

Resource Summary Report

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Field Museum of Natural History Pritzker Laboratory for Molecular Systematics and Evolution Core Facility

RRID:SCR_026354

Type: Tool

Proper Citation

Field Museum of Natural History Pritzker Laboratory for Molecular Systematics and Evolution Core Facility (RRID:SCR_026354)

Resource Information

URL: <https://sites.google.com/fieldmuseum.org/research/laboratories/core-laboratories/dna?authuser=0>

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Description: Interdepartmental, multiuser core facility dedicated to genetic analysis and preservation of the world's biodiversity. Provides equipment used in molecular biology enabling to pursue study of genetic diversity throughout tree of life and at all taxonomic levels, from populations to classes to phyla of organisms.

Synonyms: Field Museum of Natural History Pritzker Laboratory for Molecular Systematics and Evolution, , Pritzker Laboratory for Molecular Systematics and Evolution, Field Museum Pritzker Laboratory for Molecular Systematics and Evolution

Resource Type: core facility, access service resource, service resource

Keywords: ABRF, genetic analysis and preservation, biodiversity, genetic diversity, tree of life, taxonomic level,

Resource Name: Field Museum of Natural History Pritzker Laboratory for Molecular Systematics and Evolution Core Facility

Resource ID: SCR_026354

Alternate IDs: ABRF_5944

Alternate URLs: https://coremarketplace.org/RRID:SCR_026354/?citation=1

Record Creation Time: 20250131T053327+0000

Record Last Update: 20260808T190740+0000

Ratings and Alerts

No rating or validation information has been found for Field Museum of Natural History Pritzker Laboratory for Molecular Systematics and Evolution Core Facility.

No alerts have been found for Field Museum of Natural History Pritzker Laboratory for Molecular Systematics and Evolution Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#) .

Bassani S, et al. (2012) The X-linked intellectual disability protein TSPAN7 regulates excitatory synapse development and AMPAR trafficking. Neuron, 73(6), 1143.