REVIEW



Deep brain stimulation for monogenic Parkinson's disease: a systematic review

Tomi Kuusimäki^{1,2} • Jaana Korpela^{1,2} • Eero Pekkonen^{3,4} • Mika H. Martikainen^{1,2} • Angelo Antonini⁵ • Valtteri Kaasinen^{1,2}

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Abstract

Deep brain stimulation (DBS) is an effective treatment for Parkinson's disease (PD) patients with motor fluctuations and dyskinesias. The key DBS efficacy studies were performed in PD patients with unknown genotypes; however, given the estimated monogenic mutation prevalence of approximately 5–10%, most commonly *LRRK2*, *PRKN*, *PINK1* and *SNCA*, and risk-increasing genetic factors such as *GBA*, proper characterization is becoming increasingly relevant. We performed a systematic review of 46 studies that reported DBS effects in 221 genetic PD patients. The results suggest that monogenic PD patients have variable DBS benefit depending on the mutated gene. Outcome appears excellent in patients with the most common *LRRK2* mutation, p.G2019S, and good in patients with *PRKN* mutations but poor in patients with the more rare *LRRK2* p.R1441G mutation. The overall benefit of DBS in *SNCA*, *GBA* and *LRRK2* p.T2031S mutations may be compromised due to rapid progression of cognitive and neuropsychiatric symptoms. In the presence of other mutations, the motor changes in DBS-treated monogenic PD patients appear comparable to those of the general PD population.

Keywords Parkinson's disease · Monogenic · Genetic · Deep brain stimulation

Introduction

Deep brain stimulation (DBS) provides symptomatic motor benefit for patients with advanced Parkinson's disease (PD) [1–4]. The benefit of symptom control through DBS surpasses that of optimal medical treatment in patients with

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☐ Tomi Kuusimäki tomi.kuusimaki@utu.fi

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- Division of Clinical Neurosciences, Turku University Hospital, Hämeentie 11, POB 52, 20521 Turku, Finland
- Department of Neurology, University of Turku, Turku, Finland
- Department of Neurology, Helsinki University Hospital, Helsinki, Finland
- Department of Clinical Neurosciences (Neurology), University of Helsinki, Helsinki, Finland
- Department of Neurosciences, University of Padua, Padua, Italy

motor fluctuations and dyskinesias, and it is a relatively safe treatment option for motor complications of idiopathic PD [1–5]. DBS is often performed in relatively early-onset PD, a population in which it has been estimated that at least 5–10% of cases are not sporadic, but may carry genetic mutations [6, 7]. Genetic cases often are phenotypically different compared to sporadic patients, and this factor may influence clinical outcome [6, 8].

Though DBS has demonstrated efficacy, randomized studies have been performed in PD patients without genetic characterization raising questions of suitability of various monogenic forms and their relevance in DBS outcome. It is known that medication effects may vary between different mutations. For example, patients with *PRKN* mutations generally are particularly prone to levodopa-induced dyskinesias, whereas patients with *LRRK2* mutations tend to show a normal sustained benefit for levodopa [8–11]. The effects of other antiparkinsonian drugs, such as rasagiline, may also be modulated by the genotype [12]. Given the variability in medication effects, it is conceivable that there are also differences in the treatment response to DBS in advanced monogenic PD. There are several case reports and small case series of DBS outcomes in patients with genetic



PD, but due to a lack of information synthesis, we performed a systematic review on the effects of DBS in genetic PD.

Methods

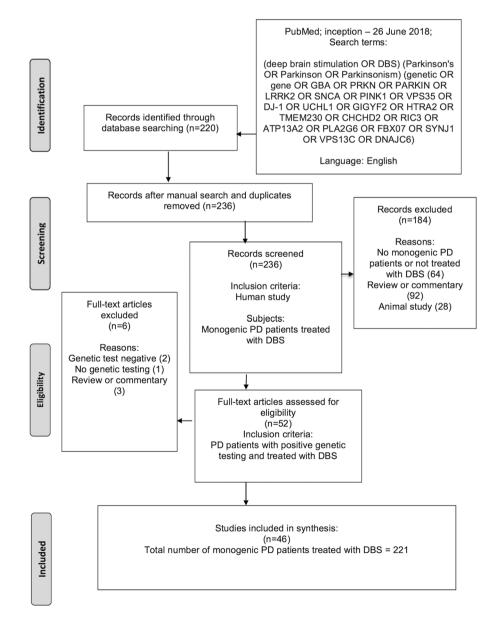
Search strategy

The Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) statement was followed [13]. We performed a PubMed search from inception to June 26, 2018 with keywords "deep brain stimulation or DBS", "Parkinson's or Parkinson or Parkinsonism" and "genetic or gene or GBA or PRKN or PARKIN or LRRK2 or SNCA or PINK1 or VPS35 or DJ-1 or UCHL1 or GIGYF2 or HTRA2 or

TMEM230 or CHCHD2 or RIC3 or ATP13A2 or PLA2G6 or FBX07 or SYNJ1 or VPS13C or DNAJC6". All original English language articles concerning genetic PD patients treated with DBS were included. Animal studies and review articles were excluded.

The initial search identified 220 articles, and we included an additional 16 relevant studies found in the manual search of reference lists (Fig. 1). All abstracts of these studies were screened, and 184 studies were excluded in the first round (no monogenic PD patients or not treated with DBS n = 64, review or commentary article n = 92, animal study n = 28). The remaining 52 studies were assessed fully for eligibility and six more studies were excluded in the second round (genetic test negative n = 2, no genetic testing n = 1, review or commentary article n = 3). Finally, 46 studies of these

Fig. 1 Flow chart of study inclusion and exclusion





236 studies met all selection criteria and were included in the systematic review (Table 1). A summary of the included studies is presented in Table 2. The included studies reported 221 genetic PD patients who were treated with DBS. However, two studies reported partially the same patients [14, 15].

Specific aims

This review of evidence aimed to systematically investigate DBS outcome in monogenic PD compared to the general PD population. The primary aim was to evaluate the motor benefit of the DBS operation in each monogenic PD type. An additional aim was to evaluate effects on non-motor symptoms, including possible cognitive and neuropsychiatric symptoms.

Selection criteria

Search terms and the PubMed search were planned by two authors (T.K. and V.K.). All titles and abstracts were reviewed by one investigator (T.K.). Studies were excluded if the title and/or abstract were not suitable for the aim of the review. Full texts were obtained for appropriate studies or if the relevance of an article was uncertain. The inclusion criteria for the selected studies were as follows: (1) a human study, (2) genetic PD patients treated with DBS, and (3) English language. The data extracted from each study were study year, first author's family name, number of patients, mutated gene, specific mutation, patient age at disease onset and DBS implantation, target nucleus of DBS, more specific lead positioning, pre- and postoperative UPDRS-III scores, follow-up time and outcome (Table 1). UPDRS-III scores of control cohort's (mutation non-carriers, NC) are also reported in Table 1 if the information was available. In the outcome evaluation, an improvement of 30% or more in the UPDRS-III motor score was considered to indicate favourable outcome; 20-30%, moderate outcome; and <20%, poor/mild outcome [58–60].

Quality control

The quality of the included studies was evaluated according to the Newcastle-Ottawa Scale (NOS) [61]. NOS includes selection, comparability, and exposure or outcome. The scale ranged from 0 to 11 stars, with the highest rating representing the greatest quality. Six months or more was a limit for the adequate follow-up time. Pre- and postoperative evaluation was thought to be accomplished if the outcome was reported properly with percentage improvement of the UPDRS-III score or verbally. A total score of 0–3 was considered to indicate to poor quality; 4–7, moderate quality; and 8–11, good quality. The NOS total score is presented in

Table 1 and the scale is presented more accurately in Supplementary Table 1. A summary of the assessed quality of the studies is presented in Supplementary Table 2.

Results

A summary of the primary results is presented in Table 2. Altogether, 46 studies and 221 monogenic PD patients treated with DBS were included in the systematic review (Table 1).

LRRK2

Seventeen studies [9, 15–30] reported 87 patients (target: subthalamic nucleus (STN) n=79, not available (NA) n=8). The outcome was reported in 73 patients (83.9% of patients); with percentage improvement of the UPDRS-III score in 49 patients and verbally in 24 patients. The motor outcome was mostly favourable in patients with LRRK2 mutation. Only five studies with ten patients reported poor/mild/moderate outcomes. Both patients with the p.T2031S (c.6091A>T) mutation (n=2) developed neuropsychiatric problems 5–7 years after implantation. The outcome appeared poor in patients with p.R1441G (c.4321C>G) mutations whereas it appeared excellent in patients with p.G2019S (c.6055G>A) mutations.

PRKN

Eighteen studies [11, 15, 16, 19, 21, 31–43] reported 67 patients (STN n=51, globus pallidus interna (GPi) n=5, zona incerta n=1, NA n=10). The outcome was reported in 57 patients (85.1%); UPDRS-III percentage improvement was reported in 45 patients and the outcome was described verbally in 12 patients. Fifty-one patients (76.1%) had favourable long-term motor outcomes. Six patients in three different studies were reported to have modest or poor outcomes.

GBA

Five studies [14, 15, 19, 44, 45] reported 50 patients (STN n = 33, GPi n = 4, ventral intermediate nucleus (VIM) n = 1, NA n = 12). Samples partially consisted of same patients in two studies [14, 15]. The outcome was reported in 30 patients (60.0%); UPDRS-III percentage improvement in 28 patients and the outcome was described verbally in 2 patients. Eighteen patients were reported to have favourable, three patients moderate and nine patients poor long-term motor outcomes. One study reported better outcomes with STN-DBS and VIM-DBS than with GPi-DBS



 Table 1
 The data extracted from the included studies

			Miliallo	AAO."	AAD.	larger	1	PKE-UPDKS III	FOST-UPDRS III	9/	<u> </u>	S	Outcome
-	18	LRRK2	p.G2019S	NA	NA ^A	STN°	NA A	NA	NA	NA	NA	4	Good or excellent $(n=8)$, moderate $(n=2)$, poor $(n=2)$ and NA $(n=6)$
_	15	LRRK2	p.G2019S	40.1 ± 9.4	NA	STN bilat.	+	55.8 ± 16.4 M-, 25.0 ± 13.2 M+ (NC: 51.7 ± 14.4 M-)	27.3±20.6 M-S+, 19.7±18.8 M+S+ (NC: 38.5±16.6 M- S+)	51.1 (NC: 25.5)	2	10	Favourable and better outcome compared to patients without mutation
	13	LRRK2	p.G2019S	49.5 ± 6.8	61.1 ± 6.6	STN bilat.	+	42.5±11.8 M-, 19.5±13 M+ (NC: 43.4±12.3 M-)	Short FU 28.5±13.1 M—S+, 17.4±12.9 M+S+ Long FU 30.5±12.8 M—S+, 21.2±9.2 M+S+ (NC: Short FU 27.2±14.1 M— S+, Long FU 33.9±16.1 M—S+)	Short FU 32.8±31.1 Long FU 28.5±32.9 (NC: Short FU 35.6±25.3, Long FU 17±37.1)	0.5-1 $(n = 13),$ 3 $(n = 11)$	10	Favourable and comparable to patients without mutations. One patient reported new/worse psychiatric symptoms at 3-year follow-up
	6	LRRK2	p.G2019S $(n = 7)$ p.G2019S + het. <i>PRKN</i> mutation $(n = 1)$, p.T2031S $(n = 1)$	33-48	38–65	STN bilat.	NA A	41.4±12.4 M-, 8.2±4.6 M+ (NC: 43.4±17.0 M-)	47.7±13.1 M-S-, 17.8±9.6 M-S+, 11.8±4.5 M+S-, 6.2±3.9 M+S+ (NC: 15.7±9.0)	50±36 (NC: 64)	9–10 (Long- term FU for two patients)	10	Favourable and comparable to patients without mutations, but cognitive, behavioral and psychotic problems in the patient with p.T2031S mutation after 5 years
	'n	LRRK2	₹X	(n=4) $(n=4)$	60.8 ± 9.0 $(n = 4)$	₹ _Z	NA A	₹z	$30.8 \pm 11.7 \text{ M+S+}$ $(n=4)$	∀ Z	3.5 ± 2.4 $(n = 4)$	9	The outcome is not reported. Clinical data before DBS is not available, but UPDRS III score was higher in LRRK2 -patients compared to patients without mutations at follow-up
Angeli et al. [15]	ς.	LRRK2	p.G2019S (n = 4), p.G2019S + GBA-E326K (n = 1)	35–55	NA^B	STN^c	NA A	65.4 ± 14.9 M-, 10.8 ± 5.1 M+ (NC: 47.6 ± 14.8 M-)	69.2 ± 12.4 M-S-, 30.6 ± 16.1 M-S+ (24.6 ± 11.3 M-S+)	53 (NC: 48)	<u>7</u>	6	Favourable and comparable to patients without mutations. No reported cognitive problems
ómez- Esteban et al. [20]	4	LRRK2	p.R1441G	29–55	41–65	STN bilat.	+	48.5 ± 18.5 M-, 18.0 ± 7.4 M+ (NC: 42.5 ± 10.6 M-)	39.7 ± 17.7 M–S+, 16.0 ±/-7.7 M+S+ (NC: 26.1 ± 8.4 M– S+)	18 (NC: 39)	0.5	10	Poorer response compared to patients without mutation
	8	LRRK2	p.G2019S	43–57	50–69	STN bilat.	+	NA for individual genes (NC: $35.7 \pm 6.7 \text{ M}-$)	NA for individual genes NA (NC: 44.8) (NC: 19.7±5.5 M – S+)	NA (NC: 44.8)	ς.	6	Favourable and comparable to patients without mutations



lable	(contint	ied)											
Study	N	Gene	Mutation ^a	AAOa	AAD ^a	Targeta	LP	PRE-UPDRS III ^a	POST-UPDRS III ^a	q%	FU	SON	Outcome
Lesage et.	al. 3	LRRK2	p.G2019S ($n=2$), p.T2031S	34-45	41–66	STN°	NA	NA 14 M+ $(n=1)$, NA 27 M-S+ $(n=1)$,	27 M $-$ S+ (n = 1),	NA	7 (Long-	6	Favourable to motor

Gene			Mutation ^a	AAOa	AAD^a	Target ^a	LP	PRE-UPDRS III ^a	POST-UPDRS IIIª	9%	FU	NOS	Outcome
3 LRRK2 p.G2019S ($n=2$), p.T2031S 3. ($n=1$)	p.G2019S ($n = 2$), p.T2031S ($n = 1$)		ý,	34-45	41–66	STIN°	NA	14 M + (n=1), NA $(n=2)$	27 M-S+ (n=1), 17 M-S+ and 32 M-S- (n=1)	NA A	7 (Long- term FU for one patient)	6	Favourable to motor symptoms, but depression and psychosis in the patient with p.T2031S mutation
3 LRRK2 p.G2019S 33-	p.G2019S		33-	33–62	NA	STN bilat.	NA	NA	NA	NA	NA	S	Favourable to motor symptoms
3 LRRK2 p.G2019S NA	p.G2019S		NA		NA	NA	NA	NA	NA	NA	NA	2	NA
1 LRRK2 p.R1441G and p.G2385R 28	p.R1441G and p.G2385R		28		39	STN bilat.	+	NA	€ Z	NA	7	٢	Poor motor response with severe psychiatric problems at 1 year after operation
1 LRRK2 Het. p.G2019S 49	Het. p.G2019S		49		56	STN bilat.	+	27 M-, 12 M+	25 M-S-, 8 M-S+, 5 M+S+	70.4	0.25	∞	Favourable outcome
1 LRRK2 p.N1437H (c.4309A > C) 50	p.N1437H (c.4309A > C)		20		3	STN bilat.	+	NA A	65 M-S+	ę K	0.5	∞	Poor motor outcome. Patient had also severe depression and suicidality and she finally committed suicide 6.5 months after DBS implantation
1 LRRK2 p.Y1699C 43		p.Y1699C 43	43		84	STN bilat.	X	54 M-, 32 M+	26 M-S+, 15 M+S+	52 M-, 53 M+	2.5	10	Favourable outcome. No changes in neuropsychological test parameters 6 months postoperatively
1 <i>LRRK2</i> p.R793M 42	p.R793M		42		09	STN bilat.	NA	NA	NA	64 (1 year), 56 (8 year)	∞	∞	Favourable outcome
1 LRRK2 p.Asn1437His NA	p.Asn1437His		NA		NA	STN^c	NA	NA	NA	NA	NA	4	Favourable outcome
14 PRKY One mutation: ex6hetdupl, 14–52 ex6hetdel, Arg256Cyshet [n=2], Ala398Thrhet, ex7hetdupl, and exhet3del; Hom. or compound het.: ex5hetdel—prom-ex lhetdel, ex24hetdupl—ex3hetdel, Cys289Glylon, ex5hetdel— Cys4Harghet, ex2hetdel— ex3hetdel— ex3hetde	One mutation: ex6hetdupl, ex6hetdel, Arg256Cyshet [n=2], Ala398Thrhet, ex7hetdupl, and exhet3del; Hom. or compound het.: ex5hetdel—c.253delAhet, ex3hetdel—prom-ex1hetdel, ex24hetdupl—ex3hetdel—Cys289Glyhom, ex5hetdel—Cys241 Arghet, ex2hetdel—ex3hetdel—ex3hetdel—ex3hetdel—ex3hetdel—fys241 Arghet, ex2hetdel—ex3hetdel—fys241 Arghet, ex2hetdel—fys7-1GC	t t, t, del, l, el—	14-52		32–67	STN bilat.	NA	One mutation 54.3 ± 13.9 M-, 11.6 ± 12.7 M+ Two mutations 55.4 ± 17.3 M-, 14.5 ± 10 M+ (NC: 51.9 ± 18.3 M-)	One mutation 38.4±16.8 M-S- 12.7±11.2 M+S-, 17.8±11.2 M-S+, 10.8±10.1 M+S+ Two mutations 47.7±12.8 M-S-, 17±10.9 M-S+, 14.5±12.5 M- S+, 9.3±8.6 M+S+ (NC: 17.9±15.1 M- S+)	One mutation 69±15 Two mutations 77±14 (MC: 65.5)	1–2 except 3 years for one patient with two PRKN muta- tions	10	Motor response was favourable and comparable to patients without mutations, but more cognitive problems in homozygous and compound heterozygous patients compared to patients without mutations



Table 1 (continued)	ontinue	(þ.											
Study	N	Gene	Mutation ^a	AAO^a	AAD^a	Target ^a	LP	PRE-UPDRS IIIª	POST-UPDRS IIIª	%p	FU	SON	Outcome
Moro et al. [32]	11	PRKN	One mutation: delEx6, duplEx5, 867C > T, 1306G > C, delEx5-12; Hom. or compound het:: 202delA [<i>n</i> =2], delEx3-4, delEx3+1142-3delGA, delEx2-5+duplEx8, delEx7-9	15–40	31–66	STN bilat.	NA	35–66 (MV = 49.5)	NA	Short FU 36 Long FU 42 (NC: Short FU 56, Long FU 44)	3-6	6	Favourable and comparable to patients without mutations in long-term follow-up
Pal et al. [19]	10	PRKN	₹	30.6±9.1	47.0±11.5	₹ Z	NA	∀ Z	m = 6 M+S+ $m = 6$	⋖ Z	4.0±4.2	9	The outcome is not reported. Clinical data before DBS is not available but UPDRS III score was higher in PRKN -patients compared to patients without mutations at follow-up
Angeli et al. [15]	v	PRKN	Hom: c.101_102delAG, c.1289G> A p.G430D and c.823C> T, p.Arg275Trp, c.337_376del and c.465— 46del, Hom. deletion of exon 3 and 4, c.823C> T; p.Arg275Trp and het. duplica- tion of exon 6	7–36	ZA ^B	GPi $(n=3)$, STN° $(n=2)$	ę z	All 57.0±11.2 M-, 21.0±6.4 M+ GP ₁ GP ₂ 33.3±13.9 M- STN 62.5±3.5 M- (NC: STN: 47.6±14.8 M-GP ₁ : 40.5±13.4 M-)	GPi 43.3±16.4 M-S- 42.0±19.0 M-S+ 27.3±17.6 M+S+ STN 84.0±22.6 M-S- 43.0±0.0 M-S+ 23.5±6.4 M+S+ (NC: STN: 24.6±11.3 M-S+, GPi: 51.0±7.1 M-S+)	GPi 21 STN 31 (NC: STN: 48, GPi: -28)	2.	0.	Good to motor symptoms without cognitive problems. The percentage improvement in the UPDRS III score was better with STN-DBS than with GPi-DBS
Romito et al. [11]	ς.	PRKN	G828A and Dupl ex1, DelAG 202-203, C1101T, G535A, Dupl ex1	27–45	42–63	STN bilat.	+	57.3 ± 9.3 M – 22.8 ± 7.3 M + (NC: 59.7 ± 11.3 M –)	25.2±10.0 M-S+ 21.8±7.5 M+S+ (NC: 29.0±12.3 M- S+)	56 (NC: 51.4)	1–3	10	Favourable and comparable to patients without mutations
Johansen et al. [21]	4	PRKN	Het. c.delEx3, Het. p.R275W, Het. c.duplEx7, Hom. c.delEx5 (GPi)	35–46	50–59	STN bilat. $(n=3)$, GPi unilat. $(n=1)$	+	NA for individual genes (NC: $35.7 \pm 6.7 \text{ M}-)$	NA for individual genes NA (NC: 44.8) (NC: 19.7±5.5 M— S+)	NA (NC: 44.8)	5-7	6	Favourable and comparable to patients without mutations
Kim et al. [33]	ю	PRKN	٧×	21.7 ± 8.5	49.7 ± 16.2	STN bilat.	N A	49.8±24.5 M-, 18.3±7.8 M+ (NC: 38.3±10.6 M-)	24.7±14.0 M-S+, 22.2±14.9 M+S+ (NC: 17.2±5.5 M- S+)	37.1 ± 45.4 (NC: 54.6 \pm 13.9)	2–5	10	Favourable and comparable to patients without mutations
Hassin-Baer et al. [34]	ю	PRKN	Hom. 202 A deletion	15–28	31–54	STN°	NA A	27–64 M–, 20–48 M+	NA	Ą V	NA A	L	Modest outcome with improvement in appendicular symptoms, but no change in axial features



Table 1 (continued)

Study	Ν	Gene	Mutation ^a	AAO^a	AAD^a	Target ^a	LP	$\rm PRE-UPDRS~III^a$	POST-UPDRS IIIa	9%	FU	NOS	Outcome
Sayad et al. [16]	2	PRKN	Het. c. 458C > G	48	NA	STN bilat.	+	46 M-, 28 M+	51 M-S+, 30 M+S+	-10.1	2	10	Poor response
			Het. c. 1204C > T	48			+	49 M-, 32 M+ (NC: 51.7±14.4 M-)	51 M-S+, 47 M+S+ (NC: 38.5 ± 16.6 M- S+)	-4.1 (NC: 25.5)			
Thompson et al. [35]	7	PRKN	Hom., specific mutation NA	26 (Gpi), 30 (STN)	NA	STN bilat. $(n=1)$, GPi bilat. $(n=1)$	NA	GPi 57 M-, 50 M+ STN 47 M-, 21 M+	NA A	NA	3 (STN), 8 (GPi)	9	Favourable outcome
Genç et al. [36]	-	PRKN	Het. c89G > A and large het. deletion	10	NA	STN bilat.	+	48 M-, 7 M+	7 M-S+, 4 M+S+	85.4	NA	9	Favourable to motor symptoms
Moll et al. [37]	-	PRKN	Compound het. <i>PRKN</i> mutation (delExon1 + c.924C > T)	35	45	STN bilat.	+	30 M- 5 M+	NA	NA	NA	7	Favourable to motor symptoms
Nakahara et al. [38]	-	PRKN+ PINKI	Hom. parkin mutation (p.T175PfsX2) + het. PINK1 mutation (p.R58-V59insGR)	15	09	STN bilat.	+	86 M-, 25 M+	33 M-S+, 21 M+S+	62	0.7	6	Favourable outcome
Lefaucheur et al. [39]	-	PRKN	Compound het. mutations of the <i>PRKN</i> gene, [c.101_102delAG (p.Gln34ArgfsX5)+c.155delA (p.Asn52MetfsX29)]	25	69	STN°	NA	NA	NA A	55	0.5	∞	Favourable to motor symptoms without cognitive problems
Wickre- maratchi et al. [40]	-	PRKN	Compound het. exon 2/exon 2 1 3 deletion in the <i>PRKN</i>	&	46	Zona incerta bilat.	N A	68 M-, 22 M+	NA M-S+24 M+S+	NA M-, 64.7 M+	0.5	6	Favourable outcome
Lesage et al. [41]	-	PRKN	Compound het. of the <i>PRKN</i> c.1-? $^{-7}$ +?del and c.172-? $^{-4}$ 12+?del mutations	&	39	STN bilat.	NA	46 M-, 15.5 M+	NA	NA	NA	9	Favourable outcome
Capecci et al. [42]	-	PRKN	Hom. deletion in exon 3	22	NA	STN bilat.	+	45 M-, 5 M+	7 M-S+, 3 M+S+	84.4	-	∞	Favourable outcome
Khan et al. [43]	-	PRKN	Exon 9 1101C->T (Arg334Cys), exon 7 939G->A (Asp230Asn)	30	35	STN bilat.	NA	NA	NA	NA	NA	9	Favourable outcome



±4.5 STN° NA 52.4±13.0 M-, (n=15), 18.4±14.9 GP; M+(NC: (n=2) 40.5±12.0 M-)	.4±5.8 53.5±4.5	1.4. 4.4.	Het. mutation carriers (n = 15), hom. mutation carrier (n = 1), compound het. (n = 1). Two patients also carried a mutation in another PD-associated gene; PARKIN or LRRK2	GBA Het. mutation carriers (n = 15), hom. mutation carrier (n = 1), compound het. (n = 1). Two patients also carried a mutation in another PD-associated gene; PARKIN or LRRK2	Het. mutation carriers (n = 15), hom. mutation carrier (n = 1), compound het. (n = 1). Two patients also carried a mutation in another PD-associated gene; PARKIN or LRRK2
			,		
NA All $51.3 \pm 14.0 \mathrm{M}_{-}$, $18.0 \pm 15.4 \mathrm{M}_{+}$ GPi $64.5 \pm 21.9 \mathrm{M}_{-}$ STN $50.5 \pm 12.4 \mathrm{M}_{-}$ VIM $35.0.5 \pm 12.4 \mathrm{M}_{-}$ VIM $35.0.5 \pm 12.4 \mathrm{M}_{-}$ VIM $35.0.5 \pm 13.4 \mathrm{M}_{-}$ GPi: $47.6 \pm 14.8 \mathrm{M}_{-}$ GPi: $40.5 \pm 13.4 \mathrm{M}_{-}$)	STN° ($n = 13$), GPi ($n = 2$), VIM ($n = 1$)	34–58 NA ^B STN ^c ($n=13$) GPl GPl ($n=2$), VIM ($n=1$)	NA ^B	GBA R463CR463C, L444PIE326K, 34–58 NA ^B S7 N370S, D409H, recNcil, R463C, N188S, R275Q, IVS2+1 G>A, L444P, E326KB326K, E326K (n=3), E326K and LRRK2 p.G2019S, T369M and PRKN c.1310C>T	R463C/R463C, 1444P/E326K, 34–58 NA ^B S7 N370S, D409H, recNcil, R463C, N188S, R275Q, IVS2+1 G> A, L444P, E326K/E326K, E326K (n=3), E326K and <i>LRRK</i> 2 p.G2019S, T369M and <i>PRKIV</i> c.1310C> T



lable 1	continue	(pa											
Study	N	Gene	Mutation ^a	AAO^a	AAD^a	Target ^a	LP	PRE-UPDRS III ^a	Target ^a LP PRE-UPDRS III ^a POST-UPDRS III ^a	q%	FU	NOS	NOS Outcome
Pal et al. [19]	12	GBA	p.N370S $(n=8)$, p.L444P $(n=3)$. I patient carried both GBA	41.6 ± 5.3 53.9 ± 2 $(n=11)$ $(n=9)$	53.9 ± 2.6 NA $(n=9)$	NA	NA	NA NA	$27.4 \pm 14.5 \text{ M+S+}$ (n = 11)	NA	1.6 \pm 3.0 6 ($n = 9$)	9	The outcome is not reported. Clinical

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Study	۷ ا	Gene	Mutation ^a	AAOª	AAD ⁴	Targeta	T.	PRE-UPDRS III"	POST-UPDRS III"	.%	FU	SON	Outcome
Pal et al. [19]	12	GBA	p.N370S (n=8), p.L444P (n=3). I patient carried both GBA and LRRK2 mutations and was excluded	(n=11)	53.9 ± 2.6 $(n = 9)$	₹ Z	NA	₹ Z	$27.4 \pm 14.5 \text{ M+S+}$ $(n = 11)$	₹ Z	(n=9)	9	The outcome is not reported. Clinical data before DBS is not available, but UPDRS-III score was little higher in GBA -patients compared to patients without mutations at follow-up
Weiss et al. [44]	м	GBA	p.N370S ($n = 1$) and p.L444P ($n = 2$)	47–54	65–69	SIN	e Z	26 and 53 M-, 14 and 19 M+, NA (n=1) (NC: 31–63 M-)	56-71 M-S-, 21-45 M-S+, 32-48 M+S-, 20-45 M+S+ (NC: 21-42 M-S+)	30-75 (NC: 22-54)	0-10	Ξ	Favourable outcome, but substantial increase of axial motor impairment in the long-term with declining therapeutic response in <i>GBA</i> carriers. <i>GBA</i> carriers developed also a significant cognitive impairment
Lesage et al. [45]	2	GBA	Hom. p.N370S	52	NA	STN bilat.	NA	NA	NA	NA	NA	S	Favourable outcome
			c.1263del + Rec TL	21	24						2		Some clinical benefit 2 years after DBS, but problems with postural instability
Martikainen et al. [46]	-	SNCA	Het. c.158C> A (p.A53E)	24	46	STN bilat.	₹ Z	31 M-, 8 M+	Ϋ́ Y	Ž	ج. د	6	Favourable motor outcome in the short-term but poor in the long-term follow-up. Response for motor fluctuations remained satisfactory but the cognitive and mental state of the patient deteriorated to a state of practical immobility
Perandones et al. [47]	_	SNCA	SNCA duplication	18	26	GPi bilat.	+	NA	NA	NA	0.1	9	Favourable and comparable to patients without mutations
Shimo et al. [48]	-	SNCA	SNCA duplication	35	41	STN bilat.	+	27 M-, 10 M+	13 M-S+	51.9	4	6	Favourable motor outcome without cognitive or psychiatric problems



Study	N	Gene	Mutation ^a	AAO^a	AAD^a	Target ^a	LP	$\rm PRE\text{-}UPDRS~III^a$	$ m POST-UPDRS~III^a$	9%	FU	NOS	Outcome
Antonini et al. [49]	-	SNCA	SNCA duplication at 4q22.1	14	46	STN bilat.	+	28 M-, 10 M+	16 M-S+, 10 M+S+	42.9	0	6	Favourable outcome in short-term follow-up but patient developed visual hallucinations and cognitive deterioration and died two years after operation due to metastatic breast
Ahn et al. [50]	1	SNCA	SNCA duplication	40	46	STN bilat.	X	32 M-, 6 M+	e N	N A	NA A	9	carlocal Excellent motor response but later patient's dementia worsened, requir- ing assistance in daily activities
Fleury et al. [51]	2	VPS35	p.D620N	49	09	STN bilat.	NA A	58 M-, 17 M+	32 M–S–, 18 M–S+, 18 M+S–, 15 M+S+	76 (1 year) 69 (8 years)	∞	∞	Favourable outcome
				84	55			28 M-, 15 M+	₹ Z	36 (1 year)	-		Tremor, akinesia and rigidiy improved markedly but patient's walking difficulties worsened with an increased frequency of freezing episodes and falls after surgery (problems disappeared after levodopa intake with the STN-DBS switched on)
Chen et al. [52]	1	VPS35	p.D620N	45	55	STN bilat.	+	42 M-, 15 M+	35 M-S-, 22 M-S+, 15 M+S-, 13 M+S+	37	S	6	Favourable outcome
Kumar et al. [53]	-	VPS35	p.D620N	N A	NA A	N A	Š	NA	∀ Z	NA	NA	ε	Little benefit to motor symptoms, but patient devel- oped significant dysarthria
Sheerin et al. [54]	-	VPS35	p.D620N	47	NA	NA	N A	NA	NA	NA	NA	'n	Favourable outcome. No reported cognitive problems
Borellini et al. [55]	1	PINKI	Hom. L347P	30	49	GPi	NA	44 M-	32 M+S+	27	0.1	7	Moderate outcome
Nakahara et al. [38]	-	PRKN+ PINKI	Hom. parkin mutation (p.T175PfsX2) + het. PINK1 mutation (p.R58-V59insGR)	15	09	STN bilat.	+	86 M-, 25 M+	33 M-S+, 21 M+S+	62	0.7	6	Favourable outcome



Table 1 (continued)

Study	N	Gene	Mutation ^a	AAOa	AAD ^a	Target ^a	LP	LP PRE-UPDRS III ^a	POST-UPDRS III ^a	q%	FU NO	NOS Outcome	ıtcome
Johansen et al. [21]	_	PINKI	Het. p.G411S	20	59	STN bilat. +		NA for individual genes (NC: 35.7±6.7 M-)	NA for individual genes NA (NC: 44.8) (NC: 19.7±5.5 M- S+)	NA (NC: 44.8)	6 6		Favourable and comparable to patients without mutations
Moro et al. [32]	П	PINKI	Hom. c.509T > G (p.V170G)	31	19	STN bilat. NA		35.5 M-	Ϋ́ Y	Short FU 46.5 Long FU 43.7 (NC: Short FU 56, Long FU 44)	3-6		Favourable and comparable to patients without mutations
Valente et al. [56]	-	PINKI	₹ Z	Ϋ́ Υ	e z	STN bilat. NA NA	X A	₹ Z	₹	₹	8 A	Σ	Motor outcome was not properly reported but patient developed imbalance, gait impairment, dysarthria, and behavioral changes at the age of 54 years. Mental deterioration was documented a few years later
Dufournet et al. [57]	8		22q11.2 Del. Syndrome	34–38°	NA	STN^{c} $(n=1)$ $GPi (n=2)$	NA NA	NA	NA	30–70	NA 7	茁	Favourable and comparable to patients with idiopathic PD

AAO age at disease onset (years), AAD age at DBS operation (years), LP specific lead position (reported or not), % the percentage improvement of the UPDRS III score after DBS^b, FU followup after surgery (years), NA not available, M-/+ medication OFF/ON, S-/+ stimulation OFF/ON, MV mean value, NC mutation non-carriers

^AThe mean time from PD onset to surgery was 11.4 years (SD 6.2), ^B mean duration of PD (years) at DBS assessment: PRKN = 25.2 ± 12.8, GBA = 11.2 ± 5.0, LRRK2 = 12.1 ± 1.8

^aParameters are reported in the table as in the original articles



^bIf the percentage improvement was not reported directly in the original article but UPDRS-III scores were available, we calculated the percentage improvement from the change of UPDRS-III score in the preoperative M - condition compared to the postoperative M-S+ condition (((Pre-op. UPDRS-III M-) - (Post.op. UPDRS-III M-S+))/(Pre-op. UPDRS-III M-)×100)

^cThe study did not specify whether the implantation was uni- or bilateral

^dSome patients were reported previously by Angeli et al. [15]

^eAge at PD diagnosis

Table 2 Summary of key findings according to the mutated gene

Gene	Studies (n)	Patients (n)	Target	Outcome
LRRK2	17	87ª	STN: n = 79 (90.8%) NA: n = 8 (9.2%)	Mostly favourable motor outcome. Four studies with eight patients (9.2%) reported poor motor outcomes and one study reported moderate outcomes for two patients. Both patients with the <i>LRRK2</i> p.T2031S (c.6091A>T) mutation (<i>n</i> =2) developed neuropsychiatric problems 5–7 years after implantation. The outcome appears poor in patients with <i>LRRK2</i> p.R1441G (c.4321C>G) mutations (<i>n</i> =5), whereas it appears excellent in patients with <i>LRRK2</i> p.G2019S (c.6055G>A) mutations
PRKN	18	67 ^b	STN: $n = 51$ (76.1%) GPi: $n = 5$ (7.5%) Zona incerta: $n = 1$ (1.5%) NA: $n = 10$ (14.9%)	Fifty-one patients (76.1%) had favourable long-term motor outcomes. Four patients (6.0%) were reported to have modest outcome in two different studies and one study with two patients (3.0%) reported poor benefit
GBA	5	50°	STN: <i>n</i> = 33 (66.0%) GPi: <i>n</i> = 4 (8.0%) VIM: <i>n</i> = 1 (2.0%) NA: <i>n</i> = 12 (24.0%)	Eighteen patients were reported to have favourable, three patients moderate and 9 patients poor long-term motor outcomes. One study reported better outcomes with STN-DBS and VIM-DBS than with GPi-DBS. <i>GBA</i> mutation carriers developed cognitive impairment faster than patients without mutations
SNCA	5	5	STN: <i>n</i> = 4 (80.0%) GPi: <i>n</i> = 1 (20.0%)	Favourable motor outcome but three of five patients developed cognitive or neuropsychiatric problems a few years after implantation
VPS35	4	5	STN: <i>n</i> = 3 (60.0%) NA: <i>n</i> = 2 (40.0%)	Favourable motor outcome in four cases and minor motor benefit complicated by dysarthria in one case
PINK1	5	5 ^b	STN: <i>n</i> = 4 (80.0%) GPi: <i>n</i> = 1 (20.0%)	Favourable motor outcome in three cases and moderate in one case
22q11.2.Del. Syndrome	1	3	STN: <i>n</i> = 1 (33.3%) GPi: <i>n</i> = 2 (66.6%)	Favourable motor outcome

STN subthalamic nucleus, GPi globus pallidus interna, VIM ventral intermediate nucleus, NA not available

[15]. GBA mutation carriers developed cognitive impairment faster than patients without mutations.

SNCA

Five patients were reported in five case reports [46–50] (STN n=4, GPi n=1). The motor outcome was favourable for all patients in the short-term but 3/5 patients developed cognitive and/or neuropsychiatric problems a few years after implantation. The percentage change in the UPDRS-III score was documented in two patients.

VPS35

Four studies [51–54] reported five patients (STN n=3, NA n=2). Favourable motor outcome was reported in four cases and minor motor benefit complicated by dysarthria in one case. The percentage change in the UPDRS-III score was reported in three patients.



Five case reports [21, 32, 38, 55, 56] including one patient in each report (STN n=4, GPi n=1) were reported. Favourable motor outcome was observed in three patients and moderate outcome in one case. One patient developed imbalance, gait impairment, dysarthria, and behavioral changes after operation and mental deterioration was documented a few years later.

Exclusion of poorer quality studies

Unfortunately, many studies (Table 1) lacked important information as shown in the Supplementary Table 1. Poorer quality studies have tendency for bias; therefore, in the Supplementary Table 3, data are presented after exclusion of poorer quality studies such as studies lacking the information about DBS target, pre- and postoperative evaluation, adequate follow-up time or outcome information. Furthermore, as Lythe et al. [14] and Angeli et al. [15] reported partly the same patients, we tested the conclusions also when the smaller study was excluded. Nevertheless, after the



^aOne patient had also PRKN mutation and one had GBA mutation

^bOne patient had both PRKN and PINK1 mutations

^cTwo studies reported partially same patients, but it was not possible to separate individual patients that were reported twice. One patient had also LRRK2 mutation and one had PRKN mutation

exclusion of these studies, the results remained essentially the same (Supplementary Table 4).

Discussion

We report the following key findings: (1) DBS outcome appears excellent in patients with *LRRK2* p.G2019S (c.6055G > A) mutations, good in patients with *PRKN* mutations and poor in patients with *LRRK2* p.R1441G (c.4321C > G) mutations, (2) the overall benefit of DBS in *SNCA*, *GBA* and *LRRK2* p.T2031S (c.6091A > T) mutations may be decreased due to rapid progression of cognitive and neuropsychiatric symptoms, and (3) in other mutations, the motor outcome in DBS-treated genetic PD patients appears generally comparable to that of sporadic PD patients.

A recent smaller review of 30 studies described the effects of DBS mainly in patients with LRRK2, PRKN and GBA mutations [62]. In the present PRISMA-compliant systematic review of 46 studies and 221 patients, the most comprehensive data were available for patients with LRRK2 and PRKN mutations. The combined evidence suggests that patients with LRRK2 mutations generally have a good response to DBS, and patients with the most common LRRK2 mutation, the p.G2019S mutation [7], may even have better outcome than the general PD population. However, the reported LRRK2 cases of p.R114G, p.T2031S and p.N1437H (c.4309A > C) mutation carriers appeared to have less favourable outcome. This interpretation is limited by the small number of reported DBS-treated cases of rarer LRRK2 mutations. For the PRKN mutations, the literature supports a view that patients with *PRKN* mutations are optimal candidates for DBS.

Apart from the LRRK2 and PRKN genes, the published literature concerning individual monogenic mutations and DBS is less comprehensive and the data are clearly limited with respect to both the number of patients and duration of follow-up. The available data are limited to five DBStreated patients with VPS35 mutation, and the patients have shown favourable sustained motor outcome in 4/5 cases. The available literature also suggests that most patients with mutations in GBA tend to achieve favourable long-term motor outcome from STN-DBS. Despite good motor outcome, GBA mutation carriers may develop cognitive impairment after DBS faster than patients without mutations. SNCA patients commonly develop cognitive and neuropsychiatric problems [8]. The literature supported a good motor outcome after DBS also in patients with SNCA mutations; however, 3/5 patients developed cognitive and neuropsychiatric problems a few years after DBS implantation. Indeed, the non-motor features of genetic PD may be a limiting factor in the overall benefit of DBS in some mutations, such as *SNCA* and *LRRK2* p.T2031S. While the motor benefit from DBS may initially be clear, the rapid non-motor progression may lessen the sum value for the quality of life. A recent study in *SNCA* A53T mutated rodents suggested that DBS may be neuroprotective [63]. Nonetheless, in human PD patients with *SNCA* mutations, the neuropsychiatric progression appears to be rapid despite DBS. The issue could be the level of damage at the time of implantation, and earlier DBS in these patients might possibly provide different outcomes.

Preoperative response to levodopa is the best single predictor of the postoperative outcome of DBS [64]. This indicator appears useful also in patients with monogenic mutations and the response was reported in practically all included studies. Another relevant predictor is the localization of DBS electrodes [65]. Unfortunately, there were studies, which did not report DBS targets and most studies lacked information about lead positioning. As the literature expands in the future, the effect of targets and lead positioning should be investigated in more detail. In most studies, STN was preferred over GPi as the target. Hence it remains ambiguous whether there are any relevant differences of clinical outcome between STN and GPi stimulation in monogenic PD. One study reported also a patient with VIM stimulation which is an unusual target for PD patients because VIM stimulation improves only tremor, not other PD symptoms [66, 67]. Finally, it is important to note that the genetic status may have a positive as well as a negative influence on outcome of surgery and this issue should be taken into consideration in the interpretation of DBS studies. For example, the EARLYSTIM trial was performed with young-onset PD patients [5] and there could have been an overrepresentation of PRKN patients in the sample.

In conclusion, monogenic PD patients have variable DBS outcomes depending on the mutated gene. Most patients benefit from STN-DBS, at least in the short-term; however, the current evidence does not support or is questionable for DBS implantation for patients with p.T2031S or p.R114G mutations in the *LRRK2* gene or mutations in the *SNCA* or *GBA* genes. The best outcome from DBS surgery appears to be in patients with *LRRK2* p.G2019S or *PRKN* mutations.

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Compliance with ethical standards

Research involving human participants and animals This manuscript does not contain clinical studies or patient data apart from those identified through literature search.

Conflicts of interest The authors declare that they have no conflict of interest.

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References

- Deuschl G, Schade-Brittinger C, Krack P et al (2006) A randomized trial of deep-brain stimulation for Parkinson's disease. N Engl J Med 355:896–908
- Obeso JA, Olanow CW, Rodriguez-Oroz MC et al (2001) Deepbrain stimulation of the subthalamic nucleus or the pars interna of the globus pallidus in Parkinson's disease. N Engl J Med 345:956–963
- Weaver FM, Follett K, Stern M et al (2009) Bilateral deep brain stimulation vs best medical therapy for patients with advanced Parkinson disease: a randomized controlled trial. JAMA 301:63-73
- Antonini A, Moro E, Godeiro C, Reichmann H (2018) Medical and surgical management of advanced Parkinson's disease. Mov Disord 33:900–908
- Schuepbach WM, Rau J, Knudsen K et al (2013) Neurostimulation for Parkinson's disease with early motor complications. N Engl J Med 368:610–622
- Kasten M, Marras C, Klein C (2017) Nonmotor signs in genetic forms of Parkinson's disease. Int Rev Neurobiol 133:129–178
- Deng H, Wang P, Jankovic J (2018) The genetics of Parkinson disease. Ageing Res Rev 42:72–85
- Puschmann A (2013) Monogenic Parkinson's disease and parkinsonism: clinical phenotypes and frequencies of known mutations. Parkinsonism Relat Disord 19:407–415
- 9. Healy DG, Falchi M, O'Sullivan SS, et al (2008) Phenotype, genotype, and worldwide genetic penetrance of

- LRRK2-associated Parkinson's disease: a case-control study. Lancet Neurol 7:583–590
- Alcalay RN, Mirelman A, Saunders-Pullman R et al (2013) Parkinson disease phenotype in Ashkenazi Jews with and without LRRK2 G2019S mutations. Mov Disord 28:1966–1971
- Romito LM, Contarino MF, Ghezzi D, Franzini A, Garavaglia B, Albanese A (2005) High frequency stimulation of the subthalamic nucleus is efficacious in Parkin disease. J Neurol 252:208–211
- Masellis M, Collinson S, Freeman N et al (2016) Dopamine D2 receptor gene variants and response to rasagiline in early Parkinson's disease: a pharmacogenetic study. Brain 139:2050–2062
- Moher D, Shamseer L, Clarke M et al (2015) Preferred reporting items for systematic review and meta-analysis protocols (PRISMA-P) 2015 statement. Syst Rev 4:1
- Lythe V, Athauda D, Foley J et al (2017) GBA-associated Parkinson's disease: progression in a deep brain stimulation cohort. J Parkinsons Dis 7:635–644
- Angeli A, Mencacci NE, Duran R et al (2013) Genotype and phenotype in Parkinson's disease: lessons in heterogeneity from deep brain stimulation. Mov Disord 28:1370–1375
- Sayad M, Zouambia M, Chaouch M et al (2016) Greater improvement in LRRK2 G2019S patients undergoing Subthalamic Nucleus Deep Brain Stimulation compared to non-mutation carriers. BMC Neurosci 17:6
- Greenbaum L, Israeli-Korn SD, Cohen OS et al (2013) The LRRK2 G2019S mutation status does not affect the outcome of subthalamic stimulation in patients with Parkinson's disease. Parkinsonism Relat Disord 19:1053–1056
- Schüpbach M, Lohmann E, Anheim M et al (2007) Subthalamic nucleus stimulation is efficacious in patients with Parkinsonism and LRRK2 mutations. Mov Disord 22:119–122
- Pal GD, Hall D, Ouyang B et al (2016) Genetic and clinical predictors of deep brain stimulation in young-onset Parkinson's disease. Mov Disord Clin Pract 3:465–471
- Gómez-Esteban JC, Lezcano E, Zarranz JJ et al (2008) Outcome of bilateral deep brain subthalamic stimulation in patients carrying the R1441G mutation in the LRRK2 dardarin gene. Neurosurgery 62:857–862 (discussion 862–853)
- Johansen KK, Jørgensen JV, White LR, Farrer MJ, Aasly JO (2011) Parkinson-related genetics in patients treated with deep brain stimulation. Acta Neurol Scand 123:201–206
- Lesage S, Janin S, Lohmann E et al (2007) LRRK2 exon 41 mutations in sporadic Parkinson disease in Europeans. Arch Neurol 64:425–430
- Gaig C, Ezquerra M, Marti MJ, Muñoz E, Valldeoriola F, Tolosa E (2006) LRRK2 mutations in Spanish patients with Parkinson disease: frequency, clinical features, and incomplete penetrance. Arch Neurol 63:377–382
- 24. Goldwurm S, Di Fonzo A, Simons EJ et al (2005) The G6055A (G2019S) mutation in LRRK2 is frequent in both early and late onset Parkinson's disease and originates from a common ancestor. J Med Genet 42:e65
- Hatano T, Funayama M, Kubo SI et al (2014) Identification of a Japanese family with LRRK2 p.R1441G-related Parkinson's disease. Neurobiol Aging 35:2656.e2617–2656.e2656.e2623
- Stefani A, Marzetti F, Pierantozzi M et al (2013) Successful subthalamic stimulation, but levodopa-induced dystonia, in a genetic Parkinson's disease. Neurol Sci 34:383–386
- Puschmann A, Englund E, Ross OA et al (2012) First neuropathological description of a patient with Parkinson's disease and LRRK2 p.N1437H mutation. Parkinsonism Relat Disord 18:332–338
- Perju-Dumbrava LD, McDonald M, Kneebone AC, Long R, Thyagarajan D (2012) Sustained response to deep brain stimulation in LRRK2 parkinsonism with the Y1699C mutation. J Parkinsons Dis 2:269–271



- Breit S, Wächter T, Schmid-Bielenberg D et al (2010) Effective long-term subthalamic stimulation in PARK8 positive Parkinson's disease. J Neurol 257:1205–1207
- Aasly JO, Vilariño-Güell C, Dachsel JC et al (2010) Novel pathogenic LRRK2 p.Asn1437His substitution in familial Parkinson's disease. Mov Disord 25:2156–2163
- 31. Lohmann E, Welter ML, Fraix V et al (2008) Are parkin patients particularly suited for deep-brain stimulation? Mov Disord 23:740-743
- Moro E, Volkmann J, König IR et al (2008) Bilateral subthalamic stimulation in Parkin and PINK1 parkinsonism. Neurology 70:1186–1191
- Kim HJ, Yun JY, Kim YE et al (2014) Parkin mutation and deep brain stimulation outcome. J Clin Neurosci 21:107–110
- Hassin-Baer S, Hattori N, Cohen OS, Massarwa M, Israeli-Korn SD, Inzelberg R (2011) Phenotype of the 202 adenine deletion in the parkin gene: 40 years of follow-up. Mov Disord 26:719–722
- Thompson AJ, Scholz SW, Singleton AB, Hardwick A, McFarland NR, Okun MS (2013) Variability in clinical phenotypes of heterozygous and homozygous cases of Parkin-related Parkinson's disease. Int J Neurosci 123:847–849
- Genç G, Apaydın H, Gündüz A et al (2016) Successful treatment of Juvenile parkinsonism with bilateral subthalamic deep brain stimulation in a 14-year-old patient with parkin gene mutation. Parkinsonism Relat Disord 24:137–138
- Moll CK, Buhmann C, Gulberti A et al (2015) Synchronized cortico-subthalamic beta oscillations in Parkin-associated Parkinson's disease. Clin Neurophysiol 126:2241–2243
- Nakahara K, Ueda M, Yamada K et al (2014) Juvenile-onset parkinsonism with digenic parkin and PINK1 mutations treated with subthalamic nucleus stimulation at 45 years after disease onset. J Neurol Sci 345:276–277
- Lefaucheur R, Derrey S, Guyant-Maréchal L, Chastan N, Maltête D (2010) Whatever the disease duration, stimulation of the subthalamic nucleus improves Parkin disease. Parkinsonism Relat Disord 16:482–483
- Wickremaratchi MM, Majounie E, Morris HR et al (2009) Parkinrelated disease clinically diagnosed as a pallido-pyramidal syndrome. Mov Disord 24:138–140
- Lesage S, Magali P, Lohmann E et al (2007) Deletion of the parkin and PACRG gene promoter in early-onset parkinsonism. Hum Mutat 28:27–32
- 42. Capecci M, Passamonti L, Annesi F et al (2004) Chronic bilateral subthalamic deep brain stimulation in a patient with homozygous deletion in the parkin gene. Mov Disord 19:1450–1452
- Khan NL, Graham E, Critchley P et al (2003) Parkin disease: a phenotypic study of a large case series. Brain 126:1279–1292
- Weiss D, Brockmann K, Srulijes K et al (2012) Long-term followup of subthalamic nucleus stimulation in glucocerebrosidase-associated Parkinson's disease. J Neurol 259::1970–1972
- Lesage S, Anheim M, Condroyer C et al (2011) Large-scale screening of the Gaucher's disease-related glucocerebrosidase gene in Europeans with Parkinson's disease. Hum Mol Genet 20:202–210
- Martikainen MH, Päivärinta M, Hietala M, Kaasinen V (2015) Clinical and imaging findings in Parkinson disease associated with the A53E SNCA mutation. Neurol Genet 1:e27
- Perandones C, Aráoz Olivos N, Raina GB et al (2015) Successful GPi stimulation in genetic Parkinson's disease caused by mosaicism of alpha-synuclein gene duplication: first description. J Neurol 262:222–223
- Shimo Y, Natori S, Oyama G et al (2014) Subthalamic deep brain stimulation for a Parkinson's disease patient with duplication of SNCA. Neuromodulation 17:102–103
- Antonini A, Pilleri M, Padoan A et al (2012) Successful subthalamic stimulation in genetic Parkinson's disease caused by duplication of the α-synuclein gene. J Neurol 259:165–167

- Ahn TB, Kim SY, Kim JY et al (2008) Alpha-synuclein gene duplication is present in sporadic Parkinson disease. Neurology 70:43-49
- Fleury V, Wider C, Horvath J et al (2013) Successful long-term bilateral subthalamic nucleus deep brain stimulation in VPS35 Parkinson's disease. Parkinsonism Relat Disord 19:707–708
- Chen YF, Chang YY, Lan MY, Chen PL, Lin CH (2017) Identification of VPS35 p.D620N mutation-related Parkinson's disease in a Taiwanese family with successful bilateral subthalamic nucleus deep brain stimulation: a case report and literature review. BMC Neurol 17:191
- Kumar KR, Weissbach A, Heldmann M et al (2012) Frequency of the D620N mutation in VPS35 in Parkinson disease. Arch Neurol 69:1360–1364
- Sheerin UM, Charlesworth G, Bras J et al (2012) Screening for VPS35 mutations in Parkinson's disease. Neurobiol Aging 33:838. e831–838.e835
- Borellini L, Cogiamanian F, Carrabba G et al (2017) Globus pallidus internus deep brain stimulation in PINK-1 related Parkinson's disease: a case report. Parkinsonism Relat Disord 38:93–94
- Valente EM, Salvi S, Ialongo T et al (2004) PINK1 mutations are associated with sporadic early-onset parkinsonism. Ann Neurol 56:336–341
- Dufournet B, Nguyen K, Charles P et al (2017) Parkinson's disease associated with 22q11.2 deletion: clinical characteristics and response to treatment. Rev Neurol (Paris) 173:406–410
- Stern MB, Marek KL, Friedman J et al (2004) Double-blind, randomized, controlled trial of rasagiline as monotherapy in early Parkinson's disease patients. Mov Disord 19:916–923
- Shulman LM, Gruber-Baldini AL, Anderson KE, Fishman PS, Reich SG, Weiner WJ (2010) The clinically important difference on the unified Parkinson's disease rating scale. Arch Neurol 67:64–70
- Rabie A, Verhagen Metman L, Fakhry M et al (2016) Improvement of advanced Parkinson's disease manifestations with deep brain stimulation of the subthalamic nucleus: a single institution experience. Brain Sci 6:58
- 61. Wells GA, Shea B, O'Connell D et al (2011) The Newcastle-Ottawa Scale (NOS) for assessing the quality of nonrandomized studies in meta-analysis. Available: http://www.ohri.ca/programs/clinical_epidemiology/oxford.asp
- Rizzone MG, Martone T, Balestrino R, Lopiano L (2018) Genetic background and outcome of Deep Brain Stimulation in Parkinson's disease. Parkinsonism Relat Disord. https://doi. org/10.1016/j.parkreldis.2018.08.006
- Musacchio T, Rebenstorff M, Fluri F et al (2017) Subthalamic nucleus deep brain stimulation is neuroprotective in the A53T α-synuclein Parkinson's disease rat model. Ann Neurol 81:825–836
- Charles PD, Van Blercom N, Krack P et al (2002) Predictors of effective bilateral subthalamic nucleus stimulation for PD. Neurology 59:932–934
- Okun MS, Rodriguez RL, Foote KD et al (2008) A case-based review of troubleshooting deep brain stimulator issues in movement and neuropsychiatric disorders. Parkinsonism Relat Disord 14:532–538
- Wong JK, Cauraugh JH, Ho KWD et al (2018) STN vs. GPi deep brain stimulation for tremor suppression in Parkinson disease: a systematic review and meta-analysis. Parkinsonism Relat Disord. https://doi.org/10.1016/j.parkreldis.2018.08.017
- Parihar R, Alterman R, Papavassiliou E, Tarsy D, Shih LC (2015)
 Comparison of VIM and STN DBS for Parkinsonian resting and postural/action tremor. Tremor Other Hyperkinet Mov (N Y) 5:321

