

Figure S1. Forest plot for the risk of *iceA1* gene and gastric carcinoma compared with gastritis or non-ulcer dyspepsia.

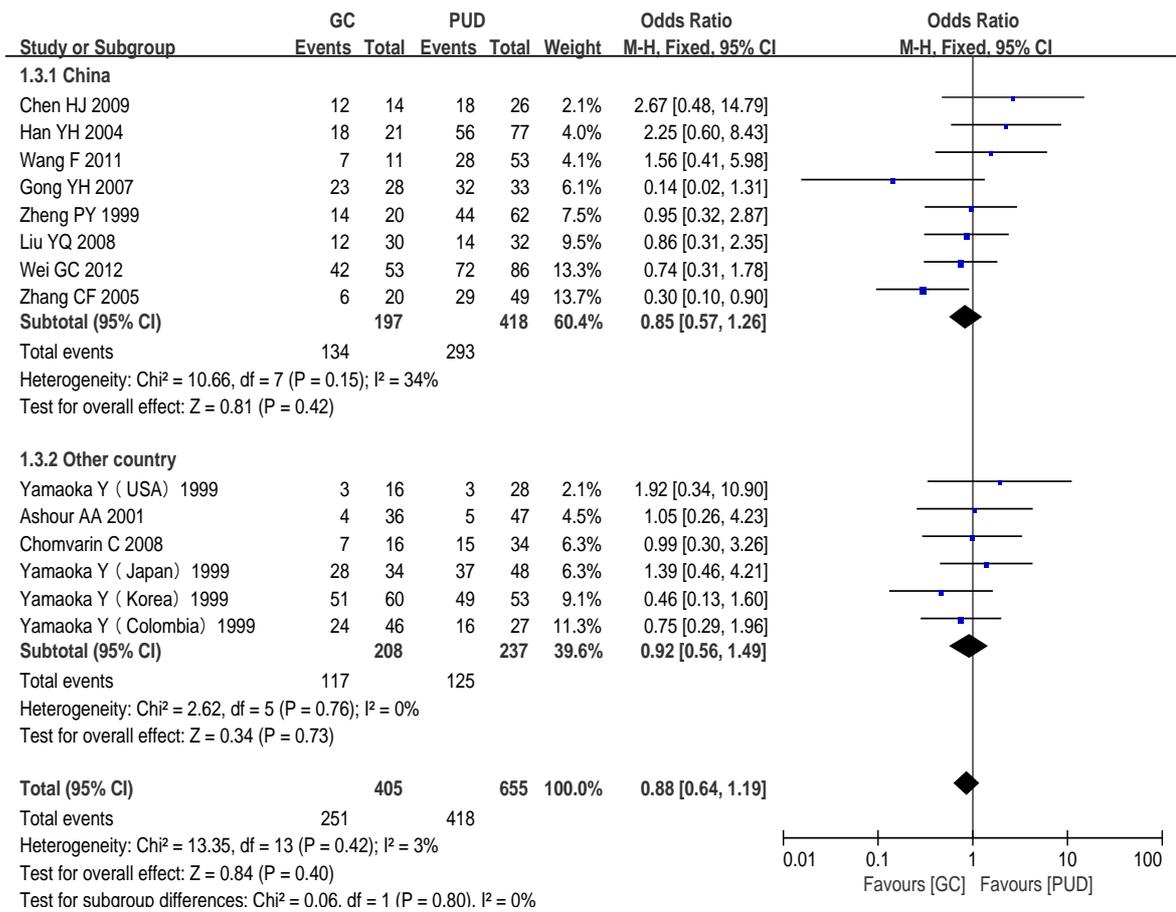


Figure S2. Forest plot for the risk of *iceAI* gene and gastric carcinoma compared with peptic ulcer disease.

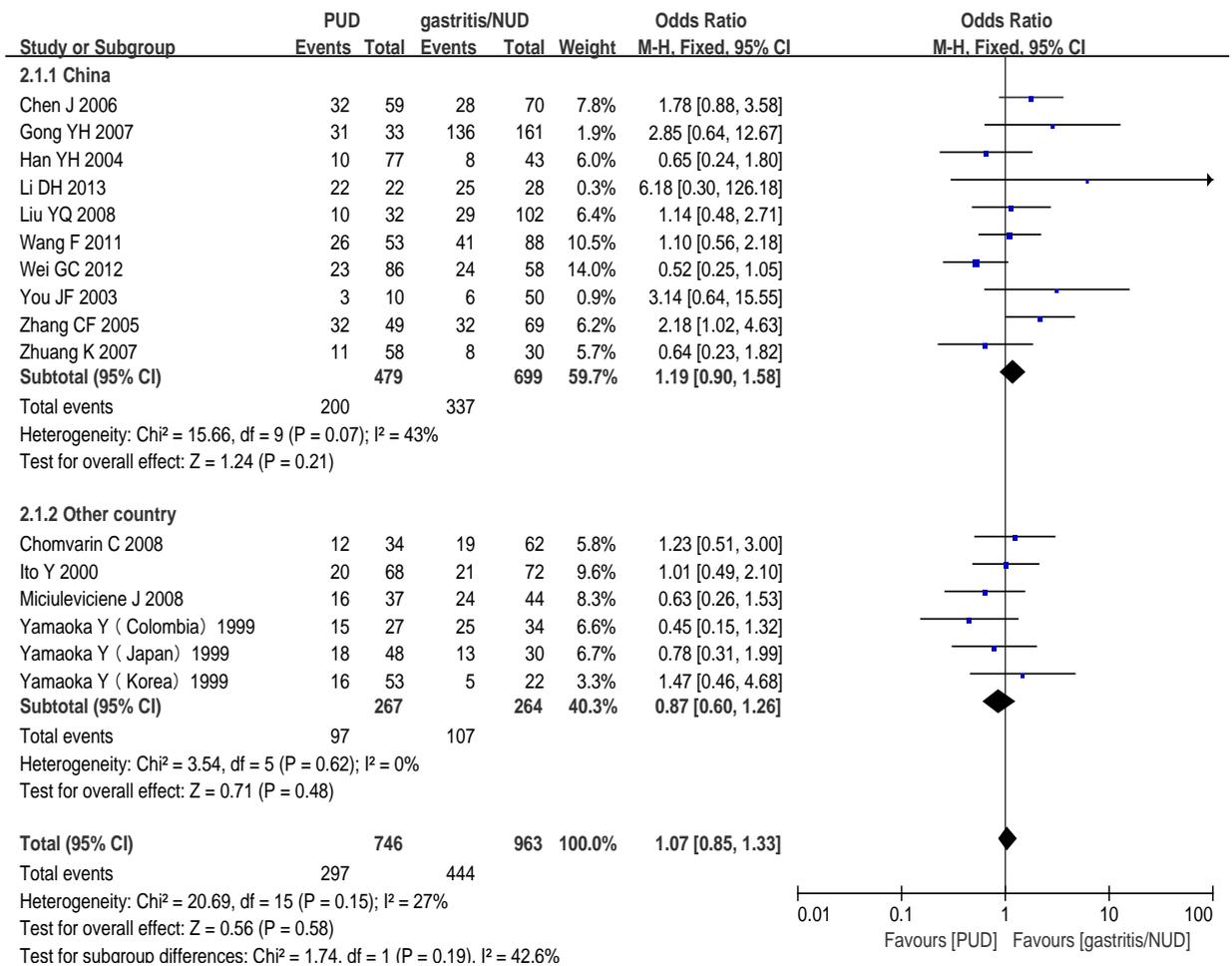


Figure S3. Forest plot for the risk of *iceA2* gene and peptic ulcer disease compared with gastritis or non-ulcer dyspepsia.

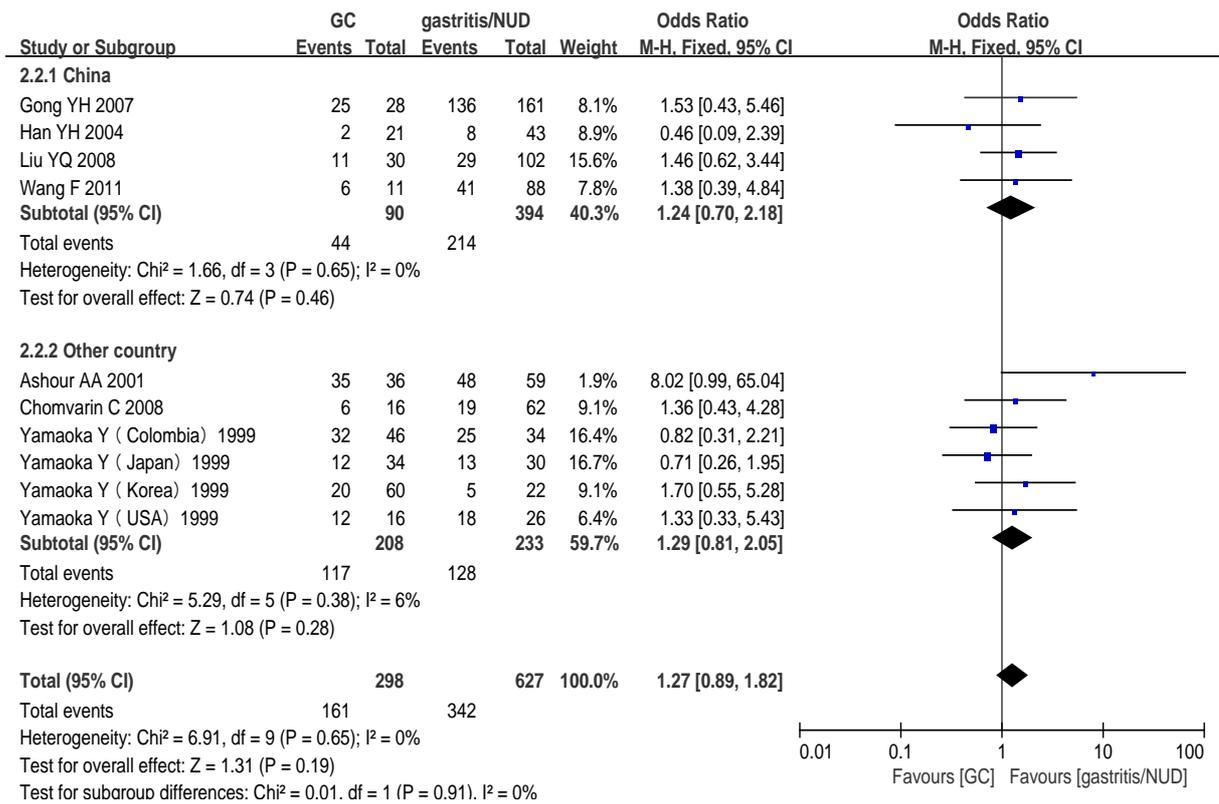


Figure S4. Forest plot for the risk of *iceA2* gene and gastric carcinoma compared with gastritis or non-ulcer dyspepsia.

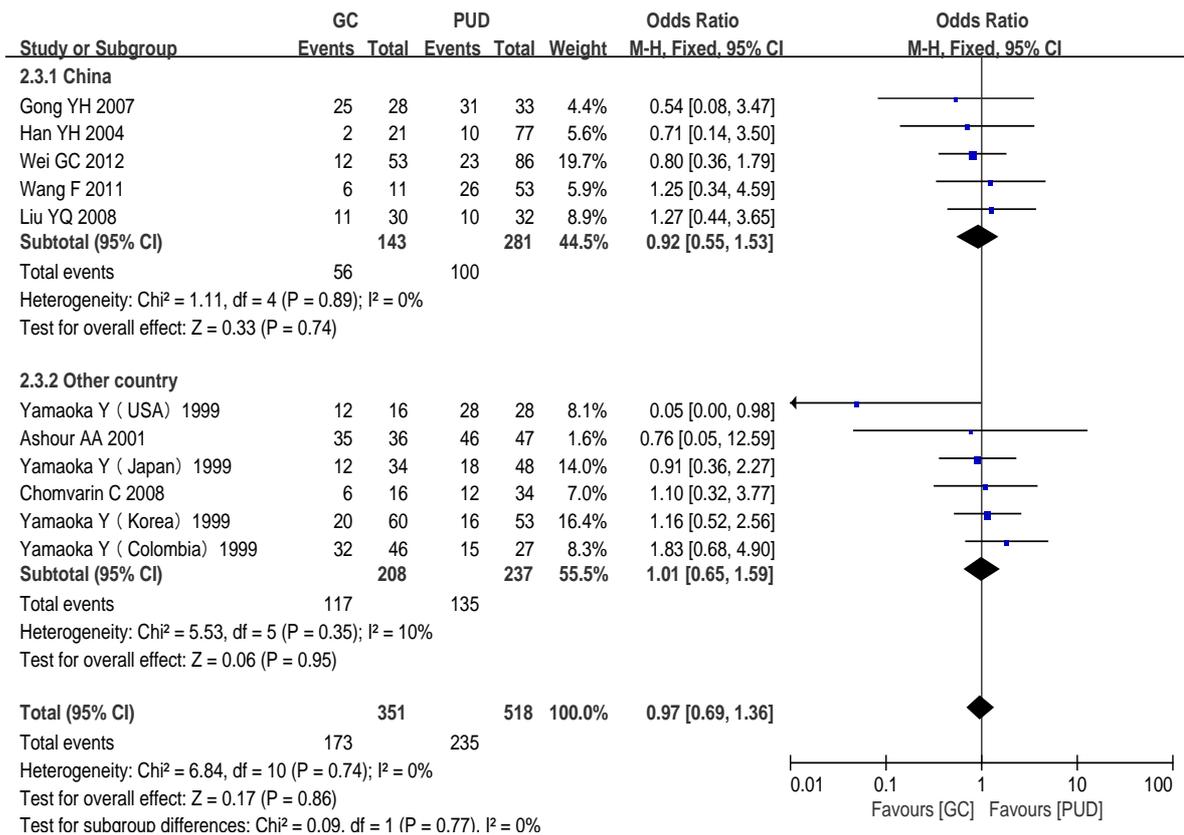


Figure S5. Forest plot for the risk of *iceA2* gene and gastric carcinoma compared with peptic ulcer disease.

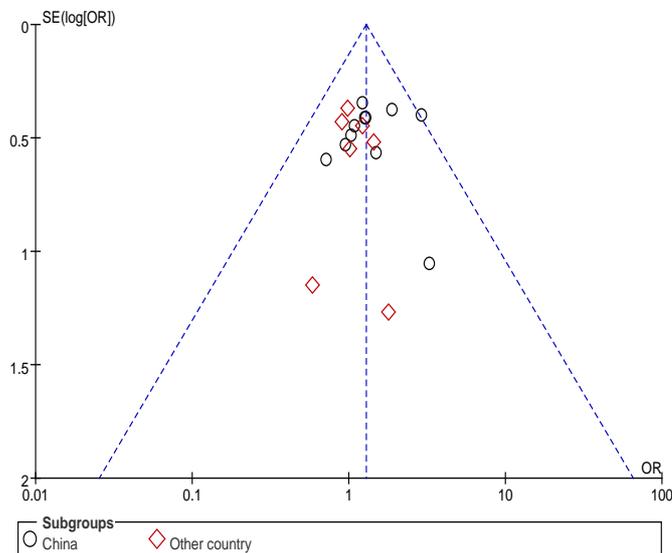


Figure S6. Funnel plot for publish bias assessment of the risk of *iceA1* gene and peptic ulcer disease compared with gastritis or non-ulcer dyspepsia.

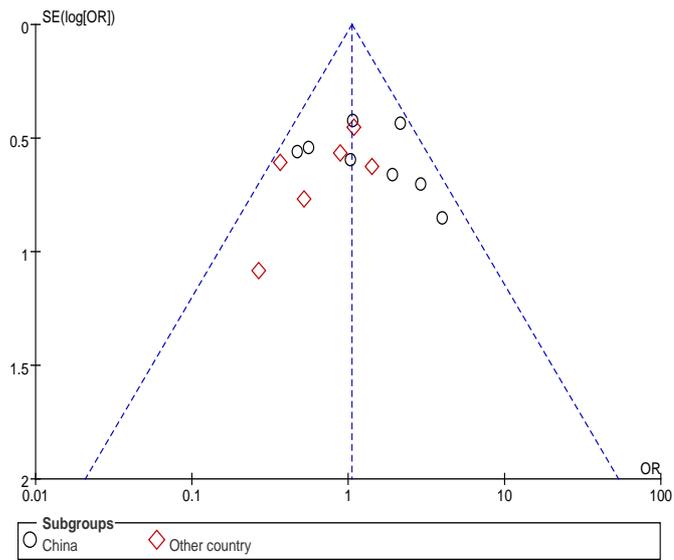


Figure S7. Funnel plot for publish bias assessment of the risk of *iceA1* gene and gastric carcinoma compared with gastritis or non-ulcer dyspepsia.

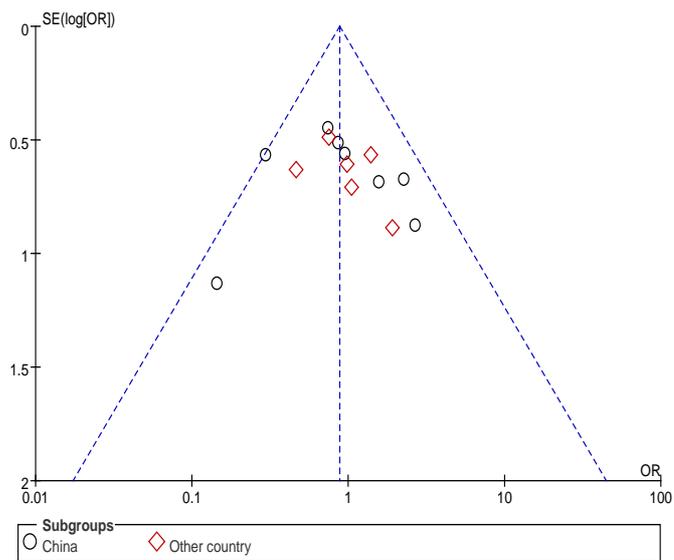


Figure S8. Funnel plot for publish bias assessment of the risk of *iceA1* gene and gastric carcinoma compared with peptic ulcer disease.

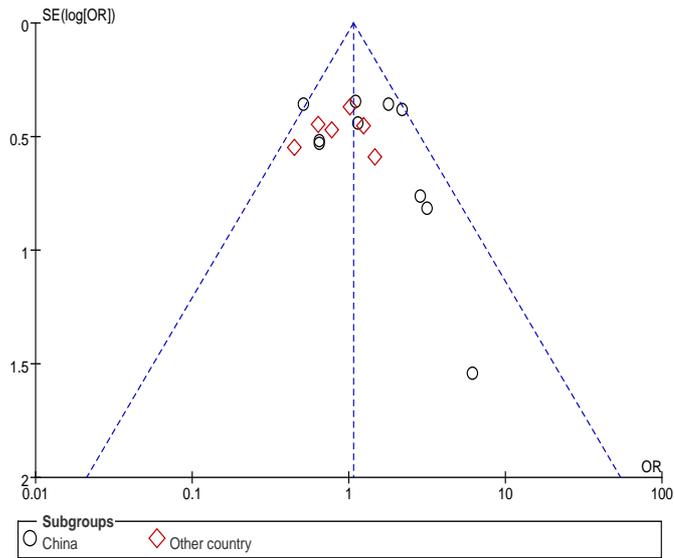


Figure S9. Funnel plot for publish bias assessment of the risk of *iceA2* gene and peptic ulcer disease compared with gastritis or non-ulcer dyspepsia.

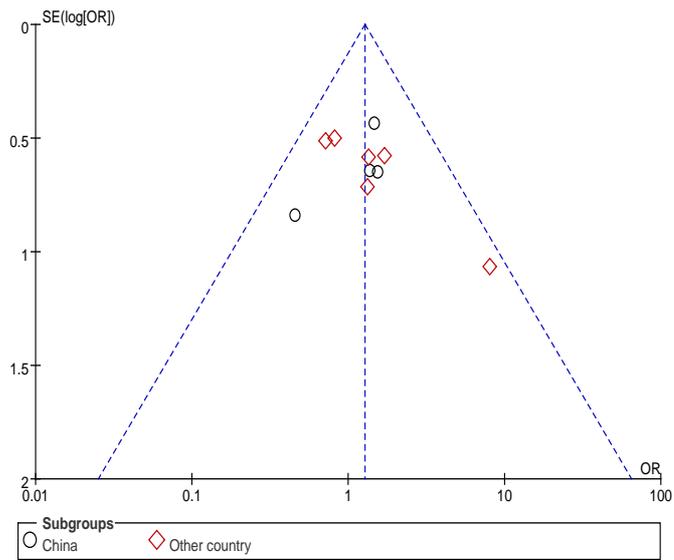


Figure S10. Funnel plot for publish bias assessment of the risk of *iceA2* gene and gastric carcinoma compared with gastritis or non-ulcer dyspepsia.

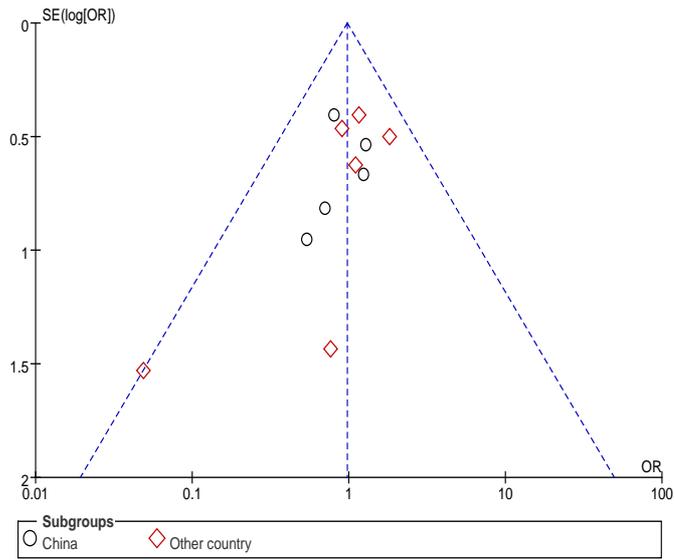


Figure S11. Funnel plot for publish bias assessment of the risk of *iceA2* gene and gastric carcinoma compared with peptic ulcer disease.