Legend:
- Original values from Schmidt et al. 2016
- Li et al. 2014
- 1. Assumes constant protein concentration and cell volumes estimated from Si *et al.*
- 2. Estimates protein concentration from experimental data of DNA, RNA-to-protein ratio, and uses cell volume estimates from Si *et al.*
- 3. Estimates total protein using measurements from Basan *et al.* protein data.

X-axis: growth rate (hr^{-1})

Y-axis: total protein mass (fg per cell)