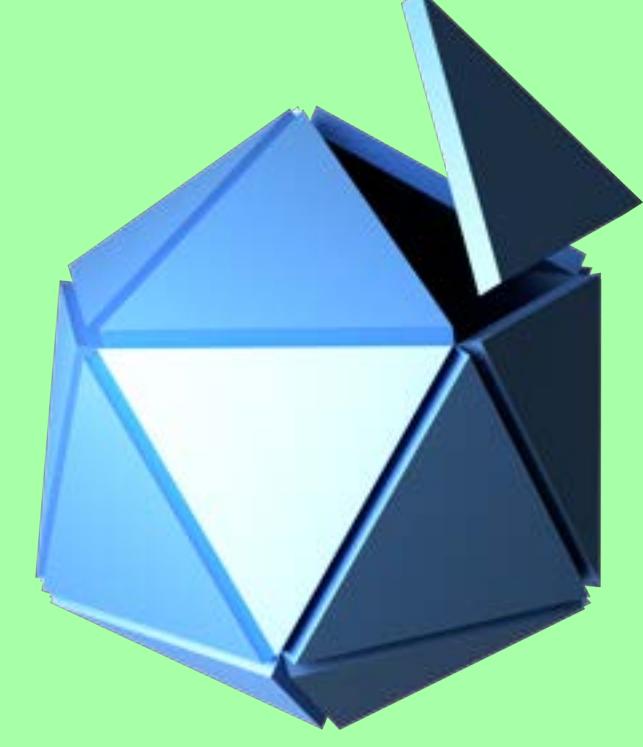


Microfluidic and CRISPRi toolkits for elucidating archaeal cytoskeleton

Brandeis
bioinspired
MRSEC

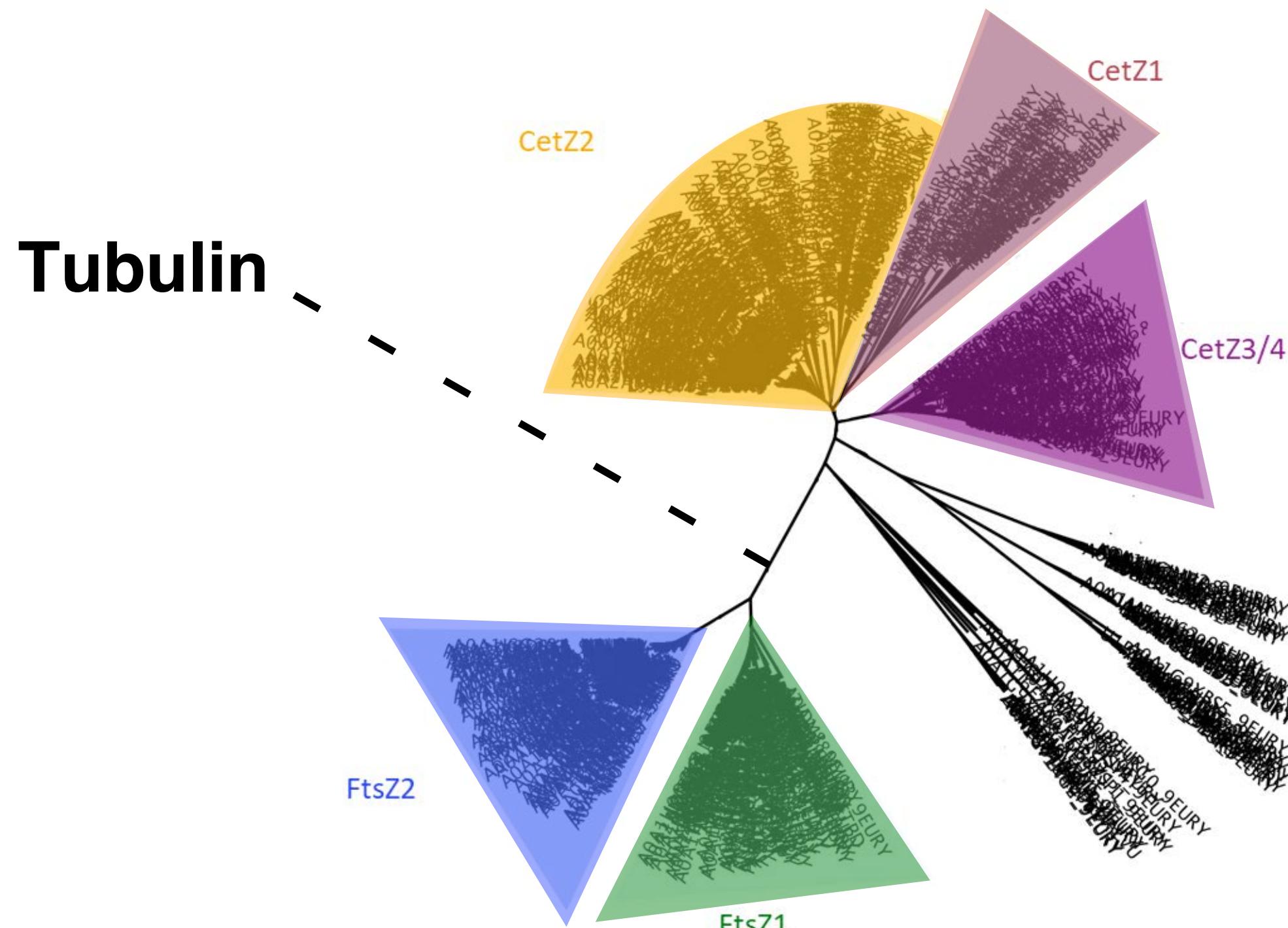


John Mallon¹, Alex Bisson¹

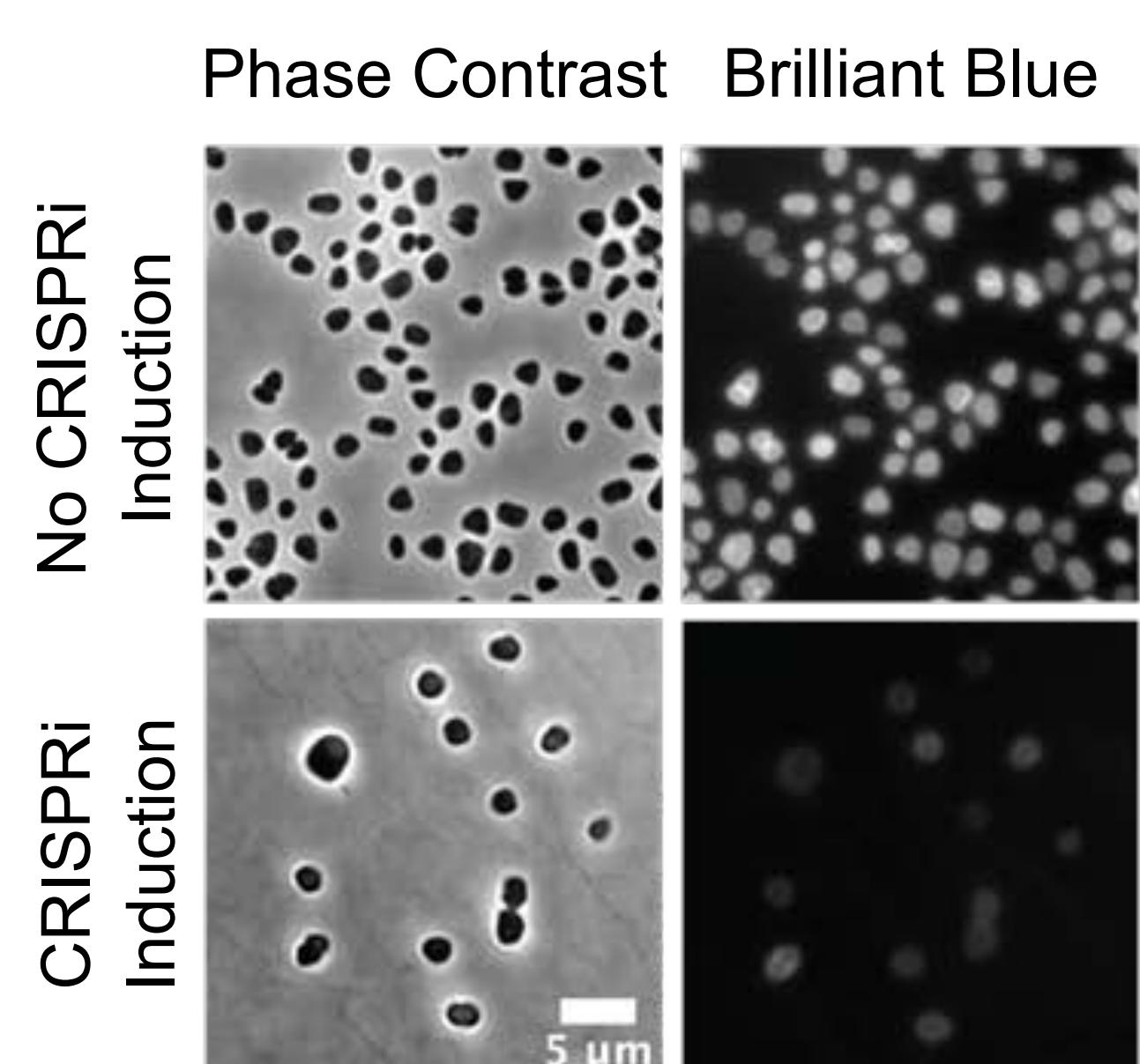
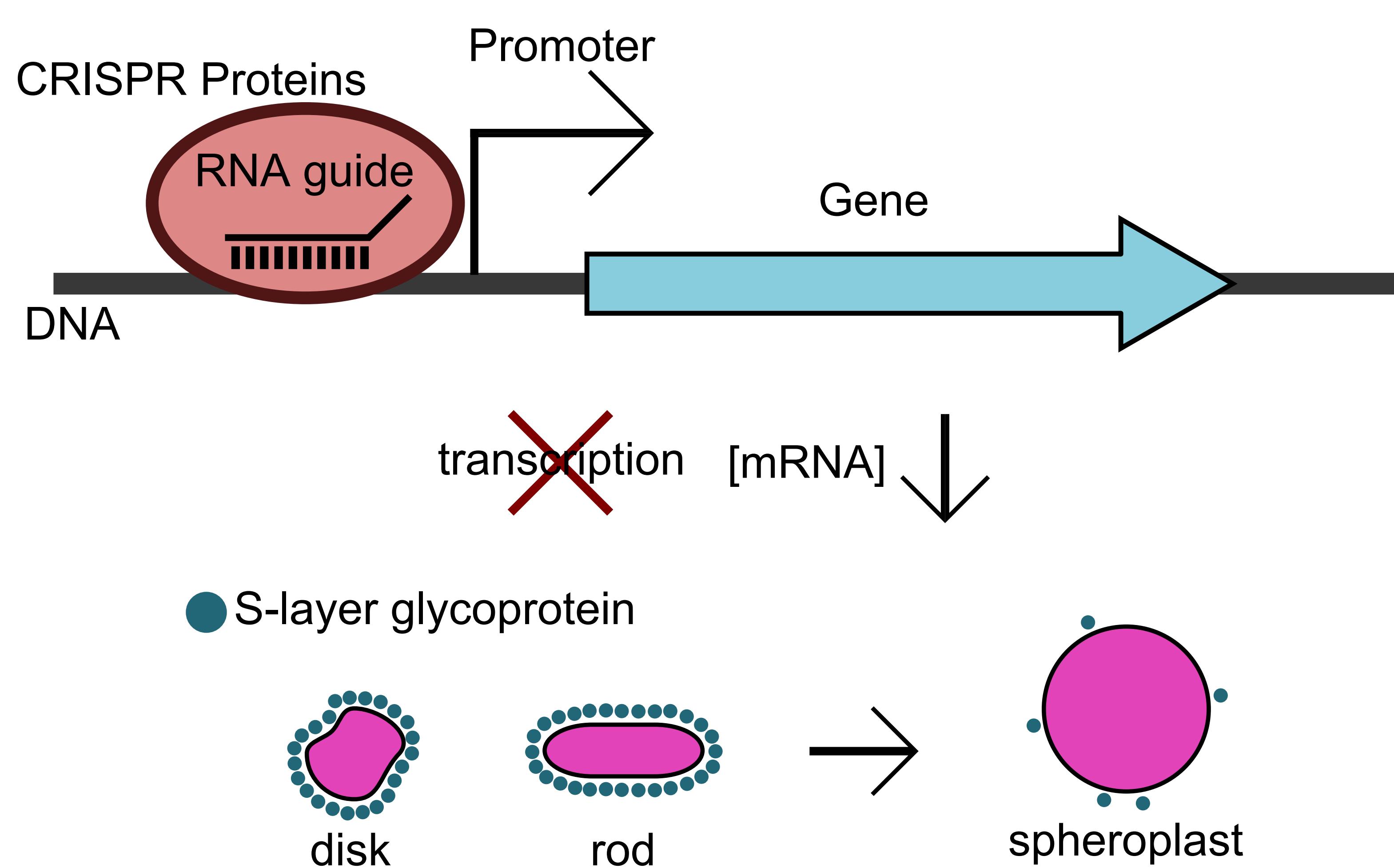
1. Department of Biology, Brandeis University, Waltham, MA

Rationale

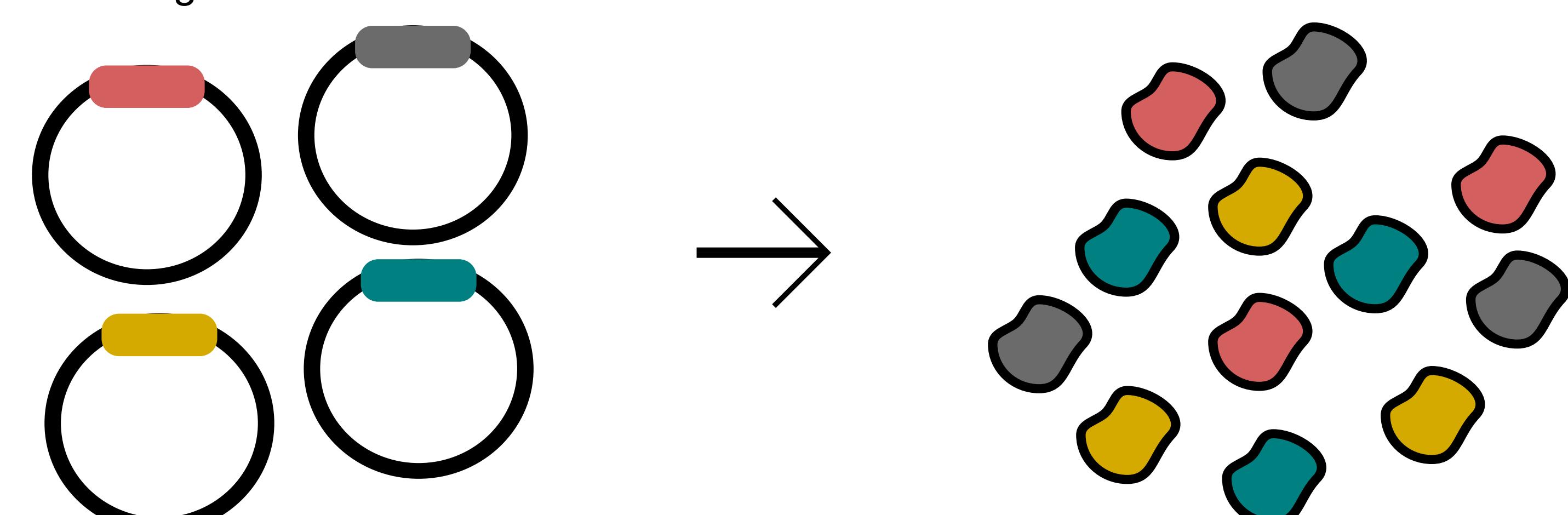
Archaea have many tubulin homologs, some of which are only found in the archaeal domain. Furthermore species will have many of these tubulins within their chromosome. To inspire new materials for active fluids research we look to develop tools for studying the properties of these archaeal tubulins



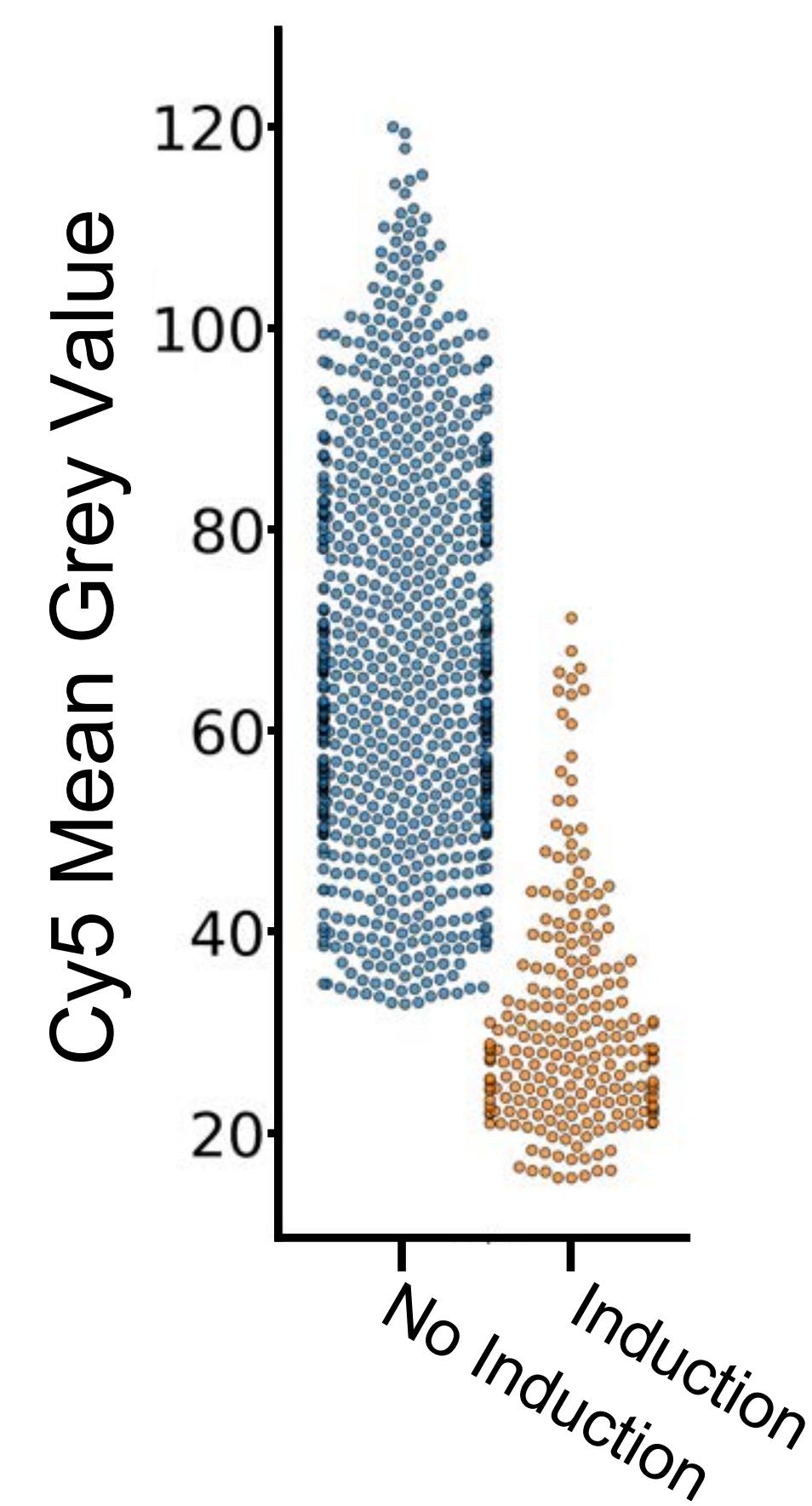
CRISPRi knockdown of S-layer



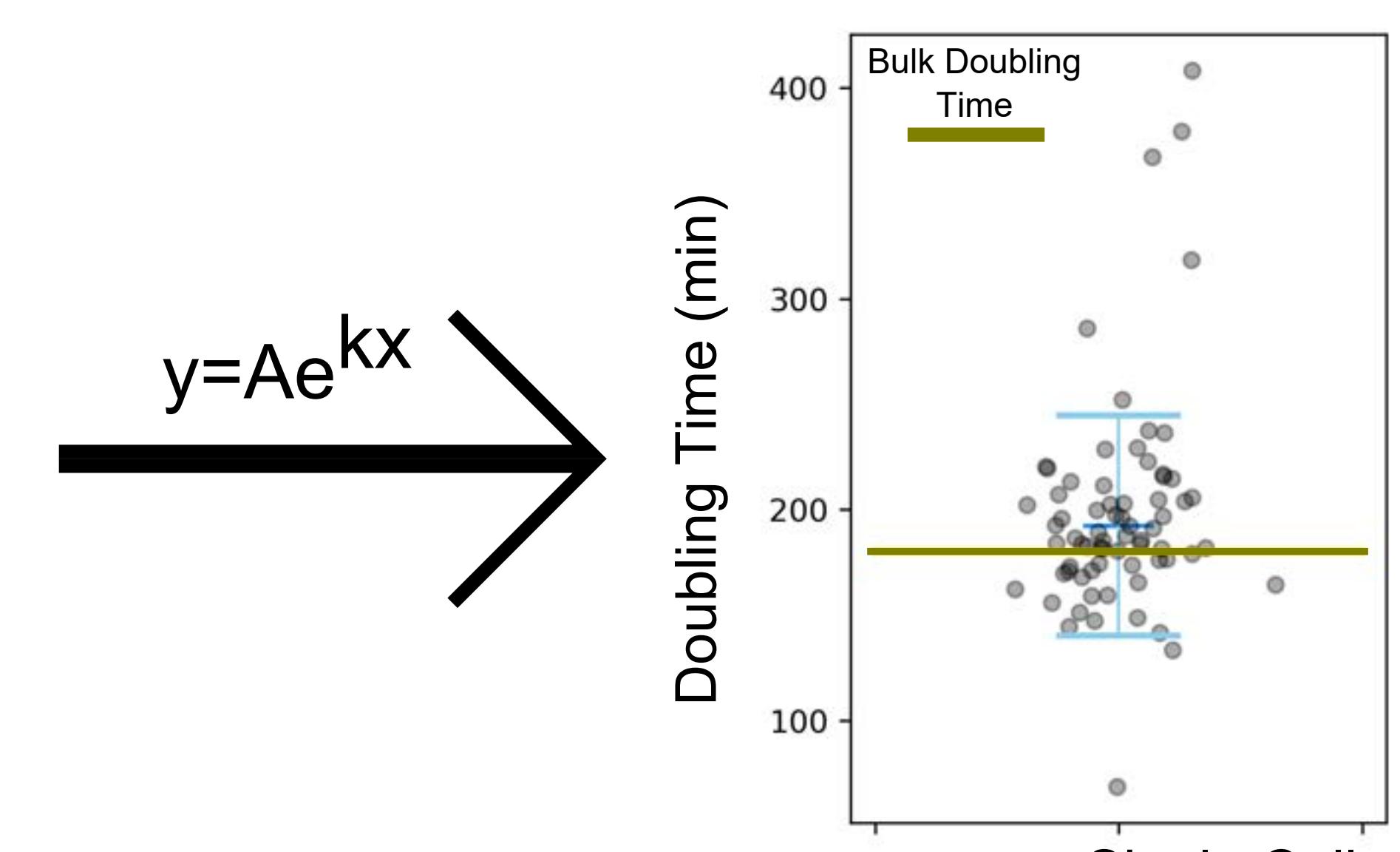
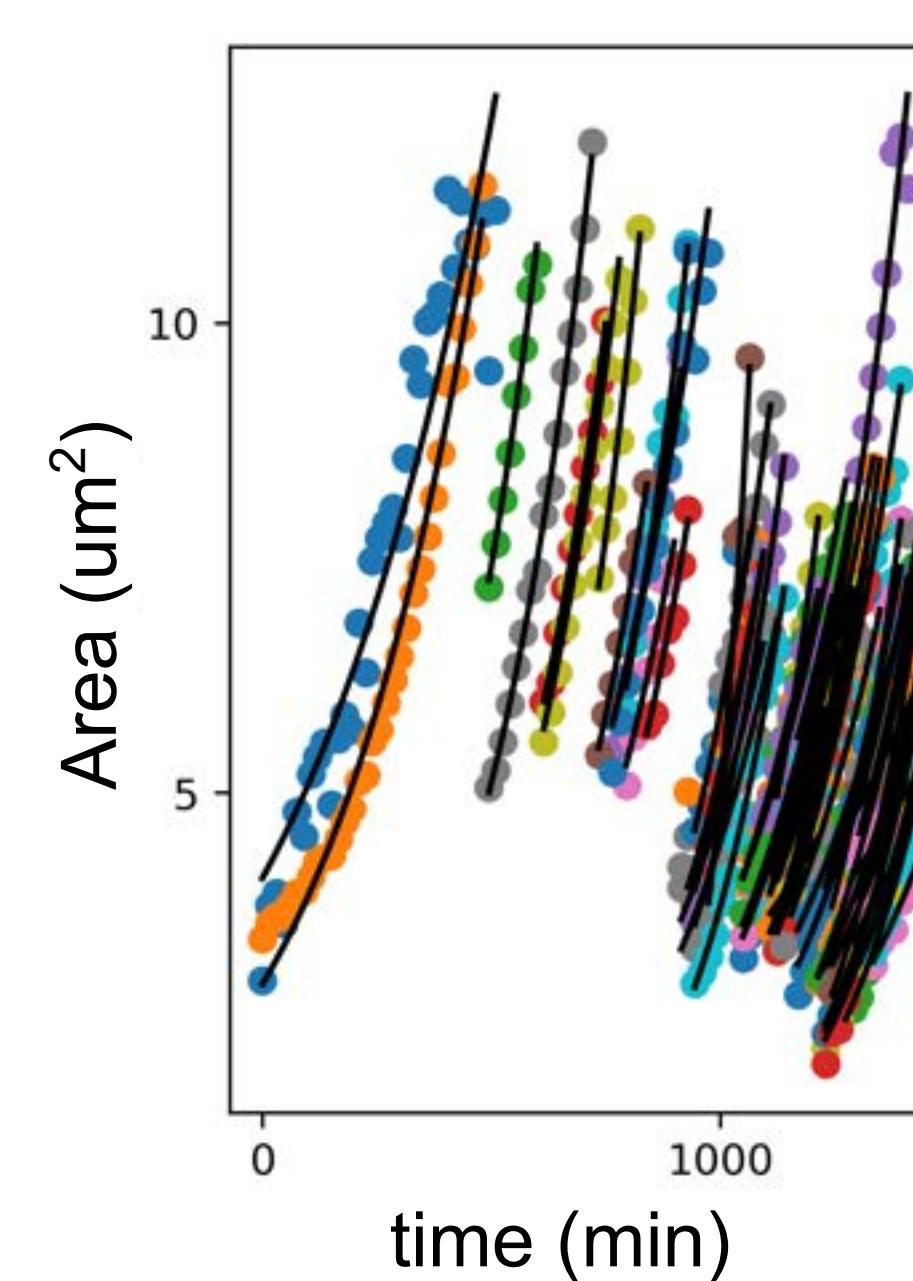
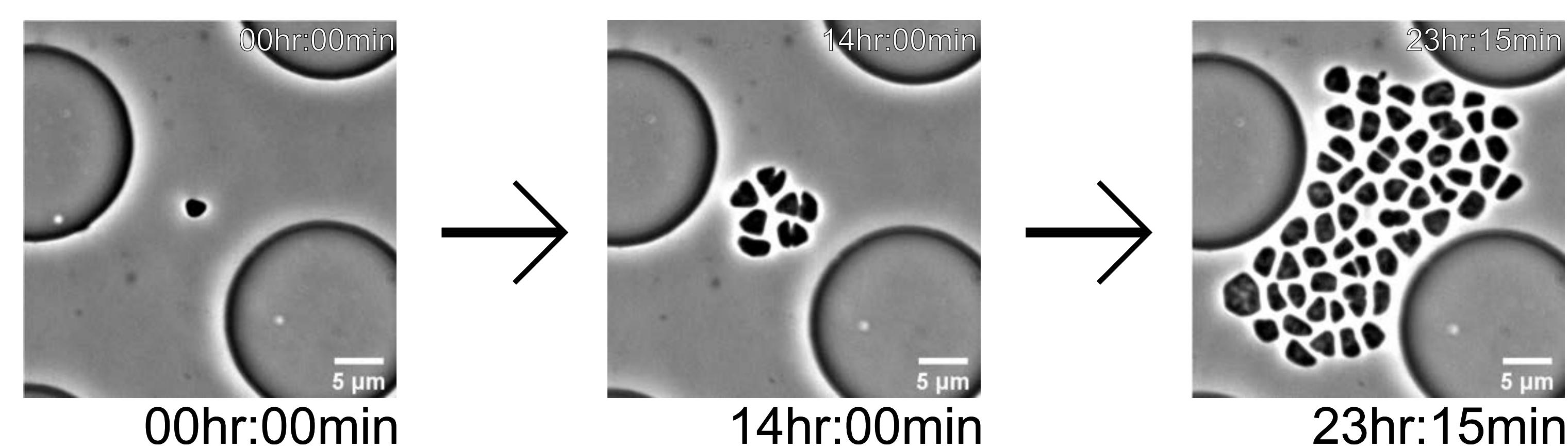
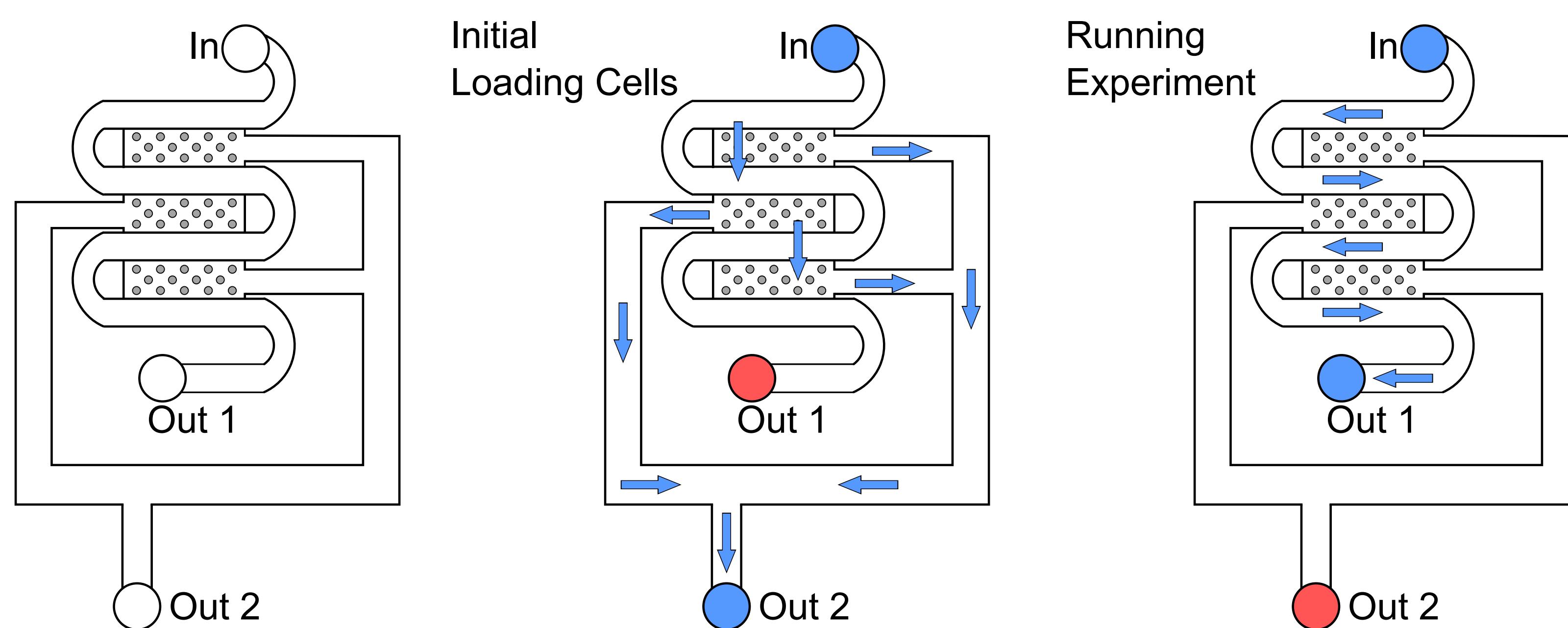
Library of ~8k RNA guides in plasmid form targeting *Haloferax volcanii* genome



Currently scaling up our transformation protocol for oversampling of library



Microfluidics is growing cells successfully



Conclusions

- Current fluidics system with two outlet design gives single cell growth rates comparable to bulk measurements
- CRISPRi strain can successfully knockdown the highest expressed gene (S-layer) in *Haloferax volcanii*

References

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2. Duggin I.G., Aylett C.H., Walsh J.C., et al. CetZ tubulin-like proteins control archaeal cell shape. *Nature*. 2015 Mar 19; 519(7543): 362-365
3. Stachler A.E., Marchfelder A. Gene Repression in Haloarchaea Using CRISPR (Clustered Regularly Interspaced Short Palindromic Repeats)-Cas-I-B System. *J Biol Chem*. 2016 Jul 15;291(29):15226-42.