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Erratum: Analysis of Dosage Mutation in *PARK2* among Korean Patients with Early-Onset or Familial Parkinson's Disease

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Authors would like to change annotation of variants in Table 1 and its legend according to HGVS nomenclature recommendation.

1. Ex7 dupl/c.101delAinsAG* → Ex7 dupl/c.106duplG

2. G284R/c.674insT → G284R/c.539_540insT

Accordingly, corrected Table 1 and its legend should be as follows:

Table 1. Demographic characteristics of patients with *PARK2* mutations, and the type and location of the mutations

Sex	Age at sample	Age at onset	Family history	Variants of <i>PARK2</i>	Zygosity
F	17	17	-	Ex1 del	Heterozygous
M	56	35	-	Ex1 del	Heterozygous
M	26	26	+	Ex1 del	Heterozygous
M	41	23	+	Ex1 del	Heterozygous
F	32	32	+	Ex1 del	Heterozygous
M	36	36	-	Ex2 del	Heterozygous
F	48	45	+	Ex2 dupl	Heterozygous
F	28	28	+	Ex3 del	Heterozygous
F	25	25	-	Ex4 del	Heterozygous
M	30	29	+	Ex4 del	Heterozygous
F	23	15	-	Ex5 dupl	Heterozygous
F	36	36	+	Ex6 del	Heterozygous
F	42	23	+	Ex7 del	Heterozygous
M	43	41	-	Ex10 del	Homozygous
F	25	6	-	Ex11 del	Homozygous
F	50	30	+	Ex4 del	Homozygous
M	34	28	-	Ex2 del/Ex4 del	Compound heterozygous
F	44	41.5	+	Ex7-10 del/Ex8-10 del	Compound heterozygous
M	24	16	-	Ex3 del/Ex1-4 del	Compound heterozygous
F	38	12	-	Ex7 dupl/c.106duplG	Compound heterozygous
F	48	12	+	G284R/c.539_540insT	Compound heterozygous
M	46	42	+	Ex2 del and Ex3 del	Phase unknown

del: deletion, dupl: duplication, Ex: exon, F: female, M: male.