

Supplement

Indoxyl Sulfate Elevated *Lnc-SLC15A1-1* Up-regulating CXCL10/CXCL8 Expression in High Glucose Endothelial Cells by Sponging MicroRNAs

Table S1. Primers of lncRNAs.

Transcript ID	Primer sequence 1	Primer sequence 2	Primer sequence 3
<i>NONHSAT</i>	F- CCTCTTGCCCTGTTGAGAGC	F- GTCCCGCTCCACTTCATAACT	
<i>238235.1</i>	R- TCCTTTCGGGGTTCAGCAAA	R- GGGTTTTCCACTGCTCCAAC	
<i>NONHSAT</i>	F- TGTAACCTCTTGGAAGCCCT	F- TGGAGTTGTGAAGGACAGTGA	
<i>231357.1</i>	R- TGGCCCCTGGAAGAACTTG	R- TGCCCTGGCTAGATTCCAC	
<i>NONHSAT</i>	F- GCCATACCTTGCAGTGTGCG	F- GCACACAGCAGGAGAGACAG	
<i>144427.2</i>	R- GAACGGGAGTCTGACCAGGTG	R- TGGGGGTCCTGAGAGTCTTC	
<i>NONHSAT</i>	F- CACCCGCTTGCTAGTGTCT	F- GCATTGGGCTAGTGTGAGGT	F- GGACCTCTGATAAGGCTGGC
<i>258517.1</i>	R- AGCTCTACCCTACACCCAG	R- AGTGGTGGGTGTCATTCCC	R- TTTGCTCTGGCTCTTGTTG
<i>NONHSAT</i>	F- AGTGTGCGGAGTTGTGTTCA		
<i>167136.1</i>	R- GGGGCTCCAGGCATAAACTC		

Table S2. Primers of microarray confirmed mRNAs.

Gene	Forward primer	Reverse primer
<i>CXCL10</i>	TCCACGTGTTGAGATCATTGC	TCTTGATGGCCTTCGATTCTG
<i>CXCL8</i>	AAGAAACCACCGGAAGGAAC	AAATTTGGGGTGGAAGGTT
<i>FAT2</i>	CACTCCCGAGATCCAAAGGG	GGCAGTAGAAGAGAAGCCCG
<i>JAM2</i>	GTCTCCTTTGTCTACTATCAAC	GGAGCCACTAATACTTCCAG
<i>TXNIP</i>	GCCACACTTACCTTGCCAAT	TTGGATCCAGGAACGCTAAC
<i>ARRDC4</i>	TGAAGAGTGGAACGAAGGCG	AGTGGGCCATTCTCCTATGC
<i>IL-6</i>	ACTCACCTCTTCAGAACG	GGCTTGTTCTCACTACT
<i>GAPDH</i>	GTCTCCTCTGACTTCAACAGCG	ACCACCCTGTTGCTGTAGCCAA

Table S3. RT stem loop primer sequences.

microRNA	Sequence
miR-150-3p	GTTGGCTCTGGTGCAGGGTCCGAGGTATTTCGCACCAGAGCCAACCTGTCCC
miR-27b-5p	GTTGGCTCTGGTGCAGGGTCCGAGGTATTTCGCACCAGAGCCAACG TTCACC
miR-297	GTTGGCTCTGGTGCAGGGTCCGAGGTATTTCGCACCAGAGCCAACCATGCAC

Table S4. Primers of miRNAs.

microRNA	Forward primer	Reverse primer
<i>miR-150-3p</i>	AGCTGGTACAGGCCTGG	GTGCAGGGTCCGAGGT
<i>miR-27b-5p</i>	GCGCAGAGCTTAGCTGATT	GTGCAGGGTCCGAGGT
<i>miR-297</i>	GGCGGATGTATGTGTGCAT	GTGCAGGGTCCGAGGT
<i>U6</i>	GCTTCGGCAGCACATATACTAAAAT	CGCTTCACGAATTTGCGTGTCAT

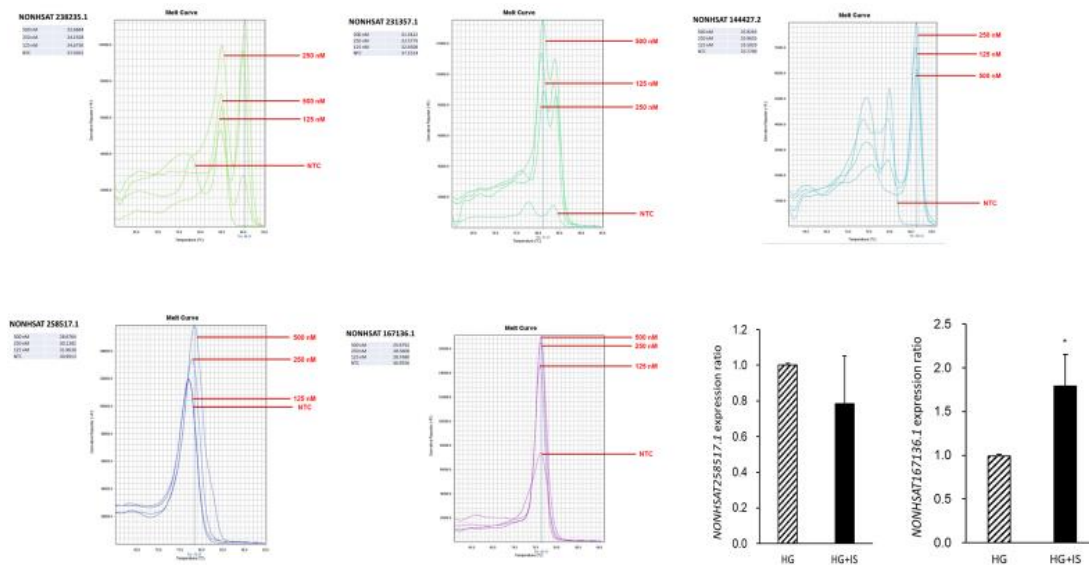
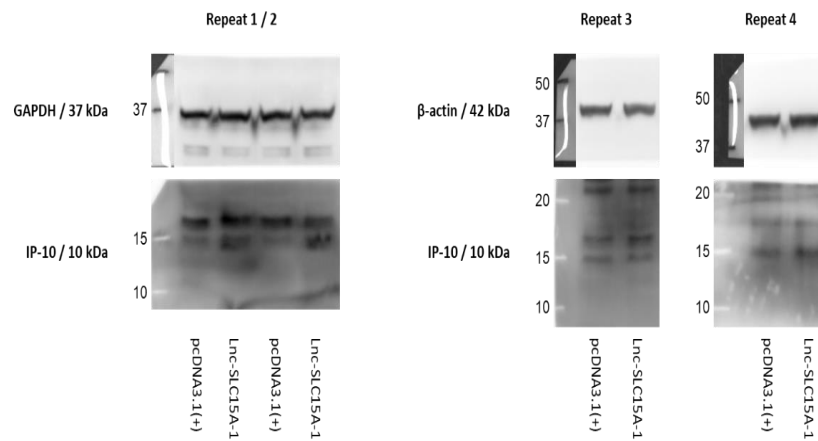


Figure S1. Primer tests of selected lncRNAs and the results of RT-PCR.

To assess the designed PCR primers against the input lncRNA sequence to determine if a product other than the desired target can be amplified, a typical denaturation (melt) curve was performed. Designed lncRNA primers with different concentrations were evaluated to determine which one could give rise to a single distinct peak in the plot of the negative derivative of fluorescence vs. temperature. The qualified designed lncRNA primers were used to perform the RT-PCR in the condition of HUVECs explored on HG and HG+IS. lncRNA of NONHSAT167136.1 expression was significantly ($p < 0.05$) increased in HUVECs in HG+IS treatment by using the designed primer in RT-PCR experiment.

(A1) Four repeats of IP-10 western blot experiments



(A2) Four repeats of IL-8 western blot experiments.

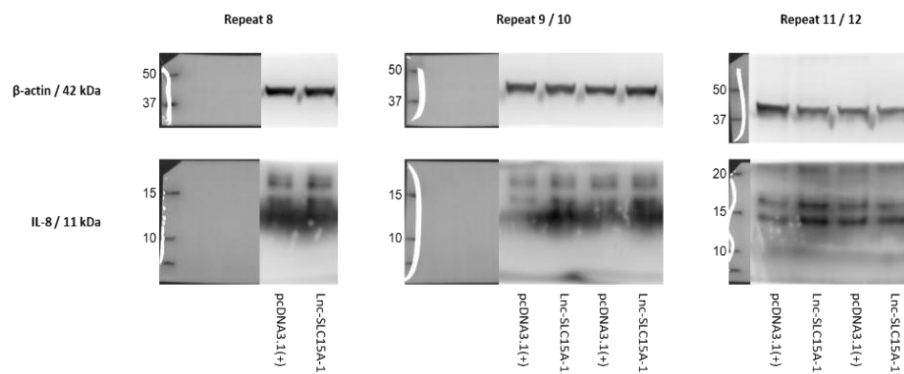


Figure S2. Original western blot gel of IP-10 and IL-8.