



What: EDAMAME is an intensive, hands-on course in microbial metagenome analysis, covering workflows from raw sequence analysis through to multivariate statistics and ecological interpretation. For more information, check out our course [website](#) and [wiki](#).

When: June 24-30, 2018

Where: Kellogg Biological Station

Why: The learning goals for the EDAMAME course are to

- Increase computing literacy
- Develop proficiency in cloud computing
- Analyze microbial amplicon sequences
- Analyze microbial shotgun metagenome sequences
- Apply ecological statistics to analyze and interpret microbial sequencing data
- Access resources provided by public sequence databases

Application: Apply [here](#) now through Friday February 2, 2018.

Questions? edamame.course@gmail.com