## **Resource Summary Report**

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# University of Washington Genomics Core Cystic Fibrosis Research Translation Center and Research Development Program

RRID:SCR\_015404

Type: Tool

## **Proper Citation**

University of Washington Genomics Core Cystic Fibrosis Research Translation Center and Research Development Program (RRID:SCR\_015404)

#### Resource Information

URL: http://depts.washington.edu/cfrtc/genomics/

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**Description:** Core provides genomics-based tools, data management and analysis tools, and creates platforms that integrate data from the Clinical and Immunology Cores for human samples and bacterial isolates. Services include consultation and experimental design assistance for using new-generation sequencing technology, data analysis, bioinformatic support, data access and storage, high throughput and new-generation whole-genome sequencing, and RNA-seq analysis of transcriptomes.

**Abbreviations:** CF Research Translation Center and Research Development Program, CFRTC

**Synonyms:** Genomics Core, Cystic Fibrosis Research Translation Center, CFRTC, Research Development Program, University of Washington

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

**Keywords:** genomics core, sequencing technology, cystic fibrosis genomics, RNA sequencing, analysis

**Related Condition:** Cystic Fibrosis

Funding: NIDDK P30 DK89507

Availability: Available to external user

Resource Name: University of Washington Genomics Core Cystic Fibrosis Research

Translation Center and Research Development Program

Resource ID: SCR\_015404

**Record Creation Time:** 20220129T080325+0000

Record Last Update: 20250423T060836+0000

## Ratings and Alerts

No rating or validation information has been found for University of Washington Genomics Core Cystic Fibrosis Research Translation Center and Research Development Program.

No alerts have been found for University of Washington Genomics Core Cystic Fibrosis Research Translation Center and Research Development Program.

#### 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.

Polte C, et al. (2019) Assessing cell-specific effects of genetic variations using tRNA microarrays. BMC genomics, 20(Suppl 8), 549.