Impaired mitochondrial accumulation and Lewy pathology in neuron-specific FBXO7-deficient mice

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Abstract
Parkinson's disease, the second most common neurodegenerative disorder, is characterized by the loss of nigrostriatal dopamine neurons. FBXO7 (F-box protein only 7) (PARK15) mutations cause early-onset Parkinson's disease. FBXO7 is a subunit of the SCF (SKP1/cullin-1/F-box protein) E3 ubiquitin ligase complex, but its neuronal relevance and function have not been elucidated. To determine its function in neurons, we generated neuronal cell-specific FBXO7 conditional knockout mice (FBXO7floxtg/fox; Nestin-Cre) by crossing previously characterized FBXO7 floxed mice (FBXO7floxtg/fox) with Nestin-Cre mice (Nestin-Cre). The resultant Fbxo7foxtg/fox; Nestin-Cre mice showed juvenile motor dysfunction, including hindlimb defects and decreased numbers of dopaminergic neurons. Fragmented mitochondria were observed in dopaminergic and cortical neurons. Furthermore, p62- and synuclein-positive Lewy body-like aggregates were identified in neurons. Our findings highlight the unexpected role of the homeostatic level of p62, which is regulated by a non-autophagic system that includes the ubiquitin–proteasome system, in controlling intracellular inclusion body formation. These data indicate that the pathologic processes associated with the proteolytic and mitochondrial degradation systems play a crucial role in the pathogenesis of PD.

Keywords: Parkinson's disease, FBXO7, Dopaminergic neuron, Mitochondria, Synuclein, p62

Main text
Autosomal recessive mutations in the FBXO7 (PARK15) gene are involved in a juvenile form of Parkinsonism with heterogeneous phenotypes characterized by either a classic Parkinson's disease (PD) phenotype, pyramidal tract signs only, or by a combination of Parkinsonism and pyramidal signs [1, 2]. FBXO7 is expressed in various types of tissues, including the gray and white matter of the brain [3].

To determine the function of FBXO7 in vivo, we generated FBXO7foxtg/fox mice. A targeting vector was constructed using 5.0- and 3-kb DNA fragments as the 5' and 3' homologous sequences, respectively (Fig. 1A). The linearized targeting vector was transfected into C57BL/6 embryonic stem (ES) cells. Selected clones were screened for homologous recombination by Southern blotting. Using the 5' external probe and a probe specific for the neo sequence, we confirmed that the clones carried the desired homologous recombination. ES cells derived from these clones were injected into C57BL/6 embryos. Chimeric offspring were crossed with C57BL/6 mice to obtain germline transmission, which was confirmed by Southern blot analysis with the 5' (Fig. 1B upper panel) and 3' (Fig. 1B lower panel) probes. Heterozygous mice were then interbred to obtain homozygous knockout and wild-type control mice. All animals were kept in a pathogen- and odor-free environment, which was maintained under a 12-h light/dark cycle at ambient temperature. Procedures were approved by the Animal Experimental Committee of the Juntendo University Graduate School of Medicine, Tokyo 113-8421, Japan
of Medicine and performed in accordance with the guidelines of the National Institutes of Health and the Juntendo University Graduate School of Medicine. Next, we generated neuronal cell-specific FBXO7\textit{\textsubscript{flox/flox}}. Nestin-Cre mice by crossing the previously characterized FBXO7\textit{\textsubscript{flox/flox}} mice with FBXO7\textit{\textsubscript{flox/flox}}. Nestin-Cre mice harboring the Cre recombinase-coding sequence downstream of a characterized fragment of the Nestin promoter, and confirmed that FBXO7 protein levels were decreased in the whole brain of Cre-expressing mice (Fig. 1C).

FBXO7\textit{\textsubscript{flox/flox}}. Nestin-Cre mice were viable at birth and indistinguishable in appearance from their littermates, but experienced gradually increasing weight loss (Fig. 1D) and had a markedly lower survival rate after birth (Fig. 1F). FBXO7\textit{\textsubscript{flox/flox}}. Nestin-Cre mice began to show impairment in motor coordination tasks and motor behavioral deficits, as determined by the footprint test (Fig. 1G) and runway test (Fig. 1H). We first conducted the footprint test, because a short stride is a characteristic of PD, including in patients with PARK15 mutations. FBXO7\textit{\textsubscript{flox/flox}}. Nestin-Cre mice had a shorter stride than FBXO7\textit{\textsubscript{flox/flox}} mice (Fig. 1G). Regarding the runway test, while FBXO7\textit{\textsubscript{flox/flox}} mice exhibited well-coordinated movement and almost no slips of the forepaw or
Fig. 1 (See legend on previous page.)
hindpaw from the beam. FBXO7flox/flox; Nestin-Cre mice could hardly move on the beam and slipped frequently (Fig. 1H). In particular, the hindpaws of FBXO7flox/flox; Nestin-Cre mice often slipped off the beam (Fig. 1I). Gait disturbance progressed, and by the terminal stage, the majority of affected mice could hardly move.

Mitochondrial damage and dysfunction in the substantia nigra have been previously reported in patients with sporadic PD [4, 5]. These observations support the notion that aberrant mitochondrial function is a critical contributor to pathological neuronal degeneration. Accumulating knowledge regarding PINK1 and Parkin, both of which are associated with mitochondria, has increased our understanding of mitochondrial quality control [6, 7].

FBXO7 and Parkin are interaction partners. FBXO7 is recruited to damaged mitochondria, and is required for the successful recruitment of Parkin. FBXO7 also binds to PINK1, and PINK1, Parkin, and FBXO7 act in concert to control the events leading to mitophagy. Interestingly, the expression of human FBXO7 in a Parkin mutant fly model rescues its phenotype, which is characterized by mitochondrial disruption and locomotor defects [8].

To characterize the damaged mitochondria in FBXO7flox/flox; Nestin-Cre mice, we performed immunohistological (Fig. 1J) and ultrastructural (Fig. 1K) analysis of cortical and dopaminergic neurons of 3-week-old mice. We observed small, round, fragmented mitochondria in these neurons in FBXO7flox/flox; Nestin-Cre mice but not in FBXO7flox/flox mice. Precise quantification revealed that the mitochondrial area was reduced in cortical and dopaminergic cells (Fig. 1L). Together, these observations suggest that mitochondrial fragmentation might be facilitated in FBXO7-deficient mice. In order for damaged mitochondria to be degraded by autophagy, they must be segregated by fission [9]. Our in vivo results are reasonable if FBXO7-mediated mitophagy contributes to mitochondrial degradation systems.

To assess the pathological contribution of damaged mitochondria, we compared tyrosine hydroxylase (TH)-immunoreactive neurons between FBXO7flox/flox; Nestin-Cre and control mice. As demonstrated in the runway test (Fig. 1H, I), FBXO7flox/flox; Nestin-Cre mice exhibited locomotor dysfunction. These FBXO7flox/flox; Nestin-Cre mice had fewer TH neurons in the central portion of the substantia nigra (SN) pars compacta (Fig. 1M, N), where the reduction in TH cell number was most prominent in other PD model mice [3, 10–13].

Finally, to identify the relationship between our model mice and PD, we conducted immunohistological and blot analysis in the brain. Interestingly, in FBXO7flox/flox; Nestin-Cre mice we identified p62-positive aggregates (Fig. 1O) that exhibited a significant increase in p62 levels (Fig. 1P) and synuclein colocalization (Fig. 1Q). These pathologies were identified in the dopaminergic neurons (Fig. 1R), including internal neurons (Fig. 1R arrows). Because p62 aggregates were not seen in endothelial cells (Fig. 1S) or microglia (Fig. 1T), FBXO7flox/flox; Nestin-Cre mice were thought to demonstrate a neuron-specific phenotype. Further research is needed, but our results suggest that loss of FBXO7 may affect p62 and synuclein proteolysis, and impaired mitochondria elimination.

Abbreviations
PD: Parkinson’s disease; EO-PD: Early-onset Parkinson’s disease; FBXO7: F-box protein only 7; UPS: Ubiquitin–proteasome system; TH: Tyrosine hydroxylase; SN: Substantia nigra; ES: Embryonic stem; VTA: Ventral tegmental area; SNpc: Substantia nigra pars compacta.

Acknowledgements
We thank Dr. Souichiro Kakuta (Laboratory of Morphology and Image Analysis, Biomedical Research Core Facilities, Juntendo University Graduate School of Medicine) for technical assistance with electron microscopy.

Author contributions
SS and NH designed the study, analyzed the data, and wrote the paper. SN, TF, US, and NT performed the experiments. All authors read and approved the final manuscript.

Funding
This work was supported by a KAKENHI Grant-in-Aid for Scientific Research from JSPS (19K07850).

Availability of data and materials
All data generated or analyzed during this study are included in this published article.

Declarations

Ethics approval and consent to participate
Not applicable.

Consent for publication
Not applicable.

Competing interests
The authors declare no competing financial interests.

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Received: 7 April 2022 Accepted: 22 May 2022

Published online: 14 June 2022

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