van der Leij et al. Figure S1 - Schematic outline of the screening procedures used in this study.

22,612 human genes

Mitochondria RNAi screen

199,901 introns

1042 SVM hits

207 manual hits

1,249 potential mitotic hits

18 cohesion candidates

8,872 potential SNW1 splice targets

572 validated mitotic genes

784 high confidence SNW1 splice targets

36 SNW1 regulated mitotic genes

ANAPC2

CDCA5

Log2fc ≥ 1.5
coverage ≥ 0.05
sum of reads ≥ 20

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