Citations:

1. Lu, Ma; <https://www.sciencedirect.com/science/article/pii/S0045653520303623>
2. Chen, Guestrin; https://www.sciencedirect.com/science/article/pii/S1470160X21010992
3. Kurokawa, Masaomi; Ying, Bei-Wen (2018). Growth data of the E. coli strains carrying the reduced genomes. figshare. Dataset. <https://doi.org/10.6084/m9.figshare.5918608.v1>
4. Article Source: [**Estimating microbial population data from optical density**](https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0276040)  
   Mira P, Yeh P, Hall BG (2022) Estimating microbial population data from optical density. PLOS ONE 17(10): e0276040. <https://doi.org/10.1371/journal.pone.0276040>
5. Pla ML, Oltra S, Esteban MD, Andreu S, Palop A. Comparison of Primary Models to Predict Microbial Growth by the Plate Count and Absorbance Methods. Biomed Res Int. 2015;2015:365025. doi: 10.1155/2015/365025. Epub 2015 Oct 11. PMID: 26539483; PMCID: PMC4619785.
6. Zwietering MH, Jongenburger I, Rombouts FM, van 't Riet K. Modeling of the bacterial growth curve. Appl Environ Microbiol. 1990 Jun;56(6):1875-81. doi: 10.1128/aem.56.6.1875-1881.1990. PMID: 16348228; PMCID: PMC184525.