

Glutathione S-transferases genes variants and chemotherapy efficacy in gastrointestinal cancer patients: a meta-analysis based on 50 pharmacogenetic studies

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Additional supporting information

Table S1 ORR, PFS, and TTP of chemotherapy in GIC patients with *GSTP1* IIe105Val variants

| Tumor types | Author | Year | ORR (Good + Poor) | | | PFS/TTP (HR) | | | OS time (HR) | | |
|-------------|----------------------|------|-------------------|-----------|-----------|-------------------|------------------|------------------|-----------------------------|------------------|------------------|
| | | | IIe/IIe | IIe/Val | Val/Val | IIe/IIe | IIe/Val | Val/Val | IIe/IIe | IIe/Val | Val/Val |
| GC | Meulendijks D | 2016 | 146(80+66) | | 27(16+11) | reference | | 0.9(0.6–1.4) | reference | | 0.9(0.6–1.4) |
| GC | Liu R | 2016 | 75(64+11) | 31(24+7) | 1(1+0) | reference | 1.24(0.78–1.96) | | reference | 1.33(0.74–2.37) | |
| GC | Liang J | 2010 | | | | 2.19(1.13–3.66)* | reference | | 2.19(1.31–2.74) | reference | |
| GC | Li QF | 2010 | 44(17+27) | 41(29+12) | | 2.36 (1.36–4.09)* | reference | | 2.27 (1.28–4.02) | reference | |
| GC | Shim HJ | 2010 | 133(46+87) | 60(26+34) | 7(5+2) | reference | 1.02 (0.73–1.42) | 0.84 (0.39–1.84) | reference | 1.12 (0.79–1.58) | 0.76 (0.33–1.77) |
| GC | Huang ZH | 2009 | | | | | | | reference | 2.13 (1.14–3.97) | |
| GC | Ott K | 2008 | 55(12+43) | 59(14+45) | 18(7+11) | | | | | | |
| GC | Goekkurt E | 2006 | 30(7+23) | 12(2+10) | 6(4+2) | | | | | | |
| GC | Seo BG | 2009 | 47(10+37) | 38(18+20) | | | | | | | |
| GC | Goekkurt E | 2009 | 64(26+38) | 47(19+28) | 22(7+15) | | | | | | |
| GC | Ruzzo A | 2006 | 87(20+67) | 70(38+32) | 18(12+6) | | | | | | |
| GC | Keam B | 2008 | 44(22+22) | 29(10+19) | | reference | 1.24(0.75–2.06)* | | reference | 0.62(0.45–1.61) | |
| GC | Ji M | 2013 | 23(3+20) | 21(4+17) | 15(11+4) | | | | | | |
| GC | JI YU ZHI | 2011 | 48(22+26) | 25(9+16) | 7(3+4) | | | | | | |
| CRC | Kap EJ | 2014 | | | | | | | reference, 0.78 (0.38–1.57) | 0.82 (0.39–1.73) | 0.60 (0.14–2.62) |
| CRC | Lai CY | 2013 | | | | | | | reference, 1.10(0.85–1.43) | 1.10(0.84–1.44) | 1.12(0.57–2.20) |
| CRC | Li HY | 2012 | | | | | | | reference | 0.85(0.58–1.28) | 0.44(0.18–0.98) |
| CRC | Fariña Sarasqueta(1) | 2011 | | | | | | | reference | 2.1(0.84–5.25) | 1.57(0.31–7.9) |

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|-----|----------------------|------|------------|------------|-----------|------------------|-------------------|-------------------|---------------------------|------------------|------------------|
| CRC | Fariña Sarasqueta(2) | 2011 | | | | | | | reference | 0.9(0.35-2.28) | 0.41(0.05-3.15) |
| CRC | Stoehlmacher J | 2004 | | | | reference | 1.22 (0.56-2.67)* | 2.13 (0.95-4.76)* | reference | 1.82 (0.71-4.66) | 2.96 (1.15-7.61) |
| CRC | Stoehlmacher J | 2002 | | | | | | | reference | 0.47 (0.27-0.81) | 0.16 (0.04-0.63) |
| CRC | Jones BA | 2009 | | | | | | | reference,0.35(0.16-0.79) | 0.50(0.20-1.23) | 0.23(0.08-0.68) |
| CRC | Bohanes P | 2015 | | | | | | | reference | 0.83(0.65-1.05) | 1.46(1.01-2.11) |
| CRC | Boige V | 2010 | | | | reference | 1.09(0.84-1.39) | 1.12(0.78-1.61) | | | |
| CRC | Huang MY | 2011 | | | | 2.77(0.34-22.48) | 1.72(0.36-8.22) | reference | | | |
| CRC | Ruzzo | 2007 | | | | reference | 0.99(0.60-2.14) | 1.15(0.70-2.01) | | | |
| CRC | Kumamoto K | 2013 | 44(14+30) | 16(11+5) | 2(1+1) | | | | | | |
| CRC | Nishina T | 2013 | 49(27+22) | 19(8+11) | | | | | | | |
| CRC | Paez D | 2011 | 45(30+15) | 47(25+22) | 8(3+5) | | | | | | |
| CRC | Zarate R | 2010 | 20(10+10) | 34(30+4) | 6(5+1) | 2.6(1.3-5.4) | reference | | | | |
| CRC | Le Morvan V(1) | 2007 | 32(15+17) | 15(7+8) | | | | | | | |
| CRC | Le Morvan V(2) | 2007 | 26(12+14) | 27(16+11) | | | | | | | |
| CRC | Chen Jian Guo | 2016 | 41(16+25) | 19(13+6) | | | | | | | |
| CRC | Dong Ning ning | 2014 | 43(20+23) | 31(20+11) | | | | | | | |
| CRC | Ying bei bei | 2009 | 54(14+40) | 48(30+18) | reference | 3.41(2.14-5.43)* | | | | | |
| CRC | HAN lei | 2015 | 51(31+20) | 20(6+14) | | | | | | | |
| CRC | Shen Dong Ya | 2015 | 80(25+55) | 47(26+21) | | | | | | | |
| CRC | Liang Jun | 2009 | 10(6+4) | 112(29+83) | | | | | | | |
| CRC | Joerger M | 2015 | 19(2+17) | 25(8+17) | 11(9+2) | reference | 0.61(0.35-1.06) | 0.24(0.12-0.49) | reference | 0.70(0.40-1.22) | 0.23(0.11-0.48) |
| CRC | Ye Chung Chen | 2009 | 125(47+78) | 41(23+18) | | | | | reference | 2.45(1.30-4.62) | |
| CRC | Hong J | 2011 | 37(16+21) | 14 (8+6) | reference | 0.47(0.21-1.04) | | | | | |

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|-----|-----------|------|------------|-----------|--------|--|--|-----------------|------------------|-----------------|
| CRC | Lamas MJ | 2011 | 35(23+12) | 37 (33+4) | | | | 2.38(0.22–3.79) | 2.12(0.41–11.03) | reference |
| EAC | Rumiato E | 2013 | 32(17+15) | 24(8+16) | 7(4+3) | | | reference | 1.36 (0.70–2.64) | |
| EAC | Wang Y | 2011 | 182(84+98) | 59(19+40) | | | | | | |
| EAC | Gui Yan | 2016 | 97(37+60) | 71(42+29) | | | | | | |
| EAC | Joerger M | 2015 | 30(12+18) | 37(18+19) | 9(7+2) | | | reference | 0.79(0.49–1.28) | 0.71(0.34–1.49) |

HR: hazard ratio; OS, overall survival; TTP, time to progression; PFS, progression-free survival; ORR: objective response rate. EC, esophageal cancer; GC, gastric cancer; CRC, colorectal cancer. * The HR of TTP.

Table S2 ORR, PFS and OS of chemotherapy in GIC patients with *GSTM1/GSTT1* variants

| Tumor types | <i>GSTT1</i> |
|-------------|--------------|
|-------------|--------------|

| | Author | Year | ORR (Good + Poor) | | PFS/HR | | OS/HR | |
|-------------|----------------|------|-------------------|------------|-----------|--------------------|-----------------|------------------|
| | | | Present | Null | Present | Null | Present | Null |
| GC | Meulendijks D | 2016 | 136(80+56) | 37(16+21) | reference | 1.1(0.7–1.6) | reference | 1.1(0.7–1.6) |
| GC | Shim HJ | 2010 | 94(37+57) | 106(40+66) | reference | 0.98(0.72–1.32) | reference | 0.77(0.57–1.06) |
| GC | Ott K | 2008 | 87(24+63) | 23(5+18) | | | | |
| GC | Goekkurt E | 2006 | 12(5+7) | 38(8+30) | | | | |
| GC | Seo BG | 2009 | 36(10+26) | 39(8+31) | | | | |
| GC | Goekkurt E | 2009 | 52(47+5) | 82(64+18) | reference | 0.99(0.69-2.88) | reference | 1.94(1.14 -3.32) |
| GC | Ruzzo A | 2006 | 154(64+90) | 21(6+15) | | | | |
| CRC | Kap EJ | 2014 | | | | | reference | 1.25 (0.89–1.75) |
| CRC | Kumamoto K | 2013 | 3(13+17) | 32(13+19) | | | | |
| CRC | Lai CY | 2013 | | | | | 1.15(0.90–1.46) | reference |
| CRC | Zarate R | 2010 | 46(34+12) | 14(11+3) | | | | |
| CRC | Stoehlmacher J | 2004 | | | reference | 0.98 (0.60, 1.60)* | reference | 1.33 (0.78-2.28) |
| CRC | Stoehlmacher J | 2002 | | | | | reference | 0.94 (0.61-1.87) |
| CRC | Ruzzo | 2007 | | | reference | 1.23(0.50-3.40) | | |
| CRC | Boige V | 2010 | | | reference | 1.26(0.92-1.72) | | |
| EC | Rumiato E | 2013 | 48(25+23) | 15(4+11) | | | reference | 1.89 (0.83–4.34) |
| Tumor types | <i>GSTM1</i> | | | | | | | |
| | Author | Year | ORR (Good + Poor) | | PFS/HR | | OS/HR | |
| | | | Present | Null | Present | Null | Present | Null |
| GC | Meulendijks D | 2016 | 84(43+41) | 89(53+36) | reference | 1.0(0.7–1.4) | reference | 1.0(0.7–1.4) |
| GC | Shim HJ | 2010 | 76(29+47) | 124(48+76) | reference | 0.98 (0.72–1.33) | reference | 1.10 (0.80–1.51) |

| | | | | | | | | |
|-----|----------------|------|-----------|-----------|-----------|---------------------|-----------|------------------|
| GC | Ott K | 2008 | 60(13+47) | 52(15+37) | | | | |
| GC | Goekkurt E | 2006 | 18(4+14) | 32(9+23) | | | | |
| GC | Huang ZH | 2009 | | | reference | 1.291 (0.774–2.154) | reference | 1.43(0.82–2.47) |
| GC | Seo BG | 2009 | 26(6+20) | 49(12+37) | | | | |
| GC | Goekkurt E | 2009 | 52(26+26) | 82(36+46) | | | | |
| GC | Ruzzo A | 2006 | 97(34+63) | 78(36+42) | | | | |
| CRC | Kap EJ | 2014 | | | | | reference | 0.79 (0.61–1.04) |
| CRC | Kumamoto K | 2013 | 23(9+14) | 39(17+22) | | | | |
| CRC | Zarate R | 2010 | 25(21+4) | 35(24+11) | | | | |
| CRC | Stoehlmacher J | 2004 | | | reference | 1.13 (0.72-1.76)* | reference | 1.14 (0.69-1.88) |
| CRC | Stoehlmacher J | 2002 | | | | | reference | 1.25 (0.68-2.30) |
| CRC | Ruzzo | 2007 | | | reference | 0.88(0.46-1.87) | | |
| CRC | Boige V | 2010 | | | reference | 0.88(0.70-1.11) | | |
| EC | Rumiato E | 2013 | 28(14+14) | 35(15+20) | | | reference | 1.05 (0.54–2.05) |

HR: hazard ratio; PFS, progression-free survival; OS, overall survival; ORR: objective response rate. * The HR of TTP.

Table S3 The association between *GSTP1* Val carriers vs. IIe/IIe and ORR of chemotherapy in GIC patients base on subgroups

| No. of studies | Study groups | Test of association | | | Models | Test of heterogeneity | | | Tau-squared |
|----------------|-----------------------|----------------------------|-------------|--------------|--------|-----------------------|------------------|--------------------|---------------|
| | | OR/HR (95% CI) | Z | P-value | | χ^2 | P-value | I ² (%) | |
| | Genotyping method | | | | | | | | |
| 8 | TaqMan assay | 1.207(0.645-2.259) | 0.59 | 0.556 | R | 34.14 | <0.001 | 76.60% | 0.6621 |
| 14 | PCR-RFLP | 1.960(1.265- 3.039) | 3.01 | 0.003 | R | 35.48 | 0.001 | 60.50% | 0.3908 |
| 3 | Sequenom-MassARRAY | 1.342(0.285- 6.322) | 0.37 | 0.71 | R | 11.97 | 0.003 | 83.30% | 1.5589 |
| 1 | HRM-SNP | 2.724(1.294-5.734) | 2.64 | 0.008 | R | - | - | - | - |
| 5 | DNA sequencing | 1.359(0.590-3.130) | 0.72 | 0.471 | R | 20.22 | <0.001 | 80.20% | 0.689 |
| | Evaluation criterion | | | | | | | | |
| 24 | RECIST | 1.662(1.136-2.431) | 2.62 | 0.009 | R | 78.54 | <0.001 | 73.30% | 0.5737 |
| 5 | NR | 1.482(0.691-3.181) | 1.01 | 0.312 | R | 14.95 | 0.005 | 73.30% | 0.5409 |
| 5 | WHO | 1.277(0.571-2.856) | 0.6 | 0.551 | R | 6.37 | 0.095 | 52.90% | 0.3559 |
| | Chemotherapy regimens | | | | | | | | |
| 29 | Platinum-based | 1.587(1.145-2.201) | 2.77 | 0.006 | R | 101.5 | <0.001 | 72.40% | 0.5538 |
| 2 | Non-platinum | 1.583(0.747-3.354) | 1.2 | 0.231 | R | 0.89 | 0.346 | 0 | 0 |
| | QS | | | | | | | | |
| 23 | High QS | 1.528(1.071-2.181) | 2.34 | 0.019 | R | 78.72 | <0.001 | 72.10% | 0.5162 |
| 8 | Low QS | 1.758(0.894-3.458) | 1.64 | 0.102 | R | 23.44 | 0.001 | 70.10% | 0.6535 |
| | Study types | | | | | | | | |
| 6 | Prospective study | 1.180(0.631-2.208) | 0.52 | 0.065 | R | 11.09 | 0.05 | 54.9% | 0.3244 |
| 25 | Retrospective study | 1.677(1.183-2.378) | 2.9 | 0.004 | R | 87.28 | <0.001 | 72.5% | 0.5471 |

OR, odds ratio; CI, confidence interval; R, random effect model; QS, quality score; PCR-RFLP, PCR-restriction fragment length polymorphism; RECIST, Response Evaluation

Criteria in Solid Tumors; WHO, World Health Organization; NR: not reported.

Figure S1 The sensitivity analysis of pooling ORs of ORR in GIC patients. (a)

ORR of *GSTP1* Val carriers vs. IIe/IIe model; (b) ORR of *GSTP1* Val/Val vs. IIe/IIe model; (c) ORR of *GSTP1* Val/Val vs. IIe carriers model; (d) ORR of *GSTT1* null vs. present; (e) OS of *GSTP1* IIe/Val vs. IIe/IIe model.

Figure S1

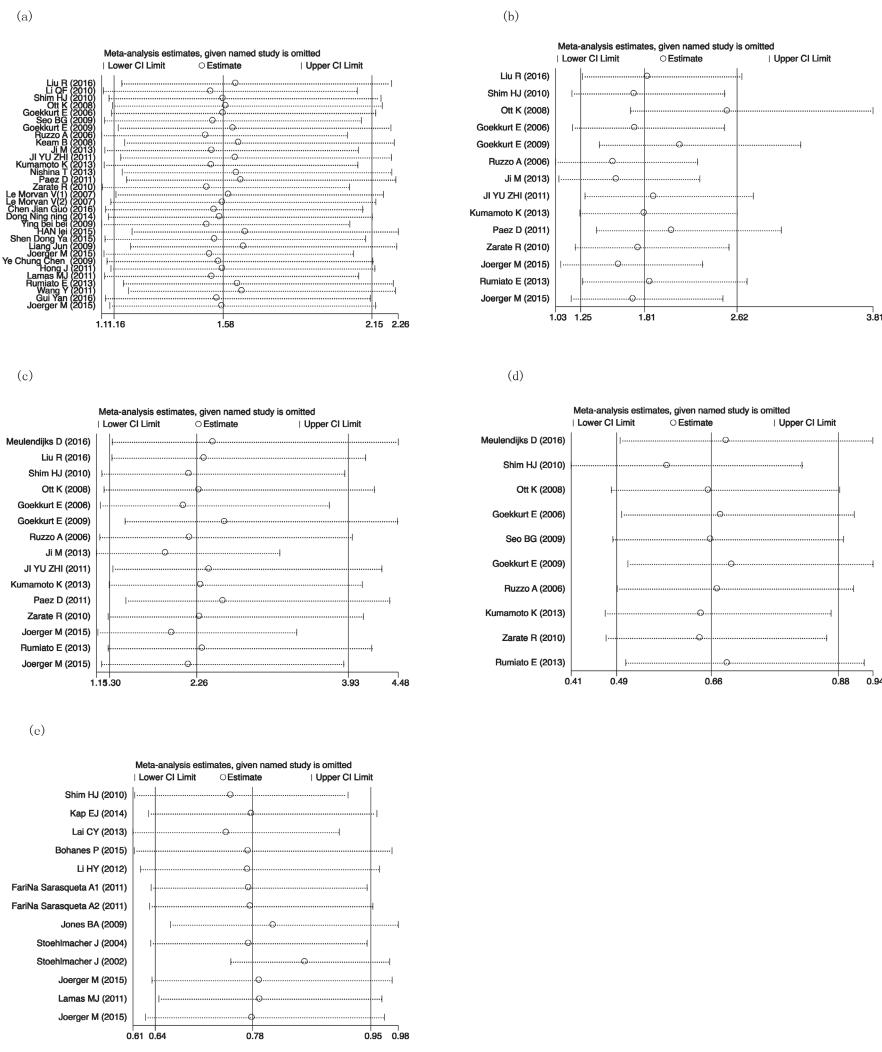


Figure S2 Forest plots of ORR in GIC patients harboring different *GSTP1* IIe105Val variants. OR: odds ratio; SE: standard error. (a) (Val/Val vs. IIe/IIe) stratified by ethnicity under random model; (b) (Val/Val vs. IIe carriers) stratified by ethnicity under random model.

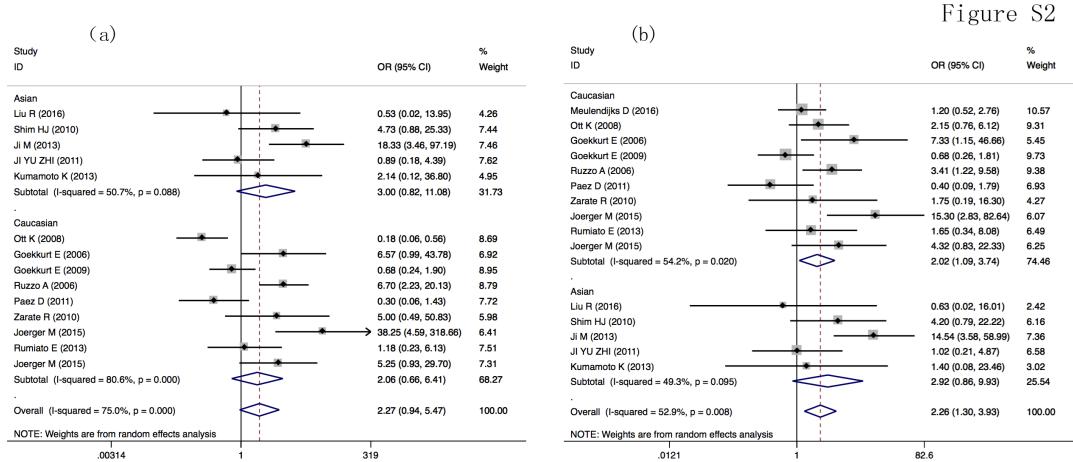


Figure S3 Begg's and Egger's bias plot for publication bias test on the association between the *GSTP1* IIe105Val variant (Val carriers vs. IIe/IIe) and ORR of GIC patients. OR: odds ratio; SE: standard error. (a) Begg's bias plot; (b) Egger's bias.

Figure S3

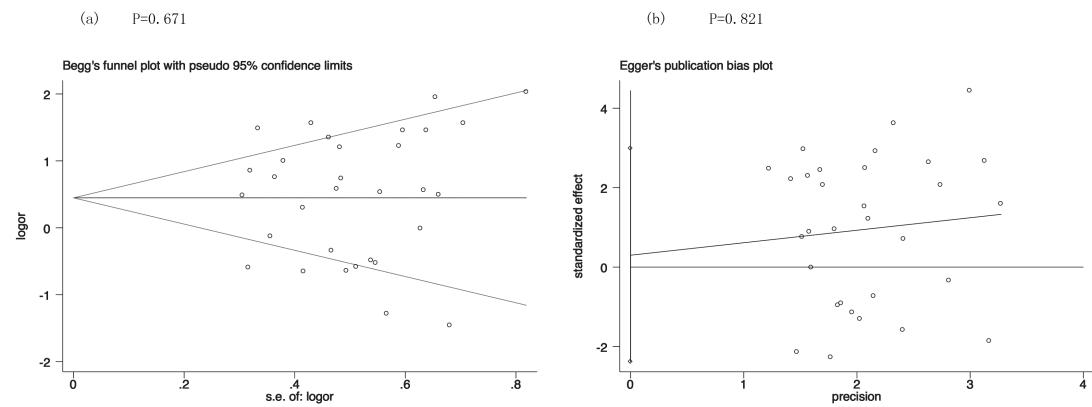


Figure S4 Begg's and Egger's bias plot for publication bias test on the association

between *GSTT1* IIe105Val variant (Val/Val vs. IIe/IIe; Val/Val vs. IIe carriers) and ORR of GIC patients. OR: odds ratio; SE: standard error. (a) Begg's bias plot under Val/Val vs. IIe/IIe model; (b) Egger's bias plot under Val/Val vs. IIe/IIe model; (c) Begg's bias plot under Val/Val vs. IIe carriers model; (d) Egger's bias plot under Val/Val vs. IIe carriers model.

Figure S4

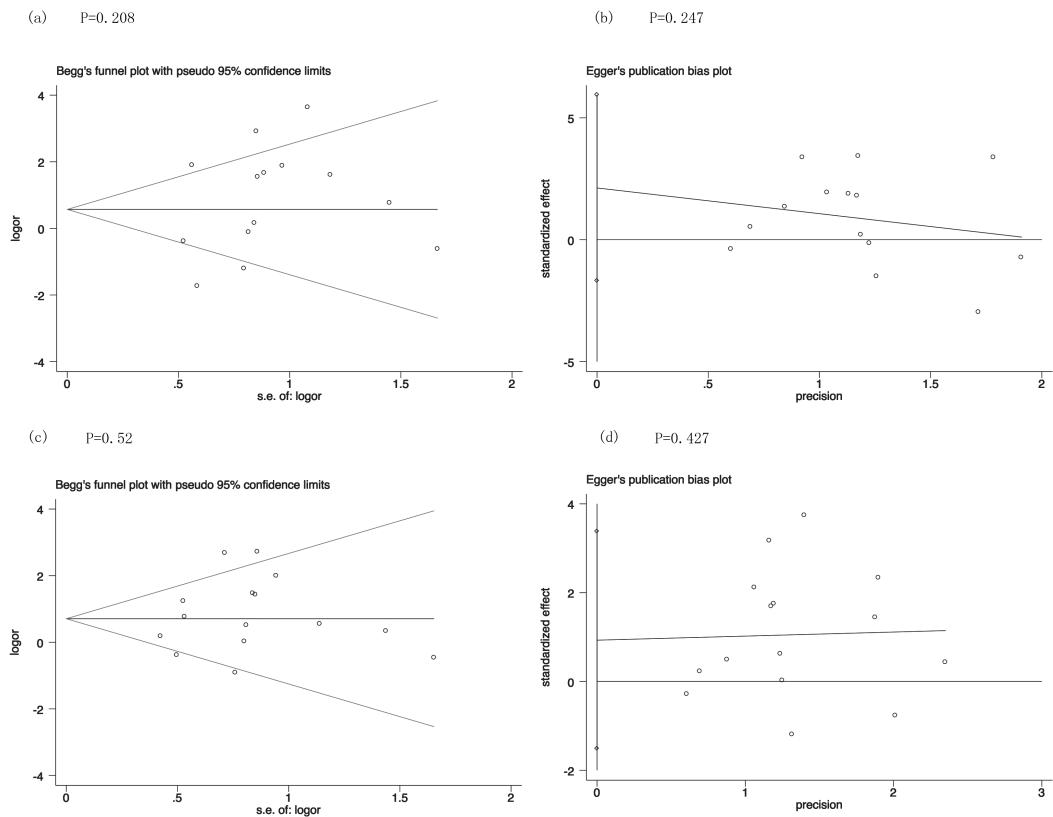


Figure S5 Begg's and Egger's bias plot for publication bias test on the association

between *GSTT1/GSTM1* null or present variant (null vs. present) and ORR of GIC patients. OR: odds ratio; SE: standard error. (a) *GSTT1* Begg's bias plot; (b) *GSTT1* Egger's bias plot; (c) *GSTM1* Begg's bias plot; (d) *GSTM1* Egger's bias plot.

Figure S5

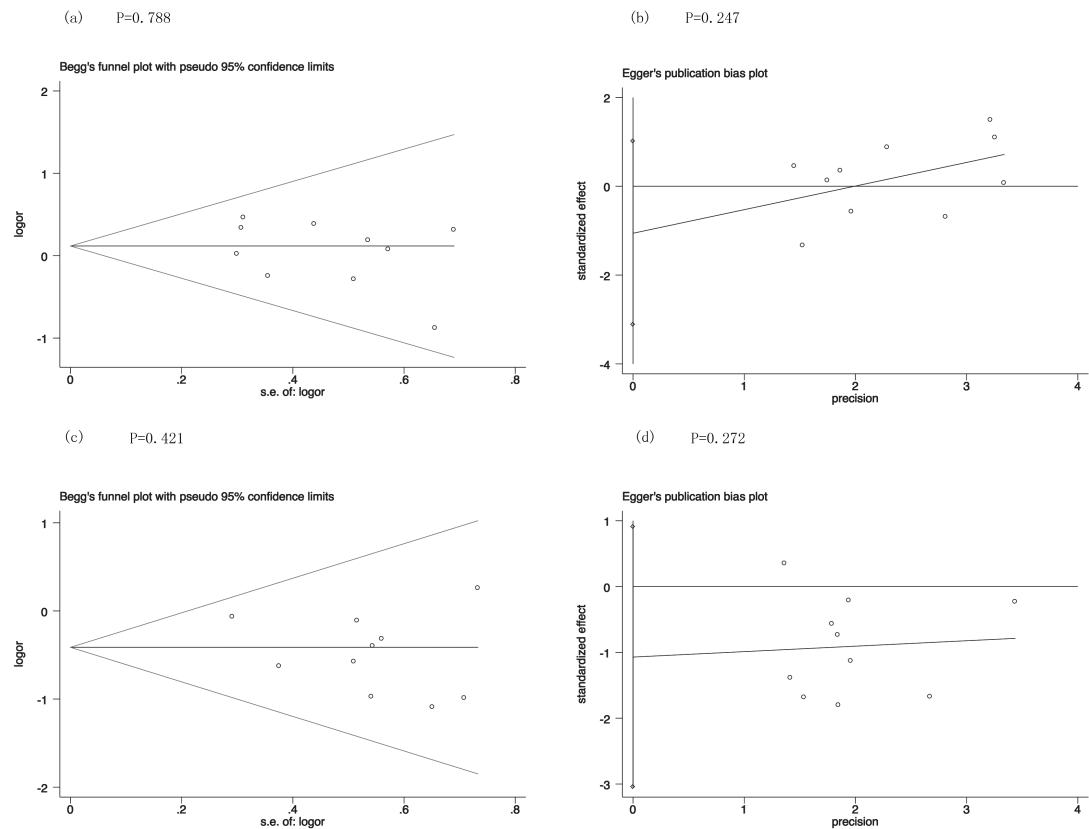


Figure S6 Begg's and Egger's bias plot for publication bias test on the association between *GSTP1* IIe105Val variant (IIe/Val vs. IIe/IIe) and the OS of GIC patients,

GSTP1 IIe105Val variant (Val carriers vs. IIe/IIe) and PFS of GIC patients. OR: odds ratio; SE: standard error. (a) Begg's bias plot under Val/Val vs. IIe/IIe model; (b) Egger's bias plot under Val/Val vs. IIe/IIe model; (c) Begg's bias plot under Val carriers vs. IIe/IIe model; (d) Egger's bias plot under Val carriers vs. IIe/IIe model.

Figure S6

