

Epistatic SNP interaction of *ERCC6* with *ERCC8* and their joint protein expression contribute to gastric cancer/atrophic gastritis risk

SUPPLEMENTARY MATERIALS

SUPPLEMENTARY TABLES

Supplementary Table 1: Effects of a single SNP on ERCC6 and ERCC8 expression

| SNP genotypes | | Positive (%) | P value |
|--|-----------------|--------------|---------|
| <i>ERCC6</i> SNP and protein expression | | | |
| rs1917799 | TT | 44(83.0) | ref. |
| | GT | 41(95.3) | 0.059 |
| | GG | 12(92.3) | 0.403 |
| | GT/GG:TT | 53(94.6) | 0.053 |
| | GG:GT/TT | 85(88.5) | 0.684 |
| <i>ERCC8</i> SNP and protein expression | | | |
| rs158572 | AA | 82(91.1) | ref. |
| | GA | 16(88.9) | 0.767 |
| | GG | 1(100.0) | 0.755 |
| | GA/GG:AA | 17(89.5) | 0.822 |
| | GG:GA/AA | 98(90.7) | 0.750 |
| rs158916 | TT | 75(91.5) | ref. |
| | CT | 21(87.5) | 0.559 |
| | CC | 3(100.0) | 0.597 |
| | CT/CC:TT | 24(88.9) | 0.688 |
| | CC:CT/TT | 96(90.0) | 0.577 |

Supplementary Table 2: The three dimensions interactions of the *ERCC6* SNP-*ERCC8* SNP- *H. pylori* infection status with the risk of GC/CAG

| SNP genotypes | | | CAG vs. CSG | | GC vs. CSG | |
|--|-------|-----|----------------------------------|-----------------|----------------------------------|-----------------|
| | | | P | OR(95%CI) | P | OR(95%CI) |
| <i>ERCC6</i> rs1917799-<i>ERCC8</i> rs158572-<i>H. pylori</i> | | | | | | |
| TT | AA | (-) | | 1(ref) | | 1(ref) |
| TT | AA | (+) | <0.001 | 5.32(3.51-8.05) | <0.001 | 4.99(3.01-8.08) |
| TT | AG+GG | (-) | 0.319 | 1.41(0.72-2.74) | 0.046 | 2.16(1.01-4.61) |
| TT | AG+GG | (+) | <0.001 | 7.74(2.80-21.4) | <0.001 | 12.5(3.99-39.2) |
| GTGG | AA | (-) | 0.507 | 0.90(0.65-1.23) | 0.090 | 1.39(0.95-2.04) |
| GTGG | AA | (+) | <0.001 | 5.14(3.62-7.30) | <0.001 | 5.49(3.58-8.40) |
| GTGG | AG+GG | (-) | 0.058 | 0.58(0.33-1.02) | 0.147 | 1.54(0.86-2.74) |
| GTGG | AG+GG | (+) | 0.001 | 2.74(1.52-4.94) | <0.001 | 3.83(1.96-7.47) |
| | | | $P_{\text{interaction}} = 0.771$ | | $P_{\text{interaction}} = 0.466$ | |
| | | | interaction index=0.80 | | interaction index=0.55 | |
| <i>ERCC6</i> rs1917799-<i>ERCC8</i> rs158916-<i>H. pylori</i> | | | | | | |
| TT | TT | (-) | | 1(ref) | | 1(ref) |
| TT | TT | (+) | <0.001 | 5.52(3.53-8.64) | <0.001 | 5.40(3.12-9.35) |
| TT | CT+CC | (-) | 0.812 | 1.07(0.62-1.86) | 0.266 | 1.44(0.86-2.72) |
| TT | CT+CC | (+) | <0.001 | 4.94(2.51-9.72) | <0.001 | 5.62(2.58-12.2) |
| GTGG | TT | (-) | 0.491 | 0.89(0.64-1.24) | 0.041 | 1.52(1.02-2.28) |
| GTGG | TT | (+) | <0.001 | 4.66(3.23-6.71) | <0.001 | 5.39(3.45-8.41) |
| GTGG | CT+CC | (-) | 0.038 | 0.59(0.36-0.97) | 0.909 | 0.97(0.56-1.68) |
| GTGG | CT+CC | (+) | <0.001 | 4.01(2.36-6.79) | <0.001 | 3.82(2.01-7.25) |
| | | | $P_{\text{interaction}} = 0.459$ | | $P_{\text{interaction}} = 0.517$ | |
| | | | interaction index=1.54 | | interaction index=1.54 | |

CSG: chronic superficial gastritis; CAG: chronic atrophic gastritis; GC: gastric cancer.