

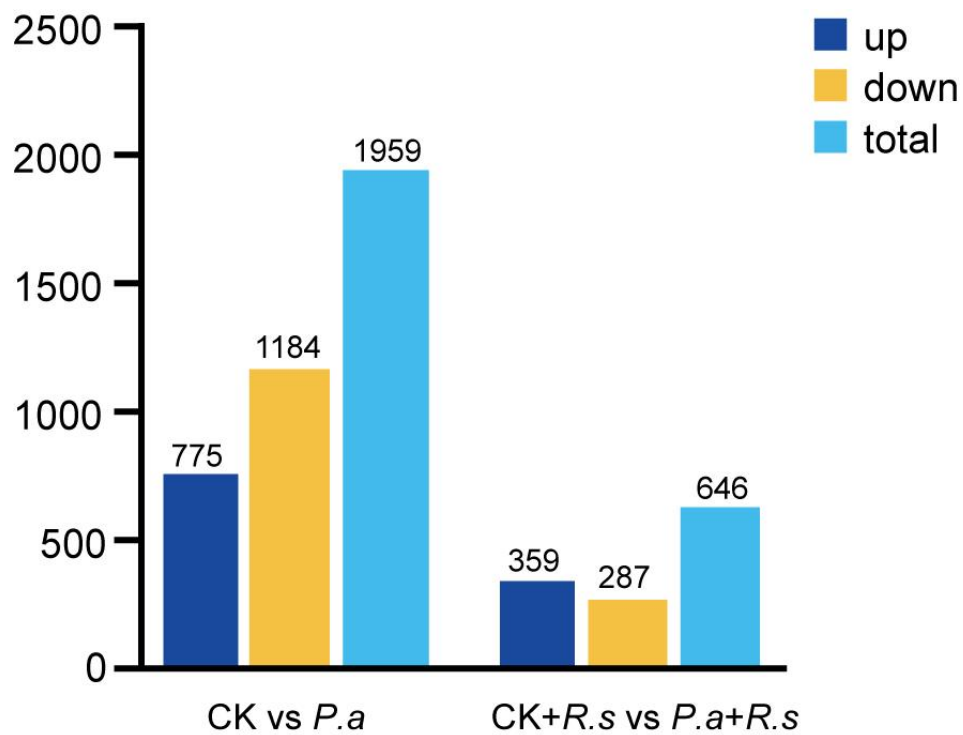


Fig. S5. Identification of differentially expressed genes (DEGs) under different treatments. DEGs were identified in the comparisons of CK vs *P.a* and CK+*R.s* vs *P.a*+*R.s*, based on $|\log_2FC| \geq 1$ and $FDR < 0.05$.