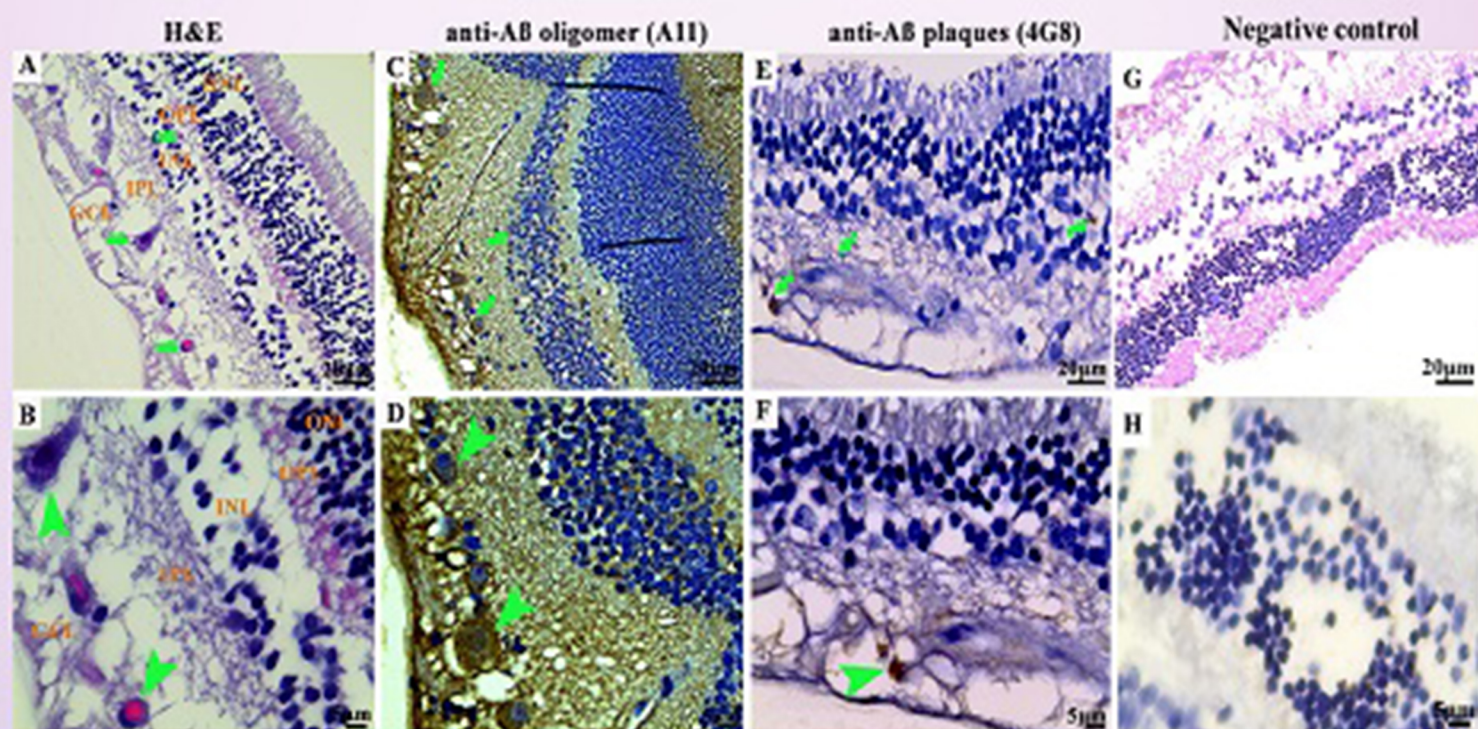


AGEING AND NEURODEGENERATIVE DISEASES

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A sequential deposition of amyloid beta oligomers, plaques and phosphorylated tau occurs throughout life in the canine retina

Umma Habiba, John Morley, Mark Krockenberger,
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Prof. Le is currently the deputy director of the Academic Committee and the director of the Institute of Neurology, Sichuan Academy of Medical Science-Sichuan Provincial People's Hospital. He is also a life-time Professor of Neurology and the director of Center for Translational Research on Neurological Diseases, the First Affiliated Hospital of Dalian Medical University. He has been engaged in clinical practice, teaching and research of neurology for almost 40 years, including 24 years of work in Baylor College of Medicine as a Professor of Neurology and Director of the Parkinson's Disease Research Center; and 10 years of working as a Professor of Neuroscience and Head of Neurogenomic Lab in the Institute of Health Science of Chinese Academy of Science jointed with Shanghai Jiao-Tong University School of Medicine.

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High-quality original articles, reviews, case reports, commentaries are welcomed from ageing and neurodegenerative diseases, particularly on the basic mechanisms of ageing and their roles in the onset and progression of neurodegenerative diseases. The journal aims to report innovative research advances on the following topics:

1. The cellular and molecular mechanisms of ageing and the pathogenesis of neurodegenerative diseases;
2. The associations between neurodegenerative diseases and the biological bases of ageing with a focus on: genomic instability, epigenetic alterations, telomere attrition, protein degradation system failure, mitochondrial dysfunction, cellular senescence, nutrient sensing deregulation, stem cell exhaustion, intercellular communication impairment, etc.;
3. Translational research into prevention and treatment of age-related neurodegenerative diseases;
4. Mechanistic bases for epidemiological observations in aging-related neurodegenerative diseases.



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GENERAL INFORMATION

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AND is a peer-reviewed and open access multidisciplinary journal that publishes high-quality original articles, reviews, case reports, commentaries, letters to editor, etc. Ageing is a major risk factor for neurodegeneration, and the prevalence of ageing-related neurodegenerative diseases such as Alzheimer's disease, Parkinson's disease, amyotrophic lateral sclerosis, etc. continues to rise with the increased ageing population. Unfortunately, there are no effective treatments available for the age-related neurodegenerative diseases. Thus, to develop successful interventions, it is important to investigate the basic mechanisms of ageing and their roles in the onset and progression of neurodegenerative diseases. Therefore, we plan to launch this new journal, which is aimed to report innovative research advances on the following topics:

1. The cellular and molecular mechanisms of ageing and the pathogenesis of neurodegenerative diseases;
2. The associations between neurodegenerative diseases and the biological bases of ageing with a focus on: genomic instability, epigenetic alterations, telomere attrition, protein degradation system failure, mitochondrial dysfunction, cellular senescence, nutrient sensing deregulation, stem cell exhaustion, intercellular communication impairment, etc.;
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Letter to Editor

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Chronic intracerebroventricular administration is a reliable method in brain studies on monkeys

Lin-Heng Zhang^{1,2}, Hong-Yi Yang³, Jing Wu³, Yun Wu³, Lu-Lu Wang³, Hai-Yang Tong³, Jin Zhang³, Wen-Chao Wang¹, Rong-Yao Huang^{1,4}, Jiang-Lei Xu^{1,4}, Jing Su¹, Xun-Ran Luo¹, Yong Yin⁵, Shi-Hao Wu¹, Xin-Tian Hu^{1,6,7}

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Abstract

Intracerebroventricular (ICV) administration through cannulas is a direct way to deliver large molecules and substances that are blocked by the blood-brain barrier into the central nervous system (CNS). It is widely used in brain studies on monkeys. However, this method is invasive, as it requires guide cannulas to be implanted into the brain. Whether the long-term implantation of the cannula and the administration of molecule-delivering vehicles,



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usually saline, can affect the brain by inducing chronic CNS inflammation or even worse brain atrophy, remains an issue to be solved. To answer this question, we investigated inflammatory markers and brain structures on three vehicle-control monkeys who received cannula implantation and one-year ICV saline administration in another study. During the experiment, the monkeys' cerebrospinal fluid (CSF) samples were collected periodically, and the level of three classic inflammatory markers (IL-1 β , IL-6, and TNF- α) were measured by electrochemiluminescence immunoassay. The monkeys' brain structures were imaged *in vivo* periodically by 9.4 Tesla magnetic resonance imaging, which can provide the best-resolution magnetic resonance images of living monkeys, and the volume of the hippocampus was measured to evaluate the brain atrophy. The data reveal that, during the administering period, the long-term levels of the inflammatory markers in the CSF and the volumes of the hippocampus did not change significantly compared with the baseline. These results suggest that the long-term ICV administration of saline through cannulas did not induce chronic neuroinflammation or brain atrophy in these rhesus monkeys, suggesting chronic ICV administration via implanted cannulas is a reliable method in monkey brain research.

Keywords: Chronic ICV administration, neuroinflammation, brain atrophy

The central nervous system is protected by the blood-brain barrier (BBB), which is a semipermeable structure composed of tight endothelial junctions to provide a physical barrier for the entry of large molecules^[1]. It restricts circulating toxic particles but facilitates nutrients to be delivered into the brain. Meanwhile, some therapeutic substances and test articles are also blocked by the BBB^[2]. Intracerebroventricular (ICV) injection physically breaches the BBB and enables the direct administration of large molecules, including chemicals, virus vectors, and therapeutic substances, into the central nervous system (CNS) through pre-implanted guide cannulas. Studies on the distribution of injected large molecules revealed that large molecules including adenovirus vectors and proteins can spread into the subarachnoid space, meninges, and choroid plexus along with the flow of the CSF and diffuse into the brain parenchyma adjacent to these sites^[3-5]. Although this method is an effective way to deliver molecules into the CNS, it is invasive and could affect the brain in many ways.

One major concern in chronic cannula-guided ICV administration comes from the implantation of the guide cannulas. To ease the repeated ICV injection, one or two guide cannulas must be implanted into the lateral ventricles. Drugs or test articles are injected into the lateral ventricles via the guide cannulas without repeated penetration of the brain parenchyma. The implantation of guide cannulas causes acute brain injury. It is well established that an acute brain injury induces acute inflammation in the brain, which is one important step in the repair of the injury. However, it is not clear if long-term implantation of cannulas can induce chronic neuroinflammation.

Chronic neuroinflammation is a persistently activated immune response in the CNS, which is always marked by sustained release of pro-inflammatory cytokines, including interleukins, tumor necrosis factor- α (TNF- α), transforming growth factor- β (TGF- β), and interferon- γ (IFN- γ)^[6,7]. Pro-inflammatory cytokines exert various functions in neuroinflammation, including induction of chemokines and adhesion molecules and recruitment of the peripheral immune cells into the CNS, leading to neuronal apoptosis and activation of the glial cells^[6]. Increasing evidence suggests that cytokine-mediated neuroinflammation plays an important role in initiating or deteriorating neurodegenerative diseases^[8-10]. Clinical evidence reveals that the pro-inflammatory cytokines are elevated in the body fluid of individuals with neurodegenerative diseases, including Alzheimer's disease, Parkinson's disease, and amyotrophic lateral sclerosis^[11-13]. Therefore, if cannula-based ICV administration can induce chronic neuroinflammation, the results of studies on brain diseases and mechanisms would be seriously confounded, and the application of this method would be seriously limited.

Non-human primates are phylogenetically closer to humans than rodents. They share with humans many brain regions, circuits, and neural cells, which are critical for higher cognitive capacity. Thus, monkeys are widely used in the study of brain diseases and mechanisms, including in CNS disease modeling with ICV administration^[14-16]. Unfortunately, no study on whether cannula-based ICV administration can induce chronic neuroinflammation in monkeys has been reported thus far.

To investigate if long-term cannula implantation and ICV saline administration can induce chronic neuroinflammation in monkeys, the levels of three most studied pro-inflammatory cytokines (IL-1 β , IL-6, and TNF- α) in the CSF of three vehicle-control monkeys used as vehicle-control in another study were continuously monitored during one-year cannula-guide ICV saline administration [see Supplementary Materials for more method details]. The measurement of IL-1 β , IL-6, and TNF- α in the CSF was carried out by an electrochemiluminescence immunoassay system, which has an extremely low detection limit in the measurement of biomarkers in body fluid.

The levels of CSF IL-1 β , IL-6, and TNF- α were measured at 12 weeks before the administration (the baseline) and the 4, 9, 14, 19, 24, 29, 34, and 39 weeks after the administration [Figure 1A]. Kruskal-Wallis H test revealed the effect of ICV saline administration on the average levels of IL-1 β and IL-6 was not significant ($P = 0.597$ for IL-1 β , $P = 0.112$ for IL-6, Figure 1B and C). The levels of CSF TNF- α were around the low detection limit, and no meaningful statistics could be applied. The data for TNF- α are not shown. In summary, the results reveal that there was no elevated CSF IL-1 β , IL-6, or TNF- α during the administration, which suggests there was no significant neuroinflammation during the ICV saline administration.

In addition, we also investigated whether there was any brain atrophy caused by long-term cannula implantation and ICV administration, considering reports on brain atrophy induced by traumatic brain injury and neuroinflammation^[17-19]. The structures of the monkeys' brains were imaged periodically by magnetic resonance imaging (MRI) (see Supplementary Materials for more method details). To increase the accuracy in the measurement of hippocampus volumes, the monkeys' brains were scanned by a 9.4 Tesla MRI system, which provides the highest resolution structural images currently available on living monkeys.

The measurement was carried out one week before the administration (the baseline) and 4, 9, 14, 19, 25, 31, 37, 43, and 49 weeks after administration [Figure 2A]. First, longitudinal MR images from the same subject were realigned to examine general structural changes in the brain. No significant brain atrophy was observed on MR images from each time point compared with the baseline [Figure 2B]. Then, the volume of the hippocampus was calculated to evaluate the atrophy in the brain because it is one of the most vulnerable structures in the brain and has a clear contour on MR images. Kruskal-Wallis H test revealed that the effect of ICV saline administration on the average volumes of total hippocampus was not significant ($P = 0.775$, Figure 2C). We further compared the average volumes of the left and right hippocampus independently. Kruskal-Wallis H test also revealed that the difference between each time point in the volumes of the left and right hippocampus were both not significant ($P = 0.770$ for the left hippocampus, $P = 0.606$ for the right hippocampus, Figure 2D and E). In summary, the results from the ultra-high-resolution MRI scan revealed that there was no reduction in the volume of the hippocampus, which suggests there was no significant atrophy in the brain.

Here, we demonstrate that, compared with the baseline, there was no elevated IL-1 β or IL-6 in the CSF of the three cannula implanted monkeys during 1-year ICV saline administration, suggesting that the cannula-guided long-term ICV administration technique did not induce chronic neuroinflammation in the monkeys. The other finding that there was no significant decrease in hippocampus volumes supports this

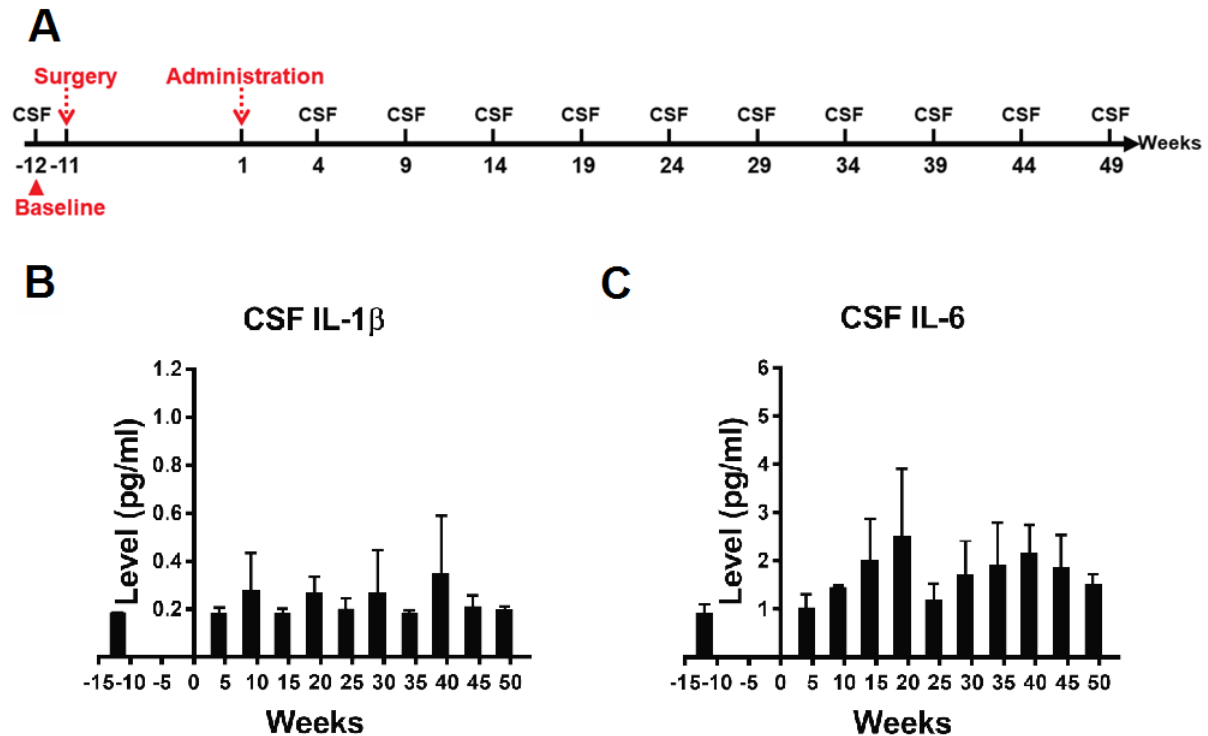


Figure 1. (A) The schedule of CSF sample collection. Cannula implantation was carried out in Week 11. ICV administration began in Week 1. CSF samples were collected in Weeks 12, 4, 9, 14, 19, 24, 29, 34, 39, 44, and 49. CSF sample collected at Week -12 was taken as the baseline. (B,C) Kruskal-Wallis H test revealed no significant difference between different time points in the average levels of CSF IL-1 β ($P = 0.597$) and IL-6 ($P = 0.112$). Data are presented as mean \pm standard deviation (SD). CSF: Cerebrospinal fluid; ICV: intracerebroventricular.

conclusion.

To increase the accuracy of the measurement of inflammatory cytokines in the body fluid, an advanced detection method was employed in this study. Inflammatory cytokines in the body fluid exert their physiological functions at low levels, and many of them are usually below the low detection threshold under physiological conditions using common enzyme-linked immunosorbent assay kits^[20]. Advanced techniques are developed to detect the cytokines in the body fluids, with much higher detection sensitivity and smaller specimen volumes. Among the newly developed assay techniques, the MSD multiplex immunoassay platform represents a combination of electrochemiluminescence and patterned arrays with an ultra-low detection limit, which is also one of the most used methods in the measurement of cytokines in body fluids nowadays. The detection sensitivity can reach 0.1 pg/mL for some cytokines using this method. The reliability of this method was validated in three recent studies, which proved that this method had a high detection rate for most of the cytokines and displayed a moderate-to-excellent intra-individual variability^[21-23]. In our experiment, the detection rate was 100% for IL-1 β and IL-6, but only 60% for the TNF- α , which indicates that the level of TNF- α in the CSF of monkeys under physiological conditions was extremely low.

The hippocampus is one of the most vulnerable structures in the brain under many conditions such as injury, aging, and neurodegenerative diseases^[24], and it is lying in the inferior horn of the lateral ventricle, directly affected by ICV administration. Due to the vulnerability and clear contour in MR images, the hippocampus volume was taken as an indicator of brain atrophy in this study. To investigate if ICV saline

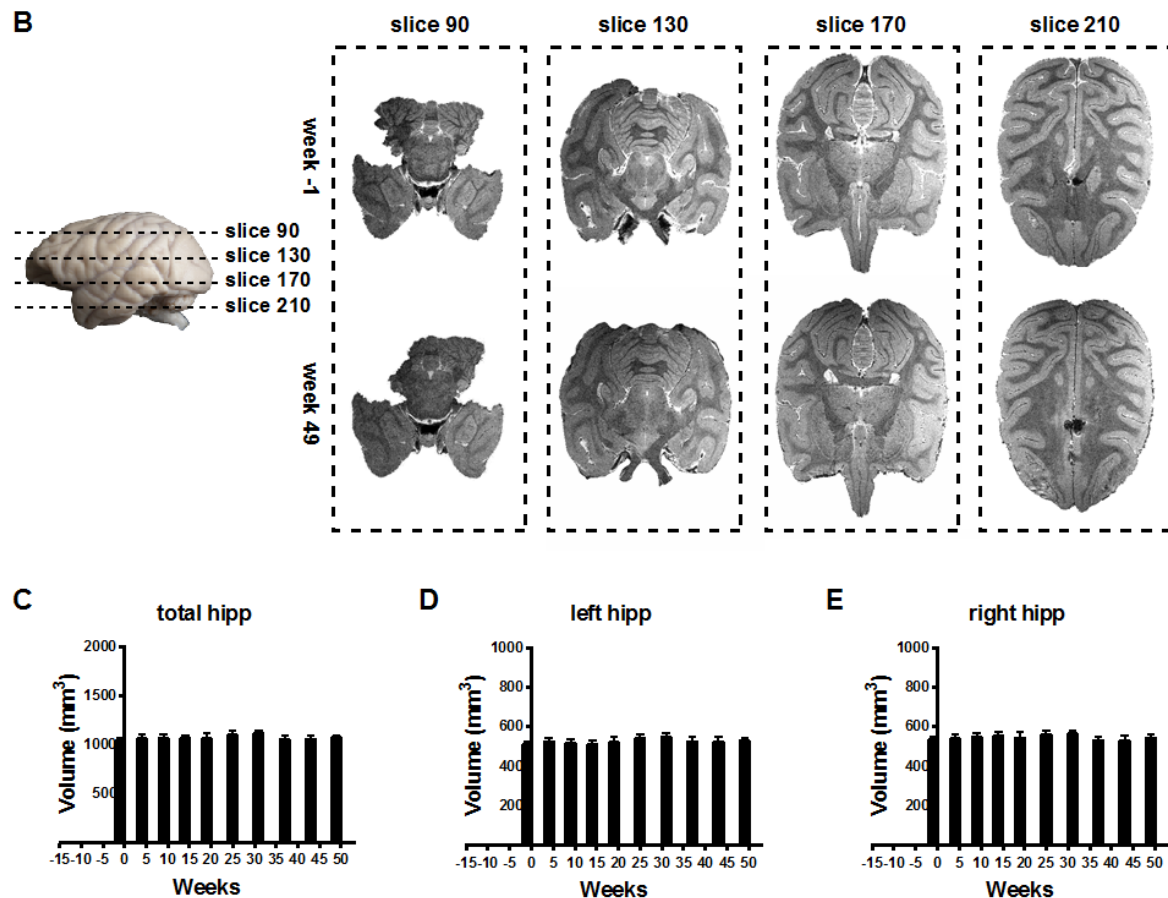


Figure 2. (A) Schedule of MRI scans. Cannula implantation was carried out in Week -11. ICV administration began in Week 1. MR images were acquired in Weeks 1, 4, 9, 14, 19, 25, 31, 37, 43, and 49. MR images acquired in Week -1 were taken as the baseline. (B) Representative MR images show there was no significant brain atrophy in Week 49 when compared with the baseline (Week 11). (C-E) Kruskal-Wallis H test revealed no significant difference between different time points in the average total hippocampus volumes ($P = 0.775$), the average left hippocampus volumes ($P = 0.770$), and the average right hippocampus volumes ($P = 0.606$). Data are presented as mean \pm standard deviation (SD). MRI: Magnetic resonance imaging; ICV: intracerebroventricular.

administration can induce atrophy, ultra-high magnetic field MRI was employed to image the brain. The fact that we did not find any decrease in the hippocampus volumes strongly suggests no brain atrophy in the monkeys' brains, which also supported the conclusion that there was no significant inflammation in the brain.

Taken together, our results reveal that long-term cannula implantation and ICV administration of drug-vehicle do not induce chronic neuroinflammation and brain atrophy in rhesus monkeys, which suggests that chronic ICV administration is a reliable method for brain studies on monkeys.

DECLARATIONS

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scanning.

Author's contributions

Designed the experiment: Hu XT, Yin Y, Wu SH, and Zhang LH

Carried out the cannula implantation surgery: Wu J, Wang WC, Huang RY, Xu JL, and Zhang LH

Carried out the MRI scanning: Yang HY, Wu Y, Wang LL, and Zhang LH

Carried out the CSF sample collection: Tong HY, Zhang J, Zhang LH, Su J, and Luo XR

Carried out the data analysis: Zhang LH

Wrote the manuscript: Zhang LH

Revised the manuscript: Hu XT and Yin Y

All authors read and approved the final version of the manuscript.

Availability of Data and Materials

Not applicable.

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Conflicts of Interest

All authors declared that there are no conflicts of interests.

Ethical Approval and Consent to Participate

All experimental procedures were carried out following “Guide for the Care and Use of Laboratory Animals”^[20] and approved by the Animal Care and Use Committee (IACUC) of the Kunming Institute of Zoology, Chinese Academy of Sciences (IACUC No.: IACUC19001).

Consent for Publication

Not applicable.

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Selective expression of neurodegenerative diseases-related mutant p150^{Glued} in midbrain dopaminergic neurons causes progressive degeneration of nigrostriatal pathway

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Abstract

Aim: Missense mutations of dynactin subunit p150^{Glued} have been associated with multiple neurodegenerative diseases, including Perry syndrome, characterized by inherited parkinsonism, depression, weight loss, and hypoventilation. The current study investigated how the pathogenic mutant p150^{Glued} affects the integrity and function of the nigrostriatal dopaminergic (DA) pathway *in vivo*.

Methods: Using a tetracycline-controlled transcriptional regulation system, transgenic mouse models were generated with selective overexpression of wild-type, motor neuron disease-related G59S mutant, or Perry syndrome-related G71R mutant human p150^{Glued} in midbrain DA neurons. A series of behavioral, neuropathological, neurochemical, electrochemical, and biochemical studies were performed on the mice to examine and compare the pathogenic impact of the two mutant p150^{Glued} on the survival and function of midbrain DA neurons.



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Results: Compared with non-transgenic control mice, transgenic mice overexpressing wild-type human p150^{Glued} showed neither motor phenotypes nor pathological, functional, or biochemical abnormalities of the nigrostriatal DA pathway. Transgenic mice overexpressing G59S mutant p150^{Glued} displayed weight loss, motor deficits, early-onset defects in dopamine transmission, and early-onset loss of DA neurons and axons. Transgenic mice overexpressing G71R p150^{Glued} mutant exhibited hyperactivities, impaired motor coordination, early-onset dysfunction of dopamine uptake, and late-onset loss of DA neurons and axons. In addition, overexpression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons preferentially downregulated the expression of dopamine transporter at dopaminergic axon terminals. Furthermore, G59S mutant p150^{Glued} rather than G71R mutant p150^{Glued} formed aggregates in midbrain DA neurons *in vivo*, and the aggregates trapped dynein/dynactin, co-localized with lysosomes, and upregulated ubiquitination.

Conclusion: These findings demonstrate that selective expression of either G59S or G71R mutant p150^{Glued} in mouse midbrain DA neurons leads to progressive degeneration of the nigrostriatal DA pathway and indicate that G59S and G71R mutant p150^{Glued} exhibit differential pathogenic impact on the survival and function of midbrain DA neurons *in vivo*.

Keywords: Perry syndrome, motor neuron disease, *DCTN1* gene, dynactin subunit p150^{Glued}, missense mutation, nigrostriatal pathway, midbrain dopaminergic neuron, neurodegeneration

INTRODUCTION

Cytoplasmic dynein and its cofactor dynactin mediate microtubule minus-end-directed trafficking, performing critical roles in mitosis, intracellular positioning and movement of organelles, retrograde axonal transport, and synapse formation and stability^[1-3]. Dynactin is a macromolecular protein complex consisting of more than ten types of subunits, with the largest subunit p150^{Glued} binding directly with dynein and microtubule^[4]. Missense mutations of *DCTN1*, the gene encoding p150^{Glued}, have been linked to different neurodegenerative diseases, in which the G59S missense mutation was linked to autosomal dominant motor neuron disease, and 13 more mutations (F52L, K56R, G67D, K68E, G71A/R/E/V, T72P, Q74P, Y78C/H, and Q93H) were associated with Perry syndrome^[5-8]. Perry syndrome is an autosomal dominant neurodegenerative disorder characterized by a series of symptoms (parkinsonism, depression, central hypoventilation, and weight loss) and pathology of severe neuronal loss in the *substantia nigra* and *locus coeruleus*^[9,10]. However, the underlying pathogenic mechanism and potential therapeutic target of neurodegeneration in Perry syndrome remain elusive.

P150^{Glued} contains two microtubule-binding domains (MTBDs) at its N-terminus: the cytoskeleton-associated protein and glycine-rich (CAP-Gly) domain, which interacts with both tubulin and microtubule end binding proteins, and the adjacent serine-rich basic domain, which has an affinity for tubulin^[11,12]. The MTBDs of p150^{Glued} are required to initiate retrograde transport and maintain microtubule stability in axons^[13-15]. Deleting the MTBDs in p150^{Glued} upregulates microtubule acetylation, increases the surface level of ionotropic glutamate receptors, disrupts neuromuscular junctions, and leads to selective degeneration of spinal motor neurons^[16]. Motor neuron disease-related G59S mutation of p150^{Glued} and all the Perry syndrome-related mutations are point mutations clustered within or close to the CAP-Gly domain, resulting in loss-of-function of the MTBD^[6-8]. Furthermore, G59S mutant p150^{Glued}, as well as G71A, G71E, G71R, T72P, Q74P, and F52L mutant p150^{Glued}, form cytoplasmic aggregates *in vitro* in a time- and concentration-dependent manner^[17-19], suggesting both loss-of-function and toxic-gain-of-function mechanisms might be involved in *DCTN1* mutation-induced neurodegeneration. Additionally, G59S mutant p150^{Glued} is less stable^[20], and both G59S and Perry syndrome-related mutant p150^{Glued} compromise normal functions of dynactin p150^{Glued} in a dominant-negative fashion when overexpressed^[13,14,21], suggesting

a dominant-negative mechanism for *DCTN1* mutation-induced neurodegeneration. Movement disorder and the underlying neuropathology, including loss of spinal motor neurons and degeneration of axon and neuromuscular junctions, have been reported in G59S mutant p150^{Glued} knock-in and transgenic mice^[20-22]. Behavioral defects have also been found in G71A mutant p150^{Glued} knock-in and transgenic mice^[23,24]. However, how the pathogenic mutant p150^{Glued} affects the integrity and function of the nigrostriatal DA pathway has not been critically examined in animal models.

Here, we investigated and compared the pathogenic impact of two different neurodegenerative diseases-related mutant p150^{Glued} in the survival and function of midbrain DA neurons *in vivo*. Transgenic mouse models with selective overexpression of wild-type, G59S mutant, or G71R mutant human p150^{Glued} in midbrain DA neurons were generated using a tetracycline-controlled transcriptional regulation system. A series of experiments were performed to identify any behavioral, neuropathological, neurochemical, electrochemical, and biochemical abnormalities in the transgenic mice. Our findings highlight the importance of p150^{Glued} and its MTBDs in maintaining the survival and function of midbrain DA neurons and reveal the pathogenic similarity and difference between G59S and G71R mutant 150^{Glued}.

METHODS

Animals

The *tetO*-human *DCTN1* WT transgenic mice (for simplicity, referred to as tetO-WT) carrying the wild-type (WT) human p150^{Glued} gene (*DCTN1*) under the control of tetracycline operator (*tetO*) were generated as described previously^[25,26]. The tetO-WT mice were mated with *Pitx3-IRE2-tTA* mice (referred to as tTA, a “Tet-Off” tool, JAX stock #021962), which predominantly express tetracycline-controlled transactivator protein (tTA) in midbrain DA neurons^[27]. This breeding strategy produced *Pitx3-IRE2-tTA/tetO*-human *DCTN1* WT transgenic mice (referred to as WT Tg), which had the selective expression of wild-type human p150^{Glued} in midbrain DA neurons, and littermate controls [non-transgenic (nTg), tetO-WT, and tTA mice].

Following the same protocol, *Pitx3-IRE2-tTA/tetO*-human *DCTN1* G59S transgenic mice (referred to as G59S Tg), which had the selective expression of G59S mutant human p150^{Glued} in midbrain DA neurons, and littermate controls (nTg, tetO-G59S, and tTA mice) were obtained.

Following the same protocol, *Pitx3-IRE2-tTA/tetO*-human *DCTN1* G71R transgenic mice (referred to as G71R Tg), which had the selective expression of G71R mutant human p150^{Glued} in midbrain DA neurons, and littermate controls (nTg, tetO-G71R, and tTA mice) were obtained.

The mice were housed in a 12 h light/dark cycle and fed a regular diet *ad libitum*. All mouse work followed the guidelines approved by the Institutional Animal Care and Use Committees of National Institute on Aging, NIH.

Genotyping

Genomic DNA was extracted from tail biopsy using DirectPCR Lysis Reagent (Viagen Biotech) and used for PCR amplification. Genotyping was performed with specific sets of PCR primers for *Pitx3-IRE2-tTA* transgene (GACTGGCTTGCCCTCGTCCCA and GTGCACCGAGGCCCCAGATCA)^[27] and *tetO*-human *DCTN1* transgene (AGGAGATGGCTGATACTGCTGATG and TTCATCGCTTCCAAGTCTCCCACA).

Behavior tests

Cohorts of WT Tg mice and littermate controls [nTg ($n = 10$), tTA ($n = 10$), tetO-WT ($n = 10$), and WT Tg ($n = 10$)] were repeatedly weighed and assessed for motor function at 1, 6, 12, and 18 months of age.

Cohorts of *G59S Tg* mice and littermate controls [nTg ($n = 10$), tTA ($n = 11$), tetO-*G59S* ($n = 10$), and *G59S Tg* ($n = 10$)] were repeatedly weighed and assessed for motor function at 1, 6, 12, and 18 months of age.

Cohorts of *G71R Tg* mice and littermate controls [nTg ($n = 11$), tTA ($n = 10$), tetO-*G71R* ($n = 10$), and *G71R Tg* ($n = 10$)] were repeatedly weighed and assessed for motor function at 1, 6, 12, and 18 months of age.

Open-field test. Mice were placed in the open-field apparatus of the Flex-Field Activity System (San Diego Instruments). Ambulatory, rearing, and fine movements of mice were traced and quantified with infrared photobeams sensors and Flex-Field software in the unit as the number of beam breaks per 30 min.

Rotarod test. Mice were placed onto a rotating rod with auto-acceleration from 0 to 40 rpm within 1 min (San Diego Instruments). The time length that each mouse stayed on the rotating rod was measured across three trials.

Histology, immunostaining, and light microscopy

Mice were sacrificed and perfused with 4% paraformaldehyde (PFA) in phosphate-buffered saline (PBS). Mouse brains were dissected, post-fixed in 4% PFA/PBS solution overnight, and submerged in 30% sucrose in PBS for at least 72 h. The 40 μm -thick brain sections were prepared using CM1950 cryostat (Leica) and stained with antibodies specific to human and mouse p150^{Glued} (amino acid 3-202 at the N-terminus of p150^{Glued}, 1:200, BD Biosciences, Catalog # 610474, recognizing wild-type p150^{Glued} but not *G59S* mutant p150^{Glued}, *G71R* mutant p150^{Glued}, or p135+), human and mouse p150^{Glued} and p135+ (amino acid 1266-1278 at the C-terminus of p150^{Glued} and p135+, 1:500, Abcam, Catalog # ab11806, recognizing wild-type, *G59S* mutant and *G71R* mutant p150^{Glued}, and p135+), tyrosine hydroxylase (TH, 1:2500, Pel-freez, Catalog # P40101-150; 1:500, ImmunoStar, Catalog # 22941), dynactin subunit p50 (1:1000, Millipore, Catalog # AB5869P), dynactin subunit actin-related protein 1 (ARP1, 1:1000, Millipore, Catalog # AB6058), cytoplasmic dynein light intermediate chain (DLIC, 1:500, Abcam, Catalog # ab123901), lysosomal associated membrane protein 2 (LAMP2, 1:500, Abcam, Catalog # ab13524), and ubiquitin (1:500, Enzo Life Sciences, Catalog # BML-PW0930), as suggested by manufacturers. Fluorescent signals were visualized with Alex Fluor-conjugated secondary antibody (1:500, Invitrogen, Catalog # A-11055, A-21202, A-21206, A-21208, A-21447, A10036, and A10040) and examined using LSM 880 laser-scanning confocal microscope (Zeiss). The paired images in all figures were collected using identical gain and offset settings, processed uniformly, and displayed as either a single optic layer or the maximum-intensity projections of the z-series confocal stack at 1.0 μm intervals.

Stereology

The number of midbrain dopaminergic neurons was estimated by unbiased stereology, as described previously^[27,28]. According to Paxinos and Franklin's *The Mouse Brain in Stereotaxic Coordinates* (4th edition), a series of coronal sections (40 μm thick) across the midbrain (every fourth section from Bregma -2.54 to -4.24 mm, 10 sections per animal) were stained with an antibody specific to TH (1:2500, Pel-Freez, Catalog # P40101-150). After subsequent staining with Vectastain Elite ABC Kit, sections were visualized with DAB Kit (Vector Laboratories). Bright-field images were captured, and the number of TH-positive neurons was assessed using the optical fractionator function of Stereo Investigator 10 (MicroBrightField) under the Axio microscope Imager A1 (Zeiss). Counters were blinded to the genotypes of mice, and the sampling scheme was designed to have a coefficient of error less than 10% for the reliability of results.

Image analysis

Quantitative analysis of images was performed with ImageJ (NIH). After converting image type to 8-bit, areas of interest were selected with Freehand tools, and then mean optical intensities or area fractions were measured. The mean intensity for the background area was subtracted from the selected area to determine the net mean intensity. For quantitative analysis of DA neurite dystrophy, 5 tile images per animal (5 sections per animal and 1 tile image per section) from the *substantia nigra pars reticulata* (SNr) were taken with a 20 \times lens. Dystrophic DA neurites were defined as neuritic varicosity $\geq 25 \mu\text{m}^2$. The area of dystrophic TH-positive DA neurites and the area of the SNr were quantified with ImageJ (NIH). For quantitative analysis of DA axon degeneration, 50 images per animal (10 sections per animal and 5 images per section) from the dorsal striatum were taken with a 63 \times lens. The area fraction of TH-positive DA axon terminals in each image was quantified with ImageJ (NIH). For quantitative analysis of G59S mutation-induced p150^{Glued} aggregates, aggregates were defined as puncta $\geq 1 \mu\text{m}^2$.

High-performance liquid chromatography (HPLC)

The dorsal striatum of mice was dissected, weighed, and homogenized by sonication in 0.1 N perchloric acid containing 100 μM EDTA (500 μl buffer per 100 mg tissue). After centrifugation, the supernatant was collected and assayed for dopamine and DOPAC content by HPLC with electrochemical detection, as described previously^[27,28]. Briefly, the mobile-phase solution containing octanesulfonic acid was used as an ion-pairing agent and pumped isocratically through a reversed-phase liquid chromatographic column. DA and DOPAC were measured by the current produced after exposure of the eluate to a flow-through electrode set to oxidizing and then reducing potentials in series, with recordings from the last electrode reflecting reversibly oxidized species.

Fast-scan cyclic voltammetry (FSCV)

The kinetics of striatal dopamine release evoked by electrical stimulation was measured using FSCV, as described previously^[27,28]. The 400 μm -thick slices of dorsal striatum were prepared and transferred into 32 °C oxygenated artificial cerebrospinal fluid (aCSF: 126 mM NaCl, 2.5 mM KCl, 1.2 mM NaH_2PO_4 , 2.4 mM CaCl_2 , 1.2 mM MgCl_2 , 25 mM NaHCO_3 , 11 mM glucose, 20 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid, and 0.4 mM L-ascorbic acid). Cylindrical carbon-fiber microelectrode prepared with T650 fibers (Goodfellow) was held at -0.4 V, and the potential was increased to 1.2 V and back at 400 V/s every 100 ms using a triangle waveform. Dopamine release in striatal slices was evoked by rectangular, electrical pulse stimulation (100-400 μA , 0.6 ms per phase, biphasic) applied every 5 min. Data were collected and analyzed with the Demon Voltammetry and Analysis software suite^[29]. Ten cyclic voltammograms of charging currents were recorded as background before stimulation, and the average of these responses was subtracted from data obtained during and after stimulation. Maximum amplitudes of extracellular dopamine transients were calculated from input/output function (I/O) curves. I/O curves were plotted with stimulus current against the concentration of dopamine response amplitude over a range of stimulus intensities. The time constant of the slope decay (τ) was used as an uptake kinetic indicator of evoked dopamine release. Following experiments, electrodes were calibrated using 1 and 10 μM dopamine solutions in aCSF.

Western blot analysis

Mouse brain tissues were homogenized and sonicated in 1% SDS lysis buffer (50 mM Tris-HCl, 150 mM NaCl, 2 mM EDTA, pH 7.5, and 1% SDS) supplemented with Protease/Phosphatase Inhibitor Cocktails (ThermoFisher Scientific). Lysates were centrifuged at 15,000 g for 15 min at 4 °C. Supernatants were collected, and protein content was quantified with the bicinchoninic acid (BCA) assay kit (ThermoFisher Scientific). Equal amounts of total proteins were size-fractionated with NuPage 4%-12% Bis-Tris gel in MES or MOPS running buffer (ThermoFisher Scientific). After transferring to nitrocellulose membranes with the

Trans-Blot Turbo Transfer System (Bio-Rad), proteins were immunoblotted with specific primary antibodies to human p150^{Glued} (1:1000, Covance, custom-made), human and mouse p150^{Glued} and p135+ (1:1000, Abcam, Catalog # ab11806), dynactin subunit DCTN4 (1:1000, Abcam, Catalog # ab170107), dynactin subunit p50 (1:1000, BD Biosciences, Catalog # 611002), dynactin subunit actin-related protein 1 (ARP1, 1:1000, Sigma-Aldrich, Catalog # A5601), tyrosine hydroxylase (TH, 1:1000, Sigma-Aldrich, Catalog # T1299), dopamine transporter (DAT, 1:1000, Millipore, Catalog # MAB369), and β -actin (1:5000, Sigma-Aldrich, Catalog # A1978). Protein signals were visualized using the Odyssey System with IRDye secondary antibodies (LI-COR Biosciences, Catalog # 926-32212, 926-32219, 926-3221, 926-32411, and 926-68023) and quantified with NIH ImageJ software.

Statistical analysis

Statistical analysis was performed using GraphPad Prism 9 (GraphPad Software). Data are presented as mean \pm SEM. Statistical significance was determined by comparing means of different groups using unpaired t-test, one-way, and two-way ANOVA with post hoc analysis. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$.

RESULTS

Generation of mouse models with selective expression of wild-type or mutant human p150^{Glued} in midbrain DA neurons

To investigate the involvement and mechanism of p150^{Glued} in neurodegeneration *in vivo*, three lines of *tetO*-human *DCTN1* mice (*tetO*-WT, *tetO*-G59S, or *tetO*-G71R) with inducible expression of wild-type or mutant human p150^{Glued} under the transcriptional control of tetracycline operator (*tetO*) were generated. The *tetO*-human *DCTN1* mice were then crossed with *Pitx3*-*IRE2*-*tTA* knock-in mice to obtain *Pitx3*-*IRE2*-*tTA*/*tetO*-human *DCTN1* WT (WT Tg), *Pitx3*-*IRE2*-*tTA*/*tetO*-human *DCTN1* G59S (G59S Tg), or *Pitx3*-*IRE2*-*tTA*/*tetO*-human *DCTN1* G71R (G71R Tg) bigenic mice [Figure 1A]. Since the tetracycline transactivator (tTA) is specifically expressed in midbrain DA neurons under the endogenous *Pitx3* promoter^[27], it turned on the expression of human wild-type or mutant p150^{Glued} selectively in midbrain DA neurons [Figure 1A]. The target overexpression of human p150^{Glued} in midbrain DA neurons of one-month-old WT mice was visualized by immunostaining [Figure 1B]. Using a human p150^{Glued}-specific antibody, Western blot confirmed the expression of human p150^{Glued} in the midbrain homogenates of one-month-old WT Tg, G59S Tg, and G71R Tg mice but not *nTg* mice [Figure 1C and D]. The steady level of G59S human p150^{Glued} was significantly less (approximately 30%) compared with WT or G71R human p150^{Glued}. Western blot further revealed substantially increased levels of total (human and mouse) p150^{Glued} (approximately two-fold) and other dynactin subunits, including DCTN4, p50, and ARP1, but markedly reduced level of p135+ (alternative splicing isoforms of *Dctn1* lacking the N-terminus) in the midbrain homogenates of WT Tg, G59S Tg, and G71R Tg mice compared to the *nTg* mice [Figure 1C and D].

G59S Tg and G71R Tg mice exhibit profound motor abnormalities.

As motor symptoms and weight loss are the main clinical manifestations in familial motor neuron disease and Perry syndrome cases with *DCTN1* mutation^[5], WT Tg, G59S Tg, G71R Tg mice, and their littermate controls were weighed, tested for locomotion activity in the open-field test, and examined for motor coordination in the rotarod test at 1, 6, 12, and 18 months of age. WT Tg mice developed normally and exhibited comparable body weight and motor function with littermate single- or non-transgenic mice in the open field test and the rotarod test [Figure 2A-D]. G59S Tg mice displayed substantial weight loss starting at 12 months of age, normal ambulatory movement, dramatic reduction of rearing movement at 18 months of age, and significant impairment in rotarod test starting at six months of age [Figure 2E-H]. G71R Tg mice showed normal body weight, significant hyperactivities of ambulatory movement and rearing, and substantially impaired rotarod performance starting at six months of age [Figure 2I-L]. Together, these

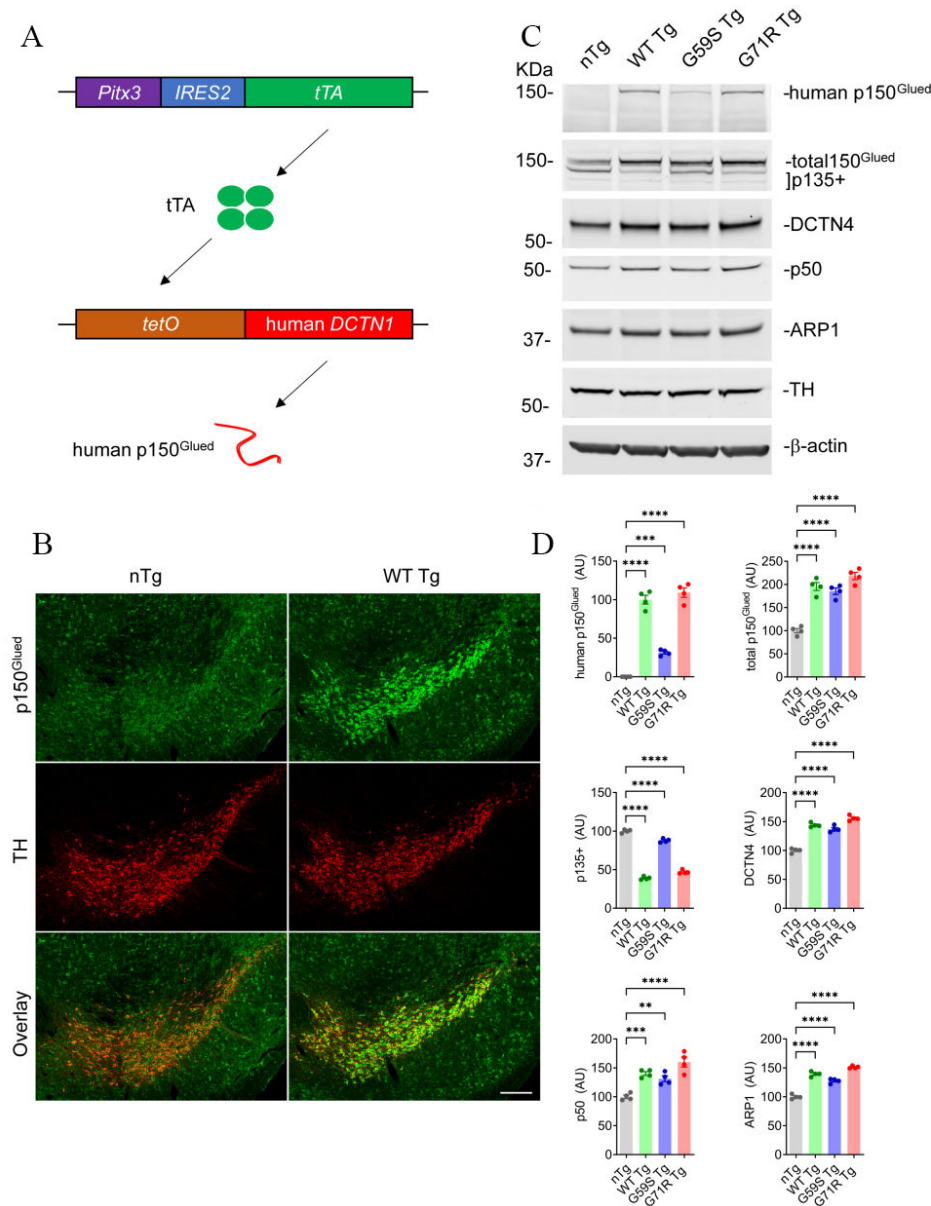


Figure 1. Selective expression of wild-type or mutant human p150^{Glued} in mouse midbrain DA neurons. (A) The schematic diagram depicts the generation of transgenic mice with target expression of human p150^{Glued} in midbrain DA neurons by crossing *Pitx3-IRES2-tTA* and *tetO-human DCTN1* mice. (B) Immunofluorescent images show the staining of p150^{Glued} (green) and TH (red) in the coronal midbrain sections of one-month-old *nTg* and *WT Tg* mice. DA neurons were visualized by TH staining. Scale bar: 200 μ m. (C, D) Western blots show the expression of human p150^{Glued}, total (human and mouse) p150^{Glued}, p135+, DCTN4, p50, and ARP1 in the midbrain homogenates of one-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice. β -actin and TH were used as the loading control. The bar graph estimates the protein level of dynactin subunits normalized with β -actin ($n = 4$ per genotype). Data are presented as mean \pm SEM. One-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$.

findings demonstrate that selective expression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons causes profound motor abnormalities in mice and suggest that both G59S and G71R mutant p150^{Glued} may impair the function and survival of midbrain DA neurons.

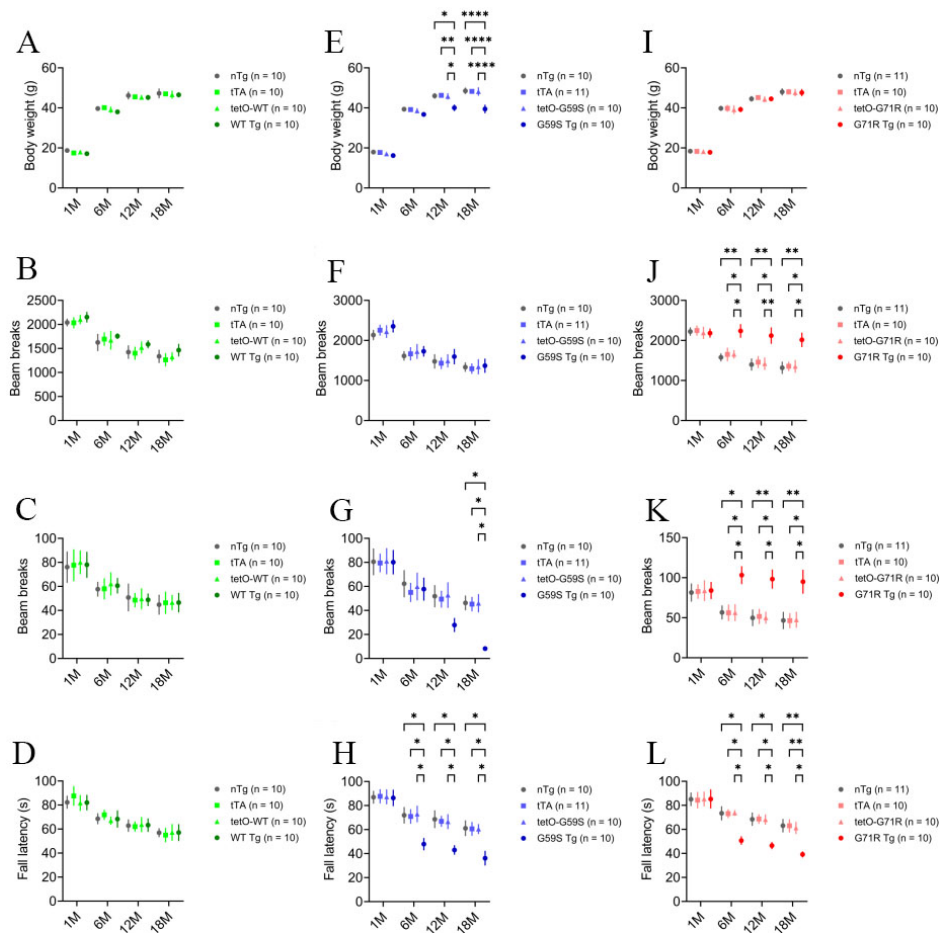


Figure 2. Profound motor abnormalities of *G59S Tg* and *G71R Tg* mice. (A-D) At 1, 6, 12, and 18 months of age, cohorts of male *nTg*, *tTA*, *tetO-WT*, and *WT Tg* mice ($n = 10$ per genotype) were repeatedly weighed (A); tested for ambulatory movement (B) and rearing (C) in the open-field test; and examined for the latency to fall in the rotarod test (D). (E-H) At 1, 6, 12, and 18 months of age, cohorts of male *nTg* ($n = 10$), *tTA* ($n = 11$), *tetO-G59S* ($n = 10$), and *G59S Tg* ($n = 10$) mice were repeatedly weighed (E); tested for ambulatory movement (F) and rearing (G) in the open-field test; and examined for the latency to fall in the rotarod test (H). (I-L) At 1, 6, 12, and 18 months of age, cohorts of male *nTg* ($n = 11$), *tTA* ($n = 10$), *tetO-G71R* ($n = 10$), and *G71R Tg* ($n = 10$) mice were repeatedly weighed (I); tested for ambulatory movement (J) and rearing (K) in the open-field test; and examined for the latency to fall in the rotarod test (L). Data are presented as mean \pm SEM. Two-way ANOVA with Tukey's multiple comparisons test was used for statistical analysis (at each time point, the mean of each genotype was compared with the mean of every other genotype). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.0001$.

***G59S Tg* and *G71R Tg* mice develop progressive degeneration of midbrain DA neurons**

Since neuronal loss in *substantia nigra* is a pathological hallmark of Perry syndrome^[9,10], the numbers of TH-positive midbrain DA neurons in the *substantia nigra pars compacta* (SNc) and ventral tegmental area (VTA) of *WT Tg*, *G59S Tg*, *G71R Tg* mice, and their control mice were counted using an unbiased stereological approach. *WT Tg* mice presented with no apparent loss of DA neurons in either the SNc or VTA at 1, 6, 12, and 24 months of age compared with control mice [Figure 3A-D]. *G59S Tg* mice displayed severe and early-onset degeneration of DA neurons in the midbrain, with approximately 13%, 28%, and 57% neuronal loss in the SNc at 6, 12, and 24 months of age, respectively, and 50% of neuronal loss in the VTA at 24 months of age [Figure 3A-D]. *G71R Tg* mice exhibited mild and late-onset degeneration of DA neurons in the midbrain, with approximately 25% neuronal loss in the SNc as well as the VTA at 24 months of age [Figure 3A-D]. Interestingly, besides neuronal loss, *G59S Tg* and *G71R Tg* mice showed remarked and progressive dystrophy of DA neurites in the *substantia nigra pars reticulata* (SNr) starting at two months of age [Figure 3A and E]. Therefore, expression of either G59S or G71R mutant p150^{Glued} in midbrain DA

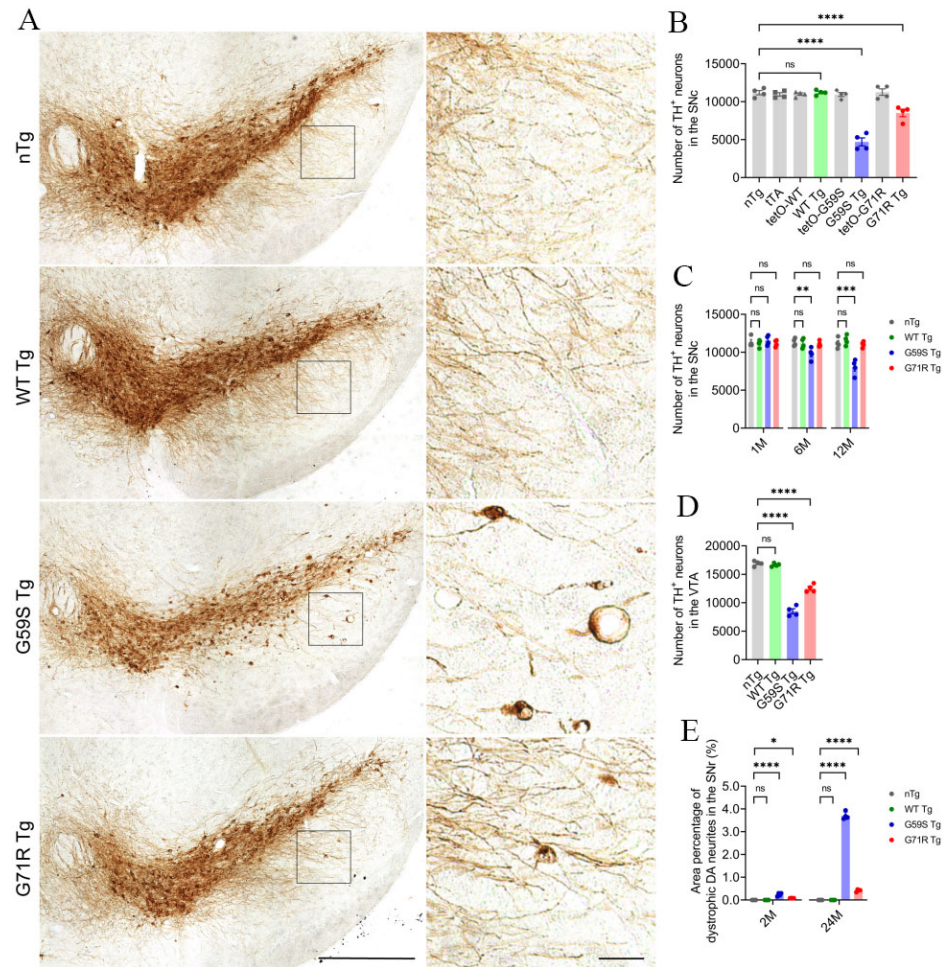


Figure 3. Progressive degeneration of DA neurons in the midbrain of *G59S Tg* and *G71R Tg* mice. (A) Immunohistochemical images show TH staining in the coronal midbrain sections of 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice. Scale bar: (left) 500 μ m (low-magnification images) and (right) 50 μ m (high-magnification images). Note the dystrophic TH-positive DA neurites in the SNr of *G59S Tg* and *G71R Tg* mice. (B) Bar graphs depict the unbiased stereological estimation of the number of TH-positive DA neurons in the SNc of 24-month-old *nTg*, *tTA*, *tetO-WT*, *WT Tg*, *tetO-G59S*, *G59S Tg*, *tetO-G71R*, and *G71R Tg* mice ($n = 4$ per genotype). Data are presented as mean \pm SEM. One-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). **** $P < 0.0001$. ns: Not significant. (C) Bar graphs depict the unbiased stereological estimation of the number of TH-positive DA neurons in the SNc of 1-, 6-, and 12-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice ($n = 4$ per genotype per time point). Data are presented as mean \pm SEM. At each time point, one-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). ** $P = 0.0099$, *** $P = 0.0001$. (D) Bar graphs depict the unbiased stereological estimation of the number of TH-positive DA neurons in the VTA of 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice ($n = 4$ per genotype). Data are presented as mean \pm SEM. One-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). **** $P < 0.0001$. (E) Bar graphs quantify the area percentage of dystrophic TH-positive DA neurites in the SNr of 2- and 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice (at each time point, $n = 4$ animals per genotype and 5 sections per animal). Dystrophic DA neurites were defined as neuritic varicosity $\geq 25 \mu\text{m}^2$. Data are presented as mean \pm SEM. At each time point, one-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). * $P = 0.0259$, **** $P < 0.0001$. TH: Tyrosine hydroxylase.

neurons leads to substantial and progressive neurodegeneration.

***G59S Tg* and *G71R Tg* mice display robust degeneration of DA axon terminals**

As axon dystrophy was reported in the striatum of patients with Perry syndrome^[30,31], the density and morphology of DA axon terminals in the dorsal striatum of *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice were

examined. Compared with age-matched nTg controls, 2-, 12-, and 24-months old *WT Tg* mice had no apparent alteration of TH-positive DA axon terminals [Figure 4A and B]. However, *G59S Tg* and *G71R Tg* mice displayed substantial loss of DA axon terminals, with approximately 15%, 33%, and 55% of axonal loss in 6-, 12- and 24-month-old *G59S Tg* mice, respectively, and 35% of axonal loss in 24-month-old *G71R Tg* mice [Figure 4A and B]. Furthermore, TH staining also revealed abnormally enlarged DA axon terminals in the striatum of *G59S Tg* and *G71R Tg* mice [Figure 4A, arrow]. These data demonstrate that expression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons causes robust degeneration of DA axon terminals and suggest dysfunctional DA transmission in *G59S Tg* and *G71R Tg* mice.

G59S Tg and G71R Tg mice show alteration of dopamine transmission

Considering the pathological findings, the steady-state level of dopamine in the dorsal striatum of *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice was assessed with HPLC. At 18 months of age, *WT Tg* and *G71R Tg* mice showed no apparent change in dopamine level compared with nTg controls, while *G59S Tg* mice displayed a significant reduction of dopamine content, with an approximately 53% of decrease [Figure 5A]. Regarding the primary dopamine metabolite 3,4-dihydroxyphenylacetic acid (DOPAC), neither its content nor the DOPAC/dopamine ratio in 18-month-old *WT Tg*, *G59S Tg*, and *G71R Tg* mice was significantly different with age-matched nTg controls [Figure 5A]. These data reveal dopamine deficiency in the striatum of *G59S Tg* mice, which might be the functional consequence of robust degeneration of midbrain DA neurons and the pathophysiological basis for the motor deficits observed in *G59S Tg* mice. By contrast, at 18 months of age, *G71R Tg* mice developed neither loss of midbrain DA neurons nor reduced dopamine content, further supporting the late-onset DA neuronal loss in *G71R Tg* mice.

Next, dopamine release kinetics in the dorsal striatum of *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice was evaluated using fast-scan cyclic voltammetry (FSCV). At six months of age, *WT Tg* mice showed comparable evoked dopamine release and dopamine uptake in response to single-pulse stimulation compared with nTg controls [Figure 5B and C]. However, *G59S Tg* mice displayed significantly decreased dopamine release and slower dopamine uptake, while *G71R Tg* mice exhibited substantially increased dopamine release and slower dopamine uptake [Figure 5B and C]. Similarly, 18-month-old *WT Tg* mice showed comparable evoked dopamine release and dopamine uptake in response to burst-pulse stimulation compared with age-matched nTg controls [Figure 5D and E]. However, *G59S Tg* mice displayed significantly decreased dopamine release and slower dopamine uptake, while *G71R Tg* mice exhibited substantially increased dopamine release and slower dopamine uptake [Figure 5D and E]. Taken together, *G59S Tg* mice showed early-onset defects in striatal dopamine transmission, consistent with the early-onset loss of DA neurons and axons. In contrast, *G71R Tg* mice showed early-onset dysfunction of dopamine release and uptake long before the substantial loss of DA neurons and axons occurred. Thus, these data indicate that expression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons profoundly compromises the dopamine transmission of the nigrostriatal DA pathway, contributing to the motor phenotypes developed in these transgenic mice.

G59S Tg and G71R Tg mice exhibit an age-dependent reduction of DA proteins

Tyrosine hydrolase (TH) is the rate-limiting enzyme in dopamine synthesis, and dopamine transporter (DAT) mediates dopamine uptake^[32,33]. Since TH and DAT play critical roles in the nigrostriatal DA pathway, the protein levels of TH and DAT in the striatal homogenates of 1- and 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice were examined. At one month of age, *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice had comparable levels of TH and DAT in the striatum [Figure 6A and B]. At 24 months of age, *WT Tg* mice showed no marked change in TH and DAT protein expression compared with nTg controls, while both *G59S Tg* and *G71R Tg* mice displayed substantial loss of TH and DAT [Figure 6A and B]. Interestingly, the percentages of DAT loss in the striatum of 24-month-old *G59S Tg* and *G71R Tg* mice (approximately

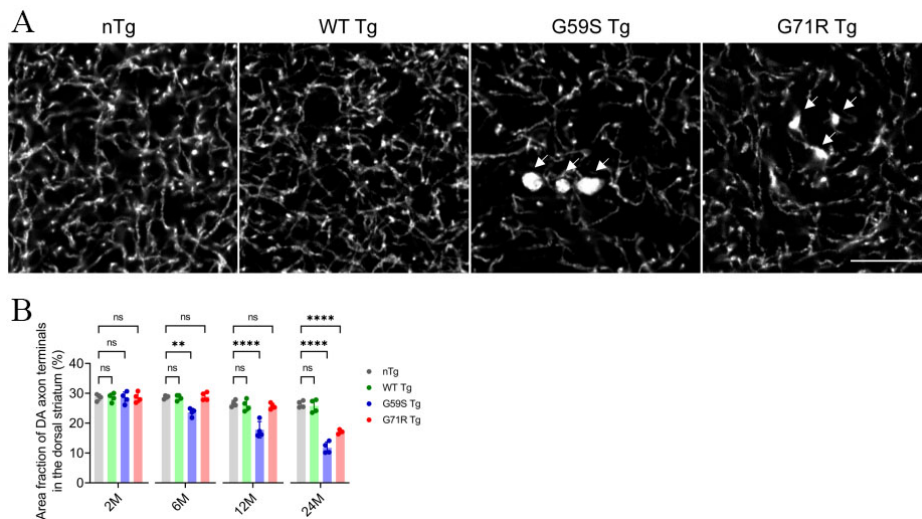


Figure 4. Robust degeneration of DA axon terminals in the striatum of *G59S Tg* and *G71R Tg* mice. (A) Representative images show TH staining in the dorsal striatum coronal sections of 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice. Scale bar: 10 μ m. Arrows point to swellings of TH-positive DA axon terminals in the striatum of *G59S Tg* and *G71R Tg* mice. (B) Bar graph estimates the area fraction of TH-positive DA axon terminals in the dorsal striatum of 2-, 6-, 12-, and 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice (at each time point, $n = 4$ animals per genotype and 10 sections per animal). Data are presented as mean \pm SEM. At each time point, one-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). ** $P = 0.0018$, **** $P < 0.0001$. ns: Not significant. TH: Tyrosine hydroxylase.

70% and 42%, respectively) were more severe than the TH loss (about 53% and 29%, respectively) [Figure 6A and B]. These data suggest expression of either *G59S* or *G71R* mutant $p150^{\text{Glued}}$ in midbrain DA neurons may preferentially downregulate the expression and function of DAT protein at DA axon terminals.

***G59S Tg* mice display age-dependent accumulation of $p150^{\text{Glued}}$ aggregates in midbrain dopaminergic neurons**

As both *G59S* and *G71R* mutant $p150^{\text{Glued}}$ were found aggregation-prone *in vitro*^[17-19], the subcellular distribution pattern of $p150^{\text{Glued}}$ in the midbrain DA neurons of 2-, 6-, 12-, and 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice was examined. TH and $p150^{\text{Glued}}$ co-staining revealed diffuse cytoplasmic localization of $p150^{\text{Glued}}$ without apparent aggregate formation in the soma and neurites of midbrain DA neurons from both young and aged *nTg*, *WT Tg*, and *G71R Tg* mice [Figure 7A and B]. By contrast, *G59S Tg* mice presented with $p150^{\text{Glued}}$ -positive cytoplasmic aggregates, mainly in the soma and in the neurites of midbrain DA neurons [Figure 7A]. Furthermore, the percentage of DA neurons with $p150^{\text{Glued}}$ aggregate in *G59S Tg* mice increased in an age-dependent manner, approximately 0%, 10.54%, 19.66%, and 82.17% at 2, 6, 12, and 24 months of age, respectively [Figure 7B]. Immunostaining further revealed the co-aggregation of dynactin subunit p50 and ARP1, as well as dynein subunit DLIC, with $p150^{\text{Glued}}$ aggregates inside the dystrophic DA neurites in the SNr of six-month-old *G59S Tg* mice [Figure 8A-C]. In addition, co-staining of coronal midbrain sections showed co-localization of lysosomes with $p150^{\text{Glued}}$ aggregates and a substantial increase of ubiquitin immunoreactivities within the dystrophic DA neurites in the SNr of six-month-old *G59S Tg* mice, compared with *nTg* controls [Figure 9A and B]. Thus, *G59S* mutation rather than *G71R* mutation induces mutant $p150^{\text{Glued}}$ to form cytoplasmic aggregates in midbrain DA neurons *in vivo*. Moreover, the aggregates trap dynein/dynactin, co-localize with lysosomes, and upregulate ubiquitination, implying detrimental effects of *G59S* mutant $p150^{\text{Glued}}$ on dynein/dynactin function and the degradative pathway in the midbrain DA neurons. These findings may explain why *G59S Tg* mice developed severe and early-onset degeneration of midbrain DA neurons, while *G71R Tg* mice showed milder and more slowly progressive neurodegeneration.

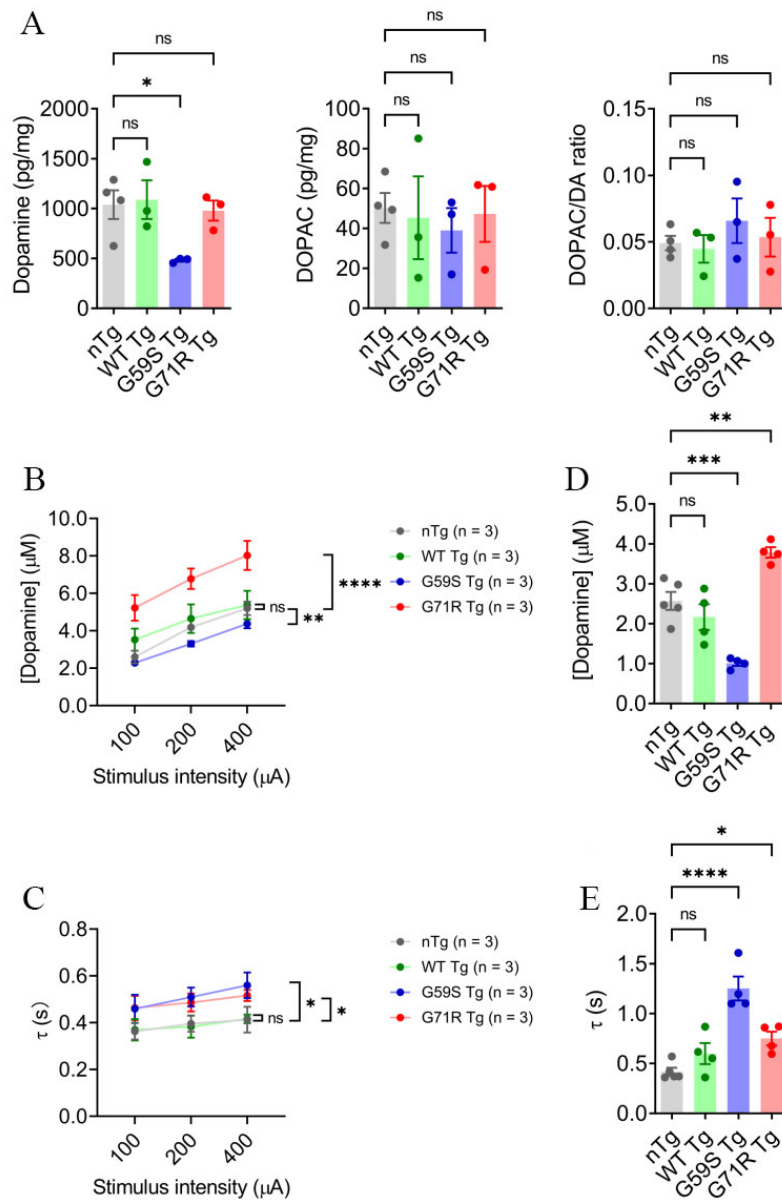


Figure 5. Alteration of dopamine content and release in the striatum of *G59S Tg* and *G71R Tg* mice. (A) HPLC measures the content of dopamine and DOPAC in the dorsal striatum of 18-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice ($n = 3$ or 4 per genotype). Data are presented as mean \pm SEM. One-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). $^*P = 0.0397$. ns: Not significant. (B, C) FSCV quantifies the peak evoked dopamine release (B) and time constant of slope decay (C) following single-pulse electrical stimulation of different stimulus intensity in the dorsal striatum of six-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice ($n = 3$ animals per genotype and 3 sections per animal). Data are presented as mean \pm SEM. Two-way ANOVA genotype factor: *nTg* vs. *G59S Tg*, $F_{(1,12)} = 25.11$, $^*P = 0.0044$; *nTg* vs. *G71R Tg*, $F_{(1,12)} = 40.19$, $^{****}P < 0.0001$ (B). Two-way ANOVA genotype factor: *nTg* vs. *G59S Tg*, $F_{(1,12)} = 9.285$, $^*P = 0.0101$; *nTg* vs. *G71R Tg*, $F_{(1,12)} = 8.491$, $^*P = 0.0130$ (C). (D, E) FSCV quantifies the peak evoked dopamine release (D) and time constant of slope decay (E) in response to burst electrical stimulation (200 μ A, 50 Hz, 5 pulses) in the dorsal striatum of 18-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice ($n = 4$ or 5 animals per genotype and 3 sections per animal). Data are presented as mean \pm SEM. One-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). *nTg* vs. *G59S Tg*, $^{***}P = 0.0004$; *nTg* vs. *G71R Tg*, $^{**}P = 0.0034$ (D). *nTg* vs. *G59S Tg*, $^{****}P < 0.0001$; *nTg* vs. *G71R Tg*, $^*P = 0.0357$ (E).

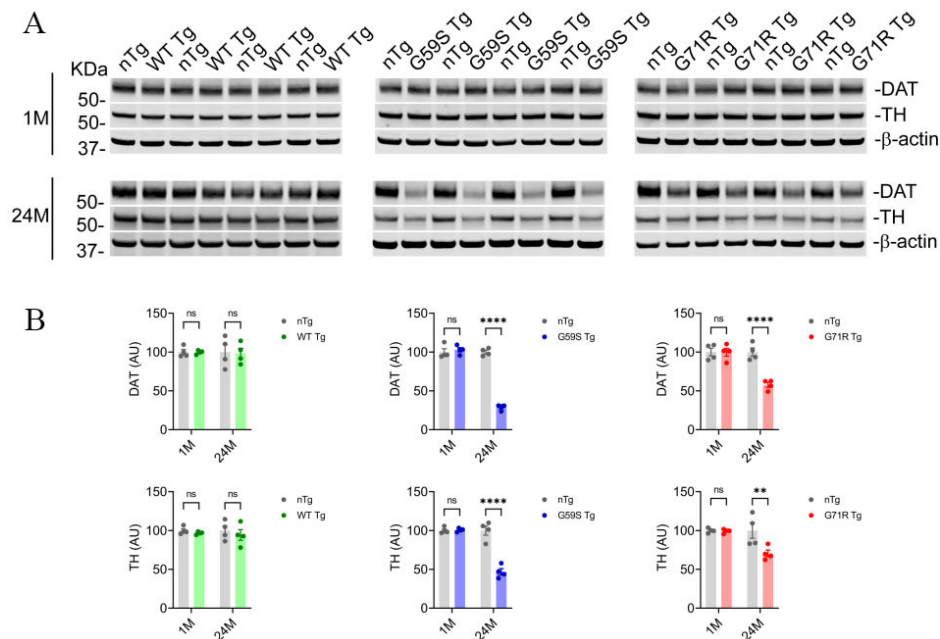


Figure 6. Reduction of dopaminergic proteins in the striatum of *G59S Tg* and *G71R Tg* mice. (A, B) Western blots show the expression of DAT and TH in the striatum homogenates of 1- and 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice. Actin was used as the loading control. The bar graph estimates the levels of DAT and TH expression normalized against β -actin ($n = 4$ per genotype). Data are presented as mean \pm SEM. Two-way ANOVA with Sidak's multiple comparisons test was used for statistical analysis. ** $P = 0.0054$, **** $P < 0.0001$. ns: Not significant.

DISCUSSION

In this study, transgenic mouse lines with selective overexpression of wild-type human p150^{Glued}, motor neuron disease-related G59S mutant p150^{Glued}, or Perry syndrome-related G71R mutant p150^{Glued} in midbrain DA neurons were generated and characterized. The *WT Tg* mice did not develop any overt motor phenotypes or midbrain DA neurodegeneration compared with age-matched *nTg* mice. The *G59S Tg* mice displayed weight loss, motor deficits, early-onset decreased dopamine transmission, and early-onset progressive loss of DA neurons and axons. The *G71R Tg* mice exhibited hyperactivities, impaired motor coordination, early-onset increased dopamine release and slower dopamine uptake, and late-onset loss of DA neurons and axons. Furthermore, both *G59S Tg* and *G71R Tg* mice showed an age-dependent reduction of DA proteins in the striatum, with a more severe loss of DAT than TH. Finally, G59S mutant p150^{Glued} rather than G71R mutant p150^{Glued} was found to form aggregates in mouse midbrain DA neurons in an age-dependent manner, and the aggregates trapped dynein/dynactin, co-localized with lysosomes, and upregulated ubiquitination.

Since the initial identification of motor neuron disease- and Perry syndrome-causative mutations in p150^{Glued}, several mouse strains of mutant p150^{Glued} have been generated to study their effects on selective degeneration of specific types of neurons. Motor neuron disease-like motor phenotypes and degeneration of spinal motor neurons have been reported in G59S p150^{Glued} transgenic and knock-in mouse models^[20-22]. Decreased exploratory activity and impaired motor coordination were observed in the transgenic mice overexpressing G71A p150^{Glued} under the control of the *Thy1* promotor^[23]. However, either the expression of G71A p150^{Glued} in midbrain DA neurons or neuronal loss in the brain tissue was detected in the transgenic mice^[23]. Recently, G71A p150^{Glued} knock-in mouse models were created^[24]. Increased immobility time in the tail suspension test, impaired motor coordination in the beam-walking test, and decreased TH staining

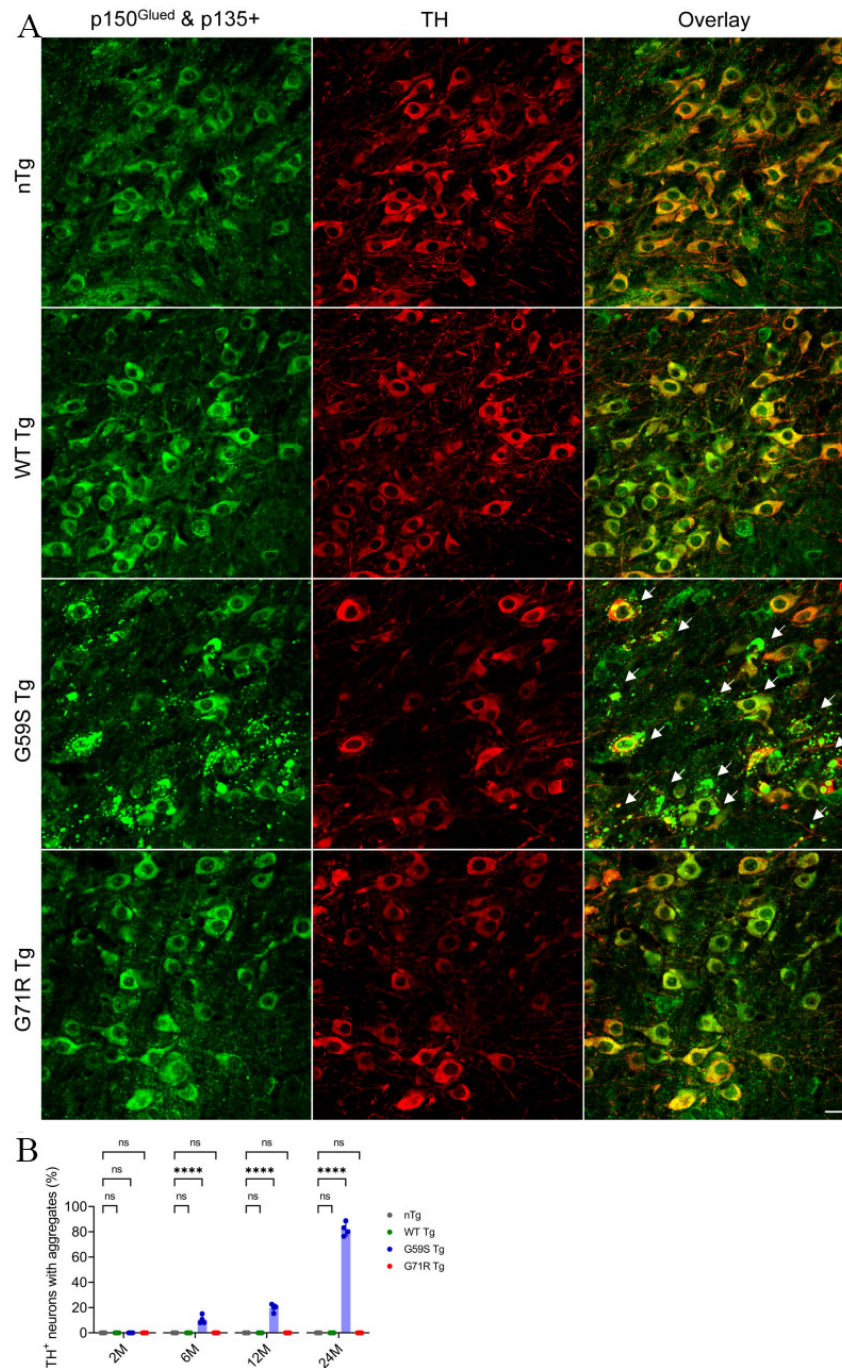


Figure 7. Formation of p150^{Glued}-positive protein aggregates in midbrain DA neurons of G59S *Tg* mice. (A) Immunofluorescent images show the staining of p150^{Glued} and p135+ (green) and TH (red) in midbrain coronal sections of 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice. Scale bar: 20 μ m. Arrows point to proteins aggregates of p150^{Glued} and p135+ in the soma and neurites of DA neurons from *G59S Tg* mice. (B) Bar graph estimates the percentage of TH-positive neurons with proteins aggregates of p150^{Glued} and p135+ in the midbrain of 2-, 6-, 12-, and 24-month-old *nTg*, *WT Tg*, *G59S Tg*, and *G71R Tg* mice (at each time point, $n = 4$ animals per genotype and 5 sections per animal). Aggregates were defined as puncta $\geq 1 \mu\text{m}^2$. Data are presented as mean \pm SEM. At each time point, one-way ANOVA with Dunnett's multiple comparisons test was used for statistical analysis (the mean of each genotype was compared with the mean of *nTg*). **** $P < 0.0001$. ns: Not significant. TH: Tyrosine hydroxylase.

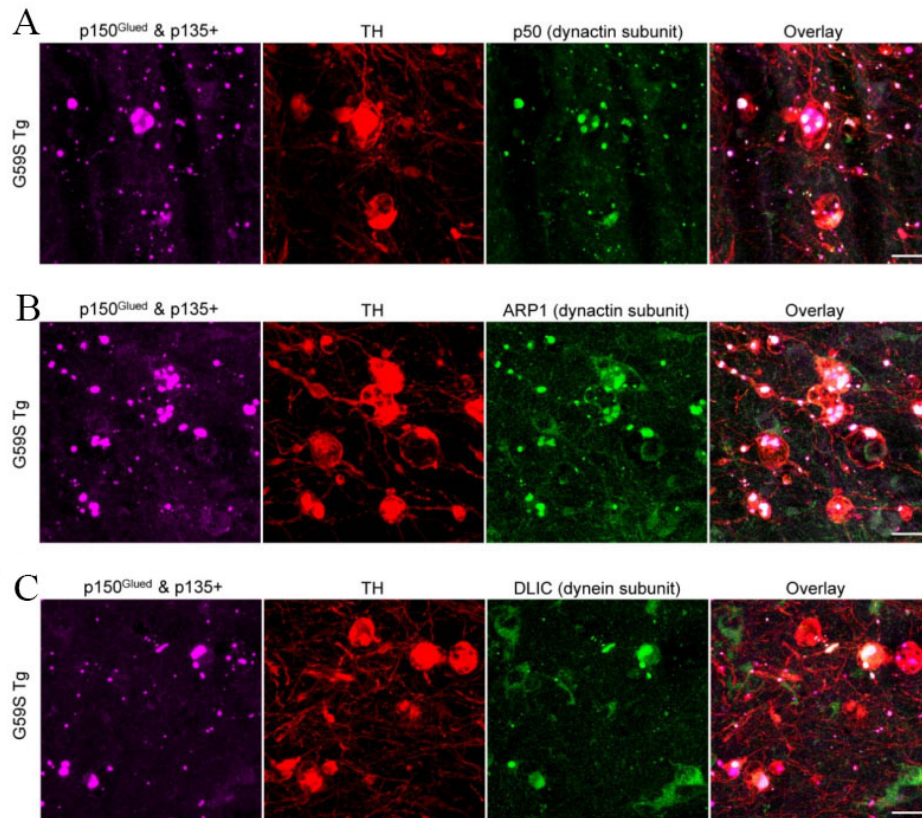


Figure 8. Co-aggregation of dynactin and dynein subunits with G59S mutation-induced p150^{Glued} aggregates in the dystrophic DA neurites of G59S Tg mice. (A) Immunofluorescent images show the staining of p150^{Glued} and p135+ (purple), TH (red), and dynactin subunit p50 (green) in the SNr of six-month-old G59S Tg mice. Scale bar: 20 μm. (B) Immunofluorescent images show the staining of p150^{Glued} and p135+ (purple), TH (red), and dynactin subunit ARP1 (green) in the SNr of six-month-old G59S Tg mice. Scale bar: 20 μm. (C) Immunofluorescent images show the staining of p150^{Glued} and p135+ (purple), TH (red), and dynein subunit DLIC (green) in the SNr of six-month-old G59S Tg mice. Scale bar: 20 μm.

intensity in the substantia nigra neurons were found in the heterozygous G71A p150^{Glued} knock-in mice^[24], recapitulating some of the clinical features of Perry syndrome. However, the pathogenic effects of mutant p150^{Glued} on Perry syndrome-related degeneration of the nigrostriatal DA pathway remain to be determined. New lines of transgenic mice (*WT Tg*, *G59S Tg*, and *G71R Tg*) with selective overexpression of wild-type, G59S mutant, or G71R mutant human p150^{Glued} in midbrain DA neurons were generated in the current study, allowing us to systematically evaluate the impact of mutant p150^{Glued} on survival and function of midbrain DA neurons. *WT Tg* mice showed neither motor phenotypes nor pathological, functional, or biochemical abnormalities of the nigrostriatal DA pathway. *G59S Tg* mice displayed substantial weight loss, dramatic reduction of rearing movement in the open field test, and significant impairment of motor coordination in the rotarod test. The pathophysiological basis for the motor phenotypes in *G59S Tg* mice is the early-onset and severe loss of DA neurons and axon terminals and the resultant deficiency of dopamine content and transmission. *G71R Tg* mice exhibited significant hyperactivities in ambulatory and rearing movements and substantially impaired rotarod performance through 6-18 months of age. In contrast to the severe and early-onset degeneration of nigrostriatal DA pathway in *G59S Tg* mice, the neurodegeneration in *G71R* mice was much milder and more slowly progressive, with no significant loss of DA neurons and axon terminals occurring before 18 months of age. Despite no early-onset DA neuron degeneration, *G71R Tg* mice showed an early-onset increase of dopamine release and slower dopamine uptake in the striatum, which may elevate the extracellular dopamine level and contribute to hyper-dopamine motor phenotypes.

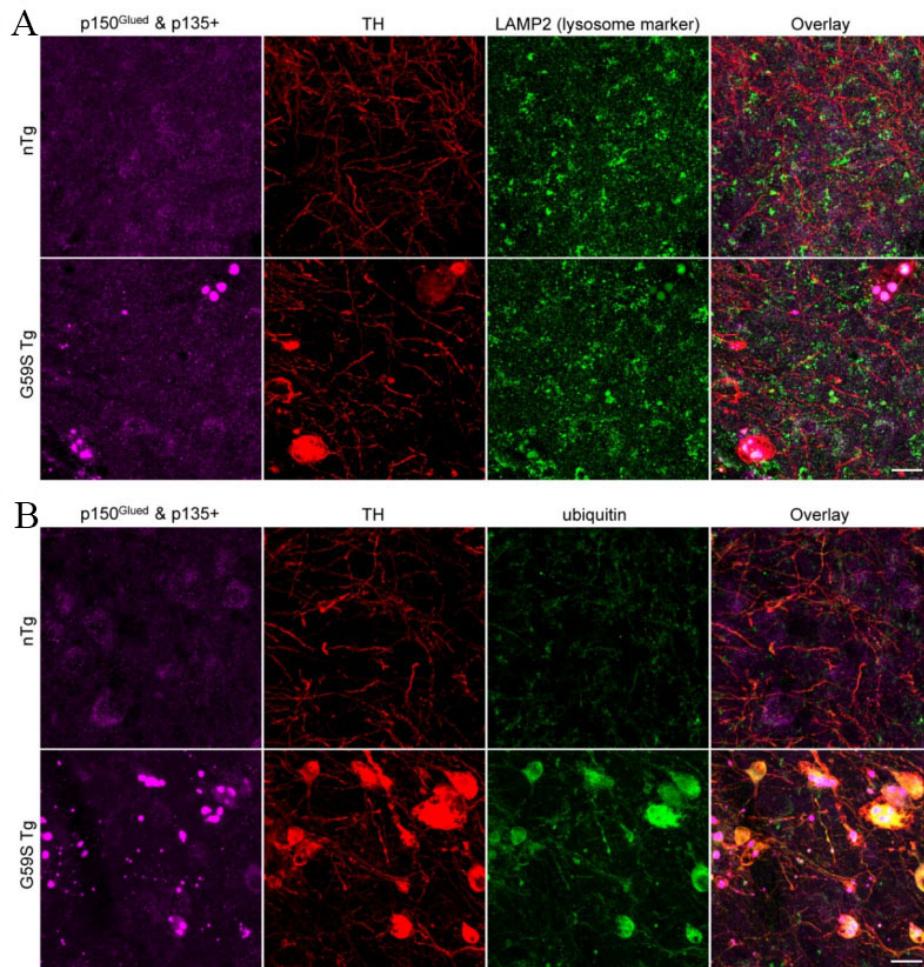


Figure 9. Co-localization of lysosomes with G59S mutation-induced p150^{Glued} aggregates and upregulated ubiquitination in the dystrophic DA neurites of G59S Tg mice. (A) Immunofluorescent images show the staining of p150^{Glued} and p135+ (purple), TH (red), and LAMP2 (green) in the SNr of six-month-old nTg and G59S Tg mice. Scale bar: 20 μm. (B) Immunofluorescent images show the staining of p150^{Glued} and p135+ (purple), TH (red), and ubiquitin (green) in the SNr of six-month-old nTg and G59S Tg mice. Scale bar: 20 μm.

Thus, our data demonstrate that both G59S and G71R mutant p150^{Glued} are pathogenic to the nigrostriatal DA pathway *in vivo*, but differentially impair the survival and functions of midbrain DA neurons. The finding on G59S mutation is unexpected since the mutation was particularly linked to a slowly progressive autosomal dominant form of lower motor neuron disease, the distal hereditary motor neuropathy 7B (HMN7B)^[6]. Our previous study showed that both homozygous *Dctn1* knock-out mice and homozygous *DCTN1* G59S knock-in mice died during early embryogenesis, indicating the loss-of-function nature of G59S mutation^[20]. Furthermore, we found that the G59S mutation disrupts the stability of p150^{Glued} protein, so we could not detect any expression and accumulation of p150^{Glued} protein in the tissues of homozygous *DCTN1* G59S knock-in embryos^[20]. By contrast, in the G59S Tg mice, an artificial overexpression of G59S mutation in DA neurons led to an extensive accumulation of misfolded p150^{Glued} proteins, which may overwhelm the protein degradation system in the DA neurons and eventually induce cell death. Therefore, the overexpression data in either cell or animal models must be interpreted with caution.

Microtubule-binding domains in p150^{Glued} are required for the initiation of retrograde axonal transport, stability of axonal microtubules, and integrity of axon terminals^[13-16]. Accordingly, degeneration of ventral root axons and disruption of neuromuscular junctions have been observed in G59S mutant p150^{Glued} transgenic mice^[21,22]. In line with these earlier findings, the current study revealed that expression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons resulted in substantial dysfunction and degeneration of DA axon terminals in the striatum. Additionally, our study showed expression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons led to the dystrophy of DA neurites in the SNr, indicating an essential role of p150^{Glued} in maintaining the structural integrity of both axons and dendrites of DA neurons. Future studies will be needed to determine the underlying mechanism of how mutant p150^{Glued} triggers the structural and functional changes of distal and proximal DA neurites.

Several mechanisms have been hypothesized to address the selective degeneration of motor neurons caused by G59S mutant p150^{Glued}, including impaired retrograde axonal transport of trophic signaling, defects of vesicular trafficking in the ER-Golgi secretory pathway, aberrant autophagosome-lysosome degradative pathway, and cytotoxicity of mutant p150^{Glued} aggregates^[17,21,22]. However, due to the lack of appropriate animal models, mechanisms for the selective degeneration of DA neurons in Perry syndrome have been poorly understood. Until recently, *DCTN1* G71A knock-in mice were generated, and homozygous knock-in mice died prenatally, suggesting a potential loss-of-function nature of the mutation^[24]. In the heterozygous *DCTN1* G71A knock-in mice, a reduction of TH immunoreactivity in DA neurons was reported^[24]. However, the effect of mutant p150^{Glued} on dopamine transmission and the related regulating proteins remains to be determined. Our HPLC and FSCV studies revealed that overexpression of human G71R p150^{Glued} in midbrain DA neurons led to substantial dopamine deficiency, significantly decreased dopamine release, and slower dopamine uptake, reflecting the severe and early-onset loss of DA neurons and axons. In contrast, overexpression of human G71R p150^{Glued} in midbrain DA neurons resulted in early-onset substantially increased dopamine release and slower dopamine uptake long before the loss of DA axons and the deficiency of dopamine content in the striatum occurred, indicating the early-onset dysfunction of DAT. Correlatively, Western blot analysis demonstrated that overexpression of either G59S or G71R mutant p150^{Glued} in midbrain DA neurons preferentially downregulated the expression of DAT at DA axon terminals. Further investigation will be required to unmask the specific cellular mechanisms leading to the DAT dysfunction and the selective cell death of DA neurons in Perry syndrome.

G59S mutation, as well as all the Perry syndrome-causative mutations, occurs within or close to the CAP-Gly domain of p150^{Glued}, causing loss of function of the MTBDs^[6-8]. G59S mutant p150^{Glued} is prone to misfold and form cytoplasmic aggregates leading to enhanced cell death both *in vitro* and *in vivo*, suggesting both loss-of-function and toxic-gain-of-function mechanisms are involved in G59S mutation-induced neurodegeneration^[17,21]. Perry syndrome-related G71A mutant p150^{Glued} has been reported to aggregate less than G59S mutant p150^{Glued} when overexpressed in HeLa cells^[18]. Moreover, no evident p150^{Glued} aggregates are observed in G71A transgenic mice^[23]. In line with these earlier findings, our study revealed that neither wild-type nor G71R mutant p150^{Glued} formed aggregates *in vivo* when overexpressed in midbrain DA neurons. In contrast, G59S mutant p150^{Glued} formed aggregates in an age-dependent way when overexpressed in mouse midbrain DA neurons. Moreover, co-aggregation of dynactin and dynein subunits with G59S mutation-induced p150^{Glued} aggregates was observed inside the dystrophic DA neurites of G59S *Tg* mice. The data suggest the sequestration of active motors in aggregates; the restricted accessibility of adaptors, cargoes, and regulatory proteins to dynein/dynactin; and thus the impairment of normal functions of dynein/dynactin in the midbrain DA neurons. In addition, co-localization of lysosomes with G59S mutation-induced p150^{Glued} aggregates and upregulated ubiquitination were found within the dystrophic DA neurites of G59S *Tg* mice, implying the disruption of the degradative pathway in the midbrain DA neurons.

These observations may explain why the degeneration of the nigrostriatal DA pathway in *G59S Tg* mice is more early-onset and severe than in *G71R Tg* mice in our study. Future studies will explore whether overexpression of exogenous *G59S* or *G71R* mutant p150^{Glued} compromises the normal functions of endogenous dynactin p150^{Glued} in the midbrain DA neurons to test if there is a dominant-negative mechanism for *DCTN1* mutation-induced degeneration of nigrostriatal DA pathway.

While the present study focused on the role of mutant p150^{Glued} in Perry syndrome-related parkinsonism and the underlying degeneration of midbrain DA neurons, the pathophysiological basis and pathogenic mechanisms of other Perry syndrome-related symptoms remain to be delineated in detail, including apathy/depression and central hypoventilation^[9,10]. Except for the loss of midbrain DA neurons, Perry syndrome patients also display marked degeneration of norepinephrine neurons in *locus coeruleus* and serotonergic neurons in the medullary raphe and ventrolateral medulla^[34,35]. Future studies will be needed to crossbreed our tetO-human *DCTN1* with suitable tTA lines and determine the pathogenic significance of mutant p150^{Glued} on norepinephrine neurons and serotonergic neurons.

In summary, our findings show selective expression of either *G59S* or *G71R* mutant p150^{Glued} in mouse midbrain DA neurons leads to parkinsonism-like motor abnormalities and degeneration of the nigrostriatal DA pathway, recapitulating some key clinical and pathological features of Perry syndrome. Moreover, *G59S* and *G71R* mutant p150^{Glued} exhibit differential pathogenic impact on the survival and function of midbrain DA neurons *in vivo*. The p150^{Glued} transgenic mouse models provide valuable tools for mechanistic insights and therapeutic targets of DA neuron degeneration in Perry syndrome.

DECLARATIONS

Acknowledgments

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Authors' contributions

Designed the study, analyzed the data, and wrote the manuscript: Cai H, Yu J, Sgobio C
Performed the experiments: Yu J, Sgobio C, Yang X, Peng Y, Chen X, Sun L, Shim H
Read and approved the final version of the manuscript: all authors

Availability of data and materials

Data and materials of the current study will be available upon reasonable request.

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Conflicts of interest

All authors declared that there are no conflicts of interest.

Ethical approval and consent to participate

All animal procedures conformed to the NIH (13-040) guide for the ethical care and use of laboratory animals. Animal protocols were approved by the Institutional Animal Care and Use Committee of National

Institute on Aging and Beijing Geriatric Hospital.

Consent for publication

Not applicable.

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Original Article

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A sequential deposition of amyloid beta oligomers, plaques and phosphorylated tau occurs throughout life in the canine retina

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Abstract

Aims: Cerebral amyloid burdens may be found in otherwise cognitively intact adults, often not showing worsening deficits with passing years. Alzheimer's transgenic rodents have been widely used to investigate this phenomenon, but a spontaneous disorder in other animals, such as dogs that cohabit with humans and thus may have some shared environmental risks, may contribute and offer opportunities not possible in the smaller laboratory animals. In animals, the spontaneous disorder most comparable to Alzheimer's disease (AD) affects mature to aged dogs and is designated canine cognitive dysfunction. Motivated by AD, many studies have revealed that amyloid progressively accumulates in the canine central nervous system, including the retina. Here, we investigated whether deposits of amyloid and/or tau can be found in the canine retina of neurologically normal animals from the first year of life to the elderly. Suppose canine ocular amyloid and tau are present from early life. In that case, that raises the question of whether similar patterns of accumulation occur in man, whether as part of aging, AD, or other.



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Methods: This study used eye tissues from 30 dogs with a variety of ophthalmic or other orbital disorders, of which 7/30 were 1-2 years old. Tissues were subdivided into dogs of three different age groups: young (1-5 years old), middle (6-10 years old), and old (≥ 11 years old).

Results: Following immunostaining of tissue sections with nanobodies against retinal $A\beta_{1-40}$ and $A\beta_{1-42}$ oligomers, and antibodies against $A\beta$ plaques ($A\beta p$) and hyperphosphorylated Tau (p-Tau), our investigations revealed that accumulation of $A\beta_{1-40}$ and $A\beta_{1-42}$ oligomers were widespread in the retina in all age groups. In contrast, $A\beta p$ were detected in the middle and old age groups but not in the young age group. Furthermore, p-Tau staining was observed in four old dogs only, while other dogs were p-Tau free. Interestingly, both $A\beta o$ and $A\beta p$ co-localized in the middle and old age groups of dogs. Moreover, diffuse granular p-Tau co-localized with intracellular $A\beta o$ in the old age group. Finally, we also observed co-localization of $A\beta o$ and $A\beta p$ in the retinal vasculature which might be similar to cerebral amyloid angiopathy associated with AD.

Conclusion: As far as we know, the presence of amyloid and tau in the canine retina is hitherto unreported. If similar, early-in-life subclinical retinal deposits occur in a human cohort perhaps identified by AD genetic risk factors, following this group may offer the prospect of preclinical therapeutic intervention in imminent dementia, a strategy recognized as likely necessary to impact this burgeoning disorder.

Keywords: Alzheimer's disease, canine cognitive dysfunction, retina, early diagnosis, $A\beta o$, $A\beta p$, p-Tau

INTRODUCTION

The principal neuropathological lesions in Alzheimer's disease (AD) brains include extracellular neuritic and/or diffuse plaques containing amyloid-beta ($A\beta$), intracellular tau protein (p-Tau) in the form of neurofibrillary tangles (NFTs) in addition to cerebral amyloid angiopathy (CAA), ubiquitin, severe synaptic loss, neuronal death, and brain atrophy^[1-4]. While the deposition of $A\beta$ in the human brain has traditionally been accepted as a major hallmark of AD^[5], its accumulation is also observed in about 20% of cognitively unimpaired, aged individuals^[6,7]. For decades, the association of $A\beta$ with AD has been demonstrated, but the significance and impact of $A\beta$ accumulation in healthy individuals are both poorly understood^[8] and a source of confusion. For example, previous studies did not establish a clear correlation with memory loss in the aging brain^[9-11]. Previous reports documented the progressive accumulation of both $A\beta$ plaques ($A\beta p$) in the brain^[12,13] and $A\beta$ oligomers ($A\beta o$) in the brain and periphery of cognitively unimpaired individuals^[14,15]. A study by Lesne *et al.* measured the levels of three $A\beta o$ "species", including $A\beta$ trimers, $A\beta^{*56}$ and $A\beta$ dimers, in brain tissues from 75 cognitively unimpaired individuals, including young children and adolescents^[14]. The authors showed that $A\beta$ trimers were present in the central nervous system (CNS) of children and adolescents, and their levels increased progressively with age, suggesting that this particular $A\beta o$ could be used to track the potential progression into AD from a very young age. Another study investigated the relationship between amyloid levels and memory performance^[15]. This study, which included 147 participants divided into three groups of adults 30-49, 50-69, and 70-89 years of age, established a clear relationship between episodic memory performance and amyloid accumulation in the youngest group^[15]. A longitudinal study by Hanseeuw *et al.* demonstrated a correlation between $A\beta$ /p-Tau and declining cognition in 60 clinically normal individuals aged between 65-85 years^[16]. The authors concluded that there was a positive correlation between $A\beta$ /p-Tau positron emission tomography (PET) and cognition, where participants with high $A\beta$ and tau were at higher risk of developing mild cognitive impairment (MCI)^[16]. These studies highlighted the importance of investigating and characterizing normal aging in cognitively unimpaired younger individuals to identify those at risk of progressive increase of AD-associated neuropathology changes and cognitive impairment and establish a time frame for early diagnostic and therapeutic intervention. However, such studies are difficult to implement in human subjects and will take decades to deliver any meaningful outcome^[8,17].

Currently available transgenic animal models of AD do not replicate the subtle clinical and pathological features of the disease, as demonstrated by their lack of reproducible therapeutic outcomes^[18,19]. As to the dog's age, there is an anatomically defined accumulation of A β peptides in the CNS, while less is known about cerebral p-Tau deposition. Some animals will develop a progressive cognitive decline with loss of learned behaviors and memory, a syndrome named canine cognitive dysfunction (CCD)^[20-24], with CNS changes similar to the neuropathological hallmarks characteristic of AD^[25-28]. In those destined to develop AD, amyloid precursor protein (APP) is sequentially cleaved by β and γ secretase leading to the production of A β peptide fragments (36-43 amino acids), which then aggregate and deposit as plaques^[4,5,29]. The canine APP and A β peptides are 98% and 100% identical to their human counterparts^[24]. In older dogs and others affected with CCD, A β deposits as diffuse plaques, while the dense core of humans' mature plaques is very rare or absent^[20,21,27,30].

There are two major isoforms of A β , A β_{1-40} (~80%-90%) and A β_{1-42} (~5%-10%) and three major assemblies^[31-34] recognized in man and animals/dogs, including monomeric A β , A β o containing 12-24 monomers which become elongated to form protofibrils and finally insoluble fibrils^[35]. Among these three stages, A β o is thought to be the most toxic to neurons and responsible for the structural and functional pathologic changes associated with AD^[36-39]. Likewise, a previous study reported that cognitive decline occurs before the accumulation of A β p in CCD, supporting the premise that earlier assembly states of A β may be the toxic species^[23]. In addition, CAA, ubiquitin, and severe synaptic loss have also been reported in CCD^[20,40-43]. Another cardinal pathological hallmark of AD is the accumulation of p-Tau^[3,4]. Several phosphorylation sites were identified on tau in aged dogs including Thr181^[42], Ser422^[43], Ser202/Thr205^[40,43], Ser396^[30,40,43], Ser189, and Ser207^[44]; however, demonstrating the presence of NFTs has been infrequent compared to the consistent demonstration of amyloid oligomers and diffuse plaques. A study by Schmidt and colleagues using anti-pT205, AT8, AT100, PHF-1, and anti-pT422 antibodies seeking the presence of tau pathology in 24 dogs aged between two and nineteen years, showed that three 13-15-year-old dogs displayed p-Tau and only one 15-year-old Pekingese dog displayed NFT-like appearance^[43].

In AD, visual disturbances are one of the early complaints and include loss of color vision, impairment of peripheral vision and object recognition, contrast sensitivity, and decreased visual memory and perception^[45-47]. A recent longitudinal study of 1349 older adults showed that poor visual acuity paralleled the development of dementia^[48], suggesting the possibility that ocular disturbances can be used as an early predictor of dementia risk in the older population^[48]. Moreover, post-mortem studies of AD and animal models of AD demonstrated a strong association between retinal accumulation of A β o^[49,50]. A β plaques^[51,52], p-Tau^[53,54] and brain depositions and cognitive decline, where retinal A β o was shown to deposit earlier than the brain and before deficits in cognition. Although, to our knowledge, age-dependent retinal deposition of A β o and/or A β p has not been investigated in AD and cognitively unimpaired individuals, including children and adolescents, our recent studies in AD mice models confirmed the conversion of cerebral and retinal A β o to A β p in an age-dependent manner, where retinal A β o was detected as early as 3-month old APP/PS1 mice, before brain pathology and cognitive decline were observed^[49,55].

In this study, we investigated the retinal accumulation of A β o, A β p, and p-Tau in three age groups of genetically diverse and neurologically intact populations of dogs. Here, following immunohistochemistry and immunofluorescence analysis, we confirmed the presence of retinal A β_{40} and A β_{42} oligomers, A β p, and p-Tau. Retinal A β_{40} and A β_{42} oligomers deposition was conspicuous and widespread and observed in all age groups, including the young 1-5-year-old neurologically intact group. Moreover, retinal co-localization of

A β_{40} /A β_{42} oligomers and A β p was observed in few middle-aged dogs and most dogs in the old neurologically intact age group, while retinal co-localization of A β_{40} /A β_{42} oligomers and p-Tau was only seen in the old neurologically intact age group of dogs. Morphologically, extracellular A β p deposits appeared as small, dot-like rounded deposits, while intracellular p-Tau deposits adopted a diffuse appearance in the retinal layers^[54]. No NFTs or neuropil threads were observed in these dogs. Taken together, these results highlight the importance of further investigations of AD-related pathology in the retina of presymptomatic children and adolescents to gain insight into disease progression and potentially identify early at-risk individuals to help implement speedy therapeutic interventions.

METHODS

Dog eye samples and animal ethics

Sections of eyes used in this study were prepared from archived cases in the Comparative Ocular Pathology Laboratory of Wisconsin (COPLOW) at the Department of Pathobiological Sciences, School of Veterinary Medicine, University of Wisconsin, Madison. Surgical enucleations had been submitted by practicing veterinarians to COPLOW for routine pathological diagnosis. These canine eyes came from 30 random, genetically diverse dogs (15 different breeds) ranging from 1 to 16 years of age with a broad variety of ocular disorders [Tables 1 and 2]. Such tissues, historically submitted for disease investigation purposes, are not subject to approval by institutional animal ethical regulations.

Tissue preparation

Enucleated eyes were submitted to the laboratory fixed in 10% neutral buffered formalin. On receipt, they were examined for gross abnormalities before trimming, followed by processing overnight using a Leica ASP300S tissue processor (Leica biosystem, Wetzlar, Germany). Tissues were then embedded in paraffin blocks and sectioned at 4 μ m thickness using a Leica RM2235 microtome (Leica biosystem, Wetzlar, Germany) and placed on charged slides. Sections were then stained with hematoxylin and eosin (H&E) and Congo red (CR). Further sections were used for immunohistochemistry (IHC) and immunofluorescence (IF) staining, and both upper and lower parts of the retina were assessed.

Hematoxylin and eosin staining

Paraffin sections were dewaxed by two changes of absolute xylene for 5 min each. Sections were rehydrated using two changes of 100% ethanol for 2 min each, 95% ethanol for 3 min, 70% ethanol for 2 min, and finally rinsed in deionized water for 2 min. H&E stain was then performed by adding Gill II Haematoxylin solution (Leica Biosystems, Wetzlar, Germany), followed by 1% acid alcohol and subsequently eosin stain (Leica Biosystems, Wetzlar, Germany). Finally, the sections were dehydrated with increasing concentrations of ethanol, from 70%, 95% to 100%, then dewaxed by two changes of xylene and finally mounted with xylene-based mounting media (Sigma Aldrich, Missouri, United States)^[56]. H&E staining was used to assess retinal morphology, the neuronal population within the eye.

Congo red staining

Initially, the CR working solution was prepared by mixing 50 ml Congo red solution and 0.5 ml potassium hydroxide solution supplied in the Congo red amyloid special stain kit (Leica Biosystems, Wetzlar, Germany). Retinal sections were placed in the working solution for 20 min and then rinsed in 5-8 changes of deionized water. This was followed by staining with Gill II Haematoxylin (Leica Biosystems, Wetzlar, Germany) for 1-3 min and rinsing in 3 changes of deionized water. Sections were then dehydrated in two changes of 95% alcohol followed by three changes of absolute alcohol for one minute each. Finally, the sections were cleared in two changes of xylene and mounted in a xylene miscible medium. Amyloid fibrils appeared as dull to red brick under light microscopy (Olympus CX 43, Shinjuku, Tokyo, Japan) and apple green birefringent under polarized light (Olympus CX 43, Shinjuku, Tokyo, Japan).

Table 1. Signalment for 30 dogs and their retinal A β oligomers and plaques IHC scores

Age groups	Age (year)	Sex	Breed	Size	Pathological A β IHC findings in the dog retina	
					A11 - A β oligomers	4G8 - A β plaques
Young (1-5 years)	1	SF	Staffordshire terrier	Medium	+	-
	1.4	M	Siberian husky	Medium	-	-
	1.8	SF	Mixed breed	ND*	+++	-
	1.10	SF	Siberian husky	Medium	++	-
	2	F	Chihuahua mix	Toy	+++	-
	2	M	Bichon frise mix	Small	+	-
	2	NM	Shih Tzu	Small	+	-
	3	NM	German shepherd	Large	+++	-
	3	F	Giant schnauzer	Large	+	-
	3.6	M	Siberian husky	Medium	+	-
Middle age (6-10 years)	7	NM	Hound mixed	Medium	-	-
	8	SF	Bedlington terrier	Small	-	-
	8	SF	Cocker spaniel	Medium	-	-
	8	M	Mixed breed	ND*	-	-
	8	M	Cocker spaniel	Medium	+	-
	8	F	German shepherd mix	Large	-	-
	8	SF	Great dane dog	Giant	-	-
	9	M	Jack Russell terrier dog	Small	-	-
	9.5	SF	Boxer dog	Large	-	-
	9.9	SF	Cocker spaniel	Medium	-	-
Elderly (11-16 years)	11	NM	Mixed breed	ND*	-	-
	11	NM	Beagle	Small	-	-
	11.9	SF	Bouvier des flanders	Large	+	-
	12	SF	German shepherd	Large	-	++
	12	SF	German shepherd mix	Large	-	-
	12	NM	Husky mix	Medium	-	-
	12.8	NM	Shih-tzu	Small	-	++
	13	NM	Border collie	Medium	-	-
	15	SF	Basset hound	Medium	-	+
	16	SF	Shih Tzu	Small	-	-

Sizes were determined according to the American Kennel Club (AKC)^[76]. Size range between 34 - \geq 54 kg is considered "giant", 24-38 kg is considered "large", 15-29 kg is considered "medium", 3-15 kg is considered "small", and 0.9-4 kg is considered "toy". A β oligomers and A β plaques staining intensity were semi-quantitatively analyzed and scored across the retinal layers under the brightfield microscope (Olympus CX 43, Shinjuku, Tokyo, Japan). The total area was examined at 40 magnification and categorized into no immunostaining "-"; low immunoreactivity found only in limited areas of the retinal layers "+", moderate immunoreactivity where A β deposits were more apparent "++"; and finally strong immunoreactivity with widespread A11 and 4G8 positive A β labeling were exhibited "+++". ND*: Not determined; SF: spayed female; F: female; NM: neutered male; M: male; IHC: immunohistochemical.

Immunohistochemical staining of amyloid-beta plaques and amyloid-beta oligomers

Eye sections that contained the retina were pre-treated using the 2100 antigen retriever (Aptum Biologics Ltd, Southampton, United Kingdom) to expose the target epitopes. The sections were then treated with 90% formic acid for 5 min at room temperature (RT) followed by cell membrane permeabilization, achieved by using 0.1% Triton X for 1 min before the addition of 0.3% H₂O₂ for 15 min to inactivate endogenous peroxidases. Sections were then blocked with protein block serum-free (Agilent, City, Country) for 15 min. The sections were then stained for 1 h with the following primary antibodies in phosphate-buffered saline (PBS): mouse purified 4G8 anti-A β ₁₇₋₂₄ (1:500; Bio legend, San Diego, CA, USA) or A11 rabbit anti-A β o (1:250; Merck Millipore, Burlington, MA, USA) antibodies. After washing with PBS, sections were

Table 2. Signalment for 30 dogs, their ophthalmological disorders, and retinal A β and p-Tau IF scores

Age groups	Age (year)	Sex	Breed	Size	Ophthalmological disorders which resulted in enucleation	Pathological IF findings in the dog retina			
						A β ₄₀ oligomers	A β ₄₂ oligomers	A β	p-Tau
Young (1-5 years)	1	SF	Staffordshire terrier	Medium	Anterior segment dysgenesis	+	+	-	-
	1.4	M	Siberian husky	Medium	Anterior segment dysgenesis	-	-	-	-
	1.8	SF	Mixed breed	ND*	Anterior segment dysgenesis	+++	+++	-	-
	1.10	SF	Siberian husky	Medium	Anterior segment dysgenesis	+++	+++	-	-
	2	F	Chihuahua mix	Toy	Anterior segment dysgenesis	+++	+++	-	-
	2	M	Bichon frise mix	Small	Proptosis	+++	+++	-	-
	2	NM	Shih Tzu	Small	Melanoma limbal	++	++	-	-
	3	NM	German shepherd	Large	Rhabdomyosarcoma orbital	+++	+++	-	-
	3	F	Giant schnauzer	Large	Phthisis bulbi	++	++	-	-
	3.6	M	Siberian husky	Medium	Anterior segment dysgenesis	++	++	-	-
Middle age (6-10 years)	7	NM	Hound mixed	Medium	Hemangiosarcoma	-	-	-	-
	8	SF	Bedlington terrier	Small	Neoplasia	-	-	-	-
	8	SF	Cocker spaniel	Medium	Conjunctivitis	+++	+++	+	-
	8	M	Mixed breed	ND*	Conjunctival melanoma	+++	+++	+	-
	8	M	Cocker spaniel	Medium	Pre glaucoma	+++	+++	+++	-
	8	F	German shepherd mix	Large	Mast cell tumor	++	++	+	-
	8	SF	Great dane dog	Giant	Complex apocrine adenocarcinoma	+	+	-	-
	9	M	Jack Russell terrier dog	Small	Sialoadenitis	++	++	-	-
	9.5	SF	Boxer dog	Large	Hemangioma	+	+	-	-
	9.9	SF	Cocker spaniel	Medium	Adenocarcinoma a orbital	+++	+++	+++	-
Elderly (11-16 years)	11	NM	Mixed breed	ND*	Squamous cell carcinoma	-	-	-	-
	11	NM	Beagle	Small	Melanoma conjunctival	+++	+++	+++	-
	11.9	SF	Bouvier des flanders	Large	Melanoma eyelid	+	+	-	-
	12	SF	German shepherd	Large	Melanoma eyelid	+++	+++	+++	+++
	12	SF	German shepherd mix	Large	Melanoma conjunctival	+++	+++	+++	-
	12	NM	Husky mix	Medium	Orbital carcinoma	+++	+++	++	+++
	12.8	NM	Shih-tzu	Small	Glaucoma	+++	+++	++	+++
	13	NM	Border collie	Medium	Melanoma skin	+++	+++	+++	+++
	15	SF	Basset hound	Medium	Conjunctival hemangiosarcoma	+++	+++	+++	-
	16	SF	Shih Tzu	Small	Squamous cell carcinoma	+	+	-	-

Existing ophthalmological conditions of the 30 neurologically intact unimpaired dogs were determined from the clinical report. Sizes were determined according to the American Kennel Club (AKC). Size range between 34 - \geq 54 kg is considered "giant", 24-38 kg is considered "large", 15-29 kg is considered "medium", 3-15 kg is considered "small", and 0.9-4 kg is considered "toy". A β oligomers A β plaques and p-Tau staining intensity were semi-quantitatively analyzed and scored across the retinal layers under the confocal microscope (LSM800, Zeiss, Oberkochen, Germany). The total area was examined at 40 magnification and categorized into no immunostaining "-"; low immunoreactivity found only in limited areas of the retinal layers "+", moderate immunoreactivity where deposits were more apparent "++"; and finally strong immunoreactivity with widespread A β , A β , and p-Tau labeling was exhibited "+++". ND*: Not determined; SF: spayed female; F: Female; NM: neutered male; M: male; IF: immunofluorescence.

incubated for 1 h at RT with the following secondary antibodies in PBS: HRP-conjugated anti-mouse IgG (Sigma-Aldrich, Missouri, United States) or anti-rabbit IgG (Sigma-Aldrich, Missouri, United States). The

sections were then washed three times with PBS before the addition of the 3,3'-Diaminobenzidine substrate chromogen system and incubated for 5-10 min. The sections were then counterstained with hematoxylin for 1 min before mounting. Sections were finally imaged using Olympus CX 43 light microscopy (Shinjuku, Tokyo, Japan).

The staining intensity was semi-quantitatively analyzed and scored across the retinal layers. Each bright field was examined at 40× magnification and categorized into no immunostaining “-”; low immunoreactivity, found in limited areas of the retinal layers “+”; moderate immunoreactivity where A β deposits were numerous “++”; and finally strong immunoreactivity with widespread A11 and 4G8 positive A β labeling “+++” [Table 1].

Immunofluorescence detection and co-localization studies

To investigate whether A β co-localized with A β p or with p-Tau, we performed immunofluorescence double labeling using camelid-derived single domain anti-A $_{40}$ (PRIOAD12) or anti-A $_{42}$ (PRIOAD13) oligomer antibodies with 4G8 anti-A β p antibody (Bio legend, San Diego, California, United States) or anti-phosphorylated tau AT8 antibody targeting amino acid residues Ser202/Thr205 (Thermo-fisher Scientific, Massachusetts, United States). Slides were processed as described above for IHC before adding the primary antibodies. The sections were double-stained with either PRIOAD 12 (1:500) or PRIOAD 13 (1:500)^[57] and 4G8 antibody overnight (1:500) or AT8 (1:500) and either PRIOAD 12 (1:500) or PRIOAD 13 (1:500). Sections were then washed in Tris-buffered saline with 0.05% Tween 20 (TBST) and incubated with secondary antibodies at a dilution of 1:500, including goat anti-llama IgG conjugated to FITC (Bethyl Laboratories, Inc, Texas, USA) or donkey-anti-mouse IgG conjugated to Texas red (Sigma-Aldrich, Missouri, United States) for 2 hours at room temperature. Other sections were used as negative control and stained with secondary antibodies with the omission of the primary antibodies. Retinal sections derived from an APP/PS1 and TAU58/2 transgenic mice models^[58,59] were used as positive control and stained with anti-A β and anti-P-Tau antibodies, respectively. The sections were covered, slipped with paramount aqueous mounting medium (Dako, Agilent, Santa Clara, California, United States), sealed, and dried overnight. Finally, the sections were visualized with LSM800 confocal microscope with a standard FITC/Texas Red double band-pass filter set (Zeiss, Oberkochen, Germany).

RESULTS

Histological assessment of the retina in 30 dogs

We performed H&E staining of the canine eye sections and examined the retinal layers to assess their general morphological appearance and identify pathological changes such as vacuolation, neuronal death, and eosinophilic deposits^[49,55]. No specific lesions were observed in the retinal layers of the young and middle age groups (data not shown). However, scattered eosinophilic deposits in the ganglion cell layer (GCL) and inner nuclear layer (INL) of the retina were noticeable in 5/10 dogs in the older age group (11-16 years old), including an 11-year-old Beagle, a 12-year-old German shepherd, a 12-year-old Husky mix, a 12.8-year-old Shih Tzu, and a 15-year-old Basset hound [Figure 1A and B]. Furthermore, CR staining used to detect any amyloid fibrils and CAA in eye sections of the dogs did not display any staining in the retina tissues from all age groups (data not shown).

Immunohistochemical detection of A β oligomers and plaques in the retina of 30 dogs

While demonstrated in the CNS of aged animals including cats and bears^[24,60], dogs have been more comprehensively studied and shown to develop human type A β deposition at a very young age. Retinal tissues stained with A11 exhibited intracellular A β o depositions in the outer nuclear layer (ONL), INL, and GCL [Figure 1C and D]. The intraneuronal A β o intensity following A11 staining ranged from low (+), moderate (++) to strong (+++) in all dogs in the young age group except a 1.4 years old male Siberian husky

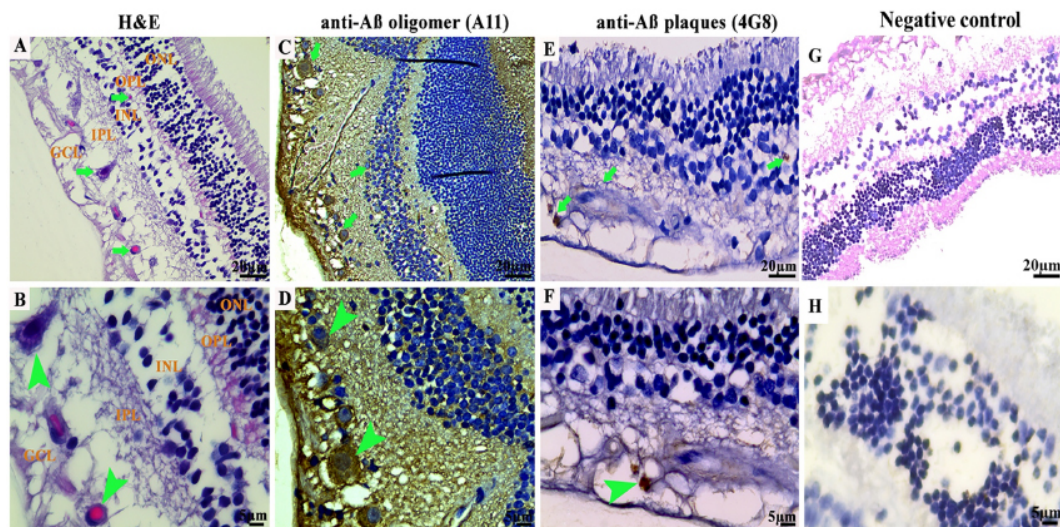


Figure 1. Photomicrographs of the microscopic lesions in the canine retina. The retina in a 12-year-old German shepherd dog. Eosinophilic deposits (green arrows) were observed in the ganglion cell layer (GCL) and inner nuclear layer (INL). Representative of 10 cases of old-aged dogs. Hematoxylin and eosin (H&E) 40 \times . (B) Higher magnification of the deposits in image (A) (green arrowhead) in the GCL. H&E 100 \times . (C) Retina of a 3-year-old German shepherd dog. Immunohistochemical staining with A11 anti-A β o IgG antibody exhibited cytoplasmic A β o depositions in the GCL and INL (green arrows). Representative of 10 cases of young age group. IHC 40 \times . (D) Higher magnification of deposits from image (C) in the GCL (green arrowheads). IHC 100 \times . (E) Retina of a 12-year-old German shepherd mix dog. Immunohistochemical staining with 4G8 anti-A β p IgG antibody exhibited extracellular A β aggregates in the GCL and INL (green arrows). IHC 40 \times . (F) Higher magnification of deposit from image (E) in the GCL (green arrowhead). (G and H) Representative retinal sections derived from dogs of all age groups stained with secondary antibodies with the omission of the primary antibody did not show any depositions (40 \times and 100 \times , respectively). The photomicrographs were taken from the peripheral region of the retina - away from the optic disc. Representative of all 30 dogs examined. IHC 100 \times . IHC: Immunohistochemistry.

that did not display any A11 positive stain [Supplementary Figure 1A]. Of note, retinal sections derived from wild-type mice were used as negative control following immunofluorescence staining [Supplementary Figure 1B and C].

A spayed female mixed breed aged 1.8 years, a female Chihuahua aged 2 years, and a neutered male German shepherd aged 3 years showed strong (+++) accumulation of A11 positive A β o [Figure 1C and D]. Two dogs displayed a low amount (+) of A11 positive A β o in the retinal layers in the middle and old age groups, including an 8-year-old male Cocker spaniel and an 11.9-year-old spayed female Bouvier des Flanders [Table 1]. In addition, 4G8-positive small rounded and dot-like extracellular A β p deposits were observed in the INL, inner plexiform layer (IPL), and GCL [Figure 1E and F] of the retina. They were structurally different from the typical large diffuse plaques normally observed in dog brains with CCD^[21,30,61]. The extracellular A β p intensity following 4G8 staining ranged from low (+) to moderate (++) in some dogs in the old age group, including a 15-year-old spayed female Basset hound, a 12.8-year-old neutered male Shih Tzu, and a 12-year-old spayed female German shepherd. Other dogs in this age group were all negative for 4G8 staining [Table 1, Figure 1E and F]. Representative retinal sections derived from dogs of all age groups stained with secondary antibodies with the omission of the primary antibody did not show any depositions [Figure 1G and H].

Immunofluorescence detection and co-localization of retinal A β_{40} , A β_{42} oligomers, and A β plaques in the retina of 30 dogs

To confirm the presence of retinal A β_{40} and A β_{42} oligomers and to determine whether A β_{40} and/or A β_{42} oligomers co-localized with A β p in the retinas of different age groups and breeds of neurologically intact dogs, we performed immunofluorescence double staining using PRIOAD12 (A β_{40} oligomers), PRIOAD13

(A β ₄₂ oligomers) camelid-derived single domain anti-A β oligomer and 4G8 anti-A β p antibodies. Morphologically, A β o appeared as globular and annular in shape^[55,62] and deposited intracellularly in the ONL, INL, and GCL [Figures 2-4]. 4G8 positive A β plaques appeared morphologically as dot-like and small rounded extracellular deposits^[50,54] in the outer plexiform layer (OPL), IPL, and GCL [Figures 2-4]. We found that both A β ₄₀ and A β ₄₂ oligomers staining was widespread in the majority of the dogs in young [Table 2, Figure 2A and B], middle [Table 2, Figure 3A and B], and old age groups [Table 2, Figure 4A and B] except four dogs including a 1.4-years-old male Siberian husky, a 7-years-old neutered male Hound mixed, an 8-years-old spayed female Bedlington terrier and an 11-years-old mixed breed. In comparison, A β p was absent in young dogs [Table 2, Figure 2C and D], but its presence in the middle age group was moderate [Table 2, Figure 3D and E] and conspicuous in the old age group [Table 2, Figure 4D and E]. Therefore, no co-localization of A β o and A β p was noticed in the retinal layers of the young age group [Figure 2E-H]. However, retinal A β p was shown to co-localize with A β ₄₀ or A β ₄₂ in the GCL, IPL & INL of the middle [Figure 3G, H, J and K] and old [Figure 4G, H, J and K] neurologically intact age groups. Co-localization of A β oligomers and 4G8 positive A β plaques were also exhibited in the retinal vessel wall, where young dogs didn't exhibit any co-localization, the middle age group showed scarce co-localization [Figure 3I and L], and the older age group exhibited conspicuous co-localization in the vessel wall [Figure 4I and L]. In the middle age group, two Cocker spaniels aged 8 years male and 9.9 years female respectively exhibited widespread co-localization of A β o and A β p, and in the old age group, a neutered male Beagle aged 11 years, two spayed female German shepherds aged 12 years, a neutered male Border collie aged 13 years, and a spayed female Basset hound aged 15 years displayed strong and widespread co-localization of A β o and A β p. This co-localization study revealed an age-dependent distribution and co-accumulation of A β o and A β p in the retina of the neurologically intact dogs. We found that the young dogs exhibited widespread accumulation of A β ₄₀ and A β ₄₂ oligomers without any plaque deposits; in the middle age group, A β ₄₀ and A β ₄₂ oligomers accumulation was less conspicuous and, in some cases, co-localized with A β p; and at the old age group, there was widespread and strong co-localization of A β o and A β p strongly distributed in most dogs [Supplementary Figures 2 and 3]. Representative retinal sections derived from dogs of all age groups stained with secondary antibodies with the omission of the primary antibody did not show any co-localization [Supplementary Figure 4A-F]. Retinal sections derived from an APP/PS1 mouse were used as positive control and confirmed the presence of A β ₄₀ and A β ₄₂ oligomers [Supplementary Figure 5A and B] and A β p [Supplementary Figure 5C]^[49,55].

Immunofluorescence co-localization of retinal A β oligomers and phosphorylated tau in the retina of 30 dogs

To confirm the presence of p-Tau and to investigate whether p-Tau co-localizes with A β o, we performed double fluorescence staining of A β o and p-Tau using PRIOAD 12 (A β ₄₀) or PRIOAD 13 (A β ₄₂) anti-oligomer^[57] and AT8 anti-p-Tau antibody, targeting Ser202/Thr205^[54]. A β o appeared as globular and annular in shape^[55,62] [Figure 5A and B] and AT8 positive p-Tau appeared morphologically as diffuse and granular intracellular deposits^[54] in the OPL, INL, IPL, and GCL [Figure 5C and D]. Overall, A β o and p-Tau did not display consistent co-localization in all age groups, as P-tau was only detected in the old age group and A β ₄₀ or A β ₄₂ oligomers were identified in all age groups. Among ten dogs in the old age group, only four dogs have displayed the presence of p-Tau including a spayed female German shepherd and a neutered male Husky aged 12 years, a neutered male Shih Tzu aged 12.8 years, and a neutered male Border collie aged 13 years, and they exhibited strong co-localization with A β ₄₀ or A β ₄₂ oligomers [Figure 5E-H]. Representative retinal sections derived from Tau 58/2 mouse were used as positive control and confirmed the presence of AT8 positive p-Tau [Supplementary Figure 5D]^[58]. Representative retinal sections derived from dogs of all age groups stained with secondary antibodies with the omission of the primary antibody did not show any co-localization [Supplementary Figure 6A and B].

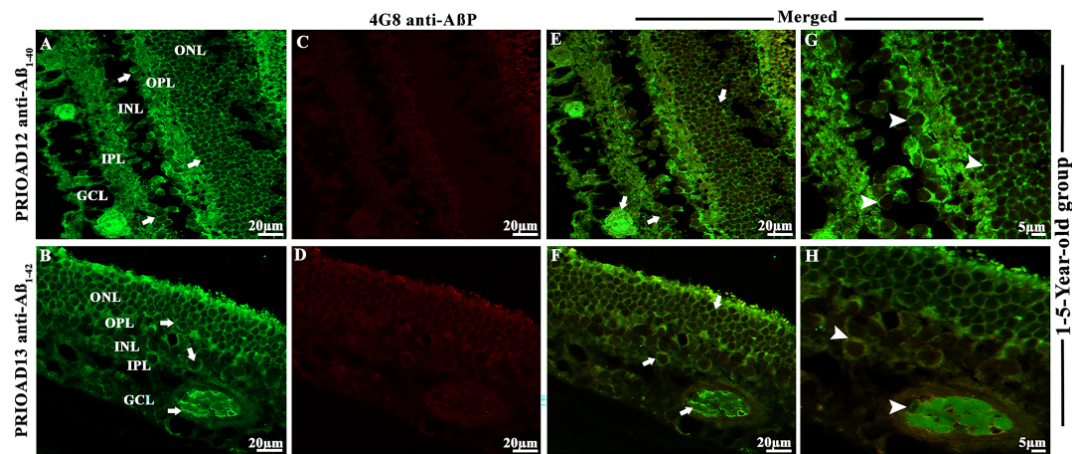


Figure 2. Immunofluorescence (IF) detection and co-localization of retinal amyloid-beta oligomers and amyloid-beta plaques in the dogs of the 1-5-year-old group. Retinal co-staining of oligomers with anti-A β_{40} (PRIOAD12) or anti-A β_{42} (PRIOAD 13) camelid-derived single domain antibodies (green) and plaques with anti-A β (4G8) antibody (red) of a 3-year-old German shepherd dog (A-H). A and B show widespread accumulation of A β_{40} and A β_{42} oligomers in the GCL, INL, and ONL (white arrows) and retinal vasculature, respectively. IF 40 \times . (C and D) No A β p was detected in the retinal layers. IF 40 \times . (E and F) Co-localization of A β o and A β p was not observed in the retinal layers of this animal (A β o was present - white arrows). G and H are higher magnification 100 \times of images (A and B) in the GCL and INL, ONL and retinal vasculature. Representative of 10 dogs in the younger age group (1-5 years). GCL: Ganglion cell layer; INL: inner nuclear layer; ONL: outer nuclear layer.

Influence of demographic factors on the retinal deposition of A β oligomers, A β plaques, and phosphorylated tau in 30 dogs

To investigate the influence of the demographic factors on retinal A β and p-Tau deposition, staining intensity was compared with the age, breed, size, and sex of the neurologically intact dogs [Table 2]. A β_{40} and A β_{42} oligomers, A β p and p-Tau fluorescence intensity, were assessed at 40 \times magnification. Immunoreactivity throughout the retinal layers was semi-quantified and categorized into no immunostaining “-”; low immunoreactivity, exhibited in limited areas of the retinal layers “+”; moderate immunoreactivity where deposits were more apparent “++” and finally strong immunoreactivity with widespread A β o, A β p and p-Tau labeling “+++” [Table 2]. After comparing the age of the dogs and immunofluorescence staining scores, we found that both A β_{40} and A β_{42} oligomers staining was high in young dogs, which slightly decreased in the middle age group, then finally, an upward trend was noticed in older dogs [Table 2, Figure 6]. In comparison, A β p was absent in young dogs, but its presence in the middle age group was moderate and high in the old age group [Table 2, Figure 6]. Finally, p-Tau deposits were not observed in the young and middle age groups, whereas old dogs exhibited widespread staining for p-Tau [Table 2, Figure 6]. However, when the size of the dog was compared with the pathological outcome, we found that six medium-sized dogs displayed strong A β o staining and only two dogs exhibited strong p-Tau staining [Table 2, Figure 7]. In addition, seven dogs revealed strong A β p staining, of which four were medium size [Table 2, Figure 7].

Influence of eye pathology on the retinal deposition of A β oligomers, A β plaques, and phosphorylated tau in 30 dogs

Further, to understand whether the presence of pre-existing underlying eye pathology in the dogs may influence retinal A β and p-Tau depositions, we compared their staining intensity with the reported underlying clinical eye conditions [Table 2]. We found that four young dogs aged 1-5 years affected with anterior segment dysgenesis, a spectrum of disorders that affect the development of the anterior segment, including the cornea, iris, ciliary body, and lens^[63-65], displayed “moderate” to “strong” intensity staining for A β_{40} and A β_{42} oligomers [Table 2]. However, another two dogs in this age group and affected with anterior segment dysgenesis showed little to no deposition of A β_{40} and A β_{42} oligomers [Table 2]. Eighteen out of

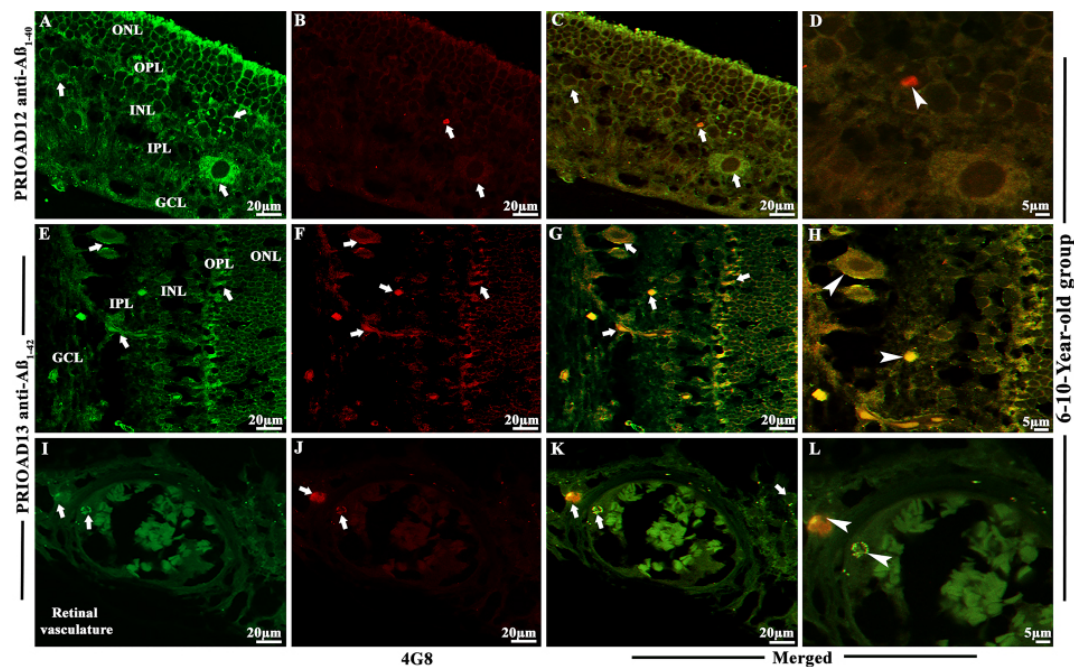


Figure 3. Immunofluorescence detection and co-localization of retinal amyloid-beta oligomers and amyloid-beta plaques in dogs of the 6-10-year-old group. Retinal co-staining with anti- $A\beta_{40}$ (PRIOD 12) and anti- $A\beta_{42}$ (PRIOD 13) camelid-derived single domain antibody (green) and 4G8 antibody (red) of a 9-year-old Cocker spaniel dog (A-L). (A) A large number of $A\beta_{40}$ and (B) $A\beta_{42}$ oligomers were found in the GCL, INL, and ONL (white arrows, 40 \times). (C) Detection of $A\beta_{42}$ oligomers in the vasculature. 4G8 positive $A\beta$ plaque-like deposits were observed in the (D and E) GCL, IPL, INL, and OPL of the retina (white arrows, 40 \times). Widespread co-localization was observed in the (G and H) retinal layers (white arrows, 40 \times). Co-localization of 4G8 positive $A\beta$ plaque with (J) $A\beta_{40}$ and (K) $A\beta_{42}$ depositions (white arrowhead) showed with higher magnification (100 \times) in the GCL and INL of the same 9-year-old Cocker spaniel dog retinal section. (C) $A\beta$ oligomers and (F) 4G8 positive $A\beta$ plaques were observed in the retinal vasculature (white arrows). (I and L) Co-localization of $A\beta$ oligomers and 4G8 positive $A\beta$ plaques were exhibited with 40 \times and with higher 100 \times magnification in the retinal vessel wall, respectively (white arrowhead). The photomicrograph was derived from the peripheral region of the retina - away from the optic disc. Representative of 10 dogs examined from middle age group (6-10 years). GCL: Ganglion cell layer; IPL: inner plexiform layer; INL: inner nuclear layer; OPL: outer plexiform layer; ONL: outer nuclear layer.

thirty dogs presented with eye neoplasms, including two in the young age group, seven in the middle age group, and nine in the old age group [Table 2]. There was no clear relation between neoplasms and the intensity of $A\beta$ or p-Tau [Table 2]. Interestingly, eye inflammation (proptosis/ phthisis bulbi in young dogs and conjunctivitis in a middle-aged dog) corresponded to $A\beta_{40}$ and $A\beta_{42}$ oligomers intensity that ranged from "moderate" to "strong" [Table 2]. Finally, a case of pre-glaucoma in the middle age group and a case of glaucoma in the old age group also matched well with $A\beta_{40}$ and $A\beta_{42}$ oligomers as well as $A\beta p$ intensity. Overall, this analysis appears to indicate that no direct influence of pre-existing eye disorders on $A\beta$ and p-Tau intensity exists; however, the small size of the cohorts used in this study did not allow to reach a substantive conclusion, and studies with much larger cohorts are needed.

DISCUSSION

Eye imaging can provide an opportunity to develop an easily accessible and point-of-care routine diagnostic testing to help predict MCI/AD early^[66,67]. A study by Ko and colleagues examined the retinal NFL thickness and cognitive status of individuals aged 40 to 69 years over three years^[68]. The authors found that individuals with thinner NFL showed a higher incidence of reduced cognition.

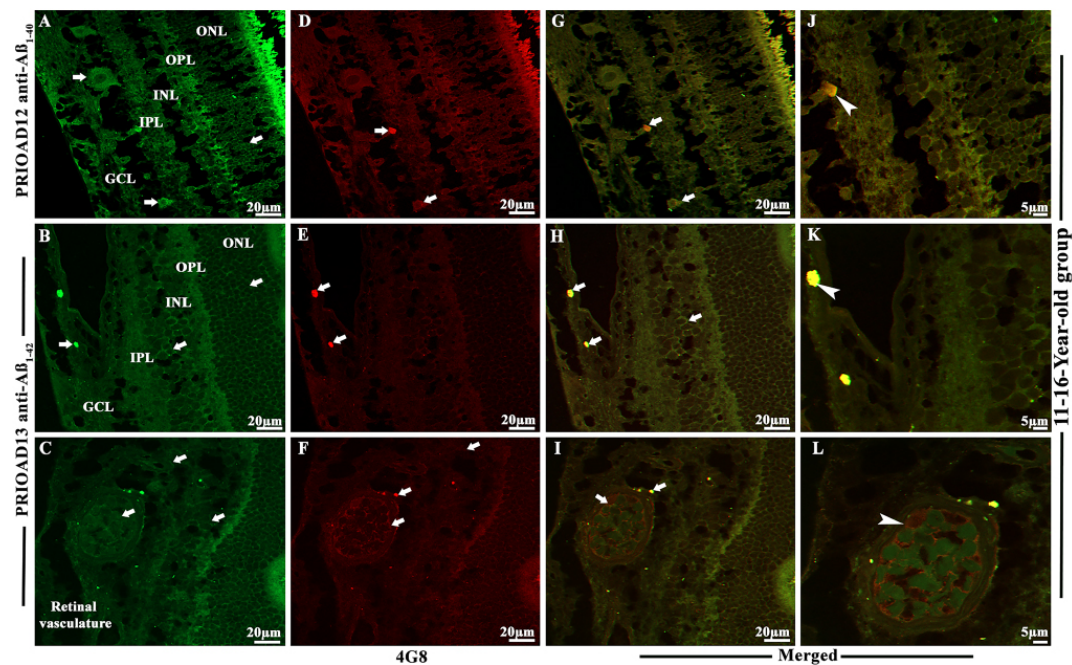


Figure 4. Immunofluorescence co-localization of retinal amyloid-beta oligomers and amyloid-beta plaques in dogs of 11-16-year-old group. Retinal co-staining with anti-A β_{40} (PRIOAD 12) and anti-A β_{42} (PRIOAD 13) camelid-derived single domain antibody (green) and 4G8 antibody (red) of a 12-year-old German shepherd dog (A-L). A large number of (A) A β_{40} and (B) A β_{42} oligomers were found in the GCL, INL, and ONL (white arrows, 40 \times). 4G8 positive A β plaque-like deposits were observed in the (D and E) GCL of the retina (white arrows, 40 \times). Widespread co-localization was observed in the (G and H) retinal layers (white arrows, 40 \times). Co-localization of 4G8 positive A β plaque with (J) A β_{40} and (K) A β_{42} depositions (white arrowhead) showed with higher magnification (100 \times) in the GCL of the same 12-year-old German shepherd dog retinal section. (C) A β oligomers and (F) 4G8 positive A β plaques were observed in the retinal vasculature (white arrows). (I and L) Co-localization of A β oligomers and 4G8 positive A β plaques were exhibited with 40 \times and also with higher 100 \times magnification in the retinal vessel wall, respectively (white arrowhead). The photomicrograph was derived from the peripheral region of the retina - away from the optic disc. Representative of 10 dogs examined from elder age group (11-16 years). GCL: ganglion cell layer; INL: inner nuclear layer; ONL: outer nuclear layer.

Despite the validation of the dog as a robust translational model for AD^[24,30,69], only very limited studies investigated AD-related changes in the young and neurologically intact dogs. One study by Stylianaki and colleagues confirmed a similar pattern to that of humans; 61 dogs were subdivided into young (0-4 years old), middle-aged (4-8 years old), aged cognitively normal (8-20 years old), and aged cognitively impaired (8-17 years-old)^[70]. The authors found that young dogs displayed the highest levels of total plasma A β_{42} and A β_{42} /A β_{40} ratio and the middle-aged dogs had the highest cerebrospinal fluid A β_{40} and A β_{42} when compared to neurologically intact aged dogs. Our first step was to investigate the distribution and morphological appearance of both A β_{40} and A β_{42} oligomers representing the canine lifetime, in the retinal layers of these neurologically intact 30 dogs. Previous studies have shown that A11 anti-oligomer antibody binds to different epitopes of the A β but also reacts with oligomeric aggregates of other proteins independent of their primary sequences. Of note, A11 cannot differentiate between A β_{40} and A β_{42} . In contrast, PRIOAD12 and PRIOAD13 nanobodies bind to A β_{40} and A β_{42} , respectively. Previous studies reported an inverse interrelationship between neurotoxicity and the size of A β , where the toxic effect of A β was shown to decrease with increased size. The general molecular weight of A β was found to range between 10-100 kDa in AD brain and included dimers to dodecamer^[71]. A study by Lambert and colleagues first describes the cytotoxic effect of the small diffusible A β on the hippocampal neurons^[72]. The small A β referred to as A β -derived diffusible ligands toxicity was tested in organotypic mouse brain slice cultures. The authors found that 17 and 22 kDa small oligomers killed hippocampal neurons at nanomolar concentration^[72]. In addition, a study by Cizas and colleagues confirmed that oligomers larger than 30 kDa have a less toxic effect on the

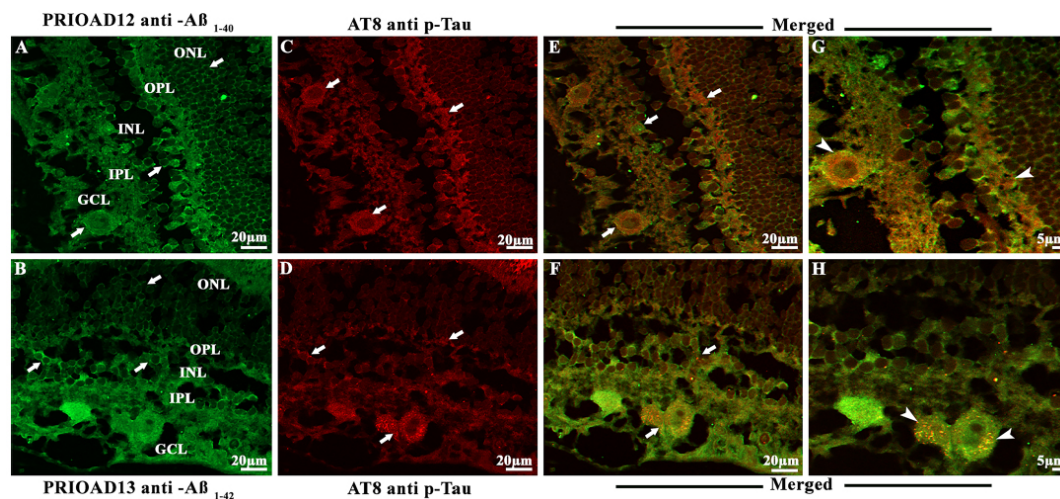


Figure 5. Immunofluorescence co-localization of retinal amyloid-beta oligomers and hyperphosphorylated tau in dogs of 11-16-year-old group. Retinal co-staining with anti-A β_{40} (PRIOAD 12) and anti-A β_{42} (PRIOAD 13) camelid-derived single domain antibody (green) and AT8 antibody (red) of a 12-year-old German shepherd dog. A large number of (A) A β_{40} and (B) A β_{42} oligomers were found in the GCL, INL, and ONL (white arrows, 40 \times). AT8 positive diffuse p-Tau-like deposits were observed in the (C and D) GCL, IPL, INL, and OPL of the retina (white arrows, 40 \times). Widespread co-localization was observed in the (E and F) retinal layers (white arrows, 40 \times). Co-localization of AT8 positive p-Tau with (G) A β_{40} and (H) A β_{42} depositions (white arrowhead) showed with higher magnification (100 \times) in the GCL and OPL of the same 12-year-old German shepherd dog retinal section. The photomicrograph was derived from the peripheral region of the retina - away from the optic disc. Representative of 10 dogs examined from elder age group (≥ 11 years). GCL: Ganglion cell layer; IPL: inner plexiform layer; INL: inner nuclear layer; OPL: outer plexiform layer; ONL: outer nuclear layer.

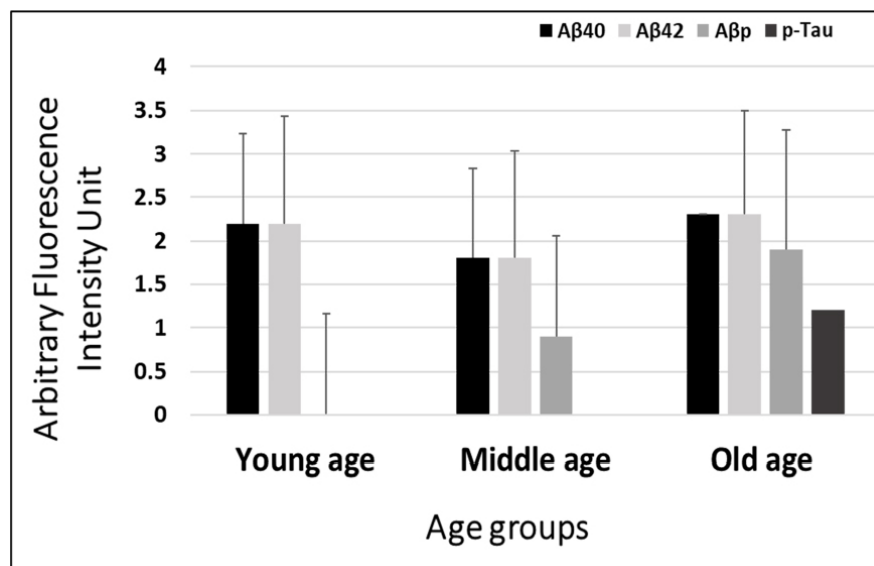


Figure 6. Semiquantitative analysis of A β_{40} and A β_{42} oligomers, A β plaques, and p-Tau in neurologically intact young (1-5-years), middle (6-10-years), and old (11-16-years) age groups of dogs. PRIOAD12 (A β_{40} oligomers), PRIOAD13 (A β_{42} oligomers), 4G8 (A β_p), and AT8 (p-Tau) fluorescence immunoreactivity and intensities were examined and quantified at 40 \times magnification under the fluorescence microscope. Semiquantitative analyses were compared with the three different age groups. A large amount of A β_{40} and A β_{42} oligomers were found in young dogs, which decreased in middle age group, then finally, an upward trend was noticed in older dogs. In comparison, A β_p were completely absent in young dogs, and then moderately found in middle, and large amounts in old age groups. Finally, p-Tau deposits were not found in young and middle age groups of dogs, whereas old dogs exhibited a considerable amount of p-Tau.

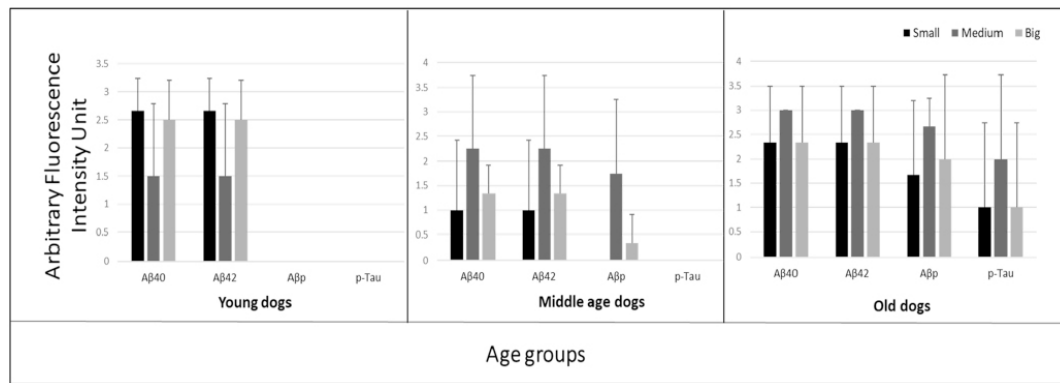


Figure 7. Influence of the size of dogs on the retinal deposition of amyloid-beta oligomers (Aβ_o), plaques (Aβ_p), and phosphorylated tau (p-Tau) in cognitively unimpaired young (1-5-years), middle (6-10-years) and old (11-16-years) age groups of dogs. PRIOAD12 (Aβ₄₀ oligomers), PRIOAD13 (Aβ₄₂ oligomers), 4G8 (Aβ_p), and AT8 (p-Tau) fluorescence immunoreactivity and intensities were examined and quantified at 40× magnification under the fluorescence microscope. In three different age groups, semiquantitative analyses were compared with the size of the dogs. In young dogs, a large number of oligomers were displayed by the small and big size dogs. The majority of medium-sized dogs displayed strong Aβ_o and Aβ_p staining intensity in the middle-aged dogs. Finally, among the different sizes of old dogs, medium-sized breeds displayed the highest amount of Aβ_o, Aβ_p, and p-Tau staining intensity.

inhibition of long-term potentiation^[73]. They also suggested the transition of sizes from small to large correlate with the high to a low toxic effect of Aβ_o^[73]. A11 specifically binds to the prefibrillar oligomers. However, studies suggested that the specificity of A11 to AD-related oligomers might vary as it recognized prefibrillar oligomers from various proteins that share a common structure including α-synuclein, islet amyloid polypeptide, polyglutamine (PolyQ), lysozyme, and prion peptide^[36]. A study by Glabe and colleagues reported the ideal band size of prefibrillar oligomer-specific antibody A11 ranged from approximately tetrameric up to ~75 kDa^[35]. In our study, we confirmed the presence of an A11-specific band at ~70KDa, confirming binding to prefibrillar oligomers in APP/PS1 mice. In comparison, camelid-derived single-domain nanobodies bind to ~15KDa band representing the small oligomers. In agreement with this current study, our previous report^[55] also suggested that nanobodies were able to detect the toxic small diffusible oligomers specific to AD, whereas A11 binds to the larger oligomers believed to be less toxic to the neurons.

Studies suggested that in the canine brain, Aβ_o may be the toxic species responsible for cognitive decline and can potentially be an early biomarker for the detection of CCD^[23,74]. Naaman and colleagues recently reported that Aβ₄₂ oligomers cause extensive retinal neurotoxicity in rats when compared to fibrillary Aβ₄₀ and Aβ₄₂^[75]. Our findings demonstrated extensive deposition of Aβ₄₀ and Aβ₄₂ oligomers in the retinal layers, including GCL, INL, and ONL in 26/30 in the young, middle, and old age groups, except for a 1.4-year-old male Siberian husky, a 7-year-old neutered male Hound mixed, an 8-year-old spayed female Bedlington terrier, and an 11-year-old neutered male mixed breed. Moreover, we did not notice any difference in the intensity of immunofluorescence (graded - to +++), the pattern of deposition, or morphology between Aβ₄₀ and Aβ₄₂ oligomers in all age groups. This presentation was unlikely influenced by the breed of the animals; for instance, out of three Siberian Huskies, one dog failed to display Aβ₄₀ and Aβ₄₂ oligomers accumulation. Remarkably, the younger age group, with an age range of 1-5 years, equivalent to humans aged 15-40 years according to the American Kennel Club^[76], has displayed extensive deposition of Aβ oligomers. Of interest, a 2013 brain study investigated the presence of three different types of Aβ oligomers, including Aβ trimers, Aβ*56, and Aβ dimers, in 75 cognitively unimpaired individuals aged 1 to 96 years^[14]. Young children and adolescents were positive for Aβ oligomers; the authors found an age-related accumulation of Aβ oligomers where the level of the amyloid-β dimer was significantly higher in subjects in their 60s, and amyloid-β trimer

in their 70s, whereas A β *56 level was significantly higher in individuals in their 40s. The investigators proposed that A β o, specifically A β *56, might trigger the pathological cascade in asymptomatic AD, a phase that may be identifiable two decades before the clinical onset^[14].

Having found that the eyes of 26/30 dogs contained A β ₄₀ and A β ₄₂ oligomers, our next step was to demonstrate the presence/deposition of fibrillar A β in the canine retina of this same cohort. We show that none of the young dogs displays 4G8-positive A β fibrils and plaques (A β p). However, A β p were observed in some of the middle age group, where three dogs exhibited "low" staining intensity and two exhibited "strong" staining intensity in the GCL, IPL & INL. A β p were also observed in the old age group, where five dogs showed "strong" signal intensity and two with "moderate" signal intensity in the GCL, IPL & INL. A pattern, not surprising, emerges; A β ₄₀ and A β ₄₂ oligomers are deposited in the retina following birth and subsequently, a fibrillar A β configuration, aggregated as plaques follow. Several studies have demonstrated the presence of A β p in the canine brain and suggested only a weak correlation with the severity of cognitive dysfunction^[77,78]. Schütt *et al.* investigated this question in further depth by studying dogs at differing ages, specifically aged 9-15 years, including neurologically intact dogs and dogs with CCD, and compared their amyloid burden with that of young dogs aged less than 6 years^[69]. The authors reported that the levels of A β deposition strongly correlated with the age of the dog but not to their cognitive capacity. Asking the same questions regarding amyloid accretion in non-CNS neuronal populations, such as the retina, has not been investigated in the canine, but some recent reports confirmed their presence in the retina of AD^[50,54,79]. In this study, we have demonstrated accumulation of retinal A β p in the mature to an aged group of these 30 dogs which supports the hypothesis that A β p might not influence the severity of cognitive deficits but can be a predictor of AD development^[11,17]. Overall, our study revealed that the A β deposition pattern in the retina was such that A β o was observed in all age groups, whereas A β p accumulation was restricted to the middle and more intensely the old age group of dogs, regardless of demographic criteria, breed and gender. Interestingly, co-accumulation of both A β o and A β p were apparent in some middle and old dogs. This is in agreement with our report this year, where 17-18 months old APP/PS1 mice showed high levels of oligomers deposits in the retinal layers that co-localized with A β p^[55]. Previous neuropathological studies in AD have shown that cerebral A β o is usually present in early disease stages and is the most cytotoxic of A β species, mostly responsible for neurotoxicity and synaptic dysfunction^[34,80-83], whereas extracellular A β p is believed to accumulate at later ages and may act as a reservoir for A β o^[84]. Moreover, the involvement of A β o in retinal degeneration in AD has also been reported^[85-87]. In the study reported herein, we noticed widespread distribution of A β o in the retinal layers of dogs of all age groups, indicating potential involvement in retinal degeneration, perhaps leading to vision impairment in some dogs^[88,89]. Of importance, a previous study by Ozawa and colleagues that focused on web and paper-based surveys of dogs aged ≥ 10 years to identify physical disturbances related to CCD showed that more than 90% of dogs affected with CCD had vision impairment^[90]. In addition to the neural retina, we observed A β deposition s in the retinal microvasculature of young, middle, and old age groups, which might imitate CAA in AD^[91-93]. Sharafi *et al.* suggested that retinal vasculature changes captured by hyperspectral imaging can differentiate cerebral amyloid status between cognitively impaired and unimpaired individuals^[94].

Central to AD pathogenesis is the intraneuronal deposition of hyperphosphorylated tau (p-Tau) in granular form and eventual organization as NFTs, one of the cardinal neuropathological hallmarks^[3,4]. In dogs with CCD, p-Tau, unlike NFTs, has been identified in the brain, but until recently, only in some cases. We speculate that, unlike human AD, p-Tau is only evident in pretangle granular form in dogs due to their shorter lifespan (Tayebi & Habiba, unpublished observation). p-Tau neuropathology was shown to develop about a decade before the formation of A β p in AD brains and was hypothesized to trigger AD^[95,96]. However, A β o accumulation was shown to precede and drive p-Tau accumulation and transneuronal spread across synapses in the parietal cortex of AD^[96].

In the retina of human AD patients, p-Tau displayed a diffuse pattern in the plexiform layers in the absence of NFTs^[54]. Schön and colleagues reported the presence of AT8-positive intracellular NFTs and diffuse p-Tau signals in the retina of 5/6 deceased AD patients^[97]. In this study, we investigated the presence of p-Tau Ser202/Thr205 to confirm the presence of retinal p-Tau in dogs. We revealed diffuse p-Tau distribution in the OPL, INL, IPL, and GCL in 4/10 dogs of the old age group, including a spayed female German shepherd and a neutered male Husky aged 12 years old, a neutered male Shih Tzu aged 12.8 years old, and a neutered male Border collie aged 13 years old. The same dogs also showed the widespread distribution of A β ₄₀ and/or A β ₄₂ oligomers and 4G8- positive A β p. Similarly, in the brains of cognitively impaired, 14-17 years old dogs, Abey and colleagues have demonstrated the presence of p-Tau Ser202/Thr205 and p-Tau Ser396 in 1/6 and 6/6, respectively^[40]. This pattern of deposition strongly supports a possible age-dependent progression as observed in AD mice models^[49,55]. Previous studies have shown that breed variance does not determine the pathological outcome associated with CCD in aged dogs^[98,99]. In agreement, our study did not reveal any breed-dependent pathological accumulation.

While previous studies have demonstrated the presence of A β p and p-Tau in the aged dog brain^[30,43], Pugleise *et al.* proposed that the acquisition of diffuse A β p and p-Tau are unrelated and independent events of aged dogs^[42]. Whether these two “ingredients” of AD dementia or CCD are closely linked (co-localized or at least act in concert) is likely crucial. In AD, it was shown that p-Tau and A β o cause neuronal toxicity and may act synergistically to trigger synaptic dysfunction^[36,37,100]. Manczak and colleagues showed that A β ₄₀ or A β ₄₂ oligomers co-localized with p-Tau in the brains of AD patients^[101]. The authors also reported that the interaction between A β o and p-Tau became more prominent with disease progression and was more pronounced at Braak stage V and VI compared to Braak stage III and IV^[102,103]. The evidence from this canine retina study would support an interaction: we also show that p-Tau co-localized with A β ₄₀ or A β ₄₂ oligomers in the OPL, INL, IPL, and GCL of the retina of 4/10 dogs in the eldest age group. A recent PET brain imaging study by Lockhart *et al.*, which demonstrated the presence of both A β and Tau pathology in cognitively normal older adults, showed a significant spatial correlation with A β and Tau deposition^[104]. Thus co-localization, while evident, does not itself necessarily imply neuronal cytotoxicity.

Several studies in the field of AD research highlighted the importance of diagnostic and therapeutic interventions in the asymptomatic phase while the individuals are cognitively intact^[14,105]. However, it is challenging to first identify and then track disease progression in humans without sensitive, specific, and cost-effective early diagnostic approaches and a lack of robust natural translational models. In that context, dogs have the potential to be an effective natural model for the study of aging and AD. They share the same environment as humans^[106,107]; there exists the option of brain CT and MRI imaging and intracranial biopsy, ease of cognitive testing which helps reduce physiological stress, and finally, a short lifespan^[108]. Canine ophthalmology is a well-established clinical and investigative discipline (inherited retinopathies, for example). The current study provides strong impetus to track how this seemingly common process evolves, with potential consequences for aging, neurodegenerative disease, and also vision.

Lateralization in retinal and cerebral A β levels was previously investigated in an AD mouse model^[109]. Some studies revealed left posterior brain-dominant lateralization in A β _{42/40}; however, no left or right brain hemisphere or retinal dominance lateralization was observed for A β ₄₀ and A β ₄₂. Although our current study does not provide information about left versus right retinal accumulation of A β , the significance of the lateralization in human Alzheimer's and CCD remains unknown. Moreover, previous reports have shown that A β accumulation was more prominent in the far-peripheral and mid-peripheral of the superior

temporal quadrant than in the central retina in AD patients^[50,79]; however, in our study, no notable differences in A β and tau deposition have been observed in different parts of the retina.

The canine eye tissues used in this study had been surgically removed in the course of managing spontaneous disorders of the globe, eyelids, or structures within the orbit which necessitated enucleation. It could be asked whether any of these conditions played any part in A β and/or p-Tau deposition in the retina. We think this unlikely, given the diverse clinical conditions involved, which ranged from developmental abnormalities, inflammatory conditions, and proptosis to various neoplasms of differing grades of the eye, while other cases were disorders arising in the conjunctiva or orbit. The span of ages in these dogs (from juvenile to quite aged) would further make this seem unlikely and some evidence of human retinal A β and/or p-Tau exists, though such explorations are largely performed only at the end of life.

DECLARATIONS

Authors' contributions

Performed experiments and revised manuscript: Habiba U

Revised manuscript: Morley J, Krockenberger M

Was a major contributor in writing and revising the manuscript: Summers BA

Conceived and designed experiments and wrote manuscript: Tayebi M

All authors read and approved the final manuscript.

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All authors declared that there are no conflicts of interest.

Ethical approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

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Clinical Observation

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Olfactory and gustatory dysfunctions in patients with COVID-19 in Wuhan, China

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Abstract

The coronavirus disease 2019 (COVID-19) is spreading all over the world. The main symptoms of COVID-19 include fever, cough, fatigue, and myalgia. However, there are few reports on olfactory and gustatory dysfunctions in patients with COVID-19. Our objective was to investigate the incidence of olfactory and gustatory dysfunctions in patients with COVID-19 infection in Wuhan, China. In this retrospective study, we collected 81 confirmed cases of COVID-19 from the Renmin Hospital of Wuhan University from February 2020 to March 2020, and analyzed the demographic characteristics, clinical manifestations (including olfactory and gustatory dysfunctions), laboratory findings, and comorbidities. In total, 81 confirmed COVID-19 patients were enrolled in this study (38 males and 43 females). The study showed that the most prevalent symptoms included cough, myalgia, and loss of appetite. On admission, 25 (30.9%) of all patients reported either olfactory dysfunction (OD) or gustatory dysfunction (GD), and 7 (8.6%) reported both OD and GD. Overall, 13.6% and 25.9% of all patients reported OD and GD, respectively. OD and GD were not associated with disease severity. Pearson correlation analysis identified some factors that are correlated with OD and GD, including headache or dizziness ($r = 0.342$, $P = 0.002$), dark urine ($r = 0.256$, $P = 0.021$), IgM titer ($r = 0.305$, $P = 0.01$), and diabetes ($r = 0.275$, $P = 0.013$). Multivariate adjusted logistic analysis showed that patients with headache or dizziness (OR = 7.382, 95%CI: 1.028-53.022) and diabetics (OR = 6.871, 95%CI: 1.096-43.086) were at higher risk of developing OD. In addition, patients with high IgM titers (OR = 1.006, 95%CI: 1.000-1.006) and diabetes (OR = 4.335, 95%CI: 1.098-17.287) were at high risk of developing GD. In 81.8% of the cases with OD and 28.6% of the cases with GD, the symptoms lasted for at least one month after



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discharge. In addition, 3.6% of inpatients without OD developed OD after discharge. These findings suggest that OD and GD are common in COVID-19. These symptoms appear early during the course of the disease and may last for one month. The incidence of OD and GD is related to neurological manifestations, diabetics, and IgM titers.

Keywords: COVID-19, SARS-CoV-2, smell, taste, infection

INTRODUCTION

Coronavirus disease 2019 (COVID-19), an infectious disease caused by the acute respiratory syndrome coronavirus-2 (SARS-CoV-2), has spread globally. As of March 20, 2022, there have been more than 464 million confirmed cases of COVID-19 globally^[1]. Early epidemiological investigations show that the basic reproduction number (R_0) of SARS-CoV-2 is estimated to be about 2.2 (range, 1.4-6.5)^[2]. The high spread and low fatality rate of COVID-19 made it a global pandemic. COVID-19 has become the largest pandemic since the H1N1 flu outbreak in 1918. Early genetic tests emphasized that the genome sequence identity of SARS-CoV-2 and SARS-CoV is 79.5%. Both viruses infect cells through angiotensin-converting enzyme 2 (ACE2)^[3,4]. The similarity of the gene sequence and infection route suggests that SARS-CoV-2 and SARS-CoV have similar clinical manifestations. Similar to SARS, the most common symptoms of COVID-19 are fever, cough, sputum, fatigue, and myalgia^[5-7]. Complications of COVID-19 involve multiple systems, such as the cardiovascular system, digestive system, and nervous system^[8-10]. Current research mainly focuses on central nervous system complications such as cerebrovascular diseases and acute encephalopathy^[10], while less is known about the peripheral nervous system complications. Symptoms in the head and neck such as dry eyes, tearing, and smell and taste disturbances were reported in COVID-19 patients and believed to be related to the neurophagy of the virus^[11-12]. It is worth noting that SARS-CoV can be detected 60-66 h after infection and is most abundant in the olfactory bulb^[13]. In addition, the virus may enter the brain through the olfactory bulb. Viral antigens can also be detected in brain areas connected to the olfactory bulb, such as the piriform nucleus and subliminal leather, basal ganglia, and midbrain^[14,15].

The purpose of this study was to investigate the incidence of olfactory and taste disorders in confirmed COVID-19 cases and to analyze the risk factors related to olfactory and gustatory dysfunctions in order to provide clues for the clinical diagnosis and treatment of COVID-19.

METHODS

Patient enrollment and data collection

This descriptive study included 81 inpatients diagnosed with COVID-19 in the Renmin Hospital of Wuhan University, Wuhan, China. The diagnosis of COVID-19 was based on clinical symptoms, computed tomography (CT), real-time RT-PCR, and next-generation sequencing. The patient's admission date was from February 1 to March 3, 2020. All patients participating in this study lived in Wuhan during the COVID-19 outbreak. Two well-trained investigators collected the data including demographic characteristics, clinical characteristics (including medical history, comorbidities, and symptoms), preliminary laboratory findings, treatment, and clinical outcomes.

According to the SARS-CoV-2 diagnosis and treatment guidelines (Versions 3-7) issued by the National Health Commission of China, the severity of COVID-19 patients was defined. Severe cases were designated when the patients fit one of the following criteria: (1) respiratory distress with respiratory rate $\geq 30/\text{min}$; (2) oxygen saturation $\leq 93\%$ at rest; (3) arterial partial oxygen pressure (PaO_2)/oxygen absorption concentration (FiO_2) $\leq 300 \text{ mmHg}$; (4) respiratory failure that needs mechanical ventilation; or (5) shock or organ failure requiring ICU care.

Procedures

Two investigators collected the information on epidemiology, clinical symptoms, laboratory test results, treatment, and clinical outcomes. Laboratory tests included routine blood examination (including white blood cells, lymphocytes, and platelets), coagulation function, lactate dehydrogenation, and immune indicators (including lymphocyte differential counts and inflammatory factors). Participants were interviewed by telephone and were asked to report their olfactory and gustatory dysfunctions at the time of hospitalization and one month after discharge. Two years after discharge, those with olfactory and gustatory dysfunctions one month after discharge were re-interviewed to report their smell and taste status. Verbal informed consent was obtained from all participants or their legal guardians. The severity of symptoms was graded as totally normal, mild, moderate, and severe. Totally normal means that respondents rated smell and taste as “Good” or “Unaffected”, mild dysfunction means the patient hardly noticed a smell and taste reduction, moderate dysfunction means there was a noticeable reduction, and severe dysfunction means the patient reported an almost complete loss of smell and taste. Patients with olfactory and gustatory dysfunctions before contracting COVID-19 were excluded, and so were patients with a diagnosis of chronic rhinitis, Alzheimer’s disease (AD), or Parkinson’s disease (PD) on admission. We also asked patients during the telephone interview whether they had a history of chronic rhinitis, AD, or PD. These patients might experience olfactory and gustatory dysfunctions, which could affect the study results. Cases with unsuccessful follow-ups were excluded.

Statistical analysis

Continuous variables were described as medians and interquartile range (IQR), and categorical variables were described as frequency rates and percentages. We used the chi-square (χ^2) test or the Fisher’s exact test to compare categorical data. Mann-Whitney-Wilcoxon test was applied to compare non-normally continuous variables. Pearson correlation analysis was used to examine the correlations between olfactory/gustatory dysfunctions and all indicators (including epidemiology, clinical symptoms, laboratory test results, treatment, and disease severity). The sample size varied due to missing data, and missing data were not imputed. The analyses regarding different factors were based on non-missing data. Multivariate adjusted logistic regression models were adopted to analyze the factors influencing olfactory and gustatory dysfunction in patients with COVID-19. All statistical analyses were performed using SPSS software (V.23.0). Two-tailed *P* values were considered statistically significant at < 0.05 . This study was approved by the Hospital Ethics Committee of the Renmin Hospital of Wuhan University (WDRY2020-K136).

RESULTS

Demographic characteristics in COVID-19

The demographic characteristics, clinical manifestations, and comorbidities are shown in Table 1. Of the 81 patients, 63 cases were diagnosed as non-severe and 18 cases were severe. The median age was 58 years (IQR, 50.0-68.5 years). The median age of the severe group was 11.5 years older than that of the non-severe group. Males accounted for 46.9% (38 of 81) of the total patients. However, males accounted for 66.7% (12 of 18) of patients in the severe group compared with 41.3% (26 of 63) of the non-severe group. These results suggest that males and older patients are more likely to develop into severe cases, which is consistent with other reports^[16]. On admission, the most common clinical manifestations were fever (72.8%) and respiratory symptoms (70.4%), while chest pain, fatigue, gastrointestinal symptoms, dark urine, and headache or dizziness accounted for 27.2%, 37%, 27.2%, 19.8%, and 14.8% of the patients, respectively. Compared with the non-severe group, the severe group had a higher proportion of dark urine (17.5% vs. 27.8%), suggesting a potential relationship between urinary system dysfunction and disease severity. Among the 81 cases, 51.9% had at least one comorbidity, including hypertension, diabetes, coronary heart disease, stroke, cancers, and chronic pulmonary diseases. Compared with the non-severe group, more patients in the severe group had a history of stroke (0% vs. 5.6%) and coronary heart disease (1.6% vs. 11.1%), suggesting that patients with

Table 1. Demographic characteristics of the 81 patients with COVID-19

Clinical characteristics and symptoms	All patients (n = 81)	Disease severity	
		Non-severe (n = 63)	Severe (n = 18)
Age, median (IQR) - years	58 (50.00-68.50)	57 (49.00-67.00)	68.5 (56.25-77.25)
Age groups - No., %			
< 65	51 (63.0)	45 (71.4)	6 (33.3)
≥ 65	30 (37.0)	18 (28.6)	12 (66.7)
Gender - No., %			
Male	38 (46.9)	26 (41.3)	12 (66.7)
Female	43 (53.1)	37 (58.7)	6 (33.3)
Symptoms or signs - No., %			
Fever on admission	59 (72.8)	46 (73.0)	12 (72.2)
Chest distress	22 (27.2)	19 (30.2)	3 (16.7)
Fatigue	30 (37.0)	23 (36.5)	7 (38.9)
Headache or dizziness	12 (14.8)	9 (14.3)	3 (16.7)
Respiratory symptoms	57 (70.4)	45 (71.4)	12 (66.7)
Gastrointestinal symptoms	22 (27.2)	17 (27.0)	5 (27.8)
Dark urine	16 (19.8)	11 (17.5)	5 (27.8)
Coexisting disorders - No., %			
Any	42 (51.9)	30 (47.6)	12 (66.7)
Hypertension	23 (28.4)	17 (27.0)	6 (33.3)
Diabetes	15 (18.5)	12 (19.0)	3 (16.7)
Coronary heart disease	3 (3.7)	1 (1.6)	2 (11.1)
Stroke	1 (1.2)	0 (0.0)	1 (5.6)
Cancer*	6 (7.4)	4 (6.3)	2 (11.1)
Chronic pulmonary disease [#]	7 (8.6)	5 (7.9)	2 (11.1)

Data are presented as medians [interquartile ranges (IQR)] and n/N (%). *Cancers referred to any malignancy. All cases had stable disease.

[#]Chronic pulmonary disease includes tuberculosis, chronic obstructive pulmonary disease, and bronchiectasis. All cases were stable and no obvious bacterial infections.

cardiovascular and cerebrovascular diseases are more likely to develop severe illness. The above findings are consistent with previous studies^[17-19].

Olfactory and gustatory dysfunctions in COVID-19

Table 2 summarizes the incidence of olfactory dysfunction (OD) and gustatory dysfunction (GD) when the patients were admitted to the hospital. On admission, 30.9% of all patients reported an altered sense of smell or taste. Overall, 13.6% and 25.9% reported OD and GD, respectively, while 8.6% reported both OD and GD. The incidence of OD in the severe group (11.1%) was not significantly different from that in the non-severe group (14.3%). Compared with the non-severe group, the incidence of GD was lower in the severe group (28.6% vs. 16.7%). However, OD and GD do not seem to be related to the severity of the disease (Fisher's exact test, $P < 0.05$).

The factors related to OD and GD are summarized in Table 3. We considered general conditions, symptoms, comorbidities, and laboratory tests and calculated the Pearson correlation coefficient. Some factors were positively correlated with OD, including headache or dizziness ($r = 0.342$, $P = 0.002$), dark urine ($r = 0.256$, $P = 0.021$), IgM titer ($r = 0.305$, $P = 0.01$), and diabetes ($r = 0.275$, $P = 0.013$). These factors were also positively correlated to GD (all $r > 0.2$, $P < 0.05$). The IgG titers were not significantly related to patients with OD ($r = 0.083$, $P = 0.496$) or GD ($r = 0.164$, $P = 0.176$). However, IgG titers were associated with the

Table 2. The incidence of olfactory and gustatory dysfunctions in patients with different disease severity

Disease severity	Olfactory dysfunction				Gustatory dysfunction				OD and GD	Any
	Total	Mild	Moderate	Severe	Total	Mild	Moderate	Severe		
Non-severe - n/%	9 (14.3)*	5 (7.9)	1 (1.6)	3 (4.8)	18 (28.6)*	8 (12.7)	7 (11.1)	3 (4.8)	5 (7.9)*	22 (34.9)*
Severe	2 (11.1)	0 (0.0)	1 (5.6)	1 (5.6)	3 (16.7)	1 (5.6)	1 (5.6)	1 (5.6)	2 (11.1)	3 (16.7)
All patients	11 (13.6)	5 (6.2)	2 (2.5)	4 (4.9)	21 (25.9)	9 (11.1)	8 (9.9)	4 (4.9)	7 (8.6)	25 (30.9)

*Compared with the severe group, $P > 0.05$. OD: Olfactory dysfunction; GD: gustatory dysfunction.

Table 3. Factors associated with olfactory and gustatory dysfunctions in COVID-19

Factors	OD		GD		OD or GD	
	r	P value	r	P value	r	P value
Headache or dizziness	0.342	0.002	0.308	0.005	0.248	0.026
Dark urine	0.256	0.021	0.273	0.014	0.273	0.014
IgM	0.305	0.01	0.238	0.046	0.251	0.035
IgG	0.083	0.496	0.164	0.176	0.269	0.024
Diabetes	0.275	0.013	0.298	0.007	0.301	0.006

OD: Olfactory dysfunction; GD: gustatory dysfunction.

incidence of patients with either OD or GD ($r = 0.269$, $P < 0.05$).

The association between various factors and the risk of olfactory and gustatory dysfunction in patients with COVID-19 is shown in Table 4. In the multivariate adjusted logistic regression model, patients with headache and dizziness (adjusted OR = 7.382, 95%CI: 1.028-53.022) were more likely to have olfactory dysfunction than those without headache and dizziness. Patients with diabetes had a higher risk of olfactory dysfunction compared to those without diabetes (adjusted OR = 6.871, 95%CI: 1.096-43.086). In the logistic regression analysis of gustatory dysfunctions, patients with high IgM titers had a higher risk of developing gustatory dysfunctions compared to those with low IgM titers (adjusted OR = 1.003, 95%CI: 1.000-1.006). In addition, patients with diabetes had a higher risk of developing GD compared to patients without diabetes (adjusted OR = 4.335, 95%CI: 1.098-17.287). Similarly, when OD and GD were treated as one symptom, patients with diabetes (adjusted OR = 7.963, 95%CI: 1.557-37.515) and patients with higher IgM titers (adjusted OR = 1.006, 95%CI: 1.001-1.010) had a higher probability of developing this symptom.

Table 5 describes the incidence of OD and GD one month after the patients were discharged from the hospital. Six patients had residual GD, and eleven patients reported OD. Of patients with OD or GD during hospitalization, 48.0% still complained of OD or GD one month after discharge. In patients with no OD or GD during hospitalization, 3.6% developed OD or GD after discharge. Continued follow-up of these patients with OD or GD showed that, in 60% (6 of 10) of the cases with OD and 20% (1 of 5) of the cases with GD, the symptoms lasted for at least two years; in 20% (2 of 10) of the cases with OD, symptoms aggravated. One OD patient and one GD patient lost in follow-up were excluded. We also analyzed the relationship between treatment and the incidences of olfactory and urinary dysfunction after discharge [Supplementary Table 1] and found that treatments had no effect on the clinical outcomes of OD and GD after patients were discharged, suggesting that the OD and GD are most likely primary symptoms caused by SARS-CoV-2, rather than the side effects of drugs.

DISCUSSION

This retrospective descriptive study included 81 confirmed cases of COVID-19 and focused on the OD and

Table 4. Multivariate adjusted logistic regression of olfactory and gustatory dysfunctions

Variables	β	SE	Wald	Adjusted OR	95%CI	P
OD						
Headache or dizziness	1.999	1.006	3.949	7.382	1.028-53.022	0.047
Dark urine	0.656	1.040	0.397	1.927	0.251-14.796	0.528
IgM	0.002	0.002	1.867	1.002	0.999-1.006	0.172
IgG	-0.001	0.898	3.671	5.591	0.961-32.515	0.055
Diabetes	1.927	0.937	4.234	6.871	1.096-43.086	0.040
Constant	-3.258	0.759	18.412			
GD						
Headache or dizziness	1.358	0.789	2.963	3.890	0.828-18.263	0.085
Dark urine	0.432	0.806	0.287	1.540	0.317-7.472	0.592
IgM	0.003	0.002	4.058	1.003	1.000-1.006	0.044
IgG	0.001	0.002	0.480	1.001	0.998-1.004	0.488
Diabetes	1.471	0.703	4.387	4.335	1.098-17.287	0.036
Constant	-2.098	0.531	15.591			
OD or GD						
Headache or dizziness	1.297	0.841	2.387	3.659	0.704-19.026	0.123
Dark urine	0.338	0.868	0.151	1.402	0.256-7.684	0.697
IgM	0.006	0.002	6.961	1.006	1.001-1.010	0.008
IgG	0.003	0.002	2.526	1.003	0.999-1.007	0.112
Diabetes	2.040	0.714	6.369	7.963	1.557-37.515	0.012
Constant	-2.479	0.589	17.727			

OD: Olfactory dysfunction; GD: gustatory dysfunction.

Table 5. The clinical outcomes of olfactory and gustatory dysfunctions in COVID-19 patients

During hospitalization	After discharged*			
	OD (n = 11)	GD (n = 6)	OD and GD (n = 3)	OD or GD (n = 14)
None - n/%	2 (3.6)	0 (0.0)	0 (0.0)	2 (3.6)
OD	9 (81.8)	3 (27.3)	3 (27.3)	9 (81.8)
GD	6 (28.6)	6 (28.6)	3 (14.3)	9 (42.9)
OD and GD	6 (85.7)	3 (42.9)	3 (42.9)	6 (85.7)
Any	9 (36.0)	6 (24.0)	3 (12.0)	12 (48.0)

*All enrolled patients' chest CT scanning signs disappeared and permitted discharge. OD: Olfactory dysfunction; GD: gustatory dysfunction.

GD in COVID-19 patients in Wuhan. On admission or during hospitalization, patients self-reporting OD, GD, and both accounted for 13.6%, 25.9%, and 8.6% of all subjects, respectively. The incidence of OD and GD in the severe cases was not significantly different from the non-severe cases, suggesting OD and GD are not related to the severity of the disease. In addition, 48.0% of patients had persistent OD or GD after the other clinical manifestations of COVID-19 disappeared. Continued follow-up showed that, in 20% of the cases with OD, symptoms even aggravated two years after infection.

There are currently few studies on the incidence of OD and GD in patients with COVID-19^[20]. One study found that 5.1% of COVID-19 patients reported hyposmia^[21]. Compared with this study, we found a higher rate of hyposmia (13.6%). A multicenter European study reported that 85.6% and 88.0% of patients suffer from olfactory and gustatory dysfunctions, respectively^[22]; this result indicates that, compared with Asian

patients, gustatory dysfunctions are more common in European patients. Notably, a recent study found that the prevalence of smell and taste dysfunction in COVID-19 patients has dropped significantly with the advent of the Omicron variant^[23]. Thus, the different strains of the virus may have distinct effects on olfactory and gustatory dysfunction.

The reasons for the different incidences of OD and GD may be as follows: (1) Quantitative measurement is more sensitive than self-report. Studies have shown that only 35% of patients are aware of their olfactory deficits^[24]. (2) The different affinity of the virus to different populations may lead to clinical differences among patients in different regions. A study showed that ACE2 mutations reduce the correlation between human ACE2 and SARS-CoV S-protein, thereby reducing the chance of infection^[25]. ACE2 polymorphisms and the differences in expression levels between Asian and European populations may explain the difference in olfactory dysfunction between Asian and European populations^[26]. (3) Different strains of the virus may cause different clinical manifestations.

Pearson correlation coefficient showed that headache or dizziness, dark urine, IgM, and diabetes all showed a positive correlation with OD and GD (all $P < 0.05$). We noted that chronic rhinitis and certain neurodegenerative diseases may directly lead to taste or smell disorders. In this study, such patients were excluded. Headache or dizziness are the most common symptoms of the nervous system. In a previous report, the incidence of headache and dizziness in COVID-19 patients were 13.1% and 16.8%, respectively^[21]. OD and GD are positively related to dizziness and headache. It has been reported that ACE2 is highly expressed in the nasal goblet and ciliated cells^[27]. The virus may infect the olfactory nerve early and cause OD before other neurological manifestations. OD and GD were also significantly related to dark urine. This may suggest that dark urine, similar to OD/GD, is a sign of early infection and damage to the urinary system. IgM is an early antibody produced by the immune system after infection. The interaction of infectivity, virulence, and immune response may explain the positive correlation between IgM and OD/GD. Consistent with other results, diabetes was significantly correlated with smell dysfunction^[28]. The infection might exacerbate potential nerve damage in diabetic patients. In addition, multivariate adjusted logistic analysis showed that patients with headache or dizziness and diabetics were at higher risk of developing olfactory dysfunction, while patients with high IgM titers and diabetes were at high risk of developing gustatory dysfunction, which is consistent with the results of the above Pearson correlation coefficient.

One month or more after discharge, some patients still suffered from OD and GD. A few patients who did not have OD or GD during hospitalization developed these symptoms after discharge. Considering the damage caused by the virus to the olfactory or gustatory nerve, this may be one of the possible sequelae of COVID-19 patients. It is worth noting that SARS-CoV can be detected 60-66 h after infection and is most abundant in the olfactory bulb^[29]. While the infection routes of SARS-CoV-2 and SARS-CoV are similar, OD may be an early symptom of COVID-19. Early screening of people with OD and GD, early detection of virus infections, and early isolation of COVID-19 patients can help prevent the spread of COVID-19.

In conclusion, we found that OD and GD are common symptoms of COVID-19. They appear early during the disease and may last from one month to even two years. Headache or dizziness, IgM titers, and diabetes are correlated with the occurrence of OD and GD. This retrospective study has some limitations. First, we collected self-reported data, which may lead to information bias. Second, the sample size was relatively small and geographically limited. We only recruited 81 confirmed cases of COVID-19 in our database since COVID-19 infectious was effectively controlled in Wuhan in early 2020. The cases reported here represent what happened to the first batch of patients in China. Third, grading bias might exist since we did not perform objective smell and taste tests. Fourth, as patients with other infectious diseases were not included

in the control group, it could not be determined whether COVID-19 had a more severe and longer-lasting impact on taste and smell than other infectious diseases.

DECLARATIONS

Authors' contributions

Full access to all of the data in the study, take responsibility for the integrity of the data and the accuracy of the data analysis, and wrote the manuscript: Zou L

The acquisition of data: Yu T, Zhang Y

Reviewed literature and gave technical support: Dai L

Conceived and supervised the project: Zhang Z, Zhang Z

All authors read and approved the submission of the manuscript.

Availability of data and materials

All analyzed data of this study are included within the article.

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Conflicts of interest

All authors declared that there are no conflicts of interest.

Ethical approval and consent to participate

This study was approved by the Hospital Ethics Committee of the Renmin Hospital of Wuhan University [WDRY2020-K136].

Consent for publication

All patients agreed to participate in this study.

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Review

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Electroneurography abnormality in Parkinson's disease: a potential biomarker to help diagnosis

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Abstract

Parkinson's disease (PD) is a common neurodegenerative disease, pathologically characterized by the progressive degeneration of dopaminergic neurons in the substantia nigra. Although various biomarkers and imaging criteria for PD have been established, objective and reliable evaluation methods are still lacking. Electroneurography, as an objective measurement of evoked compound muscle action potentials, is used to assess the integrity of the peripheral nerve and is important in the diagnosis and differential diagnosis of PD with neuromuscular injury. Moreover, it provides references for the evaluation and quantification of the motor function in PD. Here, we summarize recent advances in clinical research of electroneurography in PD, including the peripheral nerve conduction velocity, needle electromyography, surface electromyography, and motion unit number estimation. The potential values of electroneurography in PD diagnosis are also involved.

Keywords: Parkinson's disease, electroneurography, needle electromyography, surface electromyography, motor unit number estimation



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INTRODUCTION

Parkinson's disease (PD) is a common neurodegenerative disease in middle-aged and elderly people. Currently, it is believed that the main pathological change of PD is the progressive degeneration of dopaminergic neurons in the substantia nigra. The prevalence rate of people over 65 years old in China is 1.7%, and the incidence increases with aging^[1]. PD is clinically characterized by motor deficits including rest tremor, bradykinesia, myotonia, and postural balance disorders, as well as non-motor symptoms such as sensory disturbances, autonomic dysfunction, and mental and cognitive disorders. PD diagnosis is primarily based on history and physical examination^[2]. In the early stage of the disease, there is still a lack of objective or reliable diagnostic measurements. Although several auxiliary examinations have been established including olfactory test^[3], transcranial B-mode sonography of substantia nigra^[4], positron emission tomography with 18-fluorodeoxyglucose examination^[5], and autonomic nervous function examination such as ventricular variation rate detection^[6], these methods lack sensitivity and specificity in the early stage of the disease. Meanwhile, with recent advances in the understanding of the non-motor symptoms of PD, increasing lines of evidence show that peripheral neuropathy (PN) caused by neuroinflammation, oxidative stress, environmental toxins, and apoptosis may play an important role in the pathophysiological process of PD. Axonal or myelin sheath injury caused by these factors often manifests as peripheral nerve conduction velocity, including the prolonged latency, decreased compound muscle action potential amplitude, altered conduction velocity, conduction block, or waveform dispersion.

As an objective examination method, electroneurography has been widely used in the diagnosis and differential diagnosis of PN, muscular diseases, neuromuscular junction, and other diseases. This article reviews the research progress in the clinical application of electroneurography in PD.

THE MECHANISM OF PERIPHERAL NEUROPATHY IN PD

Peripheral nerve conduction velocity (NCV) examination includes motor nerve conduction velocity, sensory nerve conduction velocity, and F-wave, which can objectively reflect the integrity of myelin sheath. According to the traditional view, PD lacks epidemiological correlation with PN, and it is limited to patients with rare gene mutations such as PARK2, FMR1, ATXN3, and ATP13A2^[7] or mitochondrial disease. In recent years, it has been demonstrated that PN can also be found in idiopathic PD. Damage affecting medium to large fibers was found in PD in lower limbs, which can be confirmed by NCV as a new clinical fact, and the most common nerve found was the superficial fibular nerve, followed by the sural^[8].

Axonopathy in PD, as shown by the decreased amplitude of sural nerve^[9,10], may also be due to long-term treatment with levodopa. Levodopa is methylated by catechol-O-methyltransferase to produce S-adenosyl homocysteine in central and peripheral tissues, which is immediately cleaved to form homocysteine (HCY). Folate and vitamin B12 as cofactors are required to convert HCY back to S-adenosylmethionine (SMA). Therefore, with the increase of levodopa, vitamin B12 consumption increases, and PN associated with vitamin B12 deficiency occurs^[11]. On the other hand, HCY induces inflammatory responses by increasing brittleness to mitochondrial toxins and increasing free radicals, impinging on DNA repair mechanisms^[12,13] and RNA methylation^[14], leading to terminal degeneration of sural nerve axon. The incidence rate of large fibrous lesions in PD is 16.3%, and in small fibers, confirmed by biopsy is 56.9%^[15]. However, the literature does not exclude the possible influence of other factors such as diabetes and age-related factors on peripheral nerve damage. Levodopa is not the initial factor of PD with peripheral neuropathy, but the treatment exacerbates the progression of neuropathy.

A third mechanism of the causal relationship between PN and PD is the massive accumulation of α -synuclein in peripheral nerve terminal and damaged axon transport, resulting in small fibers^[16]. Pathological changes have also been demonstrated in animal models, i.e., the spreading of α -synuclein is associated with sensory nerve degeneration and defective nociception. α -synuclein aggregates in the afferent sensory system, including sensory neurons, axons from the dorsal roots, and spinal dorsal horn neurons, and can reduce NCV through the ultrastructural damaged small and medium myelinated fibers. Meanwhile, α -synuclein, immunoreactive in lamina I of the dorsal horn, modulates pain processing, which is related to the pain of PD^[17,18]. PN is a frequent yet underestimated feature of PD. PD patients with PN are older, have a higher levodopa dose, have worse axial motor features, and are independently associated with cognitive impairment. NCV examination can provide an objective diagnostic basis for PD with peripheral nerve damage. PN may be an independent peripheral marker of PD, indicating that there is a long course of disease accompanied by worse axial motion and impairment^[19].

NEEDLE ELECTROMYOGRAPHY

Needle electromyography (NEMG) is a method to record various electrical characteristics of muscle resting or voluntary contraction by using a concentric needle electrode to determine the localization and type of neuromuscular disease. NEMG inspection is conducted in three parts. The first is the spontaneous activity in the muscle resting state. The second is the contraction [motor unit action potential (MUAP)], which is the sum of synchronous discharges of a group of muscle fibers dominated by one anterior horn cell. The third is the electrical activity during vigorous muscle contraction (recruitment pattern). NEMG is often used in PD with camptocormia and anal electromyography.

The NEMG in PD with camptocormia

Abnormal postures are common in PD, mainly including camptocormia, antecollis, Pisa syndrome, scoliosis, and striatal deformity^[20]. According to epidemiological statistics, the incidence of PD with camptocormia is about 22.5%, also with multiple system atrophy (MSA), dementia with Lewy bodies, frontotemporal lobar degeneration, progressive supranuclear palsy, and Alzheimer's disease^[21]. While the exact mechanism underlying the pathogenesis of camptocormia is unknown, it has been recognized that both central and peripheral mechanisms may be involved, including abnormal motor function of basal ganglia-thalamic-cortical circuits^[22], sensory afferent dysfunction, muscle and joint proprioception abnormalities^[23], acetylcholine-dopamine transmitter imbalance^[24], and inflammatory infiltration in paraspinal muscles. The possible pathogenesis can be divided into the following four types^[25]: those caused by the progression of PD, manifested in the form of PD with dystonia, secondary to paraspinal muscle myogenic diseases, and antipsychotics caused by treatment.

Myopathy, myositis, neurogenic changes, and dystonia can be seen in NEMG in PD with camptocormia^[26]. NEMG of paravertebral muscles can suggest myopathy-like manifestations, with a narrow duration of multiphase potential and low amplitude, or accompanied by a small amount of spontaneous potential^[27], as well as dorsal and ventral trunk muscles (mainly about rectus abdominis, iliopsoas, internal oblique, and external oblique muscles)^[28]. Distal and proximal limb muscles may also be involved^[29].

In some patients, burst discharges were observed in the responsible muscles under electromyography, which usually suggests pathogenesis caused by dystonia. The camptocormia in PD is aggravated when walking or exercising and relieved by standing against a wall. Similar to the "sensory trick" of dystonia, camptocormia can be caused by the strong contraction of abnormal muscles^[30]. NEMG can be used to guide the treatment of botulinum toxin A when the pathogenesis suggests dystonia to identify the target muscles with the more active potentials. When botulinum toxin A was injected into rectus abdominis muscle and external oblique

muscle for treatment, the abnormal posture and abdominal pain were significantly improved 7-8 weeks after injection^[31,32].

The external anal sphincter electromyography in PD

External anal sphincter electromyography (EAS-EMG) is widely used in the differential diagnosis of neurological dysfunction of lumbosacral segment related diseases, such as lumbosacral radiculopathy, tethered cord syndrome, neurodegenerative diseases, and intraoperative monitoring of lumbosacral related operations. In the 1960s, Graham and Oppenheimer^[33] proposed that Onuf's nucleus, located in 1/3 of the ventral gray matter of S2-S3 sacral medullary, is a motor nucleus which sends out the pudendal nerve to innervate the external urethral sphincter, anal sphincter, and pelvic floor striated muscle.

Both PD and MSA are associated with the impairment of autonomic nerve function, and their pathological mechanisms are different. Since MSA may be associated with the motor neuron degeneration of Onuf's nucleus, EAS-EMG (including duration, phase, and satellite potential) could efficiently make a differentiation between MSA and PD with a sensitivity of 100% and specificity of 86%^[34]. The prolonged motor unit potentials (MUPs) duration and polyphasic MUPs usually indicate collateral reinnervation, which can be seen in early PD and MSA^[35]. Compared with PD, MSA is mainly characterized by a more prolonged mean duration and decreased amplitude during vigorous contraction^[36]. Most studies show that the abnormal rate of MSA is about 70%; muscular denervation such as fibrillation potential, positive sharp wave, and the duration of the action potential are widened; and the amplitude and polyphase wave are increased, as can be seen under the electromyography^[37].

Based on EAS-EMG, using the test of the bulbocavernosus reflex (BCR) by stimulating the pudendal nerve can also reflect the degree of damage to the autonomic nerve. The amplitude of BCR in PD is lower than in MSA. It is confirmed that PD can be accompanied with functional impairment of the autonomic nervous, but the degree is less than in MSA. Patients with MSA have a longer latent period and higher amplitude. In some cases, BCR can be used to distinguish MSA from PD^[38]. However, it is well known that the amplitude is greatly affected by the individual; thus, latency is more accurate in reflecting the degree of autonomic nerve damage, which is not mentioned in the literature above.

APPLICATION OF SURFACE ELECTROMYOGRAPHY IN PD

Surface electromyography (sEMG) is a non-invasive method to record the movement of motor units through the skin surface with electrodes^[39]. The characteristics of the signals mainly include linear (amplitude and frequency) and nonlinear (complexity and orderly) indices. It is usually used to evaluate the activity and quantify the function of neuromuscular objectively. Motor neurons are activated when muscles contract autonomously or receive an external stimulus; all recruited motor units produce continuous MUAP, which is transmitted through fat and subcutaneous tissues and finally recorded by surface electrodes^[40]. Many researchers have used sEMG technology for gait disorder evaluation, fatigue status detection, tremor analysis, central motor control, and acoustic kinematics in PD.

Application of sEMG in the assessment of gait disorder in PD

Gait disorder is a common feature of PD with the main characteristics of reduced shuffling, reduced step length and speed, difficult starting, increased rigidity, and freezing. It is one of the important factors of falls in PD. Some gait disturbances, such as bradykinesia and rigidity, have proven to be due to the decreased output of cortical motor caused by the imbalance of basal ganglia loop or supplementary motor area, as well as the changes in cortical and subcortical connectivity^[41]. Some abnormal gait with instability is not only caused by the central factor but also peripheral factors, such as the greater contraction of agonist/antagonist

muscle groups and less effective recruitment of individual muscles^[42]. sEMG can describe the walking characteristics of the patient to evaluate the motor function and formulate the next treatment and rehabilitation plan^[43] by analyzing the way different muscles in the lower body are recruited. Gait analysis by sEMG should include the phase of stance and swing, two-limb support, and one-limb support^[44]. Anterior tibial (TA) is the most frequently assessed muscle as it can receive more projections from the cortex with other limb muscles. Patients with PD always have reduced TA activity during the stance phase^[45]. Some patients with freezing have a higher activity of bilateral TA during the swing phase^[46], which reflects the impairment in motor control with the step length. The activity of medial gastrocnemius (MG) in PD is lower than in healthy controls, suggesting the defective function of proprioceptive^[47]. MG plays an important role in supporting the body in the vertical direction and stabilizing the ankle joint to maintain the balance of walking speed and posture^[48]. The asymmetric activities of TA-MG in PD are higher, which is an early feature of PD, as they have reduced muscle activity^[49]. The increased EMG variability usually indicates a high risk of falls in PD^[50], and it is an important trigger factor with frozen gait. Different walking modes, such as straight walk, U-turn, and point turn, require high energy consumption and therefore limit walking speed and flexibility^[51]. Kugler *et al.* used a classification algorithm based on data collected from sEMG and accelerometers for step segmentation^[52]. The proposed step detection method reached 98.9% sensitivity and 99.3% specificity for the classification accuracy of distinguishing between PD and healthy subjects. The reduced step length seems to be a specific feature of PD gait that can be detected early in the disease and can be a useful marker of disease progression. Co-activation is another feature of gait in PD, which can stabilize the joints and increase rigidity^[53]. Interventions for gait disorders must modify the activities and patterns of individual muscles, changing the coordinated contraction on the basis of gait impairment in PD.

Fatigue evaluation of sEMG in PD

Patients with PD often have unexplained fatigue as a sense of overwhelming tiredness, which is different from apathy, depression, dementia, or sleepiness, and it has two forms: subjective fatigue and objective fatigue^[54]. The incidence rate ranges from 33% to 50%^[55]. Some studies have demonstrated that this figure is as high as 70%. Fatigue can be observed in some but not all patients with good motor function, as well as in patients with depression, sleep disturbance, and autonomic symptoms^[56]. Basal ganglia dysfunction, such as the extrastriatal dopaminergic projections or frontal striato-thalamo-cortical loops, has been suggested as the pathophysiology, and the reduced blood perfusion in the frontal lobes can also be a contributor to fatigue^[57]. Some potential biomarkers such as C-reactive protein, cytokines, and interleukin-6 may be correlated to fatigue. Other researchers have suggested pathophysiological mechanisms including dysfunction of the hypothalamic-pituitary-adrenal system^[58]. The demographic characteristics indicate that the patients with fatigue are 1.4 years older on average, more likely to be female, and have a longer disease duration, higher UPDRS motor scores and levodopa dose, worse cognitive performances, and depression^[59]. Fatigue is usually measured through scales with no specific biomarkers. The scales are suitable for subjective fatigue. Subjective fatigue and objective fatigue are distinct and independent concepts. sEMG can detect bioelectrical activities associated with muscle contraction and reveal neural changes in objective fatigue. It can be measured based on changes during sustained muscle contraction. Longer periods of vigorous contraction indicate that more activities are required to initiate movement, leading to bradykinesia^[60]. Some researchers have indicated that the sEMG pattern of fatigue in PD is not different from that in healthy controls^[61], which might be due to the impacts from patients with subjective fatigue. sEMG can be used as a viable clinical assessment tool to detect the muscle pattern of fatigue and identify the risk during the onset.

The potential mechanism in tremor with PD by sEMG

With the development of the sEMG technology, many researchers have recently used acceleration measurement combined with sEMG to explore the tremor pattern of PD. The diagnosis of PD is

challenging; to date, sEMG is the gold standard for the analysis and diagnosis of tremors^[62]. The movement during tremor and different burst modes of a pair of antagonistic muscles can be recorded by sEMG. The frequency of most pathological tremors is 4-8 Hz, some palatal tremors are below 4 Hz, and orthostatic tremors are 13-18 Hz^[63]. Tremor generators are often classified as central or peripheral in origin^[64]. An abnormal rhythm of tremor exists in the basal ganglia of PD, while the network of ET is in the lower olive nucleus. The tremor harmonics as a valuable biomarker for tremor is important^[65].

The rest tremor as a typical and unique feature of PD has been considered for a long time, but recent studies have shown that rest tremor is also present in 15% of patients with ET^[66], causing misdiagnosis. For tremor analysis based on sEMG, parkinsonian rest tremor differs from ET as the tremor amplitude is suppressed during initiation of voluntary and target movements^[67]. The suppressed rest tremor may have delayed re-emergence when a new posture is sustained (re-emergent tremor), and the rest tremor in PD is usually unilateral in onset with 4-6 Hz^[63]. The rest tremor suppression test has demonstrated a very good sensitivity of ≥ 0.92 and a specificity of ≥ 0.69 for identifying rest tremor of PD^[67].

Thirty-nine percent of CBD patients have tremors during the disease course, including rest tremor, postural tremor, and action tremor. They are similar to low amplitude action myoclonus and differ from the typical rest tremor of PD^[68].

Shaikh *et al.*^[69] applied sEMG to differentiate PD from drug-induced secondary parkinsonian syndrome (dopamine receptor blockers such as antiemetic drugs and antipsychotics and DRBA-induced parkinsonism). Intentional tremor amplitude increased significantly in DRBA-induced parkinsonism, and the amplitude of tremor in resting and postural is similar. It may be due to the irregularity in the oscillation waveforms and the disturbance of central oscillator output. Tremor analysis is particularly useful in the diagnosis of various tremors.

The application of sEMG in central motor control of PD

Determinism (%DET), sample entropy (SampEn), and intermuscular coherence provide estimates of the signals of sEMG structure as nonlinear patterns. Higher %DET, lower SampEn, and higher theta, alpha, and beta band intermuscular coherences are observed in PD, which is correlated with MDS-UPDRS scores^[70]. This may reflect the oscillatory synchronization between the central nervous system and motor neurons, resulting in the enhancement of motor unit synchronization. The enhanced theta (4-6 Hz) and alpha (8-12 Hz) band coherences contribute to PD tremor. The amplified alpha-band drive in PD is involved in reticulospinal pathways, cortex and corticospinal output to muscles, and cerebello-thalamo-cortical circuit^[71]. The above signal analysis process is complex and requires specific mathematical models or analysis software. Further research is needed to verify the reliability and validity of the analysis method.

The use of sEMG in acoustic kinematics

The output of speech includes the cognitive, neuromotor cortex, neuromuscular, and musculoskeletal components. Gómez *et al.*^[72] used sEMG to differentiate the speech kinematic behavior by absolute velocities (dynamic and acoustic) of the joint jaw-tongue in PD, and their amplitude distributions can be used as potential biomarkers. They showed the characteristics of the interaction between the orofacial muscles by sEMG, which suggest that the facial nucleus and perioral muscles are well-preserved in PD. The lower amplitude ratio is consistent with thalamocortical activation, is limited by motor cortex activity, and contributes to bradykinesia and hypokinesia of orofacial muscles^[73].

CLINICAL APPLICATION OF MOTOR UNIT NUMBER ESTIMATION IN PD

The axon of an anterior horn motor cell and all muscle fibers it dominates are called a motor unit. Motor unit number estimation (MUNE) is a non-invasive electrophysiological test for estimating the signs of degeneration of anterior horn cells. Mes found MUNE measurements were smaller in PD with different muscles, suggesting a slight degeneration^[74]. Changes in MUNE are also associated with age, due to the degeneration of motor neurons due to physiological aging. It was also confirmed in the model of rats^[75] that the threshold of MUAP was lower and recruited less. Neurofibre degeneration and Lewy body appeared in large- and medium-sized motor neurons of spinal cord anterior horn cells, while small motor neurons were not found.

Some studies showed the increased motor unit amplitude and duration, which means reinnervation of motor units in PD^[76]. Asymptomatic motor neuron degeneration may be one of the pathological manifestations of PD, and MUNE has a certain standard deviation which needs to be confirmed by further studies with other electroneurography techniques.

CONCLUSION

PD is a common neurodegenerative disease, the diagnosis of which is based on clinical criteria only due to the lack of biomarkers. Thus, it has become important to detect the disease early. NEMG is an invasive method and poorly accepted by patients, but EAS-EMG is sensitive and specific for PD and MSA. NEMG is necessary for the auxiliary diagnosis of PD with camptocormia and striatal deformity; it can also be used as a guide for the therapy with botulinum toxin A to identify the target muscles for more precise treatment. sEMG is non-invasive with significant value in PD gait analysis and tremor. Especially, tremor analysis technology is simple and reliable in separating PD from other diseases. It may be a choice to be included in the diagnosis criteria. The application of sEMG in central motor control and acoustic kinematics is not widely used in clinical practice, but it is irreplaceable in the research on the development of quantitative biomarkers such as optimized neuromuscular performance and motor neuron activity. Electroneurogram has localization significance in PD with the diagnosis of the damage in peripheral nerve and anterior horn cells, and it needs further research. Therefore, electroneurography plays a certain role in the clinical diagnosis of PD and needs large samples to be further confirmed.

DECLARATIONS

Authors' contributions

Prepared the first draft: Hu Y

Made substantial contributions to conception and revised manuscript: Song C

Gave critical comments: Liang Z

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All authors declared that there are no conflicts of interest.

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AUTHOR INSTRUCTIONS

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Manuscript Type	Definition	Abstract	Keywords	Main Text Structure
Original Article	An Original Article describes detailed results from novel research. All findings are extensively discussed.	Structured abstract including Aim, Methods, Results and Conclusion. No more than 250 words.	3-8 keywords	The main content should include four sections: Introduction, Methods, Results and Discussion.
Review	A Review paper summarizes the literature on previous studies. It usually does not present any new information on a subject.	Unstructured abstract. No more than 250 words.	3-8 keywords	The main text may consist of several sections with unfixed section titles. We suggest that the author includes an "Introduction" section at the beginning, several sections with unfixed titles in the middle part, and a "Conclusion" section in the end.
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Systematic Review	A Systematic Review collects and critically analyzes multiple research studies, using methods selected before one or more research questions are formulated, and then finding and analyzing related studies and answering those questions in a structured methodology.	Structured abstract including Aim, Methods, Results and Conclusion. No more than 250 words.	3-8 keywords	The main content should include four sections: Introduction, Methods, Results and Discussion.
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2.3 Manuscript Structure

2.3.1 Front Matter

2.3.1.1 Title

The title of the manuscript should be concise, specific and relevant, with no more than 16 words if possible. When gene or protein names are included, the abbreviated name rather than full name should be used.

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Methods should contain sufficient details to allow others to fully replicate the study. New methods and protocols should be described in detail while well-established methods can be briefly described or appropriately cited. Experimental participants selected, the drugs and chemicals used, the statistical methods taken, and the computer software used should be identified precisely. Statistical terms, abbreviations, and all symbols used should be defined clearly. Protocol documents for clinical trials, observational studies, and other non-laboratory investigations may be uploaded as supplementary materials.

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This section should discuss the implications of the findings in context of existing research and highlight limitations of the study. Future research directions may also be mentioned.

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It should state clearly the main conclusions and include the explanation of their relevance or importance to the field.

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Journal articles ahead of print	Odibo AO. Falling stillbirth and neonatal mortality rates in twin gestation: not a reason for complacency. <i>BJOG</i> 2018; Epub ahead of print [PMID: 30461178 DOI: 10.1111/1471-0528.15541]
Books	Sherlock S, Dooley J. Diseases of the liver and biliary system. 9th ed. Oxford: Blackwell Sci Pub; 1993. pp. 258-96.
Book chapters	Meltzer PS, Kallioniemi A, Trent JM. Chromosome alterations in human solid tumors. In: Vogelstein B, Kinzler KW, editors. The genetic basis of human cancer. New York: McGraw-Hill; 2002. pp. 93-113.
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Conference paper	Christensen S, Oppacher F. An analysis of Koza's computational effort statistic for genetic programming. In: Foster JA, Lutton E, Miller J, Ryan C, Tettamanzi AG, editors. Genetic programming. EuroGP 2002: Proceedings of the 5th European Conference on Genetic Programming; 2002 Apr 3-5; Kinsdale, Ireland. Berlin: Springer; 2002. pp. 182-91.
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Diagrams with describing words (including, flow chart, coordinate diagram, bar chart, line chart, and scatter diagram, *etc.*) should be editable in word, excel or powerpoint format. Non-English information should be avoided;

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Tables should be cited in numeric order and placed after the paragraph where it is first cited;

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General italic words like *vs.*, *et al.*, *etc.*, *in vivo*, *in vitro*; *t* test, *F* test, *U* test; related coefficient as *r*, sample number as *n*, and probability as *P*; names of genes; names of bacteria and biology species in Latin.

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Numbers appearing at the beginning of sentences should be expressed in English. When there are two or more numbers in a paragraph, they should be expressed as Arabic numerals; when there is only one number in a paragraph, number < 10 should be expressed in English and number > 10 should be expressed as Arabic numerals. 12345678 should be written as 12,345,678.

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8.1 Initial check

8.1.1 Initial manuscript check

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