

<i>CDC11</i> locus	<i>SHS1</i> locus	<i>BNI5</i> locus	SD+AA	SD+AA+5-FOA
<i>CDC11-mC</i>	<i>SHS1</i>	<i>BNI5</i>		
	<i>shs1(ΔCTE)</i>	<i>BNI5</i>		
<i>cdc11(ΔCTE)-mC</i>	<i>SHS1-BNI5-eGFP</i>	<i>BNI5</i>		
	<i>SHS1-BNI5-eGFP</i>	<i>bni5Δ</i>		
	<i>shs1(ΔCTE)-BNI5-eGFP</i>	<i>BNI5</i>		
	<i>shs1(ΔCTE)-BNI5-eGFP</i>	<i>bni5Δ</i>		
	<i>SHS1-eGFP-BNI5</i>	<i>BNI5</i>		
	<i>SHS1-eGFP-BNI5</i>	<i>bni5Δ</i>		
	<i>shs1(ΔCTE)eGFP-BNI5</i>	<i>BNI5</i>		
	<i>shs1(ΔCTE)eGFP-BNI5</i>	<i>bni5Δ</i>		

Figure S4. Endogenous Bni5 is not required for the ability of Shs1-Bni5 translational fusions to rescue growth *cdc11(ΔCTE) shs1(ΔCTE)* cells. Strains GFY-160, GFY-162, GFY-911, GFY-1104, GFY-913 and GFY-1105 were grown overnight and spotted onto medium with or without 5-FOA to select for cells that has lost the covering *URA3*-marked *CDC11*-expressing plasmid. Strains GFY-160, GFY-162, GFY-888, GFY-1102, GFY-890 and GFY-1103 were also tested in the same way.