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PROPOSAL CALL DEADLINES

Choose from a variety of resources from sequencing and synthesis to metabolomics, proteomics, imaging, and more:

Community Science Program calls:

Functional Genomics: Proposals due January 31
New Investigator: Proposals due March 1
Large-Scale Annual Community Science Program (CSP) Letters of Intent due March 24

Facilities Integrating Collaborations for User Science (FICUS): Letters of intent due March 17

FICUS offers resources from EMSL, samples the National Ecological Observatory Network (NEON), and capabilities from the Bio-SANS beamline at the High Flux Isotope Reactor (HFIR) through the Center for Structural Molecular Biology.

JGI Deputy Honored with DOE’s Lawrence Award

Susannah Tringe, JGI Deputy for User Programs and Interim Director of Berkeley Lab’s Environmental Genomics and Systems Biology Division, has been awarded one of DOE's highest honors. She was recognized for “advances in sequence-based studies of microbial assemblies, revealing the roles of microbial communities in carbon cycling, in interactions with plants, and as drivers of methane production, nutrient cycling, greenhouse gas emissions, and water recycling.” Read more >
DOE Joint Genome Institute

Genomics of Energy & Environment Meeting

August 30 – September 1, 2021

All Virtual in 2021!
Free registration. View Speakers and agenda here >

Virtual 2021
Viral EcoGenomics & Applications (VEGA)

Six 1.5 hour sessions: April 8; April 15; April 22; April 29; May 6; May 13
View confirmed speakers and agenda here >
SSRL in coordination with the EMSL will host a half-day virtual workshop on January 26, 2021, 8:00 am – 2:00 pm Pacific Time to encourage the deployment of spectroscopy and computational analysis to enhance understanding of key biogeochemical processes that control the speciation, distribution and fate of contaminants and nutrients. View the agenda and register here >

Contribute to the future of FAIR Microbiome Data

The National Microbiome Data Collaborative (NMDC) seeks to address roadblocks in microbiome data science and gaps in collaboration across disciplines by making data findable, accessible, interoperable, and reusable (FAIR). Let the NMDC know about your microbiome research data needs by completing this survey >

Dive into the Data Pool with KBase

Wednesday, January 27th at 10am PT/1pm ET

Hosted by Ben Allen, Oak Ridge National Laboratory and Dakota Blair, Brookhaven National Lab. Find out how KBase can support your research. Find out why KBase is the best way to create personalized, reproducible, shareable bioinformatics workflows. Register and create a KBase account here >