

Supplementary Table 1 Summary of completed drug trials for disease-modification in Parkinson's disease that have explored the use of biomarkers.

Molecular target	Mechanism of action	Drug(s)	Biomarker supporting target engagement	Biomarker relationship with primary outcome	Biomarker result
Alpha-Synuclein	Inhibition of α -synuclein aggregation (autophagy-cAbl inhibitors)	Nilotinib ¹ NCT02954978	CSF total & oligomeric alpha-synuclein & ratio, Dopamine metabolites (DOPAC, HVA) in CSF & Plasma, CSF total tau and P-tau 181, CSF-triggered receptors on myeloid cells levels, plasma	No correlation specified	Small amounts of nilotinib detected in the CSF, plasma DOPAC, but not CSF DOPAC or HVA level increased, reduction of α -synuclein oligomers, reduction of p-tau 181, decreased P-tau 181/total tau

		<p>Nilotinib² NCT03205488</p>	<p>and CSF pan-tyrosine Abl, and plasma and CSF tyrosine 412 Abl (activity)</p>		
			<p>Nilotinib levels (CSF & serum) and dopamine metabolites (catecholamine catabolites and pre-cursors, serotonin & metabolite) in CSF</p>	<p>No correlation specified</p>	<p>Poor CSF penetration (half-maximal inhibitory concentration), No change in dopamine metabolites</p>
	<p>Anti-α-synuclein antibody</p>	<p>Prasinezumab³ NCT03100149</p>	<p>DaT-SPECT Striatal Binding Ratio (SBR) in Putamen Ipsilateral to</p>	<p>No correlation specified</p>	<p>No DaT-SPECT change, 4 had antidrug antibodies</p>

			Clinically Most Affected Side, Blood Anti-Drug Antibodies		
		Cinpanemab ⁴ NCT03318523	CSF α -synuclein seeding amplification assay to confirm presence, DaT-SPECT imaging	No correlation specified	Plasma levels demonstrated dose proportionality, No DaT-SPECT change
		Affitope (PD01A) Active alpha synuclein immunization (NCT01568099, NCT01885494,	Blood & CSF PD01A specific antibody and total α -synuclein levels.	Nil (Safety & tolerability)	Detected vaccine-triggered antibodies in blood and CSF, trend toward lower levels of alpha-synuclein in blood and CSF

		NCT02216188, NCT02618941)			
LRRK2	LRRK2 inhibitors (promoting autophagy)	DNL201 ⁵ NCT03710707	CSF & plasma drug levels of drug, LRRK2 pS935& pRab10 in peripheral blood	Nil (Safety & tolerability)	LRRK2 pathway engagement and alteration of downstream lysosomal biomarkers, robust CSF penetration
	Antisense oligonucleotide for LRRK2 inhibition	BIIB094 ⁶ NCT03976349	Serum drug concentration, whole-blood phosphorylated serine 935 LRRK2, peripheral blood	Nil (Safety & tolerability)	CSF/unbound plasma concentration ratio~1; Whole-blood phosphorylated serine 935 LRRK2 ($\leq 98\%$), peripheral blood mononuclear cell phosphorylated threonine 73 pRab10 ($\leq 93\%$), cerebrospinal fluid

			mononuclear cell phosphorylated threonine 73 pRab10, cerebrospinal fluid total LRRK2, urine bis (mono-acylglycerol) phosphate		total LRRK2 ($\leq 50\%$), and urine bis (monoacylglycerol) phosphate ($\leq 74\%$)
Lysosomal function	Modulator of lysosomal enzyme β -glucocerebrosidase (GCase) activity	Ambroxol ⁷ NCT02941822 *GBA status explored as baseline predictor	CSF drug levels, GCase activity and protein levels and α -synuclein levels	No correlation specified	Drug detected in CSF, GCase activity decreased, GCase protein levels increased, α -synuclein concentration increased *GBA status no impact
		LTI-291 ⁸ NTR6598 (SAD) and NTR6705 (MAD)	CSF levels of drug and glycosphingolipids	Nil (Safety & tolerability)	Favorable pharmacokinetics, No glycosphingolipid changes *GBA status no impact

Inflammation		*GBA status explored as baseline predictor			
	Substrate reduction	Venglustat ⁹ NCT02906020 *only GBA mutation carriers recruited	CSF & plasma drug, glucosylceramide & glucosylsphingosine levels	No correlation specified	Dose proportional drug increase, Glucosylceramide decreased in treated group in CSF & plasma
	Altered T-cell lineage	Sargramostim ¹⁰ NCT01882010 NCT03790670	Immune phenotype and function, DNA methylation status, and gene and proteome analyses	No correlation specified	Blood Treg numbers, function, and hypomethylation of upstream FOXP3 DNA elements increased
	Enzyme myeloperoxidase (MPO) inhibition	AZD3241 ¹¹ NCT01527695	Plasma drug concentration & myelo-	Nil (Safety & tolerability)	Distribution volume of 11C-PBR28 binding to translocator protein reduced

			peroxidase activity Microglial PET imaging		Sparse blood samples limited analysis
	3-hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) inhibition	Lovastatin ¹² NCT03242499	F-dopa PET scan	No correlation specified	Percentage change in the striatal 18F-dopa uptake ratio deteriorated less in the lovastatin group
Mitochondrial function	Improved mitochondrial biogenesis	Pioglitazone ^{13,14} NCT01280123	Leukocyte PGC-1 α and target gene expression, plasma interleukin 6 (IL-6) and urine 8-hydroxy-deoxyguanosine (8OHdG)	Association of concentrations of biomarkers with change in total UPDRS score	Negative study Baseline or changes from baseline in biomarker levels not associated with rate of progression, Pioglitazone did not alter biomarker levels

	Increased urate, (antioxidant)	<p>Inosine¹⁵ NCT02642393</p> <p>Inosine¹⁶ NCT00833690</p> <p>*Serum urate below population median concentration (<5.8 mg/dL) used for inclusion</p>	<p>Serum urate, striatal dopamine transporter binding</p> <p>Serum & CSF urate</p>	<p>No correlation specified</p> <p>Nil (Safety & tolerability)</p>	<p>Negative study</p> <p>Sustained elevation in serum urate from treatment</p> <p>No difference in dopamine transporter binding from treatment</p> <p>Urate levels rose in serum and CSF of treated groups</p>
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	Improved mitochondrial function	Ursodeoxycholic acid ¹⁷ NCT03840005	ATP levels using ³¹ P MR spectroscopy	No correlation specified	Increase in free energy and inorganic phosphate levels in the treatment group
Iron	Iron chelation	Deferiprone ¹⁸ NCT02655315	Iron levels in substantia nigra and striatum (MRI T2* sequence), changes in brain-structure volume (three-dimensional T1 sequence), serum ferritin levels, density of DaT quantified with SPECT DaT, & serum prolactin	No correlation specified	Clinical worsening in treatment group Nigrostriatal iron content decreased more in the deferiprone group, No inverse correlation between brain-structure volumes and iron content, Iron content outside nigrostriatal pathway similar in trial groups, no difference DaT change, plasma ferritin decreased & plasma prolactin increased with deferiprone

Insulin resistance	GLP-1 receptor agonists (reduced insulin resistance, reduced inflammation and alpha-synuclein aggregation)	Exenatide ^{19,20} NCT01971242	Measures of insulin resistance (Akt, and mTOR signalling pathway) in serum neuronal derived exosomes, CSF drug levels, DATSCAN uptake	Association with UPDRS 3	Positive clinical outcomes Therapeutic drug levels in CSF Exenatide treatment augmented insulin resistance pathways Improvements in UPDRS 3 associated with downstream insulin resistance marker levels Trend to improvement of DATSCAN signal in exenatide group
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Abbreviations α -syn / α -synuclein; alpha-synuclein, ATP; adenosine triphosphate, c-Abl; Abelson tyrosine kinase, CNS; central nervous system, CSF; cerebrospinal fluid, GCase; DAT-SPECT; Dopamine transporter single-photon emission computerized tomography imaging, DOPAC; 3,4-Dihydroxyphenylacetic acid Glucocerebrosidase, HVA; Homovanillic acid, LRRK-2; Leucine-rich repeat kinase 2, MDS-UPDRS; Movement Disorders Society-Unified Parkinson Disease Rating Scale, PD; Parkinson's disease, PGC-1 α ; Peroxisome proliferator-activated receptor-gamma coactivator.

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