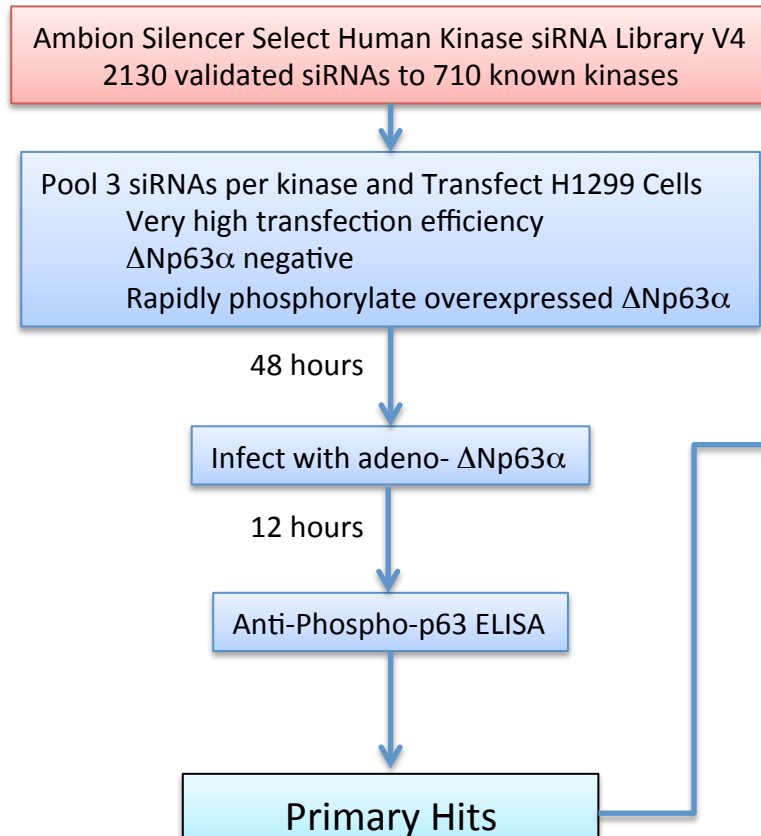


Primary Screen



Gene Symbol	Raw Kinase Data	C.V. Data
PCTK3	2.598	0.127
PIK3CD	2.698	0.149
FRK	2.983	0.127
GALK1	2.984	0.125
TGFBR1	3.039	0.143
CKB	3.053	0.129
PHKG1	3.115	0.134
SH3BP5L	3.115	0.136
NTRK2	3.133	0.13
PRKAA2	3.149	0.132
PLK3	3.197	0.121
GRK6	3.213	0.157
PRPS1L1	3.218	0.14
BTK	3.236	0.13

Secondary Screen

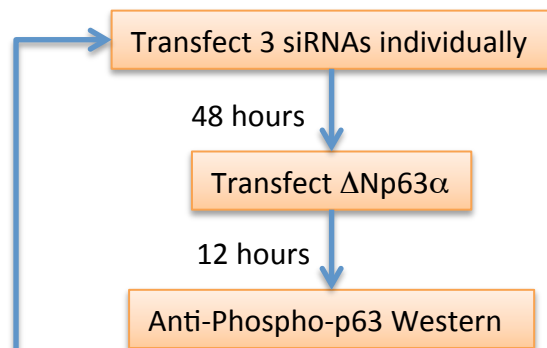


Figure S1: Schematic representation of the siRNA-based screen of the human kinome. The table at the bottom lists the top 14 hits in the screen showing the data produced from fluorescence plate readings of the IF and subsequent Abs₆₀₀ readings for crystal violet staining. Primary hits progressed to the secondary screen and kinases for which all three siRNAs repressed phosphorylation of ΔNp63α were selected.