Resource Summary Report

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Stanford Diabetes Research Center Diabetes Genomics Analysis Core

RRID:SCR_016213
Type: Tool

Proper Citation

Stanford Diabetes Research Center Diabetes Genomics Analysis Core (RRID:SCR_016213)

Resource Information

URL: https://sdrc.stanford.edu/sdrc-research-cores/dgac/home/

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Description: Core facility that offers library preparation and sequencing services on a variety of platforms - Illumina HiSeq 4000, MiSeq, HiSeq 2500 and PacBio Sequel - as well as bioinformatics analysis. It can sequence a variety of commercial sample preparation kits as well as custom workflows. DGAC provides access to high throughput sequencing and analysis to researchers at the Stanford Diabetes Research Center.

Abbreviations: GDAC, SDRC-GDAC, SDRC

Synonyms: SDRC, Genomics Analysis Core, Diabetes Genomics Analysis Core

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

Keywords: library, sequence, workflow, bioinformatic, gene, analysis, sequencing

Funding: NIDDK P30 DK116074

Availability: Available to external user

Resource Name: Stanford Diabetes Research Center Diabetes Genomics Analysis Core

Resource ID: SCR_016213

Record Creation Time: 20220129T080329+0000

Record Last Update: 20260821T194040+0000

Ratings and Alerts

No rating or validation information has been found for Stanford Diabetes Research Center Diabetes Genomics Analysis Core.

No alerts have been found for Stanford Diabetes Research Center Diabetes Genomics Analysis Core.

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 dkNET .

Shen J, et al. (2020) Paclitaxel Promotes Tumor-Infiltrating Macrophages in Breast Cancer. Technology in cancer research & treatment, 19, 1533033820945821.