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Title: Comprehensive analysis of forty yeast microarray datasets reveals a novel subset of

genes (APha-RiB) consistently negatively associated with ribosome biogenesis

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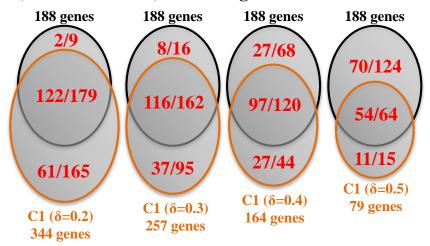
Abu-Jamous, B., Fa, R., Roberts, D. J., & Nandi, A. (2014). Comprehensive analysis of forty yeast microarray datasets reveals a novel subset of genes (APha-RiB) consistently negatively associated with ribosome biogenesis. BMC Bioinformatics, 15, Article 322. https://doi.org/10.1186/1471-2105-15-322

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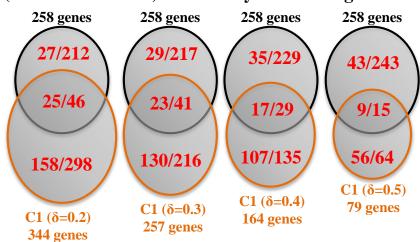
## **Supplementary Figure 2**

C1 overlap with similar clusters in the literature, and the ratios of included genes associated with the "ribosome biogenesis" GO term.

### (Wade et al. 2006) - RRB regulon



#### (Brauer et al. 2008) – Positively correlated genes with growth



### (Roy et al. 2013) – Down-regulated genes with stress

