



















































































BROAD ID	Gene deleted	30°C	37°C
KN99α	-		
CNAG_07464	<i>MBS1</i>		
CNAG_05535	<i>FHL1</i>		
CNAG_00883	<i>ECM2201</i>		
CNAG_07506	<i>FAP1</i>		
CNAG_07797	<i>CLR6</i>		
CNAG_04908	<i>CLR4</i>		
CNAG_07924	<i>MCM1</i>		
CNAG_05067	<i>CLR5</i>		
CNAG_01438	<i>SWI6</i>		
CNAG_03378	<i>CLR2</i>		
CNAG_05420	<i>USV101</i>		
CNAG_06276	<i>CEP3</i>		
CNAG_05861	<i>FKH101</i>		
CNAG_03894	<i>PDR802</i>		
CNAG_04353	<i>CLR1</i>		
CNAG_02566	<i>FKH2</i>		
CNAG_00871	<i>CLR3</i>		
CNAG_03202	<i>CAC1</i>		
CNAG_01551	<i>GAT201</i>		
CNAG_04864	<i>CIR1</i>		

BROAD ID	Gene deleted	30°C	37°C
KN99α	-		
CNAG_05431	<i>RIM101</i>		
CNAG_01626	<i>ADA2</i>		
CNAG_07680	<i>HAP5</i>		
CNAG_05222	<i>NRG1</i>		
CNAG_02215	<i>HAP3</i>		
CNAG_03849	<i>ASG1</i>		
CNAG_02774	<i>MAL13</i>		
CNAG_00031	<i>MLR1</i>		
CNAG_06352	<i>BIK1</i>		
CNAG_03279	<i>CCD4</i>		
CNAG_04093	<i>YRM103</i>		
CNAG_06252	<i>CCD6</i>		
CNAG_07435	<i>HAP2</i>		
CNAG_03902	<i>RDS2</i>		
CNAG_00440	<i>SSN801</i>		
CNAG_04345	<i>ARO8001</i>		
CNAG_00732	<i>CCD3</i>		
CNAG_01523	<i>HOG1</i>		
CNAG_02153	<i>TUP1</i>		
CNAG_00570	<i>PKR1</i>	