

Fig. S1

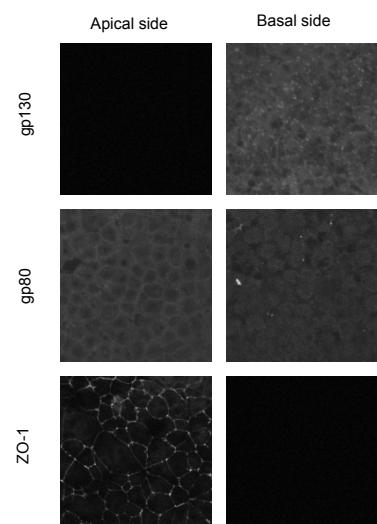


Fig. S2

1 GAATCTTGGC AACACCGAGG GCTCCTTGAA CACGGCAAA TCTTATATGG CTCTGAGATT
 61 CCAAAGCATT GACTCAGATA CCTGCCTCAT GCAAAGCCCT ATATTCTAGA GCAGTTCCC
 121 TTTCTCTGT GGCAAGACTCT TGTCCTCCCT AACAGATGGC CCAGGGATT TCAGGGCCC
NFkB
 181 CTCTCAGTCC TGGAACCCCTT GTTCCAGAGT GCTCCCTCAT CATCCAAGAG GCTGATGATG
 241 GGAGCATCTA TTAGGAGACT GGACAGGAAA TGTCCTGGCA TGTTATACAT GCAGGAGGCC
 301 TTAGACTAGG CTGCAGAGGG GGATTTGGC ATGGCTGGGA GGATCTGAAC TCTCAGAGTA
 361 TGGACAGAACGGCTAGGCC ACTACTCTCT AGGCCCTGG AGATTCAAGA GGCTCTAAC
 421 GAATGATCCA GGGCTAGGCC ACTACTCTCT AGGCCCTGG AGATTCAAGA GGCTCTAAC
 481 AAATGGAGT CCAAGACTAC ATTCTAGGAT CTGTTCTCC TGATGTAGTC TGCAGTTGG
 541 CCTCAGTCTG CAATTGAGGG GCCCTATGGC ACTGTTGCTT GGCAATGTAT TAAACAGCAG
 601 GCCTTGGAGA CTAGCACTTG AGTTAACACA GCCACCACAA CCACCACTGC CATCATCAC
 661 TTCCGGAAA GCAGCCACCT GTCTGGCTCC TGGCTTTGTC CAGCTGCCAA CCTAAGGCAT
STAT
 721 GTGCCTACGC AGGAGGCGAT GACATTGGG CTCCACGTC AAAGTTGTTT TTTTTTCCT
 781 TTCTCATGTG TTATTTCTAA AGATAACAAA GGTCAAAAGG CATCCAGCGT TTTCTGGTT
Cdx-A Cdx-B
 841 CTCATAAGCT TCTGGTCAAT ATTTAATCTG GTTTATGGAT TTTTTTAGG TCTTCTAGAT
Cdx-C Cdx-D
 901 GCCTTCTTGA GGCTGTTGT GGCCACCCAC AGACACTGT AAGGAGGAGA GAAGTCAGCC
Start of Transcription
 961 TGGCAGAGAG ACTCTGAAAT GAGGGATTAG AGGTGTTCAA GGAGCAAGAG CTTCAGCCTG
 1021 AAGACAAGGG AGCAGTCCCT GAAGACGCTT CTACTGAGAG GTCTGCCATG
Start of Translation

Fig. S3

SUPPLEMENTAL FIGURE LEGENDS

Fig. S1 IL-6 increases TJ permeability and claudin-2 expression in T84 cell monolayers. (A) TER was measured across T84 cell monolayers incubated with varying concentrations of IL-6 (0~30 ng/mL) for 48 hr. (B) Whole cell extracts of T84 cell monolayers incubated with varying concentrations of IL-6 (0~30 ng/mL) for 48 hr were immunoblotted for claudin-2 and β -actin. * $P < 0.05$ relative to the control value. Values represent the mean \pm SEM ($n = 4$).

Fig. S2 Localization of gp130 and gp80 differ in Caco-2 cell monolayers. Caco-2 cells fixed with methanol were immunolabeled for gp130, gp80, and ZO-1 and the apical and basal images of cell monolayers were collected by confocal microscopy.

Fig. S3. Sequence analysis of the 5'-flanking region of the human claudin-2 gene. Potential binding sites are underlined.

Table S1. Primers used in quantitative RT-PCR

Target gene	Forward	Reverse
hClaudin-2	5'-CTCCCTGGCCTGCATTATCTC-3'	5'-ACCTGCTACCGCCACTCTGT-3'
hCdx-1	5'-TCGGACCAAGGACAAGTACC-3'	5'-GATCTTCACCTGCCGTTCAG-3'
hCdx-2	5'-CAGTCGCTACATCACCATCC-3'	5'-TTCCCTCTCCTTGCTCTGC-3'
h β -actin	5'-GCCGAGGACTTGATTGCA-3'	5'-CTTCCTGTAACAACGCATCTCA-3'

Table S2. Primers used to delete transcriptional factor binding sites in site-directed mutagenesis of claudin-2 promoter

Deletion target	Forward	Reverse
Cdx-A	5'-AAGATAACAAAGGTCAAAAGGC-3'	5'-AACACATGAGAAAGGAAAAAAAAAAC-3'
Cdx-B	5'-GTTTCTCATAAGCTTCTGGTC-3'	5'-CGCTGGATGCCTTGACC-3'
Cdx-C/D	5'-TCTTCTAGATGCCCTCTTGAG-3'	5'-CAGATTAAATATTGACCAGAAGC-3'
STAT	5'-AGCAGCCACCTGTCTGGCTCTG-3'	5'-GGTGATGATGGCAGTGGTGGTTG-3'
NFκB	5'-GGGCCCTCTCAGCCTGG-3'	5'-TGGGCCATCTGTTAGGGGG-3'