University of Vermont Larner College of Medicine Advanced Genome Technologies Massively Parallel Sequencing Core Facility

RRID:SCR_022696
Type: Tool

Proper Citation

University of Vermont Larner College of Medicine Advanced Genome Technologies Massively Parallel Sequencing Core Facility (RRID:SCR_022696)

Resource Information

URL: https://www.med.uvm.edu/vigr/mpsf

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Description: Provides support for design, execution, and analysis of experiments involving Next Generation Sequencing through access and use of Illumina HiSeq 1500. Sequencer performs wide range of analysis including target preparation for RNAseq, Exome Seq, ChIP seq, Methyl Seq, whole genome sequencing, and small RNA sequencing. Core is involved in all aspects of Nanopore sequencing.

Synonyms: Advanced Genome Technologies Core Massively Parallel Sequencing Facility, University of Vermont Advanced Genome Technologies Core Massively Parallel Sequencing Facility

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

Keywords: USEdit, ABRF, Next Generation Sequencing, Illumina HiSeq 1500, RNAseq, Exome Seq, ChIP seq, Methyl Seq, whole genome sequencing, small RNA sequencing

Resource Name: University of Vermont Larner College of Medicine Advanced Genome Technologies Massively Parallel Sequencing Core Facility

Resource ID: SCR_022696
Alternate IDs:  ABRF_459

Alternate URLs:  https://coremarketplace.org/?FacilityID=459&citation=1

Ratings and Alerts

No rating or validation information has been found for University of Vermont Larner College of Medicine Advanced Genome Technologies Massively Parallel Sequencing Core Facility.

No alerts have been found for University of Vermont Larner College of Medicine Advanced Genome Technologies Massively Parallel Sequencing Core Facility.

Data and Source Information

Source:  SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.