

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

Children's Hospital of Philadelphia CHOP CAG Sequencing Core Facility

RRID:SCR_028416

Type: Tool

Proper Citation

Children's Hospital of Philadelphia CHOP CAG Sequencing Core Facility
(RRID:SCR_028416)

Resource Information

URL: <https://www.research.chop.edu/cag-sequencing-core>

Proper Citation: Children's Hospital of Philadelphia CHOP CAG Sequencing Core Facility
(RRID:SCR_028416)

Description: Cores Sequencing operations are integrated with Biorepository and Bioinformatics teams to keep all stages of a collaborators studies connected. Core offers genomics services in NovaSeq, Single Cell 10xGenomics, various library preparations including bulk RNAseq, Whole Genome/Whole Exome, and Sanger Sequencing including fragment analysis.

Synonyms: , CAG Sequencing Laboratory, CAG Sequencing Core

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

Keywords: ABRF, genomics services, NovaSeq, Single Cell 10xGenomics, library preparations, bulk RNAseq, Whole Genome, Whole Exome, Sanger Sequencing, fragment analysis,

Resource Name: Children's Hospital of Philadelphia CHOP CAG Sequencing Core Facility

Resource ID: SCR_028416

Alternate IDs: ABRF_5927

Alternate URLs: <https://coremarketplace.org/?FacilityID=5927&citation=1>

Record Creation Time: 20260508T053733+0000

Record Last Update: 20260808T190828+0000

Ratings and Alerts

No rating or validation information has been found for Children's Hospital of Philadelphia CHOP CAG Sequencing Core Facility.

No alerts have been found for Children's Hospital of Philadelphia CHOP CAG Sequencing 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 [SPARC SAWG](#) .

Laub S, et al. (2026) Integrative genomics and single-cell CRISPRi screening dissect Alzheimer GWAS non-coding variants regulating TSPAN14. American journal of human genetics, 113(6), 1253.