

Figure S1. The collection localities of BY, GM and OS used in this study.

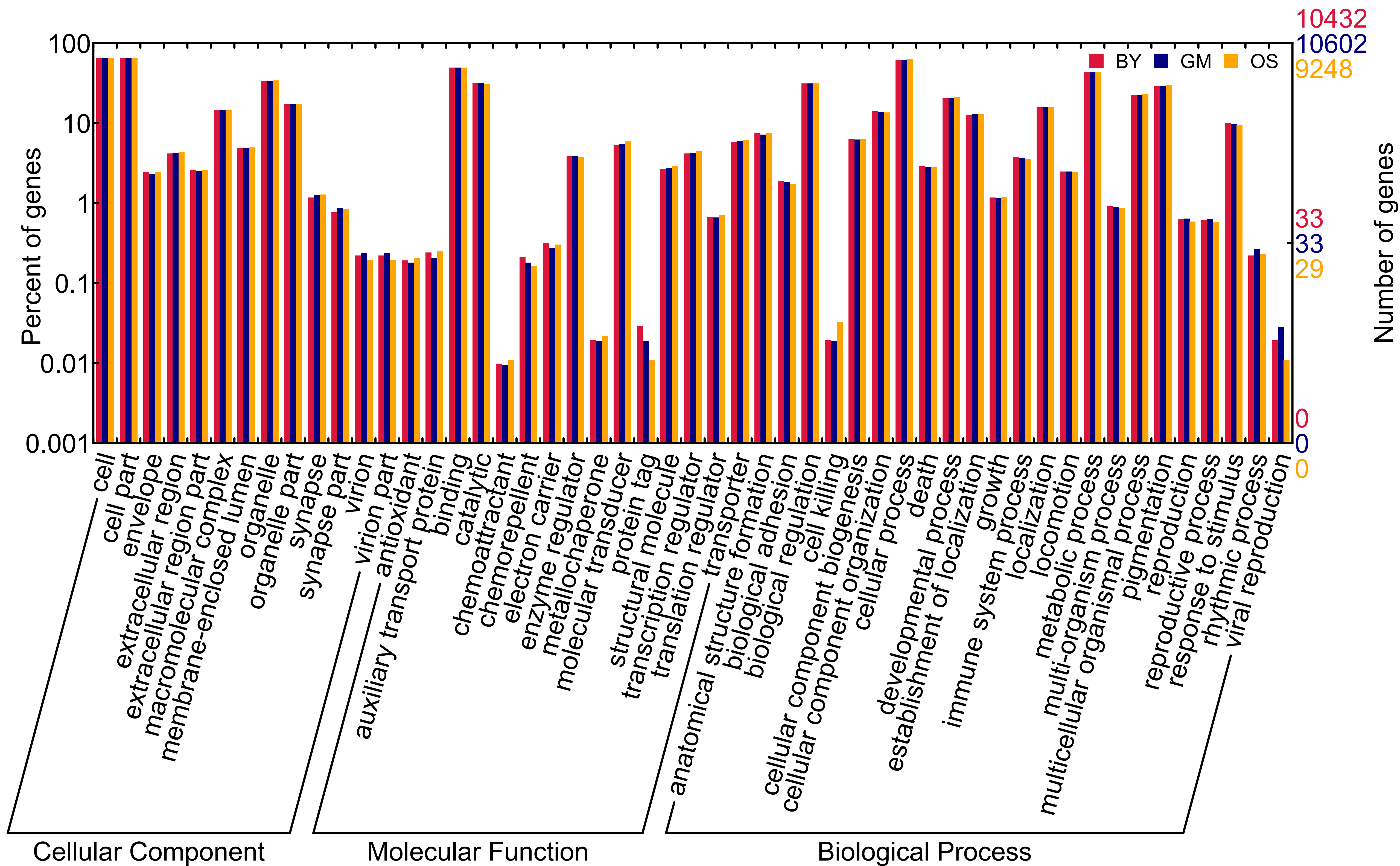


Figure S2. GO classification of annotated genes in the BY, GM, and OS groups.