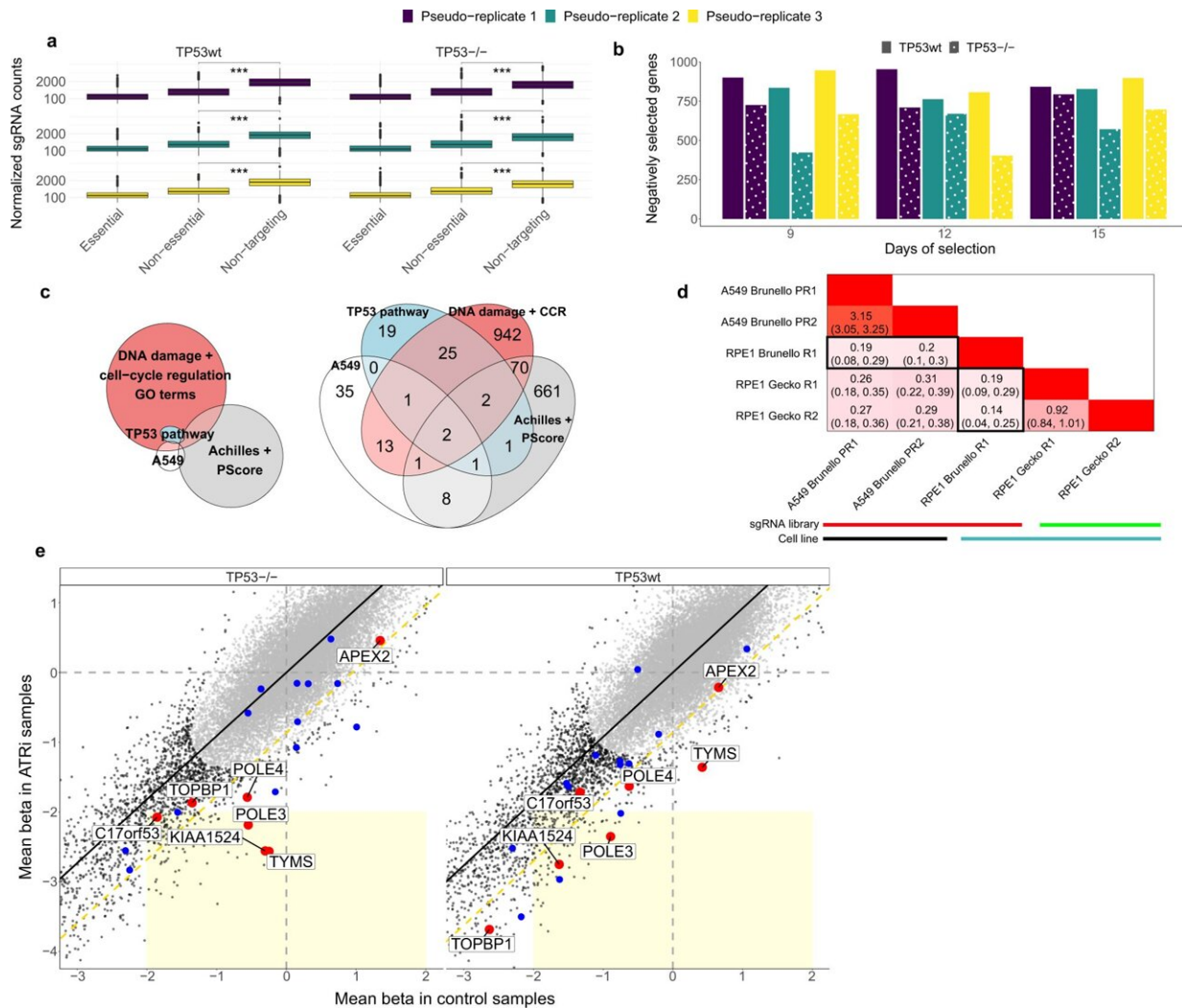


Gene editing via CRISPR/Cas9 can lead to cell toxicity and genome instability

August 9 2022



A TP53 wild-type background can confound estimates of gene selection in genetic screens. a Boxplots showing the pooled normalized sgRNA counts per sample (essential and non-essential genes, and non-targeting sgRNAs; 15 day

samples are shown). Tested using 1-tailed Mann-Whitney. *** denotes a p

Citation: Gene editing via CRISPR/Cas9 can lead to cell toxicity and genome instability (2022, August 9) retrieved 28 April 2024 from <https://phys.org/news/2022-08-gene-crispcas9-cell-toxicity-genome.html>

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