

Resource Summary Report

Generated by RRID on Aug 14, 2026

Collaborative Research Network Cloud

RRID:SCR_023923

Type: Tool

Proper Citation

Collaborative Research Network Cloud (RRID:SCR_023923)

Resource Information

URL:

<https://docs.google.com/presentation/d/13KqgDOFLbhCPRsCqf4zETdcjhZECIXD/edit#slide=id.p1>

Proper Citation: Collaborative Research Network Cloud (RRID:SCR_023923)

Description: MJFF in partnership with Aligning Science Across Parkinson's are in the process of establishing data platform, CRN Cloud, leveraging Verily Workbench to share data generated by researcher labs funded through ASAP Collaborative Research Network. The data types include mainly raw and processed single cell RNA-seq, single nucleus RNA-seq, and bulk RNA-seq from both PD and healthy control samples.

Abbreviations: CRN Cloud

Resource Type: data or information resource, organization portal, portal

Keywords: raw and processed single cell RNA-seq, single nucleus RNA-seq, bulk RNA-seq, PD and healthy control samples,

Funding: ASAP Collaborative Research Network

Availability: Free, Freely available

Resource Name: Collaborative Research Network Cloud

Resource ID: SCR_023923

Record Creation Time: 20230825T050208+0000

Record Last Update: 20260814T043051+0000

Ratings and Alerts

No rating or validation information has been found for Collaborative Research Network Cloud.

No alerts have been found for Collaborative Research Network Cloud.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Riley JF, et al. (2026) PINK1/Parkin-dependent mitophagy mediates astrocytic inflammatory responses to mitochondrial damage. bioRxiv : the preprint server for biology.

Menozzi E, et al. (2026) Microbiome signature of Parkinson's disease in healthy and genetically at-risk individuals. Nature medicine, 32(6), 2096.