

Effects of β -Methylamino-L-Alanine (BMAA) on *LC4A* Expression and Growth of *Tetrahymena thermophila*

Conclusion

- Cell counts were inconclusive, but overall, *T. thermophila* treated with BMAA were more abundant than the control cells.
- Both rounds of quantitative PCR demonstrated that BMAA exposure resulted in no major change in the expression of *LC4A*.
- There was an increase in fold change in expression of *LC4A*, as shown in Figure 3, but it was not statistically significant ($p=0.332$).
- The results disagree with our hypothesis regarding both gene expression and cell growth.

References

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Audrey Yaeger and Mackenna Landis
Department of Biology, Carroll College

Introduction

- The unicellular, eukaryotic ciliate *Tetrahymena thermophila* (*T. thermophila*) is a popular model organism in molecular biology.
- It has been proposed that β -Methylamino-L-alanine (BMAA), a non-proteinogenic amino acid secreted by Cyanobacteria, is a hyper-excitatory neurotoxin.
- If hyper-excitation of cells were to occur, a strong influx of calcium ions would induce unregulated cell signaling, which would need to be modulated by a Calmodulin protein in order to rescue homeostatic cell processes and prevent cell death.
- *LC4A* is a calmodulin homolog which sequesters calcium ions.
- **Hypothesis:** It is hypothesized that presence of BMAA in the media will decrease *Tetrahymena thermophila* growth and increase *LC4A* expression in order to help regulate calcium ion concentrations within the cell.

Methods

- **Primer synthesis:** Primers for *LC4A* were designed using Oligoanalyzer software.
- **Culturing:** *T. thermophila* cultures were maintained in NEFF media. Upon experimentation, all cultures were transferred into SPP media and experimental cultures were exposed to a single three-day dose of 1.0 mM BMAA.
- **RNA extraction:** RNA was extracted using Qiagen's RNeasy Mini Kit.
- **Reverse transcription:** cDNA was synthesized using RevertAid.
- **Quantitative PCR** was performed using PowerUp SYBR Master Mix. *BTU1* gene expression was used as positive control.
- **Cell Counts** were performed with hemocytometers.

Results

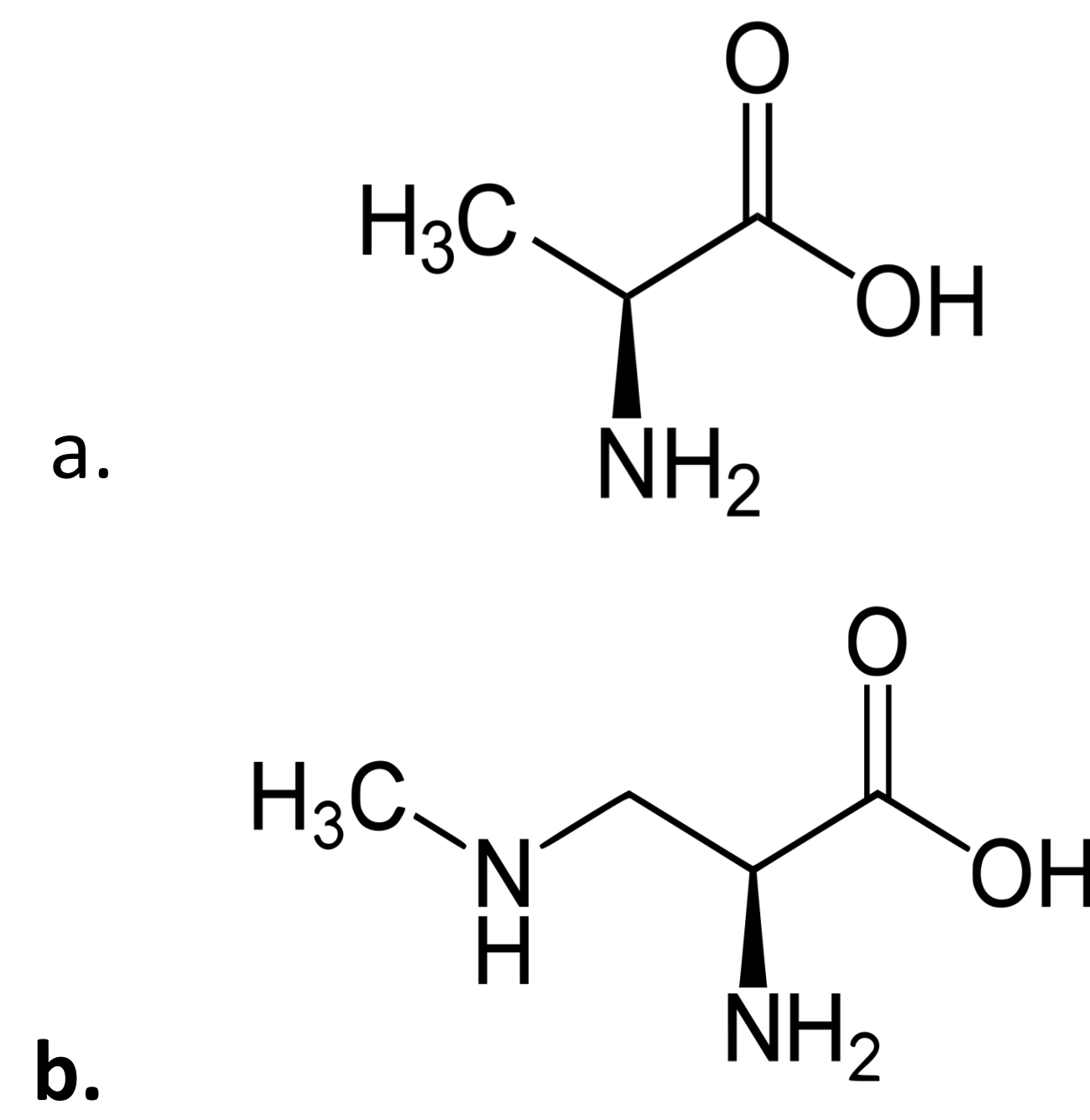


Figure 1: (a.) Chemical structure of the standard amino acid L-alanine. (b.) Chemical structure of the non-proteinogenic amino acid BMAA.



Figure 2: Cyanobacteria, which naturally produces and secretes BMAA.

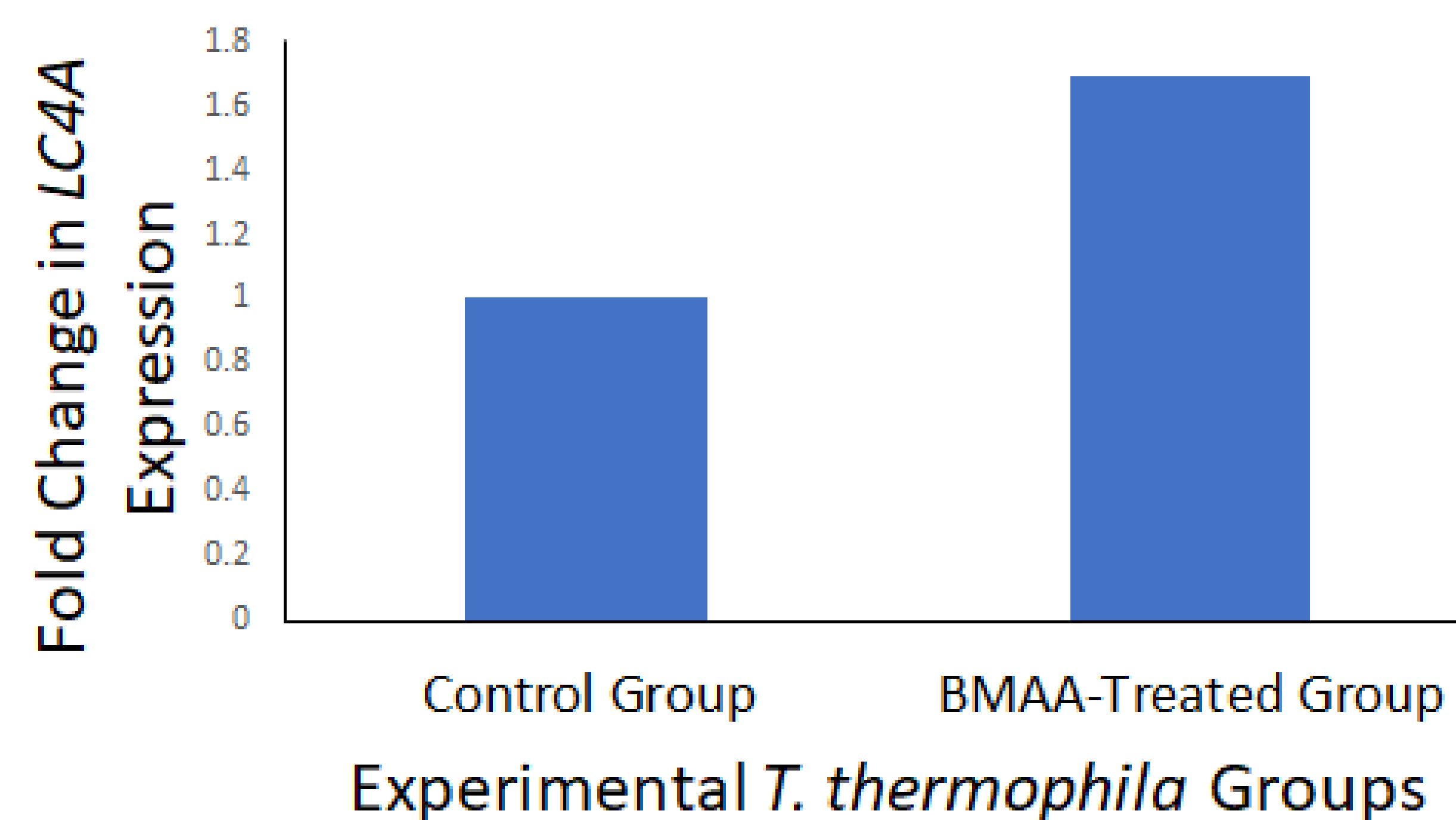


Figure 3: Fold change in expression of *LC4A*. ($p=0.332$; $n = 4$ for each group).

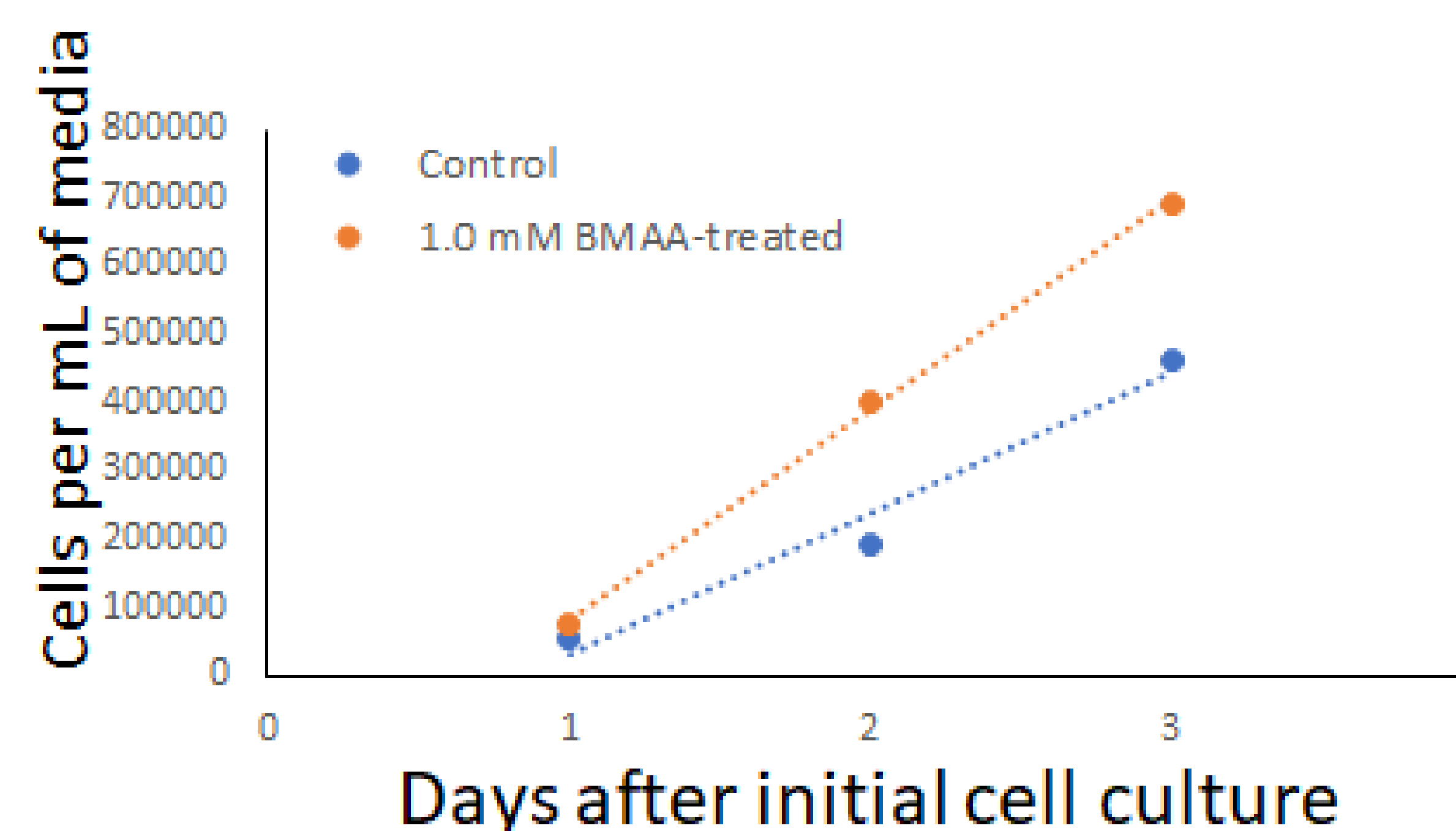


Figure 4: Number of cells per milliliter of media, averaged between round 1 and round 2. (Day 1: $p=0.346$; Day 2: $p=0.189$; Day 3: $p=0.245$, $n = 4$ for each group).

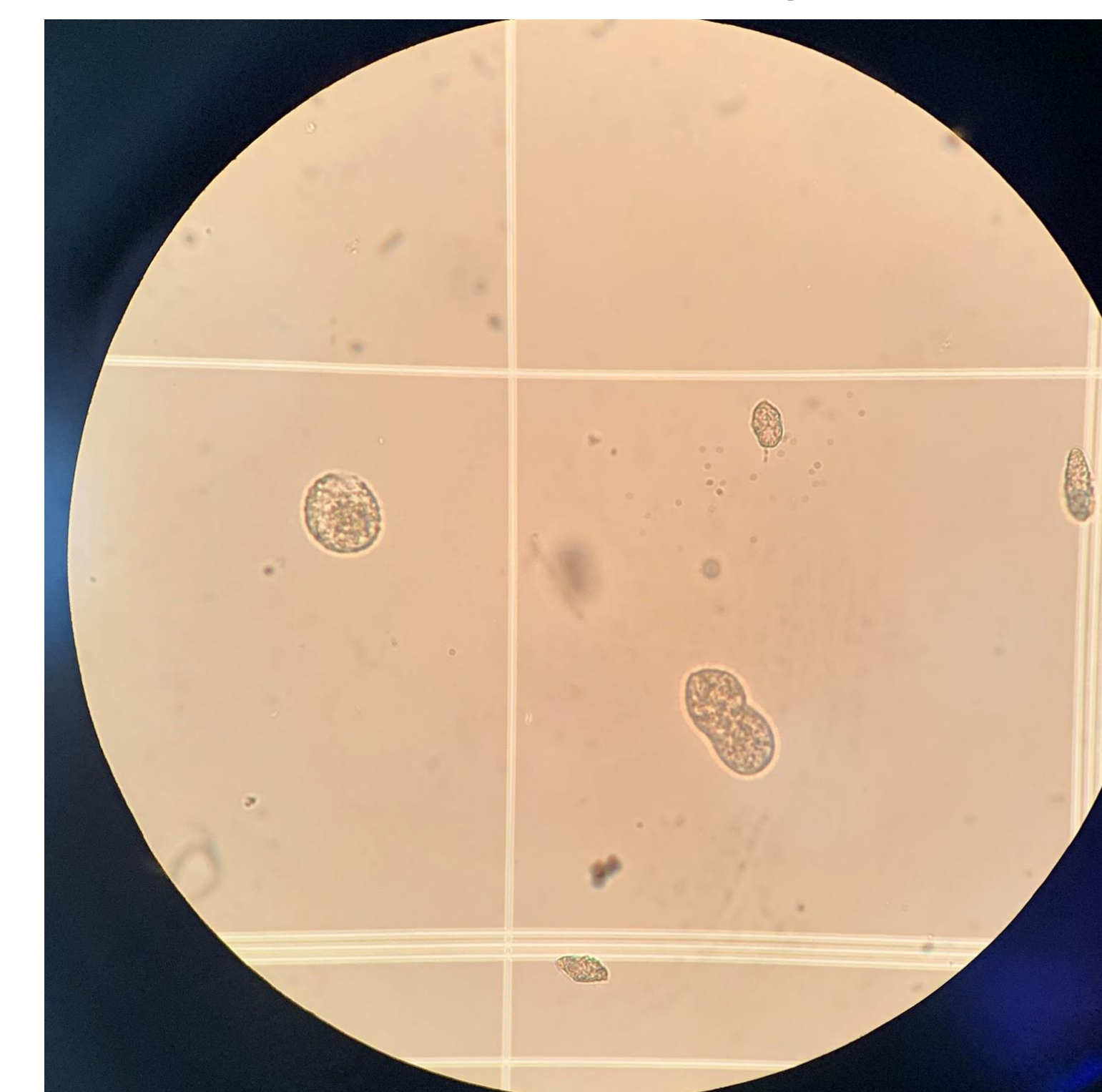


Figure 5: Image of *T. thermophila* on a hemocytometer under a microscope.