

Table S4 A survey of *rad8* genetic interaction with helicase mutants on different drugs

	genotype	growth	HU	MMS	UV	CPT
no synthetic defects	$\Delta hrp3 \Delta rad8$	-	-	-	-	-
	$\Delta tlh2 \Delta rad8$	-	-	-	-	-
	$\Delta hrp1 \Delta rad8$	-	-	-	-	-
	$\Delta swr1 \Delta rad8$	-	-	-	-	-
	$\Delta rdh54 \Delta rad8$	-	-	-	-	-
	$\Delta rrp1 \Delta rad8$	-	-	-	-	-
	$\Delta rrp2 \Delta rad8$	-	-	-	-	-
	$\Delta SPBC3B8.12 \Delta rad8$	-	-	-	-	-
	$\Delta SPBC15C4.05 \Delta rad8$	-	-	-	-	-
	$\Delta SPBC582.10C \Delta rad8$	-	-	-	-	-
	$\Delta SPAC694.02 \Delta rad8$	-	-	↑\$	-	-
	$\Delta SPCC737.07c \Delta rad8$	-	-	-	-	-
	$\Delta SPAC144.05 \Delta rad8$	-	-	-	-	-
	$cdc21-M68 \Delta rad8$	-	-	-	-	-
	$cdc21-C84 \Delta rad8$	-	-	-	-	-
Increased drug sensitivity	$\Delta rhp26 \Delta rad8$	-	-	↓↓	-	-
	$\Delta fbh1 \Delta rad8$	-	-	↓	↓	-
	$\Delta rqh1 \Delta rad8$	-	↓	↓↓	↓	ND
	$\Delta rad54 \Delta rad8$	↓	↓↓	↓↓	↓	↓
	$\Delta rad55 \Delta rad8$	↓	↓	↓↓	↓↓	↓↓
	$\Delta rad57 \Delta rad8$	↓	↓	↓↓	↓	↓
	$pfh1-R20 \Delta rad8$	-	↑	↓	-	-
	$dna2-K961T \Delta rad8$	-	↓	↓↓	↓	ND
	$dna2^{ls} \Delta rad8$	-	-	↓	↓	-
	$\Delta fml1 \Delta rad8$	-	↓↓	↓↓	↓↓	↓↓
mixed drug sensitivity	$\Delta fml2 \Delta rad8$	-	-	↓	-	-
	$\Delta fml1 \Delta fml2 \Delta rad8$	- elongated	↓↓	↓↓↓	↓↓	↑
	$\Delta chl1 \Delta rad8$	-	-	↓↓	-	↑
	$\Delta srs2 \Delta rad8$	-	↑	↓	-	↓↓
	$\Delta snf22 \Delta rad8$	-	↑	-	-	↓

The level of sensitivity is scored by the fitness on the drug plates. No difference from the either of the single mutants is labeled as "-". The level of increased drug sensitivity is scored by number of "↓". The level of reduced drug sensitivity is scored by number of "↑". ND = not determined. \$ one RNA helicase partially decreased the MMS sensitivity of *rad8*. SPBC3B8.12 = SPBC11C11.11c.