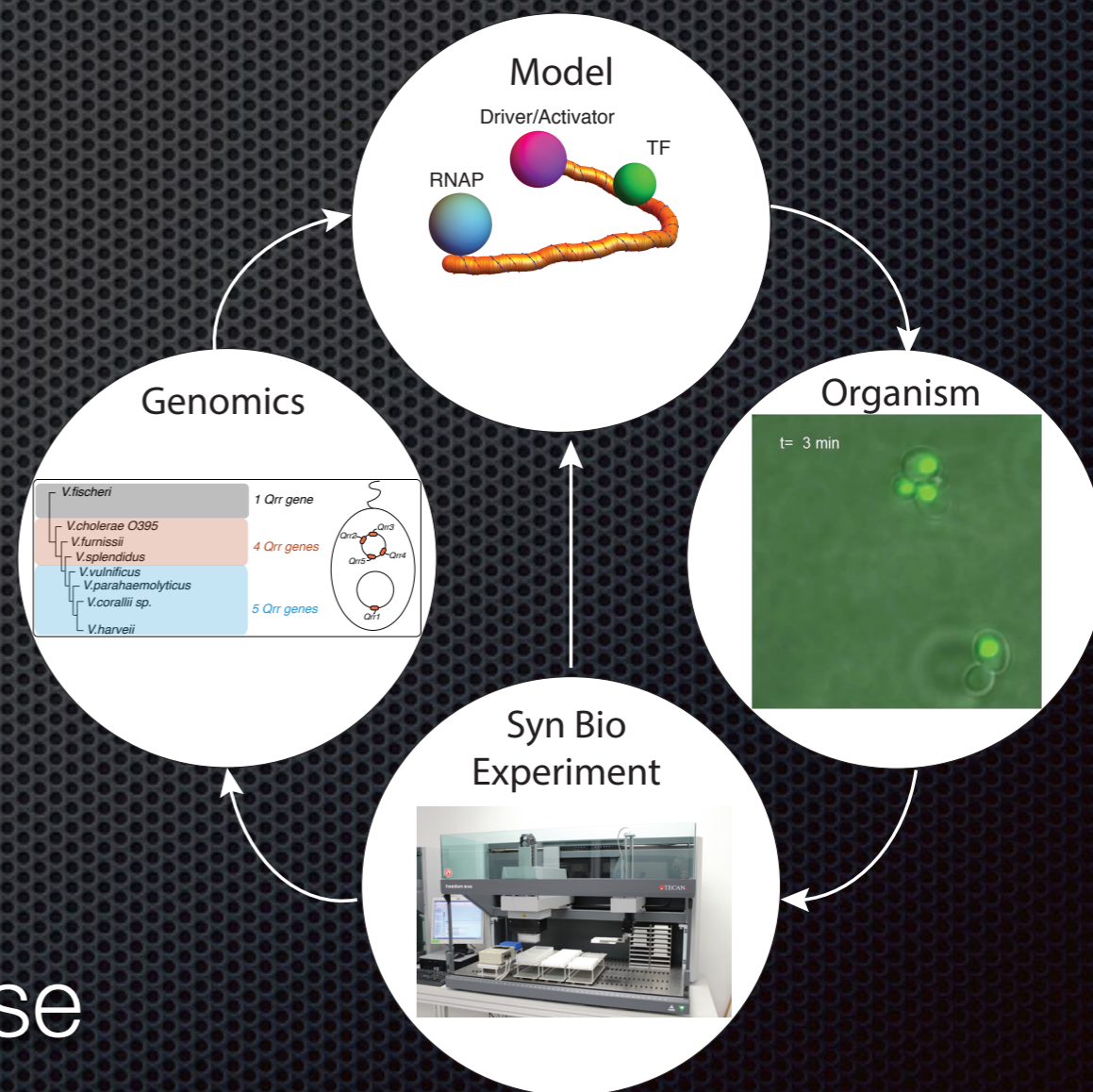


# Massive Reverse Genomics (MRG)

- A collaborative multi-disciplinary 4M € project for 2015-2018 with the Amit (Technion), Segal (WIS), Furlong (EMBL), Teichmann (EBI), Taipale (Karolniska Institute) groups, and Agilent Technologies.
- Design, analyze, and model massive libraries of synthetic enhancers in bacteria, yeast, mammalian cells, and Drosophila embryos.
- Potential postdocs or graduate students with a background in **Physics**, **Computer Science**, **Biology**, **Bioengineering**, **Bioinformatics** etc. may apply.
- Main goal will be to decipher the regulatory code, which determines when, where, under-what-conditions, and at what intensity genes get expressed.



If you are interested please  
 send C.V. to  
[roeeamit@technion.ac.il](mailto:roeeamit@technion.ac.il)