

Experiment (02) Title: Assessment of Cell Disruption by Ultrasonication

Principle: The rate of protein released by mechanical cell disruptions is usually sound to the proportional to the amount of releasable protein.

Materials required:

- i) A Sonicator
- ii) 2.0 mL centrifuge tubes
- iii) Overnight grown *E. coli* cell suspension
- iv) Spectrophotometer

Procedure:

- i) *E. coli* DH5 α cells were inoculated (3%) in 100 ml LB (Luria-Bertani) media and kept for overnight incubation at 37°C at 120 rpm
- ii) This homogenous suspension of overnight grown cells of *E. coli* DH5 α was made in sterile water
- iii) After this an aliquot of 1.0 mL of *E. coli* DH5 α cell suspension was then taken in 10 separate 2.0 mL centrifuge tubes
- iv) The cell suspension in the centrifuge tubes (Tube No. 1-10) were each subjected to ultrasonication for various time periods (as shown in the table below).
- v) For each time period, ultrasonication was accomplished with 30% amplitude setting and 10 seconds ON and 20 seconds OFF cycle.
- vi) After ultrasonication, for the respective time periods, a 20 μ L aliquot of the sample was aspirated from each tube and added to 480 μ L of water. To each sample, 0.5 mL of Bradford reagent was added and the tubes were incubated in the dark for 10 minutes.
- vii) Subsequently, the absorbance for each sample was recorded at 595 nm

Ultrasonication Duration	Sample Volume (μ L)	Diluent Volume (μ L)	Fold Dilution	Absorbance A_{595}
5.0 minutes	20	480		
3.0 minutes	20	480		
1.0 minute	20	480		

Using a standard Bradford assay (using 2.0 μ g - 18 μ g BSA), a prior calibration plot was generated in order to obtain a calibration equation for estimation of the amount of protein released from each sample. The calibration equation is: $y = 0.0316x$, where y is absorbance measured at 595 nm and x is the amount of protein in microgram (μ g). A prior experiment indicated that the maximum amount of protein released from 1.0 mL of overnight grown cells of *E. coli* DH5 α is 28575 microgram (μ g). Based on the data obtained (as shown in the table), graphically calculate and report the cell disruption constant (in min^{-1}).

Hint: Use the absorbance values, fold dilution and the calibration equation to obtain the amount of protein (in microgram) released from one mL of each sample.