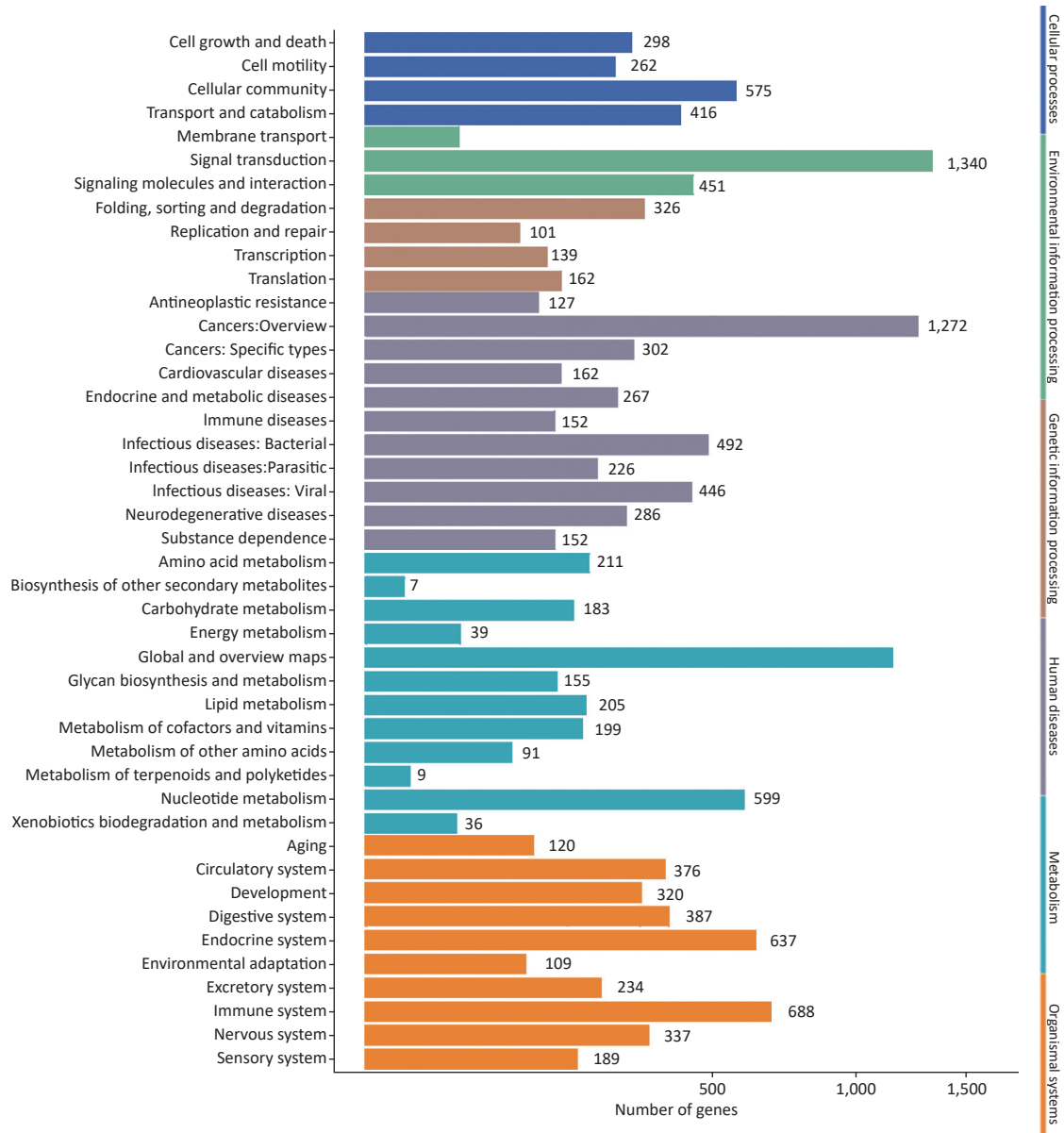


Supplementary Figure S1. GO functional classification of target genes of miRNAs induced by HRV16. The x-axis shows the means number of DEGs (the number is presented by its square root value). The y-axis shows GO terms. All GO terms are grouped into three categories: blue, biological process; green, cellular component; orange, molecular function.



Supplementary Figure S2. KEGG classification of target genes of miRNAs induced by HRV16. The x-axis shows the number of DEGs. The y-axis shows the secondary KEGG pathway terms. All secondary pathway terms are grouped in top pathway terms indicated in different colors.