

Resource Summary Report

Generated by SPARC SAWG on Sep 12, 2026

Michael J. Fox Foundation for Parkinsons Research

RRID:SCR_006183

Type: Tool

Proper Citation

Michael J. Fox Foundation for Parkinsons Research (RRID:SCR_006183)

Resource Information

URL: <http://michaeljfox.org/>

Proper Citation: Michael J. Fox Foundation for Parkinsons Research (RRID:SCR_006183)

Description: A Parkinson's research foundation dedicated to finding a cure for Parkinson's disease and to ensuring the development of improved therapies. Pipeline Programs fund investigator-initiated proposals focused on the following critical points along the translational pathway to new therapies for Parkinson's disease.

Abbreviations: MJFF

Synonyms: Michael J. Fox Foundation for Parkinson's Research, Michael J. Fox Foundation

Resource Type: data or information resource, disease-related portal, funding resource, portal, topical portal

Keywords: parkinson's disease, clinical, translational, foundational, research, funding resource

Availability: Public, Funding available to researchers

Resource Name: Michael J. Fox Foundation for Parkinsons Research

Resource ID: SCR_006183

Alternate IDs: nif-0000-00518

License URLs: <https://www.michaeljfox.org/terms-and-conditions.html?navid=footer-terms>

Record Creation Time: 20220129T080234+0000

Record Last Update: 20260912T195636+0000

Ratings and Alerts

No rating or validation information has been found for Michael J. Fox Foundation for Parkinsons Research.

No alerts have been found for Michael J. Fox Foundation for Parkinsons Research.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 76 mentions in open access literature.

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

Pourmajidian M, et al. (2026) Five energy metabolism pathways show distinct regional distributions and lifespan trajectories in the human brain. PLoS biology, 24(1), e3003619.

Lange LM, et al. (2026) Rare but Relevant? Assessing Variants in Dystonia-Linked Genes in Parkinson's Disease. Movement disorders : official journal of the Movement Disorder Society, 41(1), 247.

Farahani A, et al. (2026) Aging and metabolism contribute separately to brain-body health. PLoS biology, 24(6), e3003856.

Step K, et al. (2026) Genome-Wide Assessment Reveals Ancestral Differences in Homozygosity Patterns Potentially Linked to Parkinson's Disease Etiology. Movement disorders : official journal of the Movement Disorder Society, 41(5), 1128.

Schmidt MY, et al. (2026) Unraveling the intersection of aging and Parkinson's disease: a collaborative roadmap for advancing research models. NPJ Parkinson's disease, 12(1), 28.

Kleinz T, et al. (2025) RAB32-Linked Parkinson's Disease: Deep Phenotyping, MDSCGene Literature Review, and Application of SynNeurGe Criteria. Movement disorders : official journal of the Movement Disorder Society, 40(12), 2746.

Step K, et al. (2025) Genome-wide assessment identifies novel runs of homozygosity linked to Parkinson's disease etiology across diverse ancestral populations. medRxiv : the preprint server for health sciences.

Timmermans NA, et al. (2025) A generalizable and open-source algorithm for real-life monitoring of tremor in Parkinson's disease. NPJ Parkinson's disease, 11(1), 205.

Rissardo JP, et al. (2025) A Narrative Review on Biochemical Markers and Emerging Treatments in Prodromal Synucleinopathies. Clinics and practice, 15(3).

Kelly MP, et al. (2025) Cyclic nucleotide phosphodiesterases as drug targets. *Pharmacological reviews*, 77(3), 100042.

Bury AG, et al. (2025) Investigating the ageing-Parkinson's disease nexus: standardisation of in vitro models and techniques by the PD-AGE network. *NPJ Parkinson's disease*, 11(1), 289.

Step K, et al. (2025) Insights into X-Linked Susceptibility to Parkinson's Disease in the South African Population. *medRxiv : the preprint server for health sciences*.

Catto F, et al. (2025) Quantitative 3D histochemistry reveals region-specific amyloid- β reduction by the antidiabetic drug netoglitzzone. *PloS one*, 20(5), e0309489.

Blauwendraat C, et al. (2025) Tackling a disease on a global scale, the Global Parkinson's Genetics Program, GP2: A new generation of opportunities. *American journal of human genetics*, 112(9), 1988.

Magalhães AD, et al. (2025) Large-scale seroepidemiology uncovers nephro-urological pathologies in people with tau autoimmunity. *PLoS biology*, 23(11), e3003488.

Cox KM, et al. (2024) Detecting rhythmic spiking through the power spectra of point process model residuals. *bioRxiv : the preprint server for biology*.

Sokołowski A, et al. (2024) Longitudinal brain structure changes in Parkinson's disease: A replication study. *PloS one*, 19(1), e0295069.

Manes JL, et al. (2024) A neurocomputational view of the effects of Parkinson's disease on speech production. *Frontiers in human neuroscience*, 18, 1383714.

Sharma PK, et al. (2024) Role of the leucine-rich repeat protein kinase 2 C-terminal tail in domain cross-talk. *The Biochemical journal*, 481(4), 313.

Galper J, et al. (2024) Prediction of motor and non-motor Parkinson's disease symptoms using serum lipidomics and machine learning: a 2-year study. *NPJ Parkinson's disease*, 10(1), 123.