

# Supplementary for “Transcription Factor Driven Gene Regulation in COVID-19 Patients”

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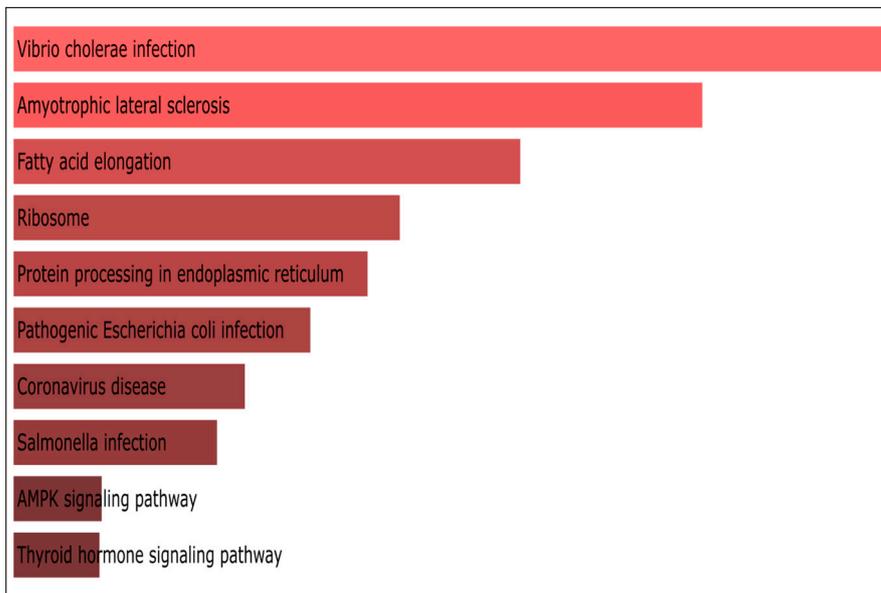
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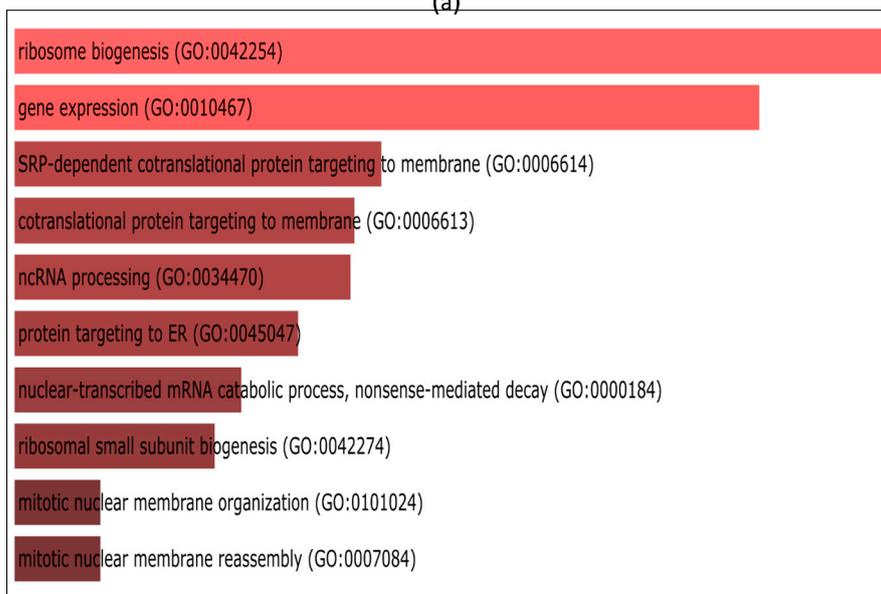
† These authors contributed equally to this work.

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(a)



(b)

Figure S1: Significant KEGG pathways and (b) Significant GO Enrichment Analysis for the 31 identified genes