Animal models of Parkinson’s disease: New models provide greater translational and predictive value

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Parkinson’s disease (PD) is a devastating neurodegenerative disease in which current treatments only provide symptomatic relief and do not alter its progression. As such, there is a great necessity for better treatment options for PD. Animal models play a key role in preclinical research, and translational and predictive models are critical to develop new treatments. This article reviews the current chemically-induced and genetic models of Parkinson’s disease, and discusses the advantages and limitations of available models. Also discussed here are the newly available and novel knockout rat models of Parkinson’s disease recently commercialized by Sigma Advanced Genetic Engineering (SAGE®) Labs. These models will potentially help elucidate the molecular mechanisms that underlie PD and facilitate drug development.

Introduction

Parkinson’s disease (PD) is a progressive neurodegenerative condition that results in movement and cognitive disorders. The incidence of PD in patients 60 years or older is ~1%. PD arises from the degeneration of dopamine (DA) neurons in the substantia nigra. Loss of striatal dopaminergic function in PD leads to resting tremor, bradykinesia, rigidity, and postural instability.

The cause of PD is unknown. Most cases are sporadic, however, 15% of patients have a first degree relative with the disease (Samii A, 2004) and up to 5% of cases have been linked to genes known to be associated with PD (Lesage S, 2009). Treatment measures have been mostly focused on augmentation of dopaminergic signaling—pharmacologic treatments include levodopa (a dopamine precursor), dopamine agonists, and monoamine oxidase B (MAO-B) inhibitors (prevent dopamine metabolism). These drugs are more focused on symptomatic rather than disease modifying treatment; there is great necessity and large research efforts to develop disease modifying agents that slow progression or even prevent or reverse PD (Savitt JM, 2006).

Animal models are a key part of preclinical research in drug discovery. Chemically-induced models of PD are most often rodents administered chemicals toxic to dopaminergic neurons. Genetic models of PD have focused on mutating or knocking out genes known to cause familial PD. As discussed below, rodent models thus far have faced limitations due to lack of strong construct (i.e. genotype or intervention) and/or face validity (i.e. phenotype), as well as species and strain limitations.

Chemically-induced models

Two of the most widely used animal models of Parkinson’s disease are the chemically induced 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) and 6-hydroxydopamine (6-OHDA) rat. Both MPTP and 6-OHDA are neurotoxic and are substrates for the dopamine transporter (DAT). 6-OHDA can also be a substrate for the norepinephrine transporter (NET), and thus when used alone, 6-OHDA is not solely specific for DA neurons. Both MPTP and 6-OHDA are typically used in rats rather than mice, due to the larger size (easier surgeries and microinjections), richer behavioral profile, and greater translational relevance.

The chemically-induced models of PD have strong face validity for both the loss of dopaminergic neurons and the elicited motor phenotype. They are useful models of dopaminergic loss, especially where the cause of the loss is not important for the study. However, these models have particularly poor construct validity (6-OHDA and MPTP poisoning are extremely rare) in that they only represent loss of dopaminergic neurons and cannot be used to investigate progression of the pathology of the disease including the mechanisms responsible for neurodegeneration. Additionally, these models do not form Lewy bodies, an important hallmark of PD, and they cannot be used to assess any of the non-dopaminergic changes known to be present in PD.

Genetic models

Genetic models of PD have focused on genes known to be implicated in familial Parkinson’s disease. In these models, the gene of interest is usually either knocked out, or a humanized version of the gene (incorporating a known human mutation) is either knocked in or transgenically expressed (Dawson TM, 2010). Historically, these models have been in mouse due to the ease of genetic manipulation in this species. Target genes in genetic models typically include alpha synuclein (Snca), leucine rich repeat kinase 2 (Lrrk2), parkin (Park2), PTEN-induced putative kinase 1 (Pink1), and DJ-1 (Park7).

The genetic models offer much higher construct validity than the chemically-induced models—these models possess defects in genes where dysfunction has been linked to PD in humans. This is especially relevant for individuals with familial PD harboring mutations in these particular genes. For sporadic PD, the construct validity is less clear. While sporadic patients, by definition, do not possess mutations in the above genes, the underlying pathology and molecular mechanisms are likely to converge at some point. Further, while the context may not be identical, the genetic models, unlike the chemically-induced models, do offer the possibility to directly examine PD progression and pathology.
While the genetic models excel in construct validity, they have thus far been disappointing when it comes to face validity. Phenotypes seen in the existing mouse models have been mild at best, with some models exhibiting no real phenotype (Dawson TM, 2010). An ideal model should display a progressive loss of dopamine neurons and formation of Lewy bodies, which is then manifested as motor deficits. Non-dopaminergic aspects of PD should also be modeled. To date, no genetic mouse model fulfills these criteria (Dawson TM, 2010).

As mentioned, traditional genetic models of PD have been made in mouse. Until very recently, mouse has been the only real mammalian option for genetic manipulation. Compared to the rat, the mouse is smaller, exhibits a simpler yet more erratic behavioral profile, and on the whole is less predictive and translational. The advent of zinc finger nucleases (ZFNs), however, now enables genetic engineering in animals other than mouse, including the more predictable rat (Geurts AM, 2009). SAGE® Labs has now harnessed the species flexibility of ZFNs to produce knockout rat models of PD. Developed in collaboration with the Michael J. Fox Foundation, SAGE Labs has commercialized knockout rats for Lrrk2, Parkin, Pink1, and DJ-1.

Unlike the genetic mouse models, early work on knockout rat PD models has been encouraging. Pink1 and DJ-1 knockout rats display a progressive loss of dopaminergic neurons in the substantia nigra, with a ~50% loss by eight months of age (Figure 1). This is the first observation of dopamine cell loss in any genetic model of PD. Coupled with the loss of dopaminergic neurons are moderate to severe motor deficits. Specifically, the Pink1 and DJ-1 knockout rats show impaired performance on the tapered balance beam and hind limb fatigue assays, as well as gait abnormalities as assessed by NeuroCube (Figure 2). A subpopulation (~30%) of Pink1 and DJ-1 knockout rats also show a marked, severe dragging of the hind limbs with an onset of around five months of age.

Initial results have demonstrated that the Pink1 knockout rats also possess non-motor and non-dopaminergic characteristics of PD. Anosmia, or the loss of the sense of smell, is frequently reported as an early symptom in human PD patients. Awake Pink1 knockout rats displayed deficits in olfactory circuits in response to almond odor (previously demonstrated as pleasurable (Kulkarni P, 2012)) as assessed by fMRI. The fMRI also revealed lack of activation of downstream centers suggesting anhedonia, or lack of pleasure, another symptom frequently seen in PD. Diffusion tensor imaging (DTI) in Pink1 KO rats further revealed altered DTI signals not only in the expected dopaminergic regions, but in non-dopaminergic regions as well. The rat is particularly well suited for imaging studies such as these, as its increased size offers higher resolution as compared to mouse.

In all, the PD rat models from SAGE Labs offer exciting new tools for PD researchers. They allow study in the more preferable and translational rat, and exhibit PD phenotypes not seen in the mouse. These models allow for unprecedented access to study PD-relevant neurodegeneration of dopaminergic neurons in an intact animal. Additionally, unlike chemically-induced models, they allow for the study of PD progression and pathology, an aspect that is absolutely critical to develop therapeutics that can be used to halt disease progression or even prevent PD outright.

Figure 1. Progressive dopamine cell loss in Pink1 and DJ-1 knockout rats.
Pink1 and DJ-1 knockout rats show a ~50% loss in tyrosine hydroxylase-stained neurons at eight months of age as determined by immunohistochemistry and stereology.
Data source: The Michael J. Fox Foundation

Reference: