

Figure S2: Chromosome 2 disomy contributes to adaptation to *BIR1* deletion by increasing the copy number of the *SLI15* gene.

- A) Schematic for the duplication of the SLI15 gene. One copy of SLI15 was added to chromosome 5.
- **B)** 10-fold dilution series of strains on 5-FOA (selecting against *URA3*) and -uracil (selecting for *URA3*) plates. Selection against *URA3* eliminates the copy of *BIR1* present on a minichromosome, resulting in *bir1* \(\triangle \) cells.
- C) Chromosome 2 copy numbers in $bir1\Delta$ -ad strains and $bir1\Delta$ -ad strains with SLI15 duplication prior to adaptation as measured by qPCR. Mean values and the standard errors are shown in red.
- **D)** Doubling times for $bir1\Delta$ -ad strains and $bir1\Delta$ -ad strains with SL115 duplication prior to adaptation as measured by optical density in rich liquid media. Mean values and the standard errors are shown in red.
- **E)** Heat-map visualization of the chromosome copy numbers for the $bir1\Delta$ -ad + SL115 duplication strains. Each of the strains is represented as a row.
- F) Schematic for the relocation of the SLI15 gene, with the sole copy of SLI15 now on chromosome 8.
- **G)** Heat-map visualization of the chromosome copy numbers for the $bir1\Delta$ -ad + SLI15 relocation strains. Each of the strains is represented as a row.
- **H)** Relative colony sizes for strains that contain an extra copy of chromosome 2 and one of the other indicated chromosomes. Values were normalized as in Figure **3E**.
- I) Median colony sizes in square millimeters of double disomic strains on YPAD plates. Normalized values are in Figure 3E.
- **J)** Scatter plot of relative colony sizes obtained from Figures **3E** and **S2H** plotted against the sum of the chromosome sizes of the two induced disomic chromosomes (r = -0.1368, p = 0.48).