

Table S1. Strains used in this study

Strain	Genotype	Parent	Reference
H99	<i>MAT</i> $\alpha$		[1]
KW137	<i>MAT</i> $\alpha$ <i>CNAG_02589</i> ( <i>BDR1</i> :: <i>NAT</i> #230)	H99	[2]
YSB3785	<i>MAT</i> $\alpha$ <i>CNAG_05216</i> ( <i>RAD53</i> :: <i>NAT</i> #184)	H99	[3]
KW1213	<i>MAT</i> $\alpha$ <i>CNAG_00720</i> :: <i>NAT</i> #43 ( <i>RAD51</i> )	H99	[2]
KW1239	<i>MAT</i> $\alpha$ <i>CNAG_00720</i> ( <i>RAD51</i> :: <i>NAT</i> # <i>pHYG-RAD51-N-HA</i> )	KW1213	This study
KW1038	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> )	H99	This study
KW1039	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> )	H99	This study
KW1040	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> )	H99	This study
KW1276	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> ) <i>pHYG-KU70</i>	KW1038	This study
KW989	<i>MAT</i> $\alpha$ <i>CNAG_03637</i> :: <i>NAT</i> #234 ( <i>KU80</i> )	H99	This study
KW990	<i>MAT</i> $\alpha$ <i>CNAG_03637</i> :: <i>NAT</i> #234 ( <i>KU80</i> )	H99	This study
KW991	<i>MAT</i> $\alpha$ <i>CNAG_03637</i> :: <i>NAT</i> #234 ( <i>KU80</i> )	H99	This study
KW1272	<i>MAT</i> $\alpha$ <i>CNAG_03637</i> :: <i>NAT</i> #234 ( <i>KU80</i> ) <i>pHYG-KU80</i>	KW989	This study
KW1258	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> ) <i>CNAG_00720</i> :: <i>NEO</i> ( <i>RAD51</i> )	KW1038	This study
KW1259	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> ) <i>CNAG_00720</i> :: <i>NEO</i> ( <i>RAD51</i> )	KW1038	This study
KW1260	<i>MAT</i> $\alpha$ <i>CNAG_04220</i> :: <i>NAT</i> #43 ( <i>KU70</i> ) <i>CNAG_00720</i> :: <i>NEO</i> ( <i>RAD51</i> )	KW1038	This study
KW1249	<i>MAT</i> $\alpha$ <i>CNAG_03637</i> :: <i>NAT</i> #234 ( <i>KU80</i> ) <i>CNAG_00720</i> :: <i>NEO</i> ( <i>RAD51</i> )	KW989	This study
KW1250	<i>MAT</i> $\alpha$ <i>CNAG_03637</i> :: <i>NAT</i> #234 ( <i>KU80</i> ) <i>CNAG_00720</i> :: <i>NEO</i> ( <i>RAD51</i> )	KW989	This study

Each *NAT-STM*# indicates the *Nat*<sup>r</sup> marker with a unique signature tag.

## References

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2. Jung, K.W.; Yang, D.H.; Kim, M.K.; Seo, H.S.; Lim, S.; Bahn, Y.S. Unraveling Fungal Radiation Resistance Regulatory Networks through the Genome-Wide Transcriptome and Genetic Analyses of *Cryptococcus neoformans*. *mBio* **2016**, *7*, doi:10.1128/mBio.01483-16.
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