

Mutational analysis in familial Alzheimer's disease of Han Chinese in Taiwan with a predominant mutation *PSEN1* p.Met146Ile

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Supplementary table S1. Mutational and clinical information of the index patients

No.	Gene	Exon	Nucleotide	Amino acid	Method	AAO (year)	mGS [†]	Sex	Initial symptoms	Additional neurological feature	Age of MRI (year)	MTA	PA	WMH
1	<i>PSEN1</i>	5	c.350C>T	p.Pro117Leu	SSCP	36	3	M	Memory	Myoclonus Emotional liability	38	0	0	0
2	<i>PSEN1</i>	5	c.392A>G	p.His131Arg	NGS	49	1	M	Memory Topo [‡]	-	61	2	1	1
3	<i>PSEN1</i>	5	c.471G>T	p.Arg157Ser	NGS	51	4	M	Memory	-	59	1	2	0
4	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	45	1	F	Memory	-	-	-	-	-
5	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	45	1	F	Topo	-	45	-	0	1
6	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	40	1	F	Memory	Seizure EPS [§]	47	-	3	0
7	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	48	1	M	Memory	-	48	1	1	0
8	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	46	1	M	Memory	-	-	-	-	-
9	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	40	1	F	Memory Topo	Myoclonus Emotional liability	46	1	0	0
10	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	43	2	M	Language Dysexecutive	-	46	1	2	0
11	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	46	2	M	Memory	Myoclonus	46	0	0	0
12	<i>PSEN1</i>	5	c.438G>A	p.Met146Ile	SSCP	44	1	M	Memory	Myoclonus Seizure Choreoathetosis	55	2	1	0

										Auditory hallucination				
13	<i>PSEN1</i>	7	c.617G>A	p.Gly206Asp	SSCP	33	3	F	Memory	Seizure	-	-	-	-
14	<i>PSEN1</i>	7	c.626G>A	p.Gly209Glu	SSCP	60	2	M	Memory	Apraxia	64	2	1	0
15	<i>PSEN1</i>	8	c.838G>A	p.Glu280Lys	NGS	55	4	F	Memory	-	-	-	-	-
16	<i>PSEN1</i>	8	c.856C>G	p.Leu286Val	SSCP	43	1	M	Memory	-	-	-	-	-
17	<i>APP</i>	16	c.2066G>C	p.Asp678His	SSCP	50	4	F	Memory	-	-	-	-	-

mGS, modified Goldman score; AAO, age at onset; MTA, medial temporal atrophy; PA, posterior atrophy; WMH, white matter hyperintensity; SSCP, single strand-conformational polymorphism with subsequent Sanger sequencing; NGS, next generation sequencing

† A modified Goldman score of 1 is defined by the presence of at least three affected people in two generations, with one person being a first-degree relative of the other two; a score of 2 is familial aggregation of three or more family members with dementia not meeting the criteria for a score of 1; a score of 3 is one other affected family member with dementia (modified to give a score of 3 only if there is a history of young-onset dementia within the family, i.e., AAO less than 65 years; with a score of 3.5 if AAO is above 65); and a score of 4 is no or an unknown family history.

‡Topo, topographic disorientation

§EPS, extrapyramidal symptoms