Experimental colitis drives enteric alpha-synuclein accumulation and Parkinson-like brain

pathology

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Abstract

Intraneuronal α-synuclein accumulation is key in Parkinson's disease (PD) pathogenesis. The

pathogenic process is suggested to begin in the enteric nervous system and propagate into the brain

already decades before diagnosis of PD. In some patients, colitis might play a critical role in this

process. Here we demonstrate that patients with inflammatory bowel disease exhibit α-synuclein

accumulation in the colon and that experimental colitis triggers α-synuclein accumulation in certain

enteric nerves of mice. The type and degree of experimental inflammation modulates the extent of

colonic α-synuclein accumulation and macrophage-related signaling limits this process. Remarkably,

experimental colitis at three months of age exacerbates the accumulation of aggregated phospho-

Serine 129 α-synuclein in the midbrain (including the substantia nigra), in 21- but not 9-month-old α-

synuclein transgenic mice. This is accompanied by loss of tyrosine hydroxylase-immunoreactive

nigral neurons. Our data suggest that intestinal inflammation might play a critical role in the initiation

and progression of PD.

Parkinson's disease (PD) is a progressively debilitating neurodegenerative disease affecting 1% of the

population above 60 years [1]. Typical symptoms are motor impairments including muscle rigidity,

tremor, and bradykinesia. Neuropathologically, PD is hallmarked by loss of dopaminergic neurons in

the substantia nigra (SN), a concomitant reduction of striatal dopaminergic signaling [2], and the

presence of intraneuronal inclusions called Lewy bodies and neurites [3]. Lewy pathology is enriched in

α-synuclein (αSyn), a presynaptic protein that tends to aggregate and become phosphorylated under

pathological conditions [2]. Rare point mutations in αSyn and gene multiplications also cause familial

forms of PD and related neurological conditions, and certain single nucleotide polymorphisms close

to the αSyn gene (SNCA) locus are associated with increased risk for sporadic PD [4]. These findings

make αSyn a focal point of biomarker and drug development programs for PD.

Several years before the first appearance of motor symptoms, many patients exhibit a variety of non-

motor symptoms including constipation, sleep disorder, depression, and hyposmia [5–7]. Indeed, co-

occurrence of some of these non-motor symptoms is coupled to elevated PD risk [8–11]. Constipation is

an important non-motor feature of prodromal PD, with 28-61% of patients having exhibited

gastrointestinal dysfunction for several years during the prodrome [7,10,12]. Notably, αSyn-

immunoreactive inclusions have been found in neurons of the submucosal plexus in people with PD

[3,13]. Taken together, this converging evidence suggests an early involvement of the enteric nervous

system (ENS) in the pathogenesis of PD. Already over a decade ago, Braak and colleagues

hypothesized that αSyn-immunoreactive inclusions first appear in the ENS and then gradually engage

the brainstem, including the dorsal motor nucleus of the vagus nerve and midbrain areas [3,13]. Several

studies in preclinical models have demonstrated that αSyn pathology in the gut is associated with the

development of αSyn pathology in the brain [14–18]. It will be critical to determine factors that regulate

αSyn accumulation in the ENS and to understand whether the process underlying αSyn accumulation

in the gut can also lead to αSyn pathology in the brain.

Inflammation can potentially trigger αSyn pathology in the ENS of the gut and in the brain. A recent

finding in children with gastrointestinal inflammation suggests an immune regulatory function of

αSyn [19]. Immune pathways are indeed activated in the brain and colon of PD cases [20,21]. Also, several

genes associated with an increased PD risk have an immune system-related function [22], and it was

recently proposed that PD heritability is not simply due to variation in brain-specific genes, but that

several cell types in different tissues might be involved [23]. Adding further genetic evidence supporting

that inflammation is involved in PD pathogenesis, a genome-wide association study identified

common genetic pathways linking PD and autoimmune disorders [24]. Most prominently, LRRK2, a

major genetic risk factor for PD also confers increased risk for developing inflammatory bowel

disease (IBD) [25] and is known to modulate the function of monocytes, macrophages and other immune

cells [26,27]. Intriguingly, IBD is associated with an increased risk for developing PD and specifically

blocking the TNF pathway reduces this risk [28–31]. This suggests that the intestinal immune

environment plays a role in triggering PD or facilitating the molecular events involved in the earliest

phases of the disease process [32].

Here we tested the hypothesis that intestinal inflammation (e.g. colitis) triggers accumulation of αSyn

in the ENS and the subsequent development of αSyn pathology in the brain. We discovered that

patients with IBD exhibited increased αSyn accumulation in the submucosa of the colon.

Experimental forms of colitis in wild type and αSyn transgenic mice demonstrated that the type and

degree of inflammation regulates the amount of αSyn accumulation in the colon. Macrophage-related

signaling limited the extent of αSyn immunoreactivity. When αSyn transgenic mice were exposed to

experimental colitis at 3 months of age and then were aged normally up to 9 or 21 months, the

accumulation of aggregated αSyn in midbrain, including the SN, was much exacerbated in the 21-

months age group, but not in the 9-months age group. These 21-month old mice also exhibited loss of

nigral tyrosine hydroxylase-immunoreactive neurons. Together, our data support a critical role for

intestinal inflammation in the initiation and progression of PD.

Results

IBD patients show αSyn accumulation in the ENS and local macrophages

Recent epidemiological data links inflammatory bowel disease (IBD) to an increased PD risk [28–30]. In

order to explore if IBD is associated with enteric αSyn accumulation we performed

immunohistochemistry for αSyn in cryo-sections from colonic biopsies of patients with ulcerative

colitis (UC, n = 11, mean age 31 years), Crohn's disease (CD, n = 11; mean age 35 years), and from

healthy subjects (HS, n = 8; mean age 51 years). We observed in eight UC cases various degrees of

αSyn accumulation, mostly in structures with the morphology of neurites (**Figure 1**). Interestingly, the

eight UC cases, and four patients with CD (images not shown) also showed marked intracellular αSyn

staining in many infiltrating monocytic cells. In contrast, only one HS showed a few cells

immunoreactive for αSyn (images not shown). This finding in human tissue suggests a potential role

of local inflammation in the development of enteric αSyn accumulation.

Experimental IBD exacerbates αSyn load in submucosal plexus of αSyn transgenic and wildtype

mice

During the process of further characterizing a (Thy1)-h[A30P]αSyn transgenic mouse line [33] we

detected human αSyn accumulation in all innervated organs that we analyzed (**Supplemental Figure**

1). This included the myenteric and submucosal plexuses of the ENS, where human αSyn co-localized

with peripherin, a specific marker for peripheral nerves (**Figure 2A)**. We observed an age-dependent

increase of baseline human αSyn inclusions (irregularly sized and shaped inclusion bodies detected by

human αSyn specific monoclonal antibody clone 211) in both plexuses between the ages of three and

twelve months (**Figure 2B**). We wanted to test whether IBD-related inflammation in the colon

exacerbates this local accumulation of αSyn acutely (e.g. within a few days or weeks) and how the

age of the αSyn transgenic mice influenced the outcome. Administration of dextran sulfate sodium

(DSS) in the drinking water in acute or chronic paradigms are well-established mouse colitis models

of IBD, exhibiting infiltration of leukocytes into the submucosa with various degrees of destruction of

the colonic mucosa and submucosa [34]. Due to awareness of the variability of the DSS model in

different genetic backgrounds of mice, we first tested DSS administration at different concentrations

and durations in the (Thy1)-h[A30P]αSyn transgenic mice (**Figure 2C**), and observed leukocyte

infiltration in a dose-dependent manner and which was similar at the age of 3 and 6 months (**Figure**

2D and 3A). In the acute paradigm with mice at the age of 3 months, 2.5%, but not 1%, DSS

triggered intracellular accumulation of αSyn in nerves of the submucosal plexus **(Figure 3A, B)**. In

the chronic DSS paradigm, which was done with mice at the age of 6 months, we observed a dose-

dependent increase of αSyn load in the submucosal plexus, but at a smaller magnitude than in the

younger mouse cohort **(Figure 3A)**. Wildtype mice also express endogenous αSyn in innervated

organs, but at much lower levels compared with the overexpressed human αSyn protein in the

heterozygous (Thy1)-h[A30P]αSyn transgenic mice (**Supplemental Figure 1**). To confirm that the

finding in (Thy1)-h[A30P]αSyn transgenic mice was independent of transgenic expression of human

αSyn, we applied acute and chronic (consistent dose) DSS paradigms also in wildtype mice. In both

treatment paradigms, we observed an elevated number of inclusion bodies of endogenous murine

αSyn in the submucosal plexus **(**detected by rodent cross-reactive αSyn-specific monoclonal antibody

Syn1/clone 42, **Figure 3C, D)**. A separate experiment also confirmed that the observed effects of DSS

could not be attributed to increased gene expression of murine or the transgenic human αSyn

(Supplemental Figure 2). Together, these results confirmed the validity of this experimental IBD

paradigm to test the effect of inflammation on αSyn accumulation in the ENS in wild type and

(Thy1)-h[A30P]αSyn transgenic mice. Because 3-month old (Thy1)-h[A30P]αSyn transgenic mice

provided more optimal conditions for visualization and quantification of αSyn inclusions in the ENS,

for the remainder of the study we focused on using this transgenic mouse model.

Colitis induced by peroral DSS but not by peritoneal administration of LPS aggravates αSyn

accumulation in colonic submucosal plexus of αSyn transgenic mice

In order to explore effects of different approaches to induce inflammation in or nearby the gut in

(Thy1)-h[A30P]αSyn transgenic mice, we compared the outcome of acute 5% DSS in drinking water

with acute 0.5 mg/kg intraperitoneal LPS administration (**Figure 2C and 4**). In order to maximize the

inflammatory response, we administered both DSS and LPS at high, but still tolerable, doses. At day

7, both agents had induced variable degrees of leukocyte infiltration in the submucosa of the colon

while a marked destruction of the mucosa was induced when giving only DSS (**Figure 2D**). As

before, the DSS-exposed mice presented with increased accumulation of αSyn in the ganglia of the

submucosal plexus (**Figure 4A**). In contrast, we detected no change in αSyn load in the myenteric

plexus, consistent with lack of leukocyte infiltration in this part of the colonic wall (**Figure 4B**).

Despite the high dose, LPS-induced inflammation did not increase αSyn accumulation in the colonic

nervous plexuses (**Figure 4C, D**). Notably, LPS and DSS resulted in a differential expression of

cytokines, and consistent with leukocyte recruitment, CCL2 was elevated in both (**Figure 4F, G**). In

the LPS paradigm, mRNA for IL-10 was markedly elevated, whereas DSS strongly increased IL-6

and also IL-1β but not IL-10. Together these results indicate that only certain types of local

inflammation increase the intracellular accumulation of αSyn in the colon.

Lack of Cx3cr1 signaling during DSS colitis aggravates αSyn load in the submucosal plexus of

αSyn transgenic mice

In both the IBD patients and the (Thy1)-h[A30P]αSyn transgenic mice that experienced acute DSS

colitis, we observed several αSyn-positive cells with a morphology consistent with them being

infiltrating leucocytes (**Figure 5**). In the mice, these infiltrating cells were positive for the

macrophage marker Iba-1 (**Figure 5C-D**). In order to explore further the role of

monocytes/macrophages in the accumulation of αSyn in our DSS model, we added an experimental

arm with (Thy1)-h[A30P]αSyn transgenic mice crossed with mice that have a deletion of Cx3cr1

induced by an insertion of GFP (Cx3cr1-GFP knock-in mice) (**Figure 4A, B**). The CX3CR1-CX3CL1

axis plays an important role in maintaining the function of the lamina propria macrophage population

of the gastrointestinal wall and lack of this signaling pathway in experimental colitis models may

either aggravate or ameliorate the induced pathology [35–37]. In our experiment, the area covered by

infiltrating leukocytes following exposure to DSS located to the mucosa and submucosa and was not

significantly higher in the Cx3cr1-deficient αSyn transgenic mice than in the Cx3cr1-competent mice

(**Supplemental Figure 2A**). However, a significantly higher level of αSyn accumulated in the

submucosal plexus in αSyn transgenic mice lacking Cx3cr1 compared to αSyn transgenic mice

expressing Cx3cr1 (p = 0.001, two-way ANOVA with Tukey HSD post-hoc analysis; **Figure 4A**). In

the myenteric plexus, we found no significant increase in αSyn accumulation in neither the αSyn

transgenic mice with normal Cx3cr1 nor the αSyn transgenic mice deficient in Cx3cr1, indicating

again a prominent role for the localization of leukocyte infiltration in the process of αSyn

accumulation in the submucosa (Figure 4B). Collectively, our results indicate a potential link

between monocyte/macrophage signaling and αSyn accumulation in ENS in this experimental IBD

model.

Systemic IL-10 ameliorates DSS-induced colitis and associated enteric αSyn accumulation in αSyn

transgenic mice

As mentioned above, LPS-induced colonic leukocyte infiltration did not result in increased

accumulation of αSyn in the ENS of the colon and that the main difference in cytokine expression

between the DSS and LPS paradigms was increased IL-10 expression in the LPS group (**Figure 4**).

Interleukin-10 is an important regulator of monocytes/macrophages, and genetic ablation of IL-10

signaling or blocking IL-10 with specific antibodies has been reported to enhance DSS colitis [38,39]. To

mimic the effect of higher levels of IL-10 in an acute model of DSS colitis (5% DSS, **Figure 2C**), we

administered intravenously murine IL-10 (mIL10), which was recombinantly engineered onto two

different murine IgG variants to extend the half-life of mIL-10 in circulation (mIgG1(v1)-mIL10 and

mIgG1(v2)-mIL10, respectively). As described above, DSS induced a marked increase in leukocyte

infiltration and αSyn accumulation, and we found this to be similar in the untreated and control IgG

treated group (**Figure 6A, B**). By contrast, both mIgG1(v1)-mIL10 and mIgG1(v2)-mIL10

significantly reduced leukocyte infiltration in mice treated with DSS (p<0.0001, one-way ANOVA

with Tukey HSD post-hoc analysis; **Figure 6A, B**). Regarding human αSyn in the submucosal plexus,

only mIgG1(v2)-mIL10 significantly reduced the levels in DSS treated mice (p=0.02, one-way

ANOVA with Tukey HSD post-hoc analysis; **Figure 6B**). Interestingly, the significantly reduced

αSyn accumulation was associated with detectable serum exposure of mIgG1(v2)-mIL10, whereas

mIgG1(v1)-mIL10 was no longer detectable at the end of the *in vivo* phase, after 7 days (**Figure 6C**).

These results underline further an important role for the IL-10 pathway in keeping αSyn accumulation

at a reduced level throughout the course of experimental IBD. Together, our results with the genetic

(e.g., CX3CR1-CX3CL1 axis) and pharmacological modulation (e.g., IL-10) of DSS colitis

corroborate an important role for monocyte/macrophage pathways in the development of αSyn

accumulations in the ENS of the colon.

DSS colitis-induced submucosal αSyn accumulation at a young age persists for months and is

exacerbated by lack of Cx3cr1 signaling

IBD increases PD risk [28–30] and our own data in colon samples from IBD patients (**Figure 1**) and

recent evidence in Crohn's disease [40] indicate that such gut inflammatory conditions are associated

with αSyn accumulation in the ENS [31]. Because longer exposure to DSS mimics more closely the

chronic nature of IBD, we next elected to explore αSyn accumulation in the submucosal plexus of

(Thy1)-h[A30P]αSyn transgenic mice that were subjected to DSS colitis in a 4-week chronic

increasing dose paradigm. In order to allow for a full recovery from the chronic inflammation we then

left the mice for two months on normal drinking water and analyzed them at the age of 6 months

(**Figure 2C**). Because we also wanted to explore the effect of modulating monocytes/macrophages in

this chronic setting, an experimental arm with (Thy1)-h[A30P]αSyn transgenic mice crossed with

Cx3cr1-deficent mice was added. As expected, the area that is usually extensively covered by

leukocytes in the submucosal plexus of the acute DSS paradigm had returned to normal levels

following the two-month recovery period (**Supplemental Figure 3A**). Remarkably, however, αSyn

accumulation in the ganglia of the submucosal plexus was still almost doubled when compared to

αSyn transgenic mice that were not exposed to DSS, and this was exacerbated in αSyn transgenic

mice deficient for Cx3cr1 (**Supplemental Figure 3B**). This suggests that accumulation of αSyn is not

a transient effect or response and that improper function of monocytes/macrophages contributes to

aggravation of this accumulation.

Experimental colitis-induced at a young age exacerbates αSyn brain pathology and dopaminergic

neuron loss in old αSyn transgenic mice

The previously highlighted hypothesis by Braak and colleagues associates αSyn brain pathology in

PD with αSyn pathology in the ENS earlier in life [3,41]. To assess development of brain αSyn pathology

and to link it again to IBD risk, we exposed 3-month old hemizygous (Thy1)-h[A30P]αSyn transgenic

mice to DSS or normal drinking water and after 23 days on this chronic increasing dose paradigm

returned all mice normal drinking water for the rest of their life (**Figure 2C**). We chose to use the

αSyn transgenic model rather than wild type mice for this study because we knew that the model

exhibits some αSyn brain pathology that develops slowly also under baseline conditions. After aging

up to 9 or 21 months (i.e. mice aged for an additional 6 or 18 months after the chronic DSS paradigm

at the age of 3 months, respectively), we analyzed brain regions for pathological αSyn (proteinase K

resistant, pSer129-αSyn immunoreactive inclusions). When we examined the αSyn transgenic mice

exposed to DSS at 3 months of age, left to live only until to 9 months, we found that they exhibited

extremely low levels of pathological αSyn inclusions in the brain, similar to the levels seen in 9 month

old hemizygous (Thy1)-h[A30P]αSyn transgenic mice never exposed to DSS (**Figure 7 and**

Supplemental Figure 4). Similarly, twenty-one-month old hemizygous (Thy1)-h[A30P]αSyn

transgenic mice that only received water during their lifetimes showed relatively low levels of

pathological αSyn in the brain (**Figure 7 and Supplemental Figure 4**), which is consistent with

previous observations [42]. In marked contrast, the 21-month-old hemizygous (Thy1)-h[A30P]αSyn

transgenic mice that were exposed to DSS at 3 months of age presented with pSer129-positive αSyn

pathology throughout various brain regions in a much exacerbated fashion than mice that were aged

up to 21 months without having experienced DSS colitis at young age. The significant aggravation of

αSyn pathology also in the substantia nigra ($p \leq 0.01$ in a negative-binomial mixed-effects model

adjusting for multiple comparisons performed over all brain areas) was accompanied by a significant

loss of tyrosine hydroxylase (TH) and Nissl positive cells at 21 months of age ($p \leq 0.05$, Student's T-

test; **Figure 8**). Together, we found that DSS colitis at a young age caused an age-dependent

exacerbation of αSyn inclusion pathology and a loss of nigral dopaminergic neurons in the brains of

αSyn transgenic mice.

Discussion

Currently, there is no therapy for PD available to slow or stop its progression and an obstacle in the

quest to develop one is that we do not understand how the disease develops [43]. Intraneuronal

accumulation of αSyn (i.e. Lewy bodies and neurites) is a key neuropathological hallmark and the

distribution of Lewy pathology in postmortem brain is used for staging in PD [2,44]. Accumulation of

αSyn has also been observed in the peripheral nervous system in PD, some individuals at risk of

developing the disease, and normal individuals [45–47]. Similar to this finding in people, αSyn-

immunoreactive inclusions have also been detected in the ENS of a transgenic mouse model prior to

changes in the brain [48]. Based on preclinical models and postmortem pathology in various organs

including the brain, it has also been suggested that αSyn pathology propagates temporospatially in a

prion-like manner [3,44,49–51]. However, the initial factors triggering αSyn aggregation are yet to be

established [43] and the involvement of peripheral stimuli in the aggregation and pathogenic spread of

αSyn is only beginning to unravel.

In this study, we provide evidence that patients with IBD have increased αSyn accumulation in the

ENS (**Figure 1**) and that DSS colitis, i.e. an experimental IBD-like inflammation, triggers αSyn

accumulation in the ENS of wildtype mice and in a transgenic mouse model of PD (**Figure 3**).

Interestingly, in IBD patients and in the mouse model of IBD, we observed macrophages filled with

αSyn in the inflamed colonic wall (**Figure 5**). We found aggravation of enteric αSyn accumulation in

αSyn transgenic mice lacking Cx3cr1 signaling and amelioration of inflammation and enteric αSyn

load by systemic IL-10, suggesting that monocytes/macrophages modulate the process (**Figure 4 and**

6). We further observed that the aggravated αSyn accumulation in the ENS persisted even after two

months of recovery from DSS colitis and was aggravated in the absence of CXCR1 signaling,

indicating that the effect is not transient and mediated by monocytes/macrophages (**Supplemental**

Figure 3). Remarkably, 18 months but not 6 months post induction of DSS colitis (thus, at ages 21

months but not 9 months, respectively), αSyn transgenic mice had developed Parkinson-like brain

pathology (**Figures 7, 8, and Supplemental Figure 4**). This included elevated proteinase K resistant

pSer129-αSyn pathology in the midbrain, including the substantia nigra, and other brain regions and

an average decrease of 30-50% of TH- and Nissl-positive cells in the nigra. We chose to perform the

long-term experiments in αSyn transgenic rather than wild type mice. These particular αSyn

transgenic mice had previously been shown to slowly develop αSyn pathology in the brain [33] making

them ideal when asking the question whether transient colonic inflammation can aggravate brain

pathology in a genetically predisposed animal. In future long-term studies, we plan to address whether

αSyn pathology develops also in the brains of wildtype mice if they sustain transient experimental

IBD at a young age. In our experiments presented here, colitis in αSyn transgenic mice recapitulated

the accumulation of enteric αSyn which is proposed to occur in humans several years before PD

diagnosis [32]. Additionally, the subsequent age-related development of αSyn pathology together with

the loss of nigral dopaminergic neurons in the brain of αSyn transgenic mice mimicked a progression

of the disease similar to what is considered to occur in PD.

We established that a mechanism by which peripheral inflammation promotes αSyn accumulation in

the colon potentially involves monocytes and macrophages. Both peroral DSS and intraperitoneal LPS

administration provoked strong local immune reactions resulting in leukocyte infiltration into the

submucosa of the colon. The region of the colon which was inflamed contains the submucosal plexus

and is anatomically separated from the myenteric plexus by a thick circular muscle (**Figure 2**). This

discrete localization of inflammation to the submucosa might explain why αSyn only accumulated in

the nerves of the submucosal plexus and not in the myenteric plexus of our mice given DSS. The

mechanism underlying how intraperitoneally administered LPS leads to submucosal leukocyte

infiltration probably involves the monocyte attractant chemokine CCL2 (**Figure 4**), but the specifics

remain to be clarified [52]. Indeed, CCL2 was upregulated in the colon of our DSS model. However, in

contrast to intraperitoneal LPS, where infiltrating macrophages were present in discrete patches in the

colonic wall, DSS-related macrophage infiltration was distributed both in small groups and larger

randomly distributed patches of cells across the entire colonic submucosa. Also, perorally

administered DSS destroys the mucosa of the colon, similar to some forms of ulcerative colitis,

resulting in the transient disintegration of the intestinal epithelial barrier. In our (Thy1)-h[A30P]αSyn

transgenic mice, the subsequent immune response to the infiltration of commensal bacteria evoked an

elevated expression of cytokines such as IL-1β and IL-6. This upregulation was absent in the LPS

paradigm in which the intestinal mucosa remained intact. By acting on tight junctions, IL-1β and IL-6

can increase intestinal barrier permeability (gut leakiness), facilitating the recruitment of additional

immune cells to the site of the inflammation, eventually culminating in widespread immune activation

[53,54]. Consistent with the breach of barrier permeability in our mouse model, some PD patients exhibit

increased colonic cytokines such as IL-1β, IL-6 and TNF, occurring together with increased intestinal

permeability [20,55]. In this context, it is also notable that Crohn's patients present with increased enteric

αSyn expression [40] and even more striking that IBD patients on anti-TNF therapy have a reduced risk

of developing PD compared to IBD patients not given this treatment [29]. Here we demonstrate that

patients with IBD present with accumulation of αSyn in the ENS, as well as in infiltrating leukocytes

nearby. Notably, mucosal macrophages with intralysosomal αSyn content were previously described

in the intact human appendix [56]. These macrophages were in close proximity to the axonal varicosities

of the vermiform appendix which showed an enriched staining for αSyn in the mucosal plexus.

Furthermore, we recently found that the vermiform appendix contains aggregated and truncated αSyn

that has the propensity to seed aggregation of recombinant αSyn *in vitro* [47]. What could be a

functional role of the αSyn species found in abundance in the gut wall? Monomeric and oligomeric

αSyn species reportedly act as chemoattractants for neutrophils and monocytes, enhancing the

maturation of dendritic cells in the ENS [19,57]. With such a role in intestinal immunity, it is possible

that the tissue destruction induced by DSS in the present study led to release of αSyn, which perhaps

served as a chemoattractant for monocytes. The increased abundance of extracellular αSyn and altered

intestinal permeability, along with the DSS-evoked inflammatory response may have provided an

enabling milieu allowing further αSyn accumulation in the ENS of the colon [58]. Macrophages and

other immune cells are also regulated by several genes including *LRRK2*, an established risk gene for

PD and IBD. It will be interesting to explore how mutations in genes that control autophagy,

including the *LRRK2* gene, influence the handling of αSyn by macrophages that invade the inflamed

colon in our DSS colitis paradigm. Despite the intriguing translational aspect of our finding in the

DSS paradigm, others have very recently reported that DSS colitis in mice down-regulates the

expression of enteric αSyn on protein levels *in vivo* [59]. This is in contrast to our immunofluorescence

(e.g. increased accumulation of αSyn in submucosal plexus upon DSS colitis; **Figures 3, 4, and 6**)

and gene expression data (e.g., no change in endogenous and transgenic αSyn upon DSS colitis;

Supplemental Figure 2) in the same paradigm and may reflect the well-known lab-to-lab variability

that can occur for the DSS models [60].

Perhaps the most striking finding in our study was that a single period of DSS-induced colitis at a

young age led to an exacerbation of αSyn pathology in the brain of αSyn transgenic mice much later

in life (**Figure 7**). How does the severe αSyn inclusion pathology develop in the brain of these mice?

One hypothesis is that the brain αSyn pathology observed in this study could be due to direct effects

of peripheral immune activation on the brain and that certain peripheral triggers can directly affect

microglial activity. For instance, short-chain fatty acids derived from gut microbiota appear to

influence function and maturation of microglia in the mouse brain [61] and inflammatory mediators

released by gut microbiota into the bloodstream have been suggested to induce brain pathology and

behavioral changes in an αSyn transgenic mouse model [62]. Moreover, rats and nematodes have been

reported to develop αSyn inclusions after exposure to the bacterial amyloid protein curli, a protein

which stimulates microgliosis, astrogliosis, and secretion of IL-6 and TNF [63]. Intriguingly, a recent

study reported that peripherally applied inflammatory stimuli induce acute immune training (that

exacerbates β-amyloid pathology) and immune tolerance in the brain that reprograms microglia, an

effect which can persist for at least six months [64]. Whether this is a relevant mechanism in the DSS

paradigm needs to be explored.

Another hypothesis is that the brain αSyn pathology observed may have accumulated following the

transfer of pathogenic αSyn seeds from the gut via the vagal nerve. Several experimental studies have

demonstrated that pathogenic αSyn seeds can be transferred from the peripheral to the central nervous

system. Aggregated recombinant αSyn injected intraperitoneally, intramuscularly or into the gastric

wall of certain mouse models of PD results in αSyn inclusions in the brain [16,65]. Data from animals

injected with αSyn protein in the gut wall or viral vectors expressing αSyn into the vagal nerve

suggest that pathogenic seeds can be transmitted via the vagal nerve [15,66]. A role for the vagal nerve in

PD was also suggested by an epidemiological study indicating that vagotomy in a Danish population

is associated with decreased PD risk [67], although this association has been challenged [68]. In the present

study, αSyn pathology was much more prominent in the reticular nucleus (including the vagal area)

and midbrain areas (compared to the rostral areas) at 18 months post DSS colitis. Although we did not

conduct the definitive experiment of cutting the vagal nerve, our data support the growing body of

evidence that the vagal nerve is involved in the accumulation of αSyn aggregates in the brain.

In summary, here we report that individuals with IBD exhibit αSyn accumulation in the colon

concomitant with infiltrating monocytes/macrophages positive for αSyn. We also show that αSyn

accumulates in the colon of αSyn transgenic and wildtype mice subjected to DSS colitis and that this

process is modulated by monocyte/macrophage-related signaling. We further demonstrate that chronic

DSS colitis in young αSyn transgenic mice leads to a markedly exacerbated accumulation of αSyn

aggregates in the brain when the mice age. In the same aged mice, the numbers of TH- and Nissl

positive neurons in the substantia nigra are reduced, suggestive of a neurodegenerative process.

Together, our findings are in consonance with studies demonstrating a link between IBD and PD

[28,29,69] and suggest a critical role for intestinal inflammation and αSyn accumulation in the initiation

and progression of PD.

Methods

Mice

Male C57BL/6 wild type mice (Jackson Laboratories, Bar Harbor, USA), hemizygous Tg(Thy1-SNCA*A30P)18Pjk ((Thy1)-h[A30P]αSyn) [33] and Tg(Thy1-SNCA*A30P)18Pjk crossed with Cx3cr1tm1Litt ((Thy1)-h[A30P]αSyn /CX3CR1-def; homozygous for Cx3cr1-GFP knock-in allele; [70] transgenic mice were used for the study. (Thy1)-h[A30P]αSyn transgenic mice express mutant human αSyn under the neuron selective Thy1 promoter. (Thy1)-h[A30P]αSyn transgenic mice were crossed to Cx3cr1-def transgenic mice which express eGFP replacing fractalkine gene expression. All mice were maintained on a C57BL/6 background for more than 10 generations and under specific pathogen-free conditions. To the extent possible, littermates were used in the experiments. Health status was monitored daily during experiments.

Human Subjects

Samples from patients with Crohn's disease (CD), ulcerative colitis (UC) or healthy subjects (HS) were provided by the tissue bank, Institute of Pathology, University of Bern. Briefly, specimens were obtained from patients who underwent surgical procedures at the University Hospital (Inselspital) in Bern, Switzerland between 2004 and 2011. Three selected male patients previously clinically diagnosed with UC with a reported disease duration > 6 year (n=3) and undergoing steroid therapy combined with either metronidazole or mesalazine. CD patients were of mixed gender and aged 22-56 years ranging from 2 months to 11 years post disease diagnosis undergoing treatment with either infliximab or mesalazine in combination with steroids. Healthy subjects were of mixed gender with no report of inflammatory bowel disease, aged 40-59. All samples contained the mucosa and submucosa regions including minor parts of the circular muscle layer. Following surgical removal, tissue samples were immediately immersed in O.C.T. compound (VWR International GmbH, Dietikon, Switzerland), frozen in liquid nitrogen and stored at -80°C. Diagnosis of disease status was made according to established criteria for histopathological analysis.

Experimental IBD in mice with DSS and LPS

Paradigms for the induction of inflammation were either 1 week (acute) or 3-4 weeks (chronic) with

or without an incubation phase of 2, 6 or 18 months post application (**Figure 2**). Acute systemic

inflammation was induced by intraperitoneal Lipopolysaccharide (LPS) application [71] of 0.5 mg/kg in

100 µl injection volume on day 1 and 4 (Sigma-Aldrich Chemie GmbH, Steinheim, Germany, LPS

055:B5). Acute colitis was induced by application of 36-50kDa Dextran Sulfate Sodium (DSS) [72]

(160110, MP Biomedicals, LLC, Illkirch, France) at 0%, 1%, 2.5% or 5% in autoclaved drinking

water for 5 continuous days respectively, followed by 2 days of water (1 DSS application cycle).

Chronic colitis was achieved by 4 repeating DSS application cycles. The DSS concentration during 4

weeks of chronic colitis was either 1% or 2.5% for 4 weeks or 2.5%-4% raised 0.5% every week for 4

weeks. Mice from same littermate group were randomized per cage into vehicle and inflammation

inducing agent.

IL-10 treatment and exposure measurement

Two different forms of mouse IgG bound murine IL-10 (mIgG(v1)-mIL10 and mIgG(v2)-mIL10)

were diluted in pre-prepared sterile formulation buffer comprised of 0.5% mouse serum supplemented

with 25mM citrate, 300mM arginine to a final concentration of 0.75 mg/ml and the pH adjusted to 6.7

on the day of application. Each mouse was treated once with 150 µg i.p concurrently with the

initiation of the acute colitis paradigm with 5% DSS. The concentrations of mIgG-mIL10 fusion

proteins in murine serum samples were determined by enzyme-linked immunosorbent assays (ELISA)

specific for the Fab moiety of the administered mIgG-mIL10 fusion protein. Biotinylated mIgG-

mIL10-specific target molecules were used for capturing, goat anti-mIg IgG-HRP conjugate and

peroxidase substrate ABTS was used for quantitative detection of mIgG-mIL10 fusion proteins.

Immunohistochemistry

Animals were injected with a lethal dose of pentobarbital (150 mg/kg). Upon full anesthesia, mice

received transcardial perfusion with room temperature phosphate buffered saline (PBS). For

biochemical and immunohistochemical analysis, one section of either the proximal colon was fresh

frozen and stored at -80°C or post-fixed in 4% paraformaldehyde (PFA) solution for 24 h. Following

post-fixation, organs were incubated in 30% sucrose/PBS at 4°C for at least 48 h before further

processing. Subsequently, enteric tissue was cryotome-sectioned to 35 μm thick longitudinal sections

(approx. 1 cm length). The brain was collected and post-fixed for 24 h in 4% PFA followed by 30%

sucrose in phosphate buffer until cryo-sectioning of floating sections at 40 μm. Histological analysis

of the colon was performed using standard hematoxylin staining. Immunohistochemical staining was

accomplished using the Vectastain Elite ABC Kits and Peroxidase Substrate Kit SK-4100 (Vector

Laboratories, Burlingame, CA, USA) or fluorescently labelled secondary antibodies (Alexa coupled

to dye 488, 555 or 647, Life Technologies, Zug, Switzerland). The following primary antibodies have

been used for overnight incubation at a dilution of 1:1000: monoclonal antibody to human α-synuclein

(clone 211, sc-12767, Santa Cruz Biotechnology, Heidelberg, Germany; used on tissue from human

αSyn transgenic mice), monoclonal antibody to rat α-synuclein but cross-reactive with murine and

human αSyn (Syn1/clone 42, BD Transduction Laboratories, Allschwil, Switzerland; used for wild

type mice and in human colon), polyclonal antibody to the peripheral neuronal marker Peripherin

(Millipore Corporation, Billerica, MA, USA), and polyclonal antibody to macrophage marker Iba1

(Wako Chemical GmbH, Neuss, Germany). To detect phosphorylated αSyn (pSer129 pathology) in

the free-floating brain sections, monoclonal antibody (ab51253, Abcam, Cambridge, USA) to human

αSyn was used at a dilution of 1:10000. Prior to the pSer129 staining, the free-floating brain sections

were incubated for 10 min at room temperature in a phosphate buffered saline solution containing 10

μg/mL proteinase K (Cat # 25530015; Invitrogen, California, USA). TH-immunoreactive cells were

detected using a polyclonal antibody (657012, Millipore Sigma) at a dilution of 1:1000. To measure

the density of Nissl-positive cells, the TH-stained cells were counter-stained with Cresyl violet. The

slides were incubated in 0.1% Cresyl violet solution for 9 min and then dehydrated in 95% and 100%

ethanol and then xylene prior to coverslipping with Cytoseal 60 mounting media (Thermo Fisher

Scientific). Quantifications of the blind-coded TH/Nissl stained slides were done using

Stereoinvestigator (version 2017.01.1; MBF Bioscience, Williams, VT, USA) on Imager M2

microscope (ZEISS) coupled to a computer. We analyzed 5-7 nigral sections per animal, and a total of

7-8 animals per treatment group. We outlined the substantia nigra pars compacta and counted every

TH-immunoreactive and Nissl-positive cell in that area and computed the number of cells per section,

generating the mean cell density per animal. We then calculated the mean density of cells per

treatment group and analyzed the data using unpaired Student's T-test after confirming normality and

homoscedasticity in Prism 7.0 (GraphPad Software).

Imaging and stereological quantification of αSyn deposits in enteric nervous system

Imaging and stereological quantification was performed on a Zeiss Axio Imager Z2 fluorescence

microscope (Carl Zeiss AG, Jena, Germany). Leica TCS SP5 confocal system using an HCX PL APO

CS 40x 1.3 oil UV or an HCX PL APO LB 63x 1.4 oil UV objective was utilized for image recording.

Accumulation of αSyn in the ENS was assessed on a random set of 3 adjacent 35 µm thick, αSyn-

immunostained sections comprising the myenteric and submucosal neuronal plexuses. Analysis was

performed with the aid of Stereologer software (Stereo Investigator 10, MBF Bioscience, Williams,

VT, USA) as described previously [73]. In the myenteric plexus ganglion volume was defined by

multiple outlined plexuses containing a range of 5-20 neuronal cells and quantified by the optical

fraction fractionator technique. In contrast to the myenteric plexus, the submucosa consists of

compact plexuses with 1-5 cells including interconnecting neurites. Therefore, the entire submucosa

was set as region of interest, analyzed with the area fraction fractionator technique. Results of the

submucosal plexus are displayed by percent area containing αSyn deposits. For the IL-10 experiment,

αSyn positive inclusions from immunofluorescence images were counted for each image. Inclusion

body-like features were filtered based on having a size between 12 and 50000 pixels and a minimal

intensity value greater than 300. The filtering step was included to exclude small background features

and macrophages (very large spots). The counts were then aggregated to the animal level by summing

the inclusion feature counts of all images per animal and then normalizing for (i.e. dividing by) the

number of images for a given animal. Upon exploratory data analysis two animals were excluded: one

mouse because it only had one image and another due it being an outlier, based on its infiltration score

and image data.

Blinding of experimenters

For analyses of colon and brain tissue on slides, a second individual assigned unique codes to

stained slides. Therefore, the experimenter conducted the analyses blinded to the identity of the

mice. For randomization of treatment groups see above.

Quantification of leukocytes infiltration

To determine the leukocyte covered area in the colon after LPS or DSS application, three adjacent

hematoxylin stained sections were quantified. Total area of colon sections and localizations of

leukocyte assemblies within the tissue architecture were identified and outlined utilizing Stereologer

Software (Stereo Investigator 6, MBF Bioscience, Williams, VT, USA). Percentage of leukocyte

covered area has been set in proportion to total area of the analyzed colon section. For the IL-10

experiment, hematoxilin stained colon slices were examined by an expert pathologist blinded to

treatment conditions. A score of 0-3 was assigned to each section for each of the 3 layers lamina

propria, submucosa and muscularis based on the degree of inflammatory infiltration. A score of 0

denoted no inflammation and a score of 3 indicated extensive infiltration. The mean of the values for

all 3 layers was taken as the final measure of leukocyte infiltration per mouse.

Quantification of αSyn/Iba1 double positive macrophages

The number of α-syn+/Iba1+ positive cells was evaluated by quantification of 10 random regions in 2

adjacent sections of the proximal colon. The region of interest was set to contain the myenteric

plexus/circular muscle layer and the submucosal plexus. Cells were assessed for positive αSyn

staining and concomitant co-localization with the macrophage marker Iba1 was quantified.

Scoring of pSer129 pathology and brain heatmap

We evaluated pSer129 pathology on a full series of immunostained coronal sections from 10 mice per

treatment group (i.e. water vs. DSS-treated groups) on blind-coded slides using a previously described

method [74]. We visualized pathology from one hemisphere of all brain sections (apart from the

olfactory area) using NIKON Eclipse Ni-U microscope and assigned scores ranging from 0 to 4 to

each brain area based on the relative abundance of PK-resistant pSer129-positive inclusions (i.e. cell

bodies and neurites). In this case, 0 = no aggregates, 1 = sparse, 2 = mild, 3= dense, 4 = very dense.

For the heatmap, we obtained the average score values of each brain area for each treatment group.

The average data for each treatment group (n=10/ group) was then represented as a heatmap in a

sagittal mouse brain background (http://atlas.brain-

map.org/atlas?atlas=2#atlas=2&structure=771&resolution=16.75&x=

7755.7470703125&y=3899.625&zoom=-3&plate=100883867&z=5).

Densitometry of pSer129 αSyn brain pathology

The density of pSer129 pathology in 12 major brain areas (reticular nucleus, pontine reticular nucleus,

periaqueductal gray, gray and white layer, reticular formation, substantia nigra, ventral tegmental

area, thalamus, hypothalamus, central amygdala, pallidum and striatum) was determined in the water

and DSS-treated animals. A NIKON Eclipse Ni-U microscope was used to acquire 20x magnification

images (without condenser lens) from all the indicated brain areas, using the same exposure time for

all images. In all cases, images were acquired on three sections separated by 420 μm intervals

(localized between Bregma). We then processed the acquired images using Image J64 [75], created a

mask (to exclude background) that redirects to the original image for analysis, measured the total area

and the mean grey value of the area that had inclusions. For brain areas such as periaqueductal gray

that do not fill the entirety of the field to be analyzed, we drew a contour of the area and the analysis

was performed only within that contoured area. We subsequently calculated the grey value of the area

per square pixels for each image (i.e. A.U./px^2 = mean grey value x area stained/total area assessed).

Based on this, we calculated the average grey value per square pixels for each brain area for each

animal ($n = 6$ mice/group), and then extended this calculation to determine the average grey value per

square pixels for each treatment group and each of the twelve brain areas of interest.

mRNA expression

To assess mRNA expression levels from the proximal colon, RNA was extracted from fresh frozen

tissue with MagnaLyser green beads (Roche Diagnostics, Mannheim, Germany) and Qiazol Lysis

(Reagent cat.no.79306, Hilden, Germany) purified on MagnaPure LC (HP Kit no.03542394001, F.

Hoffmann - La Roche AG, Rotkreuz, Switzerland) and amplified via real-time PCR (4ng

RNA/reaction; Lightcycler 480, Roche Diagnostics Corporation, Indianapolis, USA). Amplification

of mRNA was performed by using TaqMan probes for human or murine specific α-synuclein and for

selected cytokines/chemokines (Applied Biosystems Europe B.V., Zug, Switzerland). Target mRNA

was normalized to tissue-specific murine GAPDH levels and displayed as relative expression after 30

amplification cycles.

Statistics

Statistical analysis of gut pathology and inflammation was performed using GraphPad Prism 6.04 or

7.0 software (GraphPad Software, Inc. La Jolla, CA, USA). The results are expressed as mean values

± standard errors of the mean (SEM). Student's T-test (or Welch's T-test for unequal variances) was

used to compare two groups and ANOVA was used for multi-comparison of groups followed by

Tukey HSD post-hoc analysis. For the statistical analysis of the pSer129 αSyn brain pathology,

negative-binomial mixed-effects models with a random intercept for each sample were used to

analyze the dataset via the 'lme4' (http://lme4.r-forge.r-project.org/) package in R v 3.4.4. To analyze

the pSer129 αSyn cell count dataset, an offset for the total area examined was included to model the

densities. Linear contrasts with false discovery rate (FDR) adjustments were then used to test our

hypotheses and account for multiple testing (for brain area and experimental group). Like the pSer129

dataset, the Iba-1/αSyn-double positive dataset were analyzed using negative-binomial regression and

Tukey HSD adjusted contrasts to test our hypotheses.

For the statistical analysis of the mRNA expression, data quality was assessed by inspecting the

distribution of Cp values of reference endogenous genes across samples, by inspecting the level of Cp

variation between technical replicates and by exploring the samples multivariate signal distribution as

in a principal component analysis. Relative gene expression levels were expressed as $2^{-(Cpgene - CpRef)}$.

Statistical analyses to assess the effect of the experimental conditions on the log2 gene expression

levels were done with linear models using the *limma* package (Bioconductor/R, Smyth, 2005). These

analyses were implemented in R v3.1.1.

For the statistical modelling of the effects of the IL-10 treatment on αSyn counts, as well as

infiltration scores, the levels for IgG1(v1)-IL10 and IgG1(v2)-IL10 treatment were compared to the

positive (vehicle/DSS) control. Additionally, since levels of the control antibody treatment (IgG1(v1))

were very similar to the positive control, the two groups were pooled in further contrasts in which

effects of individual antibodies or control IgG was assessed. For αSyn counts, a linear model on the

treatment groups with one-degree freedom contrasts was applied. For the infiltration score a Kruskal-

Wallis test, with the same contrasts, was used.

Study approvals

The human subjects' study was conducted with the approval of the local Ethical Committee in Bern

No. 47/04. Written informed consent was obtained from each patient. The animal experiments were

approved by a Roche internal review board and the local authorities.

Author contributions

S.G., N.M. and L.S. planned and performed the in vivo experiments, colon immunostaining, analysis,

and quantification; S.G. and N.M. drafted a first version of the manuscript; E.Q. performed, imaged,

quantitated pSer129, TH and Nissl staining in the brain sections, and drafted a more advanced version

of the manuscript with J.A.S., who also provided helpful discussion. F.B. and K.O.S. supported the

image acquisition and image analysis for the colon samples; M.St. performed imaging and data

analysis of experiments with wildtype mice; G.D.P. and J.S.P. performed statistical analysis of the

DSS experiments; H.R. and M.H. performed mRNA analyses; M.Se. trained S.G. and L.S. on mouse

necropsy and supported their work; P.M. performed expert pathology staging on leukocyte

infiltration; T.E. and A.W. provided mIgG-mIL-10 fusion proteins and measured serum exposure;

Z.M. performed statistical analysis for the pSer129 αSyn immunohistochemistry data. M.L.E.G.

provided helpful discussion and project planning. A.H. co-mentored S.G. and N.M., performed expert

pathology staging on leukocyte infiltration and contributed to experimental planning. C.M. trained

S.G. on the colitis model, provided human tissue and expert input on the experimental IBD model.

M.B. and P.B. co-mentored Roche Postdoctoral Fellows S.G. and N.M., conceived and oversaw the

study, and performed experimental planning; M.B., P.B. and E.Q. wrote the final version of the

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Conflicts of Interest

At the time of the study S.G. and N.M. were Roche Postdoctoral Fellows employed by Roche and

L.S., F.B., G.D.P., J.S.P., K.O.S., H.R., M.H., M.Se. M.St., P.M., A.W., T.E., A.H. and M.B. are or

were fulltime employees or trainees at Roche and they may additionally hold Roche stock/stock

options. S.G. and L.S. are currently employees of Neurimmune AG, Schlieren, Switzerland. P.B. has

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Figures and figure legends

Grathwohl et al., Figure 1

Figure 1

Alpha-Synuclein inclusions in the enteric nervous system and in macrophages of patients with

inflammatory bowel disease. (A, B) Immunofluorescence images of αSyn inclusions (Syn1

antibody) in the submucosal region of 10 μm cryo-sections from colons of patients with colitis

ulcerosa. Arrow heads point to neuritic features indicating presence of inclusions in enteric nerves.

Scale bar 50 μm. **(C)** Close-up of a colonic region with active leukocyte infiltration in a patient with

ulcerative colitis. Immunoreactivity for αSyn (red) was observed in neuritic features (arrow heads) as

well as in individual leukocytes (arrows; identification of leukocytes based on cellular morphological

features and localization). Nuclei are shown in blue (DAPI). Scale bar 20 μm.

Figure 2

Age dependent increase of intracellular αSyn accumulation in enteric nervous system of

heterozygous (Thy1)-h[A30P]αSyn transgenic mice and setup of the experimental colitis

paradigms. (A) Confocal microscopy imaging of the inclusions of human αSyn (red, antibody clone

211; human αSyn specific) within the ganglia of the submucosal plexus (green, peripherin; blue,

DAPI/nuclei) of heterozygous (Thy1)-h[A30P]αSyn transgenic mice. Arrow head points to one of the

typical irregularly sized and shaped αSyn inclusion bodies visualized in 2D z-stacks of rotated

confocal images. Scale bar, 100 μm. **(B)** Stereological quantification of normally occurring human

αSyn inclusions in the myenteric and submucosal plexuses of 3 and 12 months old heterozygous

(Thy1)-h[A30P]αSyn transgenic mice (n = 4 per group; mean and S.E.M. are shown; Student t-test

between the two age groups in each region). **(C)** Setup of experimental colitis paradigms employing

dextran sulfate sodium (DSS, per os in drinking water). Additionally, peripheral inflammation was

induced by bacterial lipopolysaccharide (LPS, intraperitoneal injection). After some chronic DSS

paradigms mice were aged on normal water up to 6, 9 or 21 months. Mice aged up to 9 or 21 months

of age were analyzed for brain pathology **(D)** Hematoxylin staining of 35 μm thick colon sections of 3

months old heterozygous (Thy1)-h[A30P]αSyn transgenic mice. Organizational layers of the intact

colon (left panel). Representative images of various severity degrees of DSS-driven colitis from weak

leukocyte infiltration (top panel of acute DSS) to mucosal ulceration (lowest panel of acute DSS).

Note the different appearance of enteric inflammation in acute LPS-driven peripheral inflammation

compared with DSS; e.g., confined immune cell clustering and lymphoid hyperplasia; intact mucosal

layer. Scale bar 50 μm (intact colon), 100 μm (acute DSS), and 200 μm (LPS).

Figure 3

Colitis severity and duration-dependent aggravation of accumulation of αSyn inclusions in the

colonic submucosal plexus of heterozygous (Thy1)-h[A30P]αSyn transgenic and wild type mice.

(A) DSS dose-dependent increase of leukocyte infiltration in the acute and chronic paradigm. The

highest acute dose (2.5%) and the two constant chronic doses led to an increase of αSyn inclusions in

the submucosal plexus (stereological quantification of αSyn inclusions in the submucosal plexus of 3

and 6 months old heterozygous (Thy1)-h[A30P]αSyn transgenic mice; n = 5-7 per group; mean and

s.e.m. are shown). **(B)** Representative 2D z-stacks of confocal images of increasing abundance of

αSyn inclusions (red, human-αSyn specific monoclonal antibody clone 211) in a ganglion of the

submucosal plexus (green, peripherin) with cellular nuclei in blue (DAPI) in the acute DSS paradigm.

Arrow heads point to the typical irregularly sized and shaped αSyn inclusion bodies that accumulate

in the highest DSS dose. Scale bar 200 μm. **(C)** Overview of colonic region of 3-month-old wildtype

mice (top row) exposed to water or acute DSS (5%) with immunofluorescence analysis of murine

αSyn load in the colon performed immediately after colitis or exposed to constant chronic DSS (2.5%)

and analysis after aging on normal water for another 2 months. White dotted rectangles in the top row

indicate the area that was zoomed out below illustrating in more detail the murine αSyn inclusions

(red, rodent αSyn cross-reactive monoclonal antibody syn1/clone 42) in the submucosal plexus

(green, peripherin). The lower three rows shows DAPI and αSyn inclusions with and without the

peripherin channel. The white dotted circled area illustrates the peripherin-positive area that was

analyzed for αSyn inclusion bodies (arrow heads in bottom row). Scale bar for the lower three panels

200 μm. **(D)** Stereological quantification of murine αSyn inclusions in the submucosal plexus of

wildtype mice right after acute DSS colitis or after 2 months of recovery from a 4-week chronic DSS

colitis (n = 5 per group). Note the regularly arranged and smoothly distributed immunoreactivity for

the physiological αSyn with barely any inclusion bodies in the intact enteric nerves of the water

group.

Figure 4

Colitis induced by peroral DSS but not peritoneal LPS enhances αSyn accumulation in the colonic submucosal plexus of heterozygous (Thy1)-h[A30P]αSyn transgenic mice and can be aggravated by lack of Cx3cr1 signaling. Mice received in an acute paradigm either peroral 5% DSS in their drinking water or intraperitoneally 0.5 mg/kg LPS. Effects of the two agents in the ENS was compared to effects induced by vehicle (see Figure 2C for timelines). Stereological quantification of αSyn inclusions in the submucosal plexus as % area **(A, C)** and in the mucosal plexus as particle load per ganglion **(B, D)** (Two-way ANOVA with Tukey post hoc test). **(E)** Representative 2D stacks of confocal images of intracellular αSyn inclusions (red, human αSyn specific monoclonal antibody clone 211; arrow heads pointing to some selected inclusions) in a ganglion of the myenteric plexus

(green, peripherin) with cellular nuclei in blue (DAPI). Scale bar 50 µm. Gene expression analysis of

selected cytokines in the colon of (Thy1)-h[A30P]αSyn transgenic mice that received either acutely

LPS **(F)** or DSS **(G)** compared to their respective vehicle or water controls. Note the strong increase

in IL-6 and the lack of elevation of IL-10 in the DSS paradigm compared to the LPS paradigm

indicating a different inflammatory colonic milieu despite the abundant leukocyte infiltration in both

paradigms. n = 5-8 per group; mean and s.e.m.; Student's t-test between inflammatory agent and

vehicle for individual cytokines.

Grathwohl et al., Figure 5

Figure 5

Alpha-synuclein co-localizes with ENS and macrophages upon DSS colitis in αSyn transgenic

mice. (A, B) Immunofluorescence image of αSyn staining in colonic region of (Thy1)-h[A30P]αSyn

transgenic mice on water **(A)** or after acute DSS colitis (2.5%) **(B)**. Note the small dotted structures of

the typical αSyn inclusions in the submucosal plexus (arrow heads) and the large features of

immunoreactivity which localize to infiltrating leukocytes (arrows; identified by their typical cellular

morphology), similar to what was observed in IBD patients in Figure 1. Scale bar 100 µm. **(C)** 2D

stacks and close-up of confocal images co-localizing αSyn (red) with the macrophage marker Iba-1

(green) in the colon of a (Thy1)-h[A30P]αSyn transgenic mouse after DSS colitis. Note the dotted

structures of the typical αSyn inclusions in the submucosal plexus (arrow heads). Scale bar 40 μm and

13 μm for the close-up. **(D)** Quantification of numbers of Iba-1/αSyn-double positive macrophages (n

= 3 per group; mean and S.E.M.)

Grathwohl et al., Figure 6

Figure 6

Systemic IL-10 ameliorates DSS colitis and associated local αSyn accumulation in (Thy1)-

h[A30P]αSyn transgenic mice. Two different recombinantly engineered and murine IgG1-fused

forms of murine IL-10 (mIgG1(v1)-mIL10 and mIgG1(v2)-mIL10) were administered (150 μg per

mouse i.p.) at the beginning of the acute DSS paradigm (5%) in (Thy1)-h[A30P]αSyn transgenic

mice. Vehicle and the mIgG1(v1) alone served as untreated controls. **(A)** Leukocyte infiltration was

assessed by visual scoring and **(B)** inclusion features of αSyn were stereologically and semi-

automatically quantified and result log scaled for statistical analysis. Both the vehicle group and the

mIgG1(v1) group had similar levels of leukocyte infiltration and αSyn inclusions and were merged for

the statistical analysis to compare with the IL-10 treated groups. Both forms of IL-10 ameliorated

leukocyte infiltration whereas mIgG1(v2)-mIL10 also blocked the appearance of αSyn inclusions

significantly (n = 3-6 per group; mean and s.e.m.; one-way ANOVA and Tukey post hoc test). **(C)**

Persistent exposure mIgG1(v2)-mIL10 versus mIgG1(v1)-mIL10 (lower limit of detection is indicated

at <0.234 μg/ml) as measured in serum at the end of the in vivo phase corresponds with beneficial

treatment effects on αSyn readout observed above. The mIgG1(v1) was only measured in two mice.

Figure 7

A single chronic DSS colitis insult causes an age-dependent accumulation of proteinase K

resistant pSer129-αSyn in various brain regions of (Thy1)-h[A30P]αSyn transgenic mice. A 3-

week chronic increasing dose DSS paradigm was performed with 3-month old (Thy1)-h[A30P]αSyn

transgenic mice. After recovering and further aging, various brain regions were analyzed for

proteinase K resistant pSer129-αSyn immunoreactivity in 9-month **(A-F)** and 21-month old **(G-L)**

mice, respectively. The dark brown features in G-L indicate proteinase K resistant pSer129-αSyn.

They are barely visible in A-F. Densitometric quantification of pSer129-αSyn immunoreactivity in

different brain regions in 9-month **(M-O)** and 21-month old mice **(R-T)** (n=6 mice per group). The

two orders of magnitude different y-axes between M-O and R-T confirm the visual impression in

panel A-L. Statistical analyses were performed using negative-binomial mixed-effects models

adjusting for multiple comparisons. Representative heatmap of the average distribution scores of

pSer129-αSyn immunoreactivity for each treatment group in varying brain regions in all the 9-month

956 **(P-Q)** and 21-month old **(U-V)** mice was generated in a sagittal mouse brain (n=10 mice per group).

957 Scale bars: 500 µm.

Grathwohl et al., Figure 8

Figure 8

L**oss of tyrosine hydroxylase and Nissl positive cells in the substantia nigra of (Thy1)-**

h[A30P]αSyn transgenic mice at 21 months of age, 18 months post recovery from DSS colitis.

(Thy1)-h[A30P]αSyn transgenic mice that were exposed to a chronic DSS-colitis paradigm at 3

months and were aged to 21 months showed a significant loss of mean count of cells with tyrosine

hydroxylase (TH) immunoreactivity and cellular Nissl staining in the substantia nigra compared to

age-matched littermate mice in the group that did not experience DSS colitis (water). **(A)**

Representative images of two levels of the substantia nigra in one mouse per group. **(B)** Stereological

quantification of cells positive for TH or Nissl (n=7-8 mice per group). Statistical analyses of the TH

dataset were performed using Student's T-test, while Welch's T-test was used for the Nissl dataset to

adjust for unequal variances. Scale bar: 500 µm.