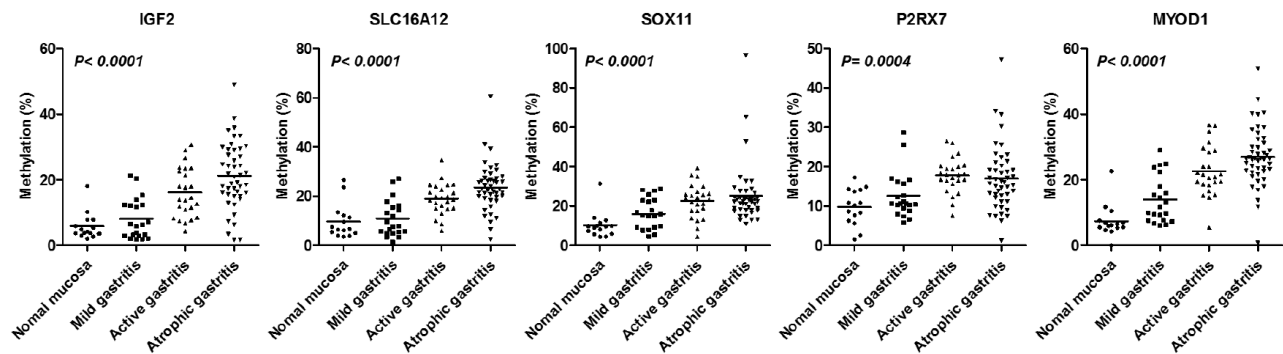
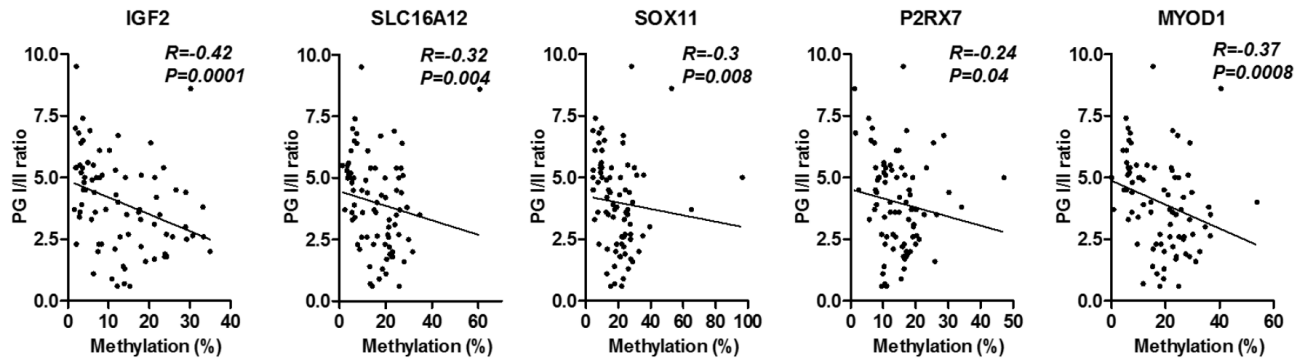


## Demonstration of potential link between *Helicobacter pylori* related promoter CpG island methylation and telomere shortening in human gastric mucosa

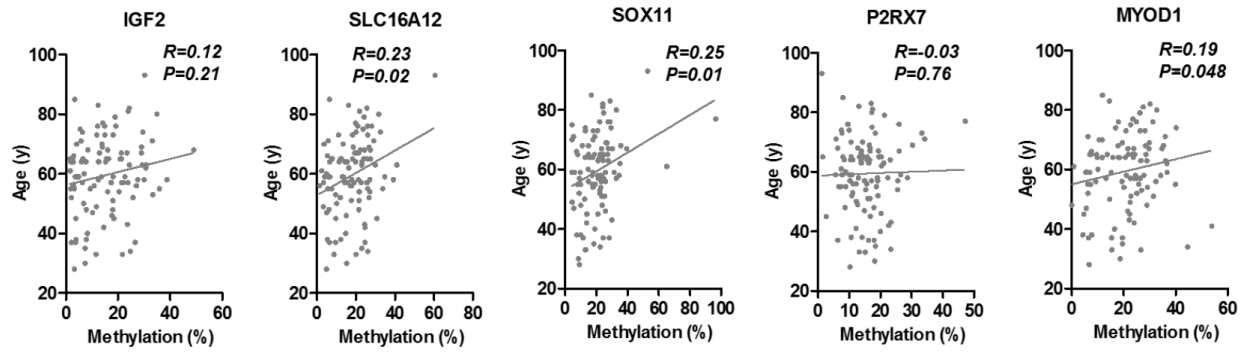
### SUPPLEMENTARY FIGURES AND TABLES



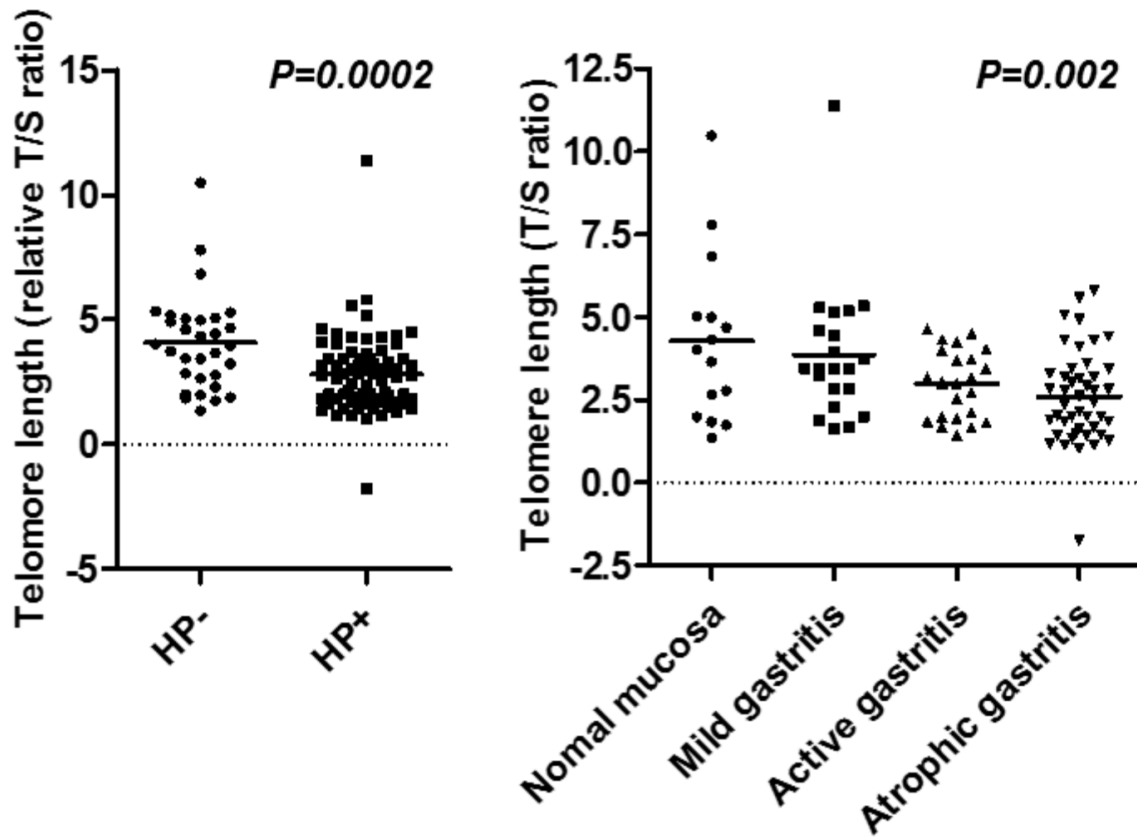
Supplementary Figure S1: Association between methylation status of five PCGIs (*IGF2*, *SLC16A12*, *SOX11*, *P2RX7* and *MYOD1*) and development of pathological state of *H. pylori* related gastritis. The statistical analysis was performed using one way ANOVA.



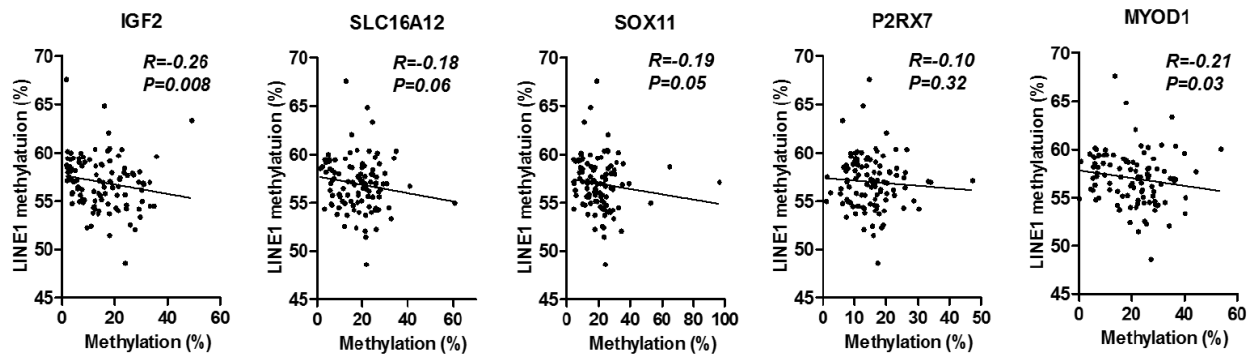
Supplementary Figure S2: Association between methylation status of five PCGIs (*IGF2*, *SLC16A12*, *SOX11*, *P2RX7* and *MYOD1*) and serum pepsinogen (PG) I/II ratio. Statistical analysis was performed using the Spearman correlation analysis.



**Supplementary Figure S3: Association between methylation status of five PCGIs (*IGF2*, *SLC16A12*, *SOX11*, *P2RX7* and *MYOD1*) and aging.** Statistical analysis was performed using the Spearman correlation analysis.



Supplementary Figure S4: Association between telomere shortening and *H. pylori* infection (left) and development of pathological state of *H. pylori* related gastritis (right). The statistical analysis was performed using Student's t-Test (left) and one way ANOVA (right), respectively.



Supplementary Figure S5: Association between methylation status of five PCGIs (*IGF2*, *SLC16A12*, *SOX11*, *P2RX7* and *MYOD1*) and *LINE1* repetitive element. Statistical analysis was performed using the Spearman correlation analysis.

**Supplementary Table S1: Primer sequences used in pyrosequencing**

See Supplementary File 1

**Supplementary Table S2: Primer sequences used in telomere length measurement**

<b>Assay name</b>	<b>Forward primer</b>	<b>Reverse primer</b>
	sequence	sequence
<i>Telomere</i>	CGGTTTGTTGGGTTTGGGTT TGGGTTTGGGTTTGGGTT	GGCTTGCCTTACCCTTACCCTTA CCCTTACCCTTACCCT
<i>Beta-globin</i>	GCTTCTGACACAACCTGTGTTCACTAGC	CACCAACTTCATCCACGTTCCACC