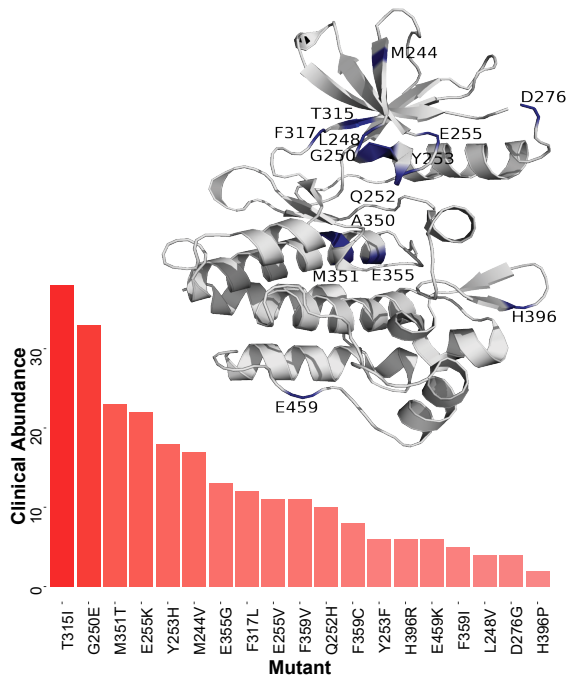
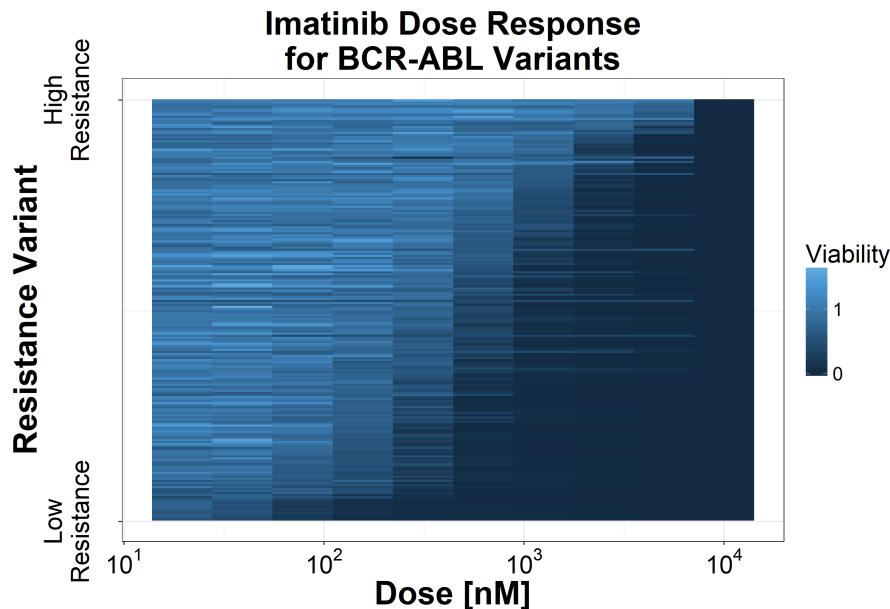


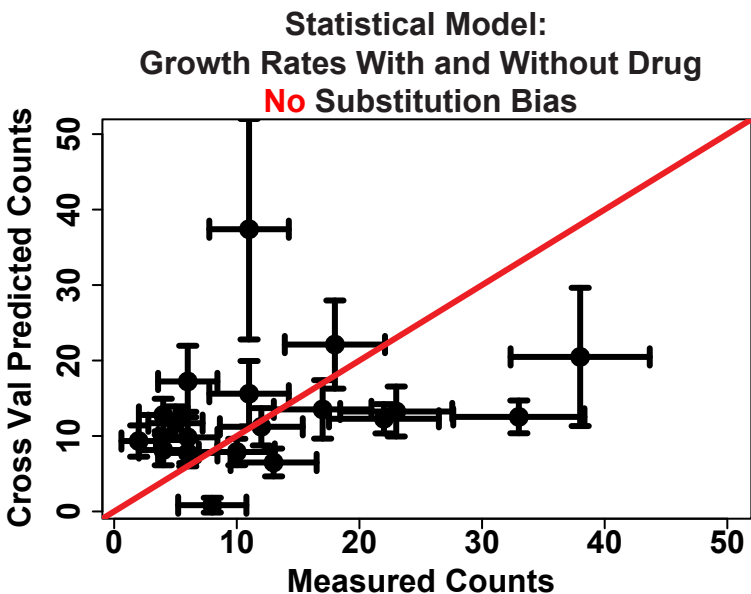
A.



B.



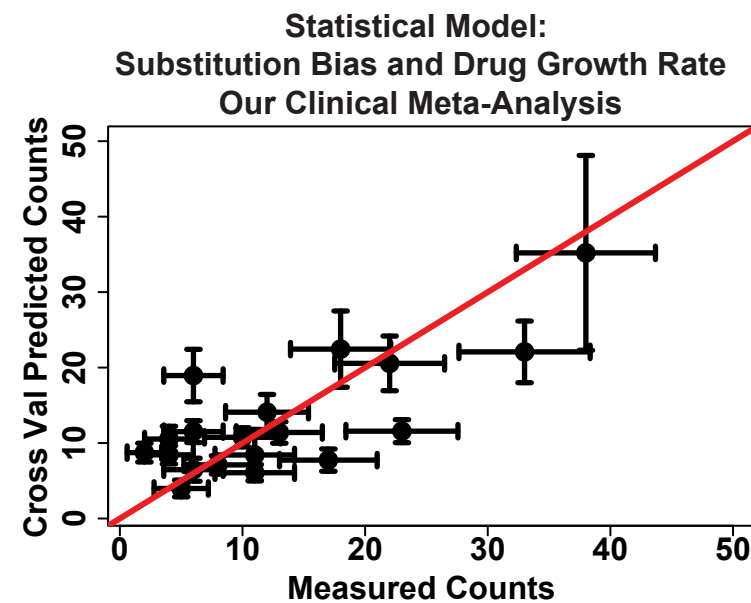
C.



D.

Mutant	IC ₅₀	Nuclueotide Change	Mutation Probability
T315I	8711	C>T	0.183
G250E	3022	G>A	0.133
E255K	1874	G>A	0.133
E459K	803	G>A	0.133
F317L	927	T>C,C>A,C>G	0.077
M351T	1022	T>C	0.055
Y253H	6298	T>C	0.055
M244V	530	A>G	0.038
E355G	1170	A>G	0.038
H396R	1037	A>G	0.038
D276G	673	A>G	0.038
Q252H	1730	G>C, G>T	0.023
L248V	1439	C>G	0.012
H396P	1734	A>C	0.011
F359V	1018	T>G	0.009
F359C	1382	T>G	0.009
E255V	8433	A>T	0.005
Y253F	2181	A>T	0.005
F359I	826	T>A	0.003

E.



F.

