

The background of the image is a photograph of a modern, multi-story glass skyscraper at dusk. The building's windows are illuminated from within, creating a warm glow against the dark blue twilight sky. A prominent feature is a curved glass facade that reflects the surrounding environment. In the foreground, there is a glass-enclosed walkway or entrance area with a white structural frame. A person in a white lab coat is walking through the glass entrance on the right side. A street lamp with a glowing globe is visible in the middle ground. The overall atmosphere is professional and modern.

**M Northwestern Medicine**  
Feinberg School of Medicine

**Accelerating Transplant  
Science: How the CTC's  
Bioinformatics Core Can  
Support Your Research**

# The Comprehensive Transplant Center's Bioinformatics Core at Northwestern University



Our mission is to advance patient-centric impact by providing high-quality data analysis, innovative computational approaches and educational support to enhance basic, pre-clinical, clinical and epidemiological transplant research.



We serve as a collaborative platform for researchers at Northwestern and beyond, providing expertise in analysis of multi-omics data—including genomics, transcriptomics, epigenetics, proteomics, and metabolomics—to drive discovery and innovation in transplantation research.

# Our Services Include (But Not Limited to):

## Data Storage and Management

Insight on storage of large data files at Northwestern through Quest High-Performance Computing Cluster

## DNA Sequencing

Genome-Wide Association Studies (GWAS) to identify genetic variants linked to transplant outcomes

HLA typing from whole genome sequencing to determine donor-recipient compatibility among cohorts of transplant patients

## RNA Sequencing

We provide tailored computational pipelines to dissect cellular heterogeneity, map gene expression within tissue architecture, and uncover transcriptional networks driving transplant outcomes

Includes single cell/nuclear RNA sequencing, spatial transcriptomics, and bulk RNA sequencing

## Epigenomics

Includes ATAC sequencing, chromatin immunoprecipitation (ChIP) sequencing analysis to provide insights on gene regulation and epigenetic modifications

## Proteomics & Immunopeptidomics

Advanced proteomics and immunopeptidomics analysis, enabling researchers to profile global protein expression, post-translational modifications, and antigen presentation dynamics

## Metabolomics

Includes comprehensive profiling, targeted metabolite identification, and metabolic flux analysis to uncover metabolic signatures

# Powering Research Success: Bioinformatics Support for Grants & Publications

## Supporting Grant Applications

- **Strengthen Proposals:** High-quality bioinformatics data for preliminary results
- **Grant Writing Support:** Assistance with computational methods and study design
- **Data Visualization:** Creation of figures, heatmaps, and network graphs
- **Collaboration & Letter of Support:** Available as co-investigators or collaborators



National Institutes  
of Health



## Enhancing Publications

- **Comprehensive Data Analysis:** From raw data processing to biological and clinical interpretation
- **Publication-Ready Figures:** High-quality visualizations for manuscripts
- **Reproducibility and Rigor:** Standardized workflows and analysis pipelines

## Looking to start a collaboration?

Please complete our Bioinformatics Intake Form located on our website:

<https://www.feinberg.northwestern.edu/sites/transplant/research/research-cores/bioinfo-core.html>



**Connor Lantz, PhD**

Lead Scientist, Bioinformatics Core  
Comprehensive Transplant Center

[connor.lantz@northwestern.edu](mailto:connor.lantz@northwestern.edu)