

A banner image for the EDAMAME workshop. It features a blue sky with white clouds at the top and green corn leaves at the bottom. The word "EDAMAME" is written in large, white, bold, sans-serif capital letters across the center of the sky. Below it, in a smaller, white, italicized font, is the text "Explorations in Data Analyses for Metagenomic Advances in Microbial Ecology (EDAMAME) Workshop".

EDAMAME

Explorations in Data Analyses for Metagenomic Advances in Microbial Ecology (EDAMAME) Workshop

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: August 6-12 2017

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: Coming soon!

Questions? edamame.course@gmail.com