	21	100	100	95	90	100	100	100	100	100	100	100	96.7	91.7	95	98.3
6	29	100	100	100	100	95	100	100	100	100	100	100	95.9	95	96.7	98.3
7	33	100	100	95	90	100	100	100	100	100	100	100	94.1	95.9	93.3	95.9
8	35	100	100	100	100	95	100	100	100	100	100	100	100	95	96.7	98.3
9	67	35	20	25	55	55	35	5	35	60	80	23.3	39.1	35	30.9	79.1
10	71	100	100	100	100	100	100	100	100	100	100	100	99.1	96.7	98.3	98.3
11	76	45	20	25	55	60	35	5	35	60	80	26.8	38.3	40.9	29.1	78.3
12	80	5	15	0	5	0	100	100	95	95	100	35	60.9	50.9	41.7	55
13	83	70	60	65	5	85	100	100	90	95	100	80	73.3	69.1	65	89.1
14	86	100	100	100	100	100	100	100	95	100	100	100	94.1	87.4	95	99.1
15	89	100	100	100	100	100	100	100	95	80	80	100	98.3	81.7	93.3	94.1
16	93	100	95	100	100	95	100	100	95	100	100	100	96.7	96.7	95.9	98.3
17	101	100	100	100	100	95	100	100	100	100	100	100	99.1	98.3	95.9	98.3
18	113	100	100	100	0	90	100	100	100	100	100	100	91.7	89.1	80	85.4
19	125	100	100	100	0	95	100	100	95	100	100	100	91.7	88.3	76.7	93.3
20	132	45	20	25	55	60	30	5	40	65	75	33.3	30.9	30.9	29.1	65.9
21	168	100	15	100	0	85	90	100	95	100	90	90	76.7	89.1	78.3	84.1
22	180	30	20	20	45	45	15	0	45	35	15	15	13.3	20.9	19.1	38.3
23	183	30	85	100	65	70	100	100	75	75	100	76.8	77.4	85.9	79.1	79.1
24	190	45	20	20	45	55	15	0	45	35	35	26.8	25	21.7	19.1	54.1
25	192	40	20	20	45	45	15	0	45	35	35	25	18.3	20	19.1	38.3
26	197	20	5	20	10	35	15	0	0	10	20	15	16.7	7.4	6.7	34.1
27	202	45	20	20	50	45	15	0	45	25	15	26.8	23.3	23.3	18.3	44.1
28	204	45	20	20	50	45	15	0	45	25	15	26.8	26.7	21.6	19.1	42.4
29	215	15	10	20	10	35	15	0	10	10	35	18.3	6.7	9.1	9.1	33.3
30	233	0	5	0	10	25	0	0	0	0	0	0	0.7	7.4	2.4	5
31	236	0	5	0	10	25	0	0	0	0	0	0	5	0	2.4	5.9
32	247	30	15	0	45	35	10	0	45	45	30	13.3	10.9	9.1	17.4	26.3
33	251	0	5	0	5	15	0	5	0	0	0	0	4.1	0	0.7	3.3
34	258	40	5	0	45	45	10	0	35	45	40	16.8	20	15	20	45
35	266	30	20	0	45	35	10	0	40	45	30	13.3	9.1	9.1	19.1	25
36	274	30	20	0	45	30	10	0	50	45	40	13.3	8.3	16.7	18.3	31.7
37	284	15	80	0	5	20	40	50	25	15	35	18.3	33.3	14.1	13.3	20
38	286	15	10	0	5	25	10	0	20	10	35	8.3	5	5.9	3.3	21.7
39	296	15	10	0	5	20	10	0	20	10	35	8.3	3.3	4.1	2.4	19.1
40	298	45	20	0	45	30	0	0	40	35	10	15	8.3	9.1	15.9	18.3
41	306	45	90	0	45	30	30	50	55	55	45	25	37.6	21.7	22.4	28.3
42	309	15	0	0	5	15	10	0	10	10	50	8.3	4.1	3.3	2.4	16.7
mean CpGs 4,9,11,20		all donors										27.6	36.7	35.5	28.5	72.5
mean CpGs 4,9,11,20		donor M4										11.2	35	28.7	32.5	81.2

Foxp3 preTSDR																								
Donor		M2					M4					M5					M6							
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5			
1	1	N/A	90	75	90	85	45	100	10	0	75	85	65	60	45	55	15	35	85	60	75			
2	30	N/A	100	80	100	100	95	100	80	75	100	100	95	85	90	90	100	100	100	100	100			
3	37	N/A	100	80	100	100	95	100	90	75	100	100	95	80	90	90	100	100	100	100	95			
4	87	N/A	100	80	95	100	95	100	80	95	95	95	95	80	90	90	100	100	100	100	95			
5	123	N/A	40	15	10	10	30	100	15	0	70	20	50	65	70	70	15	20	0	65	85			
6	187	N/A	65	75	80	100	70	0	70	70	75	80	90	75	85	80	85	45	95	30	75			
7	243	N/A	95	75	90	85	30	100	10	0	65	85	45	60	45	55	15	30	90	70	85			
8	334	N/A	50	15	10	10	30	100	10	0	80	25	75	80	80	75	15	15	5	65	85			
9	357	N/A	10	10	0	10	5	0	0	0	60	0	50	60	45	50	0	15	0	0	60			
10	524	N/A	55	10	25	45	90	100	85	80	100	30	90	85	90	95	100	75	20	95	90			
Donor		F1					F2					F3					F4							
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5			
1	1	10	70	25	0	75	10	75	30	100	90	55	25	50	45	70	75	80	65	75	60			
2	30	10	100	45	100	100	25	90	55	100	100	100	25	55	80	100	95	90	90	90	100			
3	37	20	100	45	100	100	25	90	65	100	100	100	20	50	80	95	95	90	95	90	100			
4	87	15	100	60	95	100	15	85	55	90	95	90	30	55	80	90	100	85	90	85	100			
5	123	0	95	10	0	10	0	35	25	10	20	60	25	30	45	85	15	20	15	30	70			
6	187	5	30	50	100	100	15	65	60	70	95	50	40	60	35	90	100	70	90	80	80			
7	243	10	90	35	95	75	15	90	45	100	85	60	40	50	50	95	80	90	75	75	80			
8	334	10	100	20	95	20	0	45	30	25	30	80	50	40	50	100	35	40	10	55	75			
9	357	0	25	95	5	20	0	30	15	0	35	15	40	25	15	90	20	10	5	20	55			
10	524	10	100	50	100	45	10	60	30	20	65	95	35	60	75	100	60	40	45	40	80			
Donor		F5					mean all donors								70.5	33	47.5	55.5	91.5	mean all CpGs donor F3				
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	70.5	33	47.5	55.5	91.5	mean all CpGs donor F3		
1	1	70	40	95	55	100	45.6	64.4	55	52.2	76.1													
2	30	95	75	100	80	100	77.5	86.1	76.7	90.6	98.9													
3	37	95	75	95	80	100	78.8	85.6	78.3	90.6	97.8													
4	87	85	75	90	80	95	74.4	85.6	76.7	90	95.6													
5	123	30	45	35	20	15	21.3	47.8	23.3	27.8	48.3													
6	187	80	70	60	70	95	60.6	52.8	70.6	68.9	87.8													
7	243	80	75	100	70	90	46.9	72.8	60	66.1	79.4													
8	334	35	45	55	25	30	28.8	57.8	29.4	45	56.1													
9	357	30	40	15	20	20	8.8	24.4	25	11.7	44.4													
10	524	50	50	55	30	45	55.6	67.2	48.9	61.7	73.9													
mean CpGs 5,8,9 all donors		19.6					43.3					25.9					28.2					mean all CpGs donor F3		
mean CpGs 5,8,9 donor M6		10					16.7					1.7					43.3					mean all CpGs donor F3		

Foxp3 promoter

Donor		M1					M2					M3					M4					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	N/A	15	0	15	60	N/A	0	0	0	30	N/A	0	0	0	0	0	0	0	0	0	0
2	41	N/A	15	0	5	45	N/A	0	0	0	25	N/A	0	0	0	0	0	0	0	0	0	0
3	118	N/A	15	5	10	55	N/A	0	0	0	25	N/A	0	0	0	0	0	5	0	0	5	0
4	130	N/A	15	0	10	75	N/A	5	0	0	65	N/A	0	0	0	5	0	0	0	0	0	0
5	143	N/A	15	10	20	70	N/A	5	0	0	70	N/A	0	0	0	0	0	0	0	0	0	0
6	179	N/A	10	10	20	85	N/A	15	0	0	75	N/A	0	0	5	15	5	0	5	0	0	0
7	191	N/A	15	5	15	60	N/A	15	0	0	70	N/A	0	0	5	10	0	0	0	0	0	0
8	198	N/A	10	0	25	75	N/A	5	0	0	65	N/A	0	0	0	15	0	5	0	5	0	0
9	213	N/A	0	0	15	60	N/A	5	0	0	25	N/A	0	0	0	0	0	0	0	0	0	0
10	241	N/A	15	5	15	70	N/A	5	0	0	55	N/A	0	0	0	20	0	0	0	0	0	0
Donor		M5					M6					F1					F2					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	30	0	0	0	0	5	
2	41	0	0	0	0	0	0	0	0	0	0	0	0	0	0	20	0	0	0	0	0	
3	118	0	0	0	0	0	0	0	0	0	0	0	0	0	0	20	5	0	5	0	0	
4	130	0	0	0	0	0	0	0	0	0	0	0	0	0	0	50	5	0	0	0	0	
5	143	0	0	0	0	0	0	0	0	0	0	0	0	0	0	40	5	0	0	0	5	
6	179	0	0	0	0	0	0	0	0	0	0	0	0	0	0	50	5	0	0	0	0	
7	191	0	0	0	0	0	0	0	0	0	0	0	0	0	0	50	0	0	0	0	0	
8	198	0	0	0	0	0	0	0	0	0	0	0	0	0	0	40	0	0	0	0	0	
9	213	0	0	0	0	0	0	0	0	0	0	0	0	0	0	45	0	10	0	0	0	
10	241	0	0	0	0	0	0	0	0	0	0	0	0	0	0	40	0	0	0	0	0	
Donor		F3					mean all donors					mean M1 and 2					mean M3-6 and F1-5					
CpG	nt	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	#1	#2	#3	#4	#5	
1	1	0	0	0	0	0	0	1.7	0	1.7	13.9	N/A	7.5	0	7.5	45	0	0	0	0	5	
2	41	0	0	0	0	0	0	1.7	0	0.6	10	N/A	7.5	0	2.5	35	0	0	0	0	2.9	
3	118	0	0	0	0	0	1.7	1.7	1.1	1.7	11.1	N/A	7.5	2.5	5	40	1.7	0	0.7	0.7	2.9	
4	130	0	0	0	0	0	0.8	2.2	0	1.1	21.7	N/A	10	0	5	70	0.8	0	0	0	7.9	
5	143	0	0	0	0	0	0.8	2.2	1.1	2.2	20.6	N/A	10	5	10	70	0.8	0	0	0	6.4	
6	179	0	0	0	0	0	1.7	2.8	1.7	2.8	25	N/A	12.5	5	10	80	1.7	0	0.7	0.7	9.3	
7	191	0	0	0	0	0	0	3.3	0.6	2.2	21.1	N/A	15	2.5	7.5	65	0	0	0	0.7	8.6	
8	198	0	0	0	0	0	0	2.2	0	3.3	21.7	N/A	7.5	0	12.5	70	0	0.7	0	0.7	7.9	
9	213	0	0	0	0	0	0	1.7	0	1.7	14.4	N/A	2.5	0	7.5	42.5	0	1.4	0	0	6.4	
10	241	0	0	0	0	0	0	2.2	0.6	1.7	20.6	N/A	10	2.5	7.5	62.5	0	0	0	0	8.6	
mean all CpGs							0.5	2.2	0.5	1.9	18	N/A	9	1.7	7.5	58	0.5	0.2	0.1	0.3	6.6	

Foxp3 Promoter Top/Reverse Strands									
Donor		M6			F3		F4		
subset		#2	#5	#5	#2	#5	#2	#5	#5
CpG	nt	TS	TS	RS	TS	TS	TS	TS	RS
1	1	0	60	0	60	80	60	40	0
2	41	0	50	0	70	40	0	40	0
3	118	0	60	0	90	50	60	30	0
4	130	10	100	0	80	70	60	80	0
5	143	10	100	0	80	90	60	90	0
6	179	10	100	0	0	50	0	0	0
7	191	0	90	0	80	100	60	100	0
8	198	10	100	0	80	70	60	100	0
9	213	0	90	0	50	100	60	80	0
10	241	10	90	0	40	70	60	80	0
mean all CpGs		5	84	0	63	72	48	64	0

Foxp3 TSDR

Foxp3 TSDR Top 700bp/500bp												Foxp3 TSDR Top/Reverse Strands 700bp																			
Donor		M2						F2						M2						F1						M4					
CpG	subset	#3	#3	#5	#5	#2	#2	#5	#5	#3	#3	#5	#5	#3	#3	#5	#5	#2	#2	#4	#4	#5	#5	TS	RS	TS	RS	TS	RS	TS	RS
1	1	0	0	100	90	10	20	100	90	0	0	100	100	60	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
2	6	10	0	100	100	0	20	90	80	0	0	100	100	60	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
3	47	0	0	100	90	10	20	100	100	0	0	80	100	40	100	90	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
4	55	0	0	100	100	10	20	100	100	0	0	100	100	60	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
5	65	0	0	100	100	0	20	100	100	0	0	90	100	60	100	80	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
6	68	0	0	100	90	0	20	100	100	0	0	100	100	60	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
7	74	0	0	100	100	0	20	100	100	10	0	30	100	20	100	70	100	100	100	0	100	10	100	90	100	100	100	100	100	100	
8	78	0	0	100	100	0	20	100	100	0	0	100	100	70	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
9	86	20	0	100	100	0	20	100	80	0	0	100	100	80	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
10	95	0	0	100	100	0	20	90	100	0	0	90	100	70	100	100	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
11	114	0	0	100	100	10	20	100	100	0	0	100	100	60	100	90	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
12	149	0	0	90	90	20	0	60	60	0	0	100	100	60	100	60	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
13	221	0	0	90	70	10	20	70	40	0	0	100	100	60	100	50	100	0	100	0	100	80	100	100	100	100	100	100	100	100	100
14	246	0	0	100	100	40	20	90	100	0	0	100	100	70	100	90	100	0	100	0	100	100	100	100	100	100	100	100	100	100	
15	259	0	0	100	100	20	20	100	100	10	0	100	100	70	100	80	100	0	100	0	100	90	100	100	100	100	100	100	100	100	
mean all CpGs		2	0	98.7	95.3	8.7	18.7	93.3	90	1.3	0	92.7	100	60	100	87.3	100	0	100	0	100	97.3	100								

Foxp3 TSDR Top/Reverse Strands 700bp																														
Donor		M6						F3						F4						F4										
CpG	subset	#2	#2	#4	#4	#5	#5	#2	#2	#4	#4	#5	#5	#2	#2	#4	#4	#5	#5	TS	RS	TS	RS	TS	RS					
1	1	0	100	60	100	100	100	0	100	0	100	100	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
2	6	0	100	60	100	100	100	0	100	0	100	90	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
3	47	0	100	60	100	100	100	0	100	0	100	100	100	50	100	100	100	90	100	100	100	100	100	100	100	100	100	100	100	100
4	55	0	100	60	100	100	100	0	100	0	100	90	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
5	65	0	100	60	100	100	100	0	100	0	100	100	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
6	68	0	100	60	100	100	100	0	100	0	100	90	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
7	74	0	100	50	100	100	100	20	100	30	100	80	100	50	100	70	100	50	100	100	100	100	100	100	100	100	100	100	100	100
8	78	0	100	60	100	100	100	0	100	0	100	100	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
9	86	0	100	60	100	100	100	0	100	0	100	90	100	50	100	100	100	90	100	100	100	100	100	100	100	100	100	100	100	100
10	95	0	100	60	100	100	100	0	100	0	100	100	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
11	114	0	100	70	100	100	100	0	100	0	100	100	100	50	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
12	149	0	100	70	100	90	100	0	100	0	100	80	100	0	100	50	100	100	80	100	100	100	100	100	100	100	100	100	100	100
13	221	0	100	60	100	100	100	0	100	0	100	80	100	40	100	100	100	70	100	100	100	100	100	100	100	100	100	100	100	100
14	246	0	100	70	100	100	100	0	100	0	100	100	100	40	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
15	259	0	100	70	100	100	100	0	100	0	100	100	100	40	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100
mean all CpGs		0	100	62	100	99.3	100	1.3	100	2	100	93.3	100	44.7	100	94.7	100	91.3	100											

Supplementary Table 3. Methylation percentages for *CAMTA1*, *FUT7*, *FOXP3* enhancer, preTSDR, promoter and TSDR regions in five cell populations of donors M1-6 and F1-5. Methylation percentages for individual CpG sites were calculated based on the number of ^mCs in a total of 20 sequences unless otherwise stated. CpGs in each amplicon are numbered relative to the 5'-3' direction of the coding strand with CpG 1 denoted nucleotide (nt) position 1. The five cell subsets are: #1 (CD34⁺), #2 (CD45RA⁺CD15s⁻), #3 (CD45RA⁻CD15s⁻), #4 (CD45RA⁻CD15s⁺) and #5 (Tcon). TS- top strand, RS – reverse strand

Human <i>FOXP3</i> TSDR		
Reference	Based on	Strand
(1)	Primers	Top
(2)	Primers	Reverse
(3)	Primers	Top
(4)	Primers	Reverse/Top
(5)	Primers	Top
(6)	Primers	Reverse
(7)	(1)	Top
(8)	(5)	Top
(9)	(5)	Top
(10)	(5)	Top
(11)	(5)	Top
(12)	(5)	Top
(13)	Primers	Reverse/Top
(14)	Primers	Reverse
(15)	Primers	Top
(16)	(5)	Top
(17)	(4)	Top
(18)	(5)	Top
(19)	Primers	Top
(20)	Primers	Top
(21)	Primers	Top
(22)	(1)	Top
(23)	Primers	Top
(24)	Primers	Reverse
(25)	(5)	Top
(26)	(5)	Top
(27)	EpigenDx	---
(28)	(1)	Top
(29)	(1)	Top
(30)	Primers	Reverse
(31)	Primers	Reverse
(32)	(5) and primers	Top
(33)	(5)	Top
(34)	Primers	Top
(35)	(5)	Top
(36)	Primers	Top
Murine <i>FOXP3</i> TSDR		
(37)	Primers	Reverse
(38)	(37)	Reverse
(39)	Primers	Top
(40)	Primers	Top
(41)	EpigenDx	---
(42)	EpigenDx	---
(43)	Primers	Reverse
(44)	Primers	Reverse
(45)	(40, 44)	Top, Reverse
(46)	(37)	Reverse
(47)	Primers	Top
(48)	Primers	Top

(49)	Primers	Top
(50)	Primers	Reverse /Top
(51)	(43)	Reverse
(52)	(39), (43)	Top, Reverse
(53)	Primers	Top
(54)	Primers	Top
(55)	Primers	Top
Human <i>FOXP3</i> Promoter		
(38)	Primers	Top
(56)	Primers	Top
(2)	Primers	Reverse
(3)	Primers	Top
(57)	Primers	Top
(6)	Primers	Reverse
(13)	Primers	Reverse
(19)	Primers	Top
(21)	Primers	Reverse
(25)	Primers	Top
(58)	Primers	Reverse
(59)	Primers	Top
(60)	Primers	Reverse
(61)	Primers	Top

Supplementary Table 4. Studies that used human and murine *FOXP3* TSDR and promoter regions. Template strands used to detect methylation status of the two regions were identified using primer sequences supplied in the article or *via* reference (which then had primer sequence) used in Methodology section when primer sequences were not available

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