# Resource Summary Report

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# 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/

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

**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:** 20260804T043301+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 68 mentions in open access literature.

**Listed below are recent publications.** The full list is available at ASWG .

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.

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.

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

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

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-? reduction by the antidiabetic drug netoglitazone. 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.

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

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

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.

Gorji A, et al. (2024) Machine learning for predicting cognitive decline within five years in Parkinson's disease: Comparing cognitive assessment scales with DAT SPECT and clinical biomarkers. PloS one, 19(7), e0304355.

Cox KM, et al. (2024) Detecting rhythmic spiking through the power spectra of point process model residuals. Journal of neural engineering, 21(4).

Weng JH, et al. (2023) Capturing Differences in the Regulation of LRRK2 Dynamics and Conformational States by Small Molecule Kinase Inhibitors. ACS chemical biology, 18(4), 810.

Vollstedt EJ, et al. (2023) Establishing an online resource to facilitate global collaboration and inclusion of underrepresented populations: Experience from the MJFF Global Genetic Parkinson's Disease Project. PloS one, 18(10), e0292180.

Störmer E, et al. (2023) Capturing the domain crosstalk in full length LRRK2 and LRRK2RCKW. The Biochemical journal, 480(11), 815.

Galper J, et al. (2022) Lipid pathway dysfunction is prevalent in patients with Parkinson's disease. Brain : a journal of neurology, 145(10), 3472.