### Review

# The Gut and Parkinson's Disease: Hype or Hope?

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**Abstract**. In the last two decades it has become clear that Parkinson's disease (PD) is associated with a plethora of gastrointestinal symptoms originating from functional and structural changes in the gut and its associated neural structures. This is of particular interest not only because such symptoms have a major impact on the quality of life of PD patients, but also since accumulating evidence suggests that in at least a subgroup of patients, these disturbances precede the motor symptoms and diagnosis of PD by years and may thus give important insights into the origin and pathogenesis of the disease. In this mini-review we attempt to concisely summarize the current knowledge after two decades of research on the gut-brain axis in PD. We focus on alpha-synuclein pathology, biomarkers, and the gut microbiota and envision the development and impact of these research areas for the two decades to come.

Keywords: Alpha-synuclein, prion, constipation, dysautonomia, microbiota, gut-brain-axis

## GASTROINTESTINAL PATHOLOGY AND ITS ROLE IN PARKINSON'S DISEASE ETIOLOGY

The first descriptions of alpha-synuclein (asyn) histopathology in the enteric nervous system (ENS) and peripheral autonomic ganglia date back as far as 1960 [1, 2], and several groups have now confirmed and extended these initial findings [3, 4]. By contrast to the *substantia nigra*, no overt neuronal loss is observed in the ENS in Parkinson's disease (PD) [5, 6]. Based on early pathology in the olfactory bulb and dorsal motor nucleus of the vagus, the dual-hit hypothesis posited by Braak and colleagues

proposes that initial misfolding and aggregation of asyn may occur in peripheral nerve terminals with subsequent centripetal spreading through the vagus nerve [7, 8]. This proposition gained support from two epidemiological studies showing that full truncal but not selective vagotomy seems to decrease the risk of PD [9, 10]. Also, in studies of archived tissue removed from patients up to 20 years prior to PD diagnosis, asyn pathology in the gut was reported more frequently than in matched controls [11, 12]. Several animal models, recently reviewed in detail [13], have shown cell-to-cell prion-like transmission and centripetal spreading of asyn [14, 15]. Such studies demonstrate the mechanistic plausibility of gut-to-brain transmission of asyn pathology. Nevertheless, this gut-to-brain transmission hypothesis is still widely debated and the arguments in favor or against the gut as the site of origin of PD were

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reviewed in two recent publications [16, 17]. The strongest argument against such a scenario comes from findings from the Arizona Parkinson's consortium group, which did not find a single case in which asyn pathology was present in the ENS but not in the CNS in their wide survey of 466 whole-body autopsy cases [18]. This whole-body survey does not exclude, however, that initial asyn aggregation may start in the gut, but such hypothetical initial gut pathology would have to be be highly localized, or alternatively consist of immature and potentially protease-sensitive microaggregates, which are difficult to detect by standard immunohistochemistry [16].

In the coming two decades, it is likely that major advances will be made in understanding the role of gastrointestinal asyn pathology in the etiology of PD. An important research goal is to elucidate the degree of similarity between pathophysiological processes in PD and those of true prion diseases. In variant Creutzfeldt-Jakob disease (CJD), there is little doubt about the occurrence of gut-to-brain propagation of prions [19]. The aggregates of disease-related prion protein in variant CJD and bovine spongiform encephalopathy initially form in peripheral lymphoid tissues and in the ENS and then spread through the autonomic nervous system [20, 21]. Importantly, the prion aggregates in peripheral organs of variant CJD show clear differences in glycoform ratios and protease resistance compared to prions in brain tissues [22, 23]. It has therefore been suggested that the long latency between peripheral lymphoid colonization and subsequent neuro-invasion is in part due to the need for selection of a neuro-invasive strain of prion [24]. Such mechanisms have not been considered in the context of PD, but may explain why asyn multimers or aggregates harvested in peripheral autonomic nervous tissues behave differently than those from the *substantia nigra* of PD patients [25, 26].

There is therefore an urgent need to better characterize the biochemical characteristics of both native and pathological asyn in the ENS (Fig. 1). This can best be achieved by performing a comprehensive inventory of synuclein forms present in the ENS from PD patients using two-dimensional electrophoretic analysis and mass spectroscopy or luminescent conjugated oligothiopenes, as such approaches have proven successful for characterizing the dominant pathological modification of synuclein and amyloids in the brain [27, 28]. Also, much work needs to be done within the field of animal PD models in the coming years. Although the concept of species barrier, i.e.

the inability of some prion strains to cause disease in other species, is well described in prion disease [29, 30], it has yet received little attention in PD. It has been shown, however, that sequence homology between asyn and the host protein is proportional to the rate of seeding initiation, e.g., human asyn fibrils efficiently seed monomeric human asyn, while seeding of mouse asyn is inefficient (reviewed in [31]). In addition, the importance of aging and other modulatory factors including intestinal hyperpermeability, concurrent inflammation, and microbiota alterations needs to be studied in detail before firm conclusions can be drawn (Fig. 1).

Prion pathology in variant CJD exhibit peripheral-to-central propagation, whereas the initial pathology in sporadic and familial CJD cases probably arises spontaneously within the brain. Thus, it is theoretically possible that synucleinopathies can similarly be divided into types either originating in the peripheral nervous system or arising spontaneously within the CNS [16]. Note that this does not necessarily imply that these synucleinopathies were contracted by exogenous agents, but can be explained by Braak's hypothesis that the nerve terminals of long, non-myelinated, hyperbranched axons are the most probable (but not only) place for the first occurrence of asyn aggregation [7].

One candidate for a peripheral-dominant subtype of PD may be those cases, in whom rapid eye movement sleep behavior disorder (RBD) presents during the prodromal phase. It has been shown that PD patients with RBD exhibit much higher frequencies of phosphorylated asyn pathology in the colon and in the skin compared to PD patients without RBD [32]. Also, idiopathic RBD patients exhibit marked pathology in the sympathetic and parasympathetic nervous system, but a relatively intact dopamine system [33]. In contrast, PD patients without RBD often have normal cardiac sympathetic innervation in early disease stages indicative of a sparing of the autonomic nervous system [34, 35]. An important research goal for the coming years will be to perform a thorough phenotypic characterization of PD subgroups using objective markers of neuronal dysfunction. It urgently needs to be clarified whether such distinct phenotypes can be explained by molecular characteristics of pathological asyn in the ENS and CNS (Fig. 1). An attractive hypothesis is that some specific post-translationally modified forms of asyn and/or assemblies (strains) could be associated with a more aggressive disease progression [36].

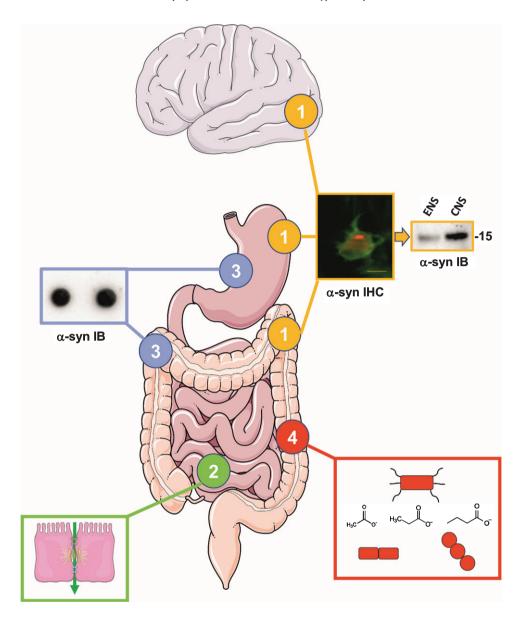


Fig. 1. The four most important issues that need to be addressed in the next 10 years regarding the gut in PD. (1) Alpha-synuclein deposits are observed in the ENS of PD patients (the microphotography shows phospho-alpha-synuclein immunostaining in the colonic myenteric plexus of a PD patient, scale bar 20 µM). However, it remains to be determined if the alpha-synuclein aggregates in the ENS are biochemically similar to the ones found in the brain as this might be critical in our understanding of the role of the gut in PD pathogenesis. The picture shows lysates from colonic biopsies (left lane, ENS) and brain samples (right lane, CNS) from two PD patients that have been analyzed by western blot using C-20R total asyn antibody. Using this simple approach, no difference is observed between the ENS and the CNS and there is therefore a critical need to perform a comprehensive inventory of synuclein forms present in the ENS from PD patients using proteomic approaches. Molecular weight in kDa is indicated on the right. (2) Triggering of initial alpha-synuclein aggregation in enteric nerve terminals through extrinsic factors could be facilitated by intestinal hyperpermeability. It remains to be definitely demonstrated that intestinal permeability is increased in PD. (3) Results of immunohistochemistry-based studies on alpha-synuclein deposits in the ENS of PD patients have provided conflicting results. There is therefore a critical need to develop alternative techniques to detect alpha-synuclein aggregates in the gut. The inset illustrates a dot blot from gastrointestinal biopsy lysates of a control subject, stained with Syn-1 alpha-synuclein antibody. (4) Alterations of gut microbiota composition in PD have been shown in multiple cross-sectional studies from diverse populations. It will be crucial to determine the mechanisms that connect gut microbiota and PD in large multicenter studies of prodromal and de novo PD patients as well as animal models employing multiomics approaches. Eventually, the clinical diagnostic and therapeutic potential of the gut microbiota must be determined. The inset illustrates differently shaped bacteria and the molecular structures of the SCFAs butyrate, acetate and propionate. Drawings were modified from Servier Medical Art, licensed under a Creative Commons Attribution 3.0 Unported License

#### **BIOMARKERS**

The observation that asyn inclusions are found in ENS of the vast majority of PD patients has led to an increasing number of immunohistochemistry-based studies aimed at developing biomarkers for disease diagnosis and progression. However, the diversity of methodology between studies, especially regarding the immunohistochemical methods used, has led to conflicting results regarding the sensitivity and specificity of gastrointestinal biopsies for the detection of asyn deposits [37]. Although several studies showed a high sensitivity of formalin-fixed paraffin embedded (FFPE) gastrointestinal biopsies for the detection of asyn inclusions in PD patients [38–41], other reports have raised concerns regarding the specificity of this approach because asyn immunoreactivity was also observed in some healthy individuals [42-45]. This lack of specificity, which likely results from technical difficulties inherent in the use of FFPE samples for the detection of asyn [46], explains why FFPE gastrointestinal biopsies are not currently used in routine clinical practice as PD biomarker. Classical biochemical approaches, such as one- and two-dimensional electrophoresis did not prove to be more efficient than immunohistochemistry for detecting aggregated asyn in gastrointestinal biopsies [47]. As a whole, these disappointing results might be explained by the low amount of aggregated/pathological asyn that is usually found in the gastrointestinal tract and especially in gastrointestinal biopsies [48]. In order to overcome this limitation, future research could take benefit from ultrasensitive techniques such as protein misfolding cyclic amplification (PMCA) and real-time quakinginduced conversion (RT-QuIC) assays, which have been recently shown to efficiently amplify aggregated asyn in cerebrospinal fluid and in formalin-fixed archived tissue [49, 50]. In addition, there is growing evidence that pathological changes in the gut in PD are not limited to enteric neurons but also involve the enteric glial cells and the intestinal epithelial cells [51, 52], two cell types that largely outnumber enteric neurons and are easily captured by routine gastrointestinal biopsies. These observations support the hypothesis that this so-called 'neuro-glio-epithelial unit' [53] might constitute an unparalleled source of biomarkers in PD beyond the sole assessment of asyn deposits/aggregates. Possible strategies to identify new enteric markers of PD might include a joint transcriptomic and proteomic analysis of the biopsies along with the analysis of biopsy supernatant [54].

Finally, the characterization of gastrointestinal dysfunction in patients has mostly relied on scoring of subjective symptoms, which is necessarily unable to capture subclinical disease and often shows poor correlation with markers of objective dysfunction [55]. Accessible and affordable methods such as radio-opaque markers to assess gastrointestinal transit times will find more widespread use in future studies. Recently, <sup>11</sup>C-donepezil PET scans have demonstrated decreased cholinergic signal in the gastrointestinal tract of both manifest and prodromal PD populations [33, 56]. Also, once an asyn-specific PET ligand becomes available it may be possible to image not only the brain but also peripheral organs and investigate longitudinally the dual-hit hypothesis. Although expensive, such techniques will be critical to improve our understanding of PD etiopathogenesis and may have potential as progression markers in upcoming trials of disease-modification.

#### GUT MICROBIOTA

The human microbiota consists of bacteria, archaea, protists, fungi, their respective viruses, and human viruses. Most research has been carried out on the bacterial component of the microbiota [57]. The number of genes encoded in the gut metagenome is approximately 150 times larger than that of the human genome [58]. Evidence is accumulating that there is an intense bidirectional interaction between the gut, its microbiota, and the brain, frequently referred to as the microbiota-gut-brain-axis, which has important implications for brain health [59].

Alterations of the gut microbiota composition in PD have been revealed in multiple case-control studies from diverse populations. While there have been variations in the reported results, in PD patients an increased relative abundance of bacteria from the genera Akkermansia, Lactobacillus, and Bifidobacterium and decreased abundances of Prevotella, Faecalibacterium, and Blautia have been reproducibly shown [60-64]. For Prevotella such reduction has also been observed in RBD patients [64]. Based on the attributed functional properties of these bacteria, such alterations could affect gut barrier integrity, short-chain fatty acid (SCFA) production, and inflammation. This would be in line with reports of a leaky gut and reduced levels of SCFAs and lipopolysaccharide binding protein in PD patients [62, 65, 66]. However, it remains to be definitely proven that the gut is hyperpermeable in PD [52] (Fig. 1). Also with respect to SCFAs, in particular butyrate, the effects on PD pathology are not unequivocally established, since beneficial and harmful effects have been reported from PD animal and *in vitro* models [67–70]. An interesting link between gut microbiota and asyn pathology could be cross-seeding of amyloid pathology induced by bacterial amyloid proteins such as curli [71]. Few studies have investigated the nasal and oral microbiota in PD. While some alterations were found in the oral microbiota, two studies failed to find alterations in the nasal compartment [64, 72]. Future research on other non-bacterial microbes such as viruses, fungi, and archaea may provide also valuable insights.

So far, human microbiome studies in PD have been carried out exclusively in medicated patients, except for one study that included also idiopathic RBD patients [64]. While the PD associated microbiome alterations have been confirmed in medication adjusted analyses, confounding effects cannot be excluded and have in fact been shown for COMT inhibitors [60, 73]. Another potential confounder is colonic dysmotility, which may independently affect microbiota composition [60, 74]. Several microbiota studies assessed constipation symptoms using questionnaires and adjusted for reported constipation in their analyses. However, questionnaires may underestimate the prevalence of objective colonic dysfunction and thus adjustment based on these responses may be insufficient [55]. It is not known, whether the observed microbiome changes play a causal role in the development of gut pathology in PD or whether they are rather a consequence of altered gut function. Also for the observed correlations between microbiome composition and motor [60, 64] and non-motor symptoms [75, 76] causality has not been established in humans. However, observations that motor symptoms, neuroinflammation, asyn pathology, and gut motility can be modulated by manipulating the gut microbiota in transgenic asyn-overexpressing mice suggest that such causal influences are possible [68].

Whether gut microbiome alterations in PD are independent of medications and constipation can be addressed by comparing microbiota and objective assessments of gut function between healthy subjects and drug naïve prodromal and/or *de novo* patients. To establish what mechanisms link microbiota alterations and PD, such studies should employ a multiomics approach involving metagenomic, metatranscriptomic, and metabolomics analyses in combination with an assessment of host factors such

as gut biopsies, permeability studies, cytokine levels and host genotype (Fig. 1). For such studies to succeed, multicenter consortia should collaborate to ensure sufficient cohort sizes and standardized methodology. Longitudinal study designs will eventually enable us to study the temporal relationship between microbiota changes, disease stages, and phenotypes. While such undertakings will be longlasting and very cost-intensive, further work in animal models may be helpful to narrow down the spectrum of potentially relevant pathways that connect microbiota and PD disease mechanisms and to test interventions.

Taken together there is good reason to envision that gut microbiota may have important implications in the future diagnostic and therapeutic landscape of PD. Analysis of the composition and/or functional aspects of the microbiome combined with clinical and genetic host factors could become a crucial element of patient phenotyping allowing more individualized treatments for PD. Using such biomarkers to select suitable patients for specifically tailored therapeutic interventions should improve odds for successful clinical trials of symptomatic or disease modifying treatments. Based on encouraging observations from RBD subjects [64], such approaches could find applications already in the prodromal phase of PD. If disease specific microbiota profiles could be established, these would be a valuable tool for the differential diagnosis of parkinsonism [77].

Therapeutic applications based on the gut microbiome are possible through a range of approaches. These include dietary interventions, application of beneficial bacteria (probiotics), substances that promote growth of beneficial bacteria (prebiotics), substances that eliminate harmful bacteria (antibiotics), and transfer of bacterial ecosystems (fecal microbiota transplantation). Furthermore, a detailed understanding of microbiome-host-interactions in PD could identify new pathways that could be targeted using more traditional pharmacological approaches. Importantly, microbiome based therapies could be effective independently of the causative role of the microbiome in PD pathogenesis. This is because the microbiome is a key regulator of neuroinflammation and could affect disease progression through modulation of microglia activity [78]. Furthermore, gut microbiota affect gut permeability and the local environment in the gut lumen and mucosa which could have an implication for the impact of environmental toxins and other PD initiating or perpetuating factors such as protein aggregation. Also,

beneficial symptomatic effects may be achieved, as has been demonstrated in a trial showing improvement of constipation in PD patients receiving a synbiotic formulation, i.e., probiotics in combination with prebiotic fiber [79]. Finally, the microbiome may also play a role in the pharmacologic treatment of PD, as has been demonstrated for *Helicobacter pylori* and small intestinal bacterial overgrowth [80]. This suggests that benefits from traditional treatments of PD, such as levodopa, might be enhanced by microbiome based interventions.

#### CONCLUSION

In his "Essay on the shaking palsy" [81], James Parkinson noted that "Although unable to trace the connection by which a disordered state of the stomach and bowels may induce a morbid action in a part of the medulla spinalis [...] little hesitation need be employed before we determine on the probability of such occurrence." In line with his statement, our understanding and appreciation of the importance of the gut-brain connection in PD has grown rapidly in recent years. We are confident that the coming two decades of microbiome-gut-brain-axis research will see an even accelerated development in this area that will reshape our understanding of the pathogenesis of PD. While there is a hype, there is definitely also hope that this will translate into improved diagnostic and therapeutic approaches and eventually disease modifying treatments for PD patients.

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#### CONFLICT OF INTEREST

FS is founder and CEO of NeuroInnovation Oy, has received consulting fees from LivaNova and Illumina Ventures, lecture fees from Abbvie and Zambon, travel support from Abbvie, LivaNova, and Zambon, is member of the scientific advisory boards of Axial

Biotherapeutics and LivaNova, and is listed as inventor on the patent applications FI20145492A, PCT/FI2015/050374, EP20150798909, and US15314240.

PD has nothing to disclose.

PB has nothing to disclose.

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