MiR-155 regulates IL-10-producing CD24^{hi}CD27⁺ B cells and 1 impairs their function in patients with Crohn's disease 2 Yingxia Zheng, ^{1, 2} Wensong Ge, ³ Yanhui Ma, ¹ Guohua Xie, ¹ Weiwei Wang, ¹ Li 3 Han, ¹Bingxian Bian, ¹Li Li, ¹Lisong Shen ^{1,*} 4 5 6 7 **Supplementary information** miRNA Microarray 8 Human CD19⁺ B cells were isolated and were stimulated with CpG 9 oligonucleotides for 48 h. Total RNA was quantified by the NanoDrop ND-2000 10 (Thermo Scientific) and the RNA integrity was assessed using Agilent Bioanalyzer 11 12 2100 (Agilent Technologies). The sample labeling, microarray hybridization and washing were performed based on the manufacturer's standard protocols. Briefly, 13 total RNA were dephosphorylated, denaturated and then labeled with Cyanine-3-CTP. 14 After purification the labeled RNAs were hybridized onto the microarray. After 15 washing, the arrays were scanned with the Agilent Scanner G2505C (Agilent 16 17 Technologies). Feature Extraction software (version10.7.1.1, Agilent Technologies) was used to analyze array images to get data. 18

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20 *Cell viability*

Isolated B cells were electroporated with control or miR-155 mimic and miR-155
stimulated with 100 nM CPG and cultured for another 48 h. The viability of the cells
was analyzed by CellTiter-Glo luminescent assay (Promega, Madison, WI);
the luminescence values were measured with microplate computer software (Bio-Rad
Laboratories).



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Supplementary figure 1: Heat map and unsupervised hierarchical clustering of
miRNAs in human CD19⁺ B cells stimulated with or without 100nM CpG
oligonucleotides for 48 h. The microRNA arrays were scanned with the Agilent
Scanner G2505C. Differentially expressed miRNAs were then identified through fold
change. Hierarchical Clustering was performed to show the distinguishable miRNAs
expression pattern among samples.



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Supplementary figure 2: Flowcytometric gating strategy for B cell subsets in a
representative healthy control. B cells defined as CD19⁺ lymphocytes were further
classified into four subsets (CD24^{hi}CD27⁺; CD24^{lo}CD27⁺; CD24^{hi}CD27⁻;
CD24^{lo}CD27⁻) according to the CD24 and CD27 markers.



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Supplementary Figure 3: (A) Isolated CD19⁺ B cells were electroporated with control 40 or miR-155 mimic; then, the cells were cultured in a 96-well microplate (2 x 10^5 cells 41 per well), stimulated with 100 nM CPG and cultured for another 48 h. The viability of 42 the cells was then analyzed. (B) FACS was used to sort CD24^{hi}CD27⁺ B cells, which 43 were electroporated with control or miR-155 inhibitor; then, the cells were cultured in 44 a 96-well microplate (2×10^5) cells per well), stimulated with 100 nM CPG and 45 cultured for another 48 h. The viability of the cells was then analyzed. NS: no 46 significant difference. 47



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- 49 Supplementary Figure 4: CD19⁺B cells isolated from HCs and CD patients; Jarid2
- 50 expression levels were measured by Q-PCR. NS: no significant difference.
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