



FIGURE S3. Density plots of cell size distributions from deletions of components of the indicated transcription-related complexes (A-D). The deletion mutants of components from each complex are the viable ones queried by (Jorgensen et al. 2002). On the y-axis are the density frequency values, while on the x-axis are the cell size bins, encompassing the cell size values shown in the corresponding spreadsheet associated with this plot (see File S1). The wild type is shown in black in every panel. The density plots shown in grey in (D), are from deletions of genes encoding components of the Mediator (MED) complex that had distributions very similar to wild type, and they corresponded to the following genes: *MED1*, *MED2*, *PGD1*, *NUT1*, *CSE2*, *SSN2*, *GAL11*, *SIN4*, *SRB2*, *SOH1*, *SSN8*.