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Figure S4 HIV exposure alters transcriptomic profile of gut homing Treg cells. A) Flow plot showing sorting of $\alpha 4\beta$ 7- and $\alpha 4\beta$ 7+ Treg cells. B) NMDS plots showing Bray Curtis distances between infant samples.

Antibody	Clone	Supplier
CD3 Apc-Cy7	UCHTI	Biolegend
CD4 AlexaFluor 700	SK3	Biolegend
CD8 BV570	RPA-T8	Biolegend
CD127 PE-CF594	hIL-7R-M21	BD
CD25 PE-Cy7	2A3	BD
Foxp3 AlexaFluor 647	2594/C7	BD
CD39 BV786	Tu66	BD
TIGIT FITC	MBSA43	eBioscience
CCR7 PerCP-Cy5.5	G043H7	Biolegend
CCR6 BV711	G034E3	eBioscience
CCR4 BV605	L291H4	Biolegend
CD161 PE-Cy5	DX12	BD
$\alpha 4\beta 7 PE$	A4B7R1	NIH AIDS reagents
FACS sorting		
CD3 FITC	OKT3	Biolegend

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Table S2 Demographic characteristic of HIV infected-exposed and HIV uninfectedunexposed mother-infant pairs

Demographics	HUU (n=16)	HEU (n=20)	P value
Median maternal age at delivery, years (range)	26 (19-36)	27 (19-39)	0.49
Median gestational age, weeks (range)	39 (37-41)	39 937-41)	0.89
Median maternal CD4 count cells/ μ L (IQR)	-	429.5	
		(344.5-516.50)	
Initiating ART, n (%)		10 (50)	
Infant			
Median birth weight, kg (IQR)	3.0 (2.9-3.3)	3.3 (2.9-3.5)	0.31
Gender, female (%)	9 (56.3)	11 (55.0%)	0.94
Exclusively breastfeeding			
7 weeks, n (%)	9 (56.3)	10 (50.0)	
15 weeks, n (%)	6 (37.5)	6 (30.0)	
36 weeks, n (%)	0 (0.0)	0 (0.0)	

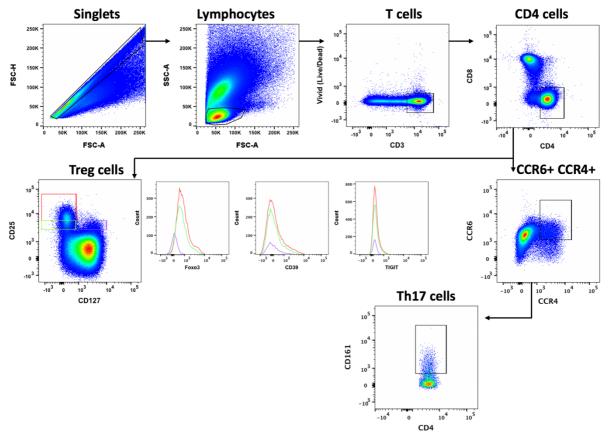


Figure S1 Flow cytometry gating strategy to identify T regulatory and Th17 cells.

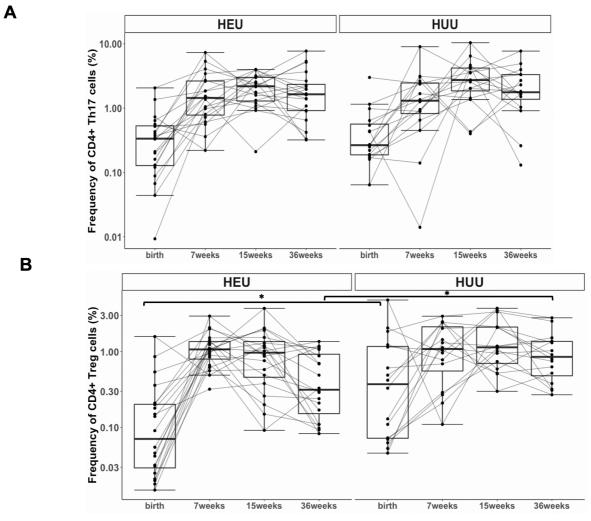


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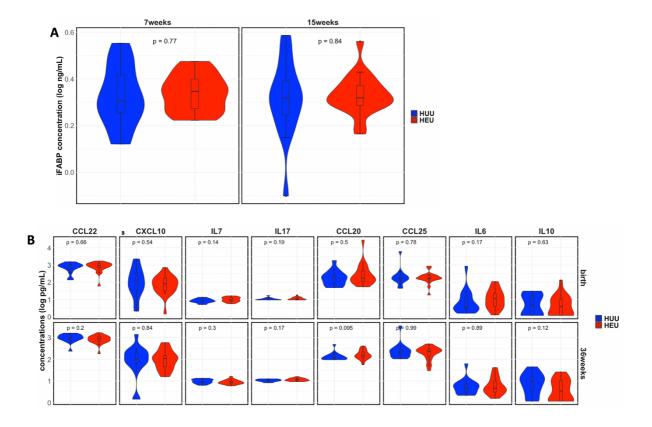


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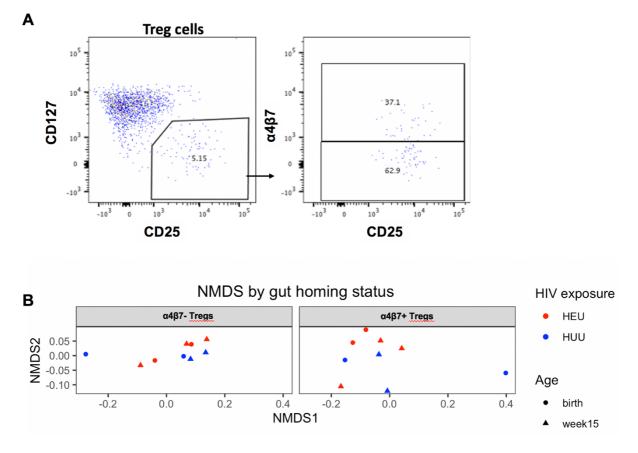


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