Supplemental Tables

Supplemental Table S1: ZIKV molecular target analysis.

The canSAR bioinformatics tool was batch analyzed with the ZIKV database of genes. Number of druggable structures, druggable ligands, activities and compounds summary are shown for each gene. Enzyme nature and RNAi data is indicated.

Supplemental Table S: Bioactivity data for the ZIKV database of genes.

The bioactivity data for the ZIKV associated genes is shown. The canSAR compound ID and IC_{50} values are shown for each compound.

Supplemental Table S3: Bioinformatics analysis of the ZIKV associated genes.

The GeneCards Suite was used to batch analyze the ZIKV database of genes. Details on gene summary and function, diseases and disorders and pathways are shown.

Supplemental Table S4: ZIKV lead compounds.

The canSAR compounds tool was used to batch analyze the ZIKV lead targets. The canSAR compound ID and IC_{50} values are shown for each compound.