## Supplementary Figure 1



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[A] Representative dot plots illustrating purity of CD71<sup>+</sup> erythroid cells. [B] Representative dot plot showing percent CD71<sup>+</sup> erythroid cells in cord blood of an infant born to a CD mother following an emergency C-section, also showing % activated immune cells (CD71<sup>+</sup>CD235a<sup>-</sup> cells). [C] Representative zebra plots showing % CD69 expression on CD4 T cells among placenta cells from HC versus UC and CD patients. [D] Representative zebra plots showing % CD69 expression on CD4 T cells among CBMCs of infants born to HC versus UC and CD mothers. [E, F] Data showing percentages of CD4 and CD8 T cells in gut tissues and spleen in control versus treated mice with anti-CD71 antibody, respectively.



1 Increased intestinal permeability

## Supplementary Figure 2

CD71<sup>+</sup> erythroid cells expand in the 2<sup>nd</sup> and 3<sup>rd</sup> trimesters of pregnancy. Profound changes of gut microbiota take place at the 3<sup>rd</sup> trimesters which coincides with the expansion of CD71<sup>+</sup> erythroid cells. Depletion of CD71<sup>+</sup> erythroid cells in a mouse model of pregnancy which mimics lower frequency of CD71<sup>+</sup> erythroid cells in IBD patients was associated with pro-inflammatory response as illustrated by upregulation of TLRs, IL-6, TNF-a and CXCL-1 expression and downregulation of TGF-b in the gut tissues.

Supplementary Table 2. Primers used for microbiota analysis.

Target bacterial group	Primers sequence (5'→3')	Tm (≌C)	Amplico n size (bp)	Reference
Total bacteria	F: CGGYCCAGACTCCTACGGG	63	200	120
	R: TTACCGCGGCTGCTGGCAC			
Bacteroides group	F: GGTGTCGGCTTAAGTGCCAT	60	140	20
	R: CGGAYGTAAGGGCCGTGC			
Lactobacillus group	F: AGCAGTAGGGAATCTTCCA	63	341	20
	R: CACCGCTACACATGGAG			
Enterobacteriaceae family	F: CATTGACGTTACCCGCAGAAGAAGC	63	195	20
	R: CTCTACGAGACTCAAGCTTGC			
Clostridium cluster I	F: GTGAAATGCGTAGAGATTAGGAA	58	665	20
	R: GATYYGCGATTACTAGYAACTC			
Clostridium cluster IV	F: GCACAAGCAGTGGAGT	60	239	20
	R: CTTCCTCCGTTTTGTCAA			
Clostridium cluster XIVa	F: AAATGACGGTACCTGACTAA	58	438-441	20
	R: CTTTGAGTTTCATTCTTGCGAA			