Resource Summary Report

Generated by dkNET on Apr 21, 2025

Harvard PCPGM Genotyping Facility

RRID:SCR 009846

Type: Tool

Proper Citation

Harvard PCPGM Genotyping Facility (RRID:SCR_009846)

Resource Information

URL: http://harvard.eagle-i.net/i/0000012e-021d-e404-b2b9-4d8780000000

Proper Citation: Harvard PCPGM Genotyping Facility (RRID:SCR_009846)

Description: Core facility that provides the following services: ABI Prism Taqman Allelic Discrimination Assay, OpenArray SNP genotyping, Illumina genotyping, Custom Illumina GoldenGate genotyping, Illumina Infinium genotyping analysis, Illumina methylation analysis. The Partners Genotyping Facility, part of the Partners HealthCare Center for Personalized Genetic Medicine (PCPGM), provides flexible, high quality, high-throughput SNP genotyping to the Harvard-Partners research community, including Harvard Medical School, hospitals in the Partners HealthCare network, investigators in the Dana-Farber-/ Harvard Cancer Center, and the Harvard School of Public Health. The portfolio of Genotyping methods at PCPGM now includes Illumina, TaqMan and TaqMan OpenArrays. Note: DF/HCC members will receive the DF/HCC discount on both genotyping and sequencing services from our facility

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

Keywords: genotyping assay, single-nucleotide polymorphism analysis, snp interrogation genotyping, dna methylation profiling assay

Funding:

Resource Name: Harvard PCPGM Genotyping Facility

Resource ID: SCR_009846

Alternate IDs: nlx_156328

Alternate URLs: http://pcpgm.partners.org/research-services/genotyping/

Record Creation Time: 20220129T080255+0000

Record Last Update: 20250421T053744+0000

Ratings and Alerts

No rating or validation information has been found for Harvard PCPGM Genotyping Facility.

No alerts have been found for Harvard PCPGM Genotyping Facility.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.