

Supplementary Materials

Decoding hereditary spastic paraplegia pathogenicity through transcriptomic profiling

Nicolas James Ho¹, Xiao Chen^{2,3,4,5,6}, Yong Lei^{7,8,#,*}, Shen Gu^{1,9,10,11,#,*}

¹School of Biomedical Sciences, Faculty of Medicine, The Chinese University of Hong Kong, Hong Kong SAR, China

²Dr. Li Dak Sum-Yip Yio Chin Center for Stem Cells and Regenerative Medicine and Department of Orthopedic Surgery of the Second Affiliated Hospital, Zhejiang University School of Medicine, Hangzhou, Zhejiang 310058, China

³Key Laboratory of Tissue Engineering and Regenerative Medicine of Zhejiang Province, Zhejiang University School of Medicine, Hangzhou, Zhejiang 310058, China

⁴Zhejiang University-University of Edinburgh Institute & School of Basic Medicine, Zhejiang University School of Medicine, Hangzhou, Zhejiang 310058, China

⁵Department of Sports Medicine, Zhejiang University School of Medicine, Hangzhou, Zhejiang 310058, China

⁶China Orthopedic Regenerative Medicine Group (CORMed), Hangzhou, Zhejiang, 310058 China

⁷School of Medicine, The Chinese University of Hong Kong (Shenzhen), Shenzhen, Guangdong 518172, China

⁸The Chinese University of Hong Kong (Shenzhen), Shenzhen Futian Biomedical Innovation R&D Center, Shenzhen, Guangdong 518172, China

⁹Key Laboratory for Regenerative Medicine, Ministry of Education, School of Biomedical Sciences, Faculty of Medicine, The Chinese University of Hong Kong, Hong Kong SAR, China

¹⁰Kunming Institute of Zoology Chinese Academy of Sciences, The Chinese University of Hong Kong Joint Laboratory of Bioresources and Molecular Research of Common Diseases, Hong Kong SAR, China

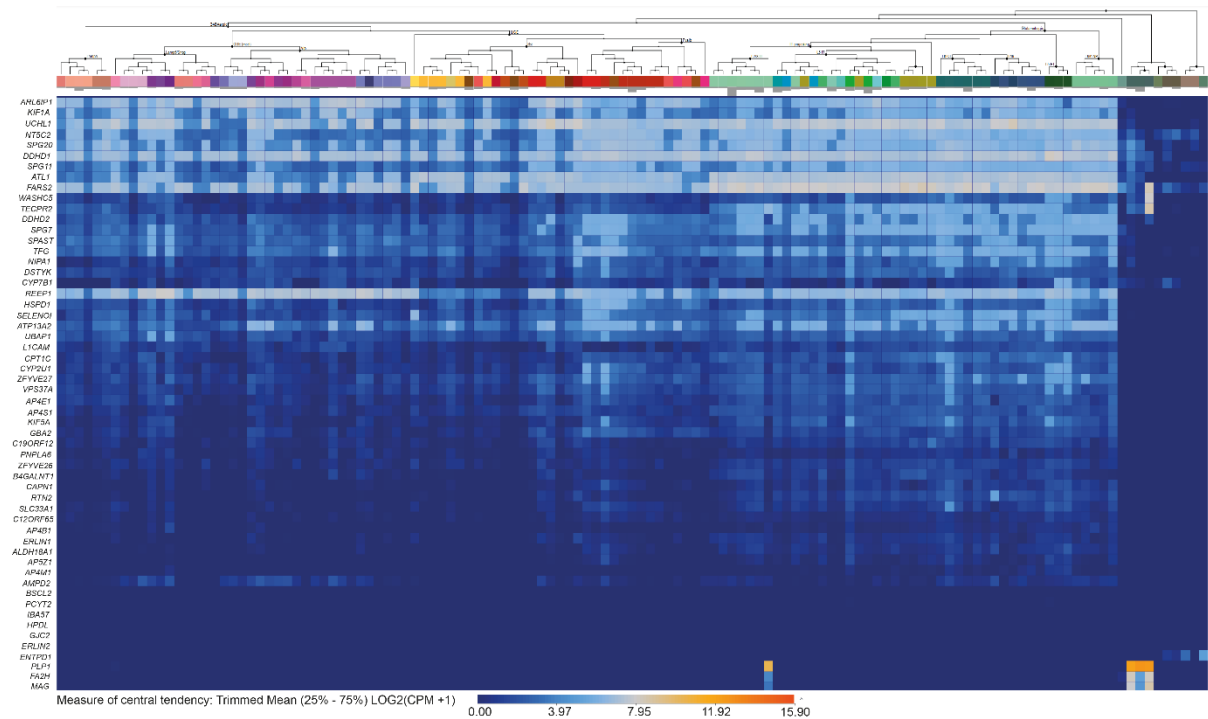
¹¹Hong Kong Branch of CAS Center for Excellence in Animal Evolution and Genetics, The Chinese University of Hong Kong, New Territories, Hong Kong SAR, China

#Authors contributed equally to this work

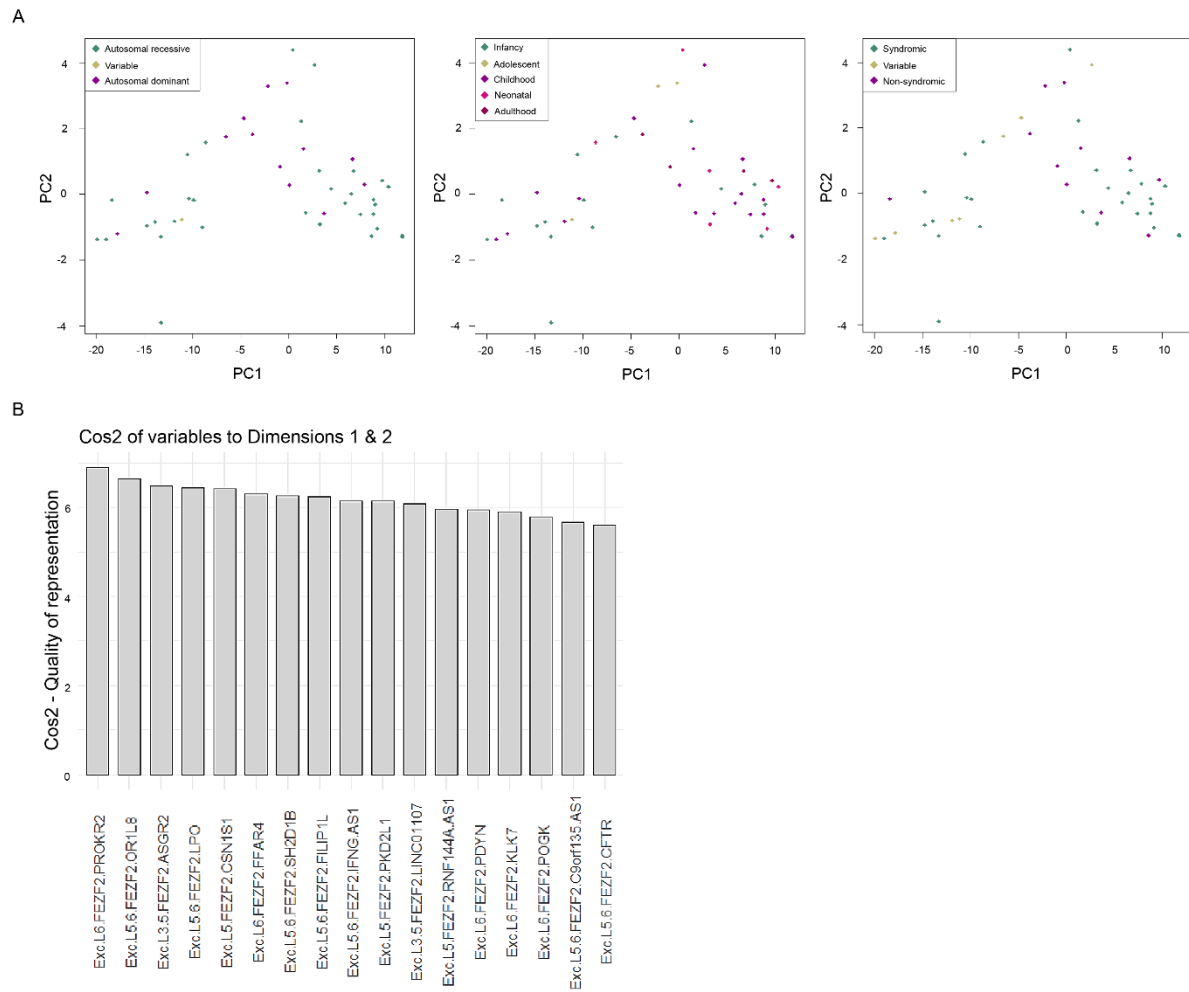
*Corresponding authors, E-mail: leiyong@cuhk.edu.cn; shengu@cuhk.edu.hk

Earliest reported onset	Neonatal										Infancy										Childhood										Adolescent										Adulthood																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																						
HSP subtype	SPG1	SPG2	SPG3	SPG4	SPG5	SPG6	SPG7	SPG8	SPG9	SPG10	SPG11	SPG12	SPG13	SPG14	SPG15	SPG16	SPG17	SPG18	SPG19	SPG20	SPG21	SPG22	SPG23	SPG24	SPG25	SPG26	SPG27	SPG28	SPG29	SPG30	SPG31	SPG32	SPG33	SPG34	SPG35	SPG36	SPG37	SPG38	SPG39	SPG40	SPG41	SPG42	SPG43	SPG44	SPG45	SPG46	SPG47	SPG48	SPG49	SPG50																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																													
Associated genes	LITAM	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1	ANKK1																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																												
Inheritance	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR	AR																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																											
Phenotype	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																											
Human forebrain (week post conception)	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																											
Human hindbrain (week post conception)	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																											
Human forebrain (postnatal)	Neonate	Infant	Toddler	School age child	Adolescent	Young adult	Adult	Older adult	Elderly																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																						

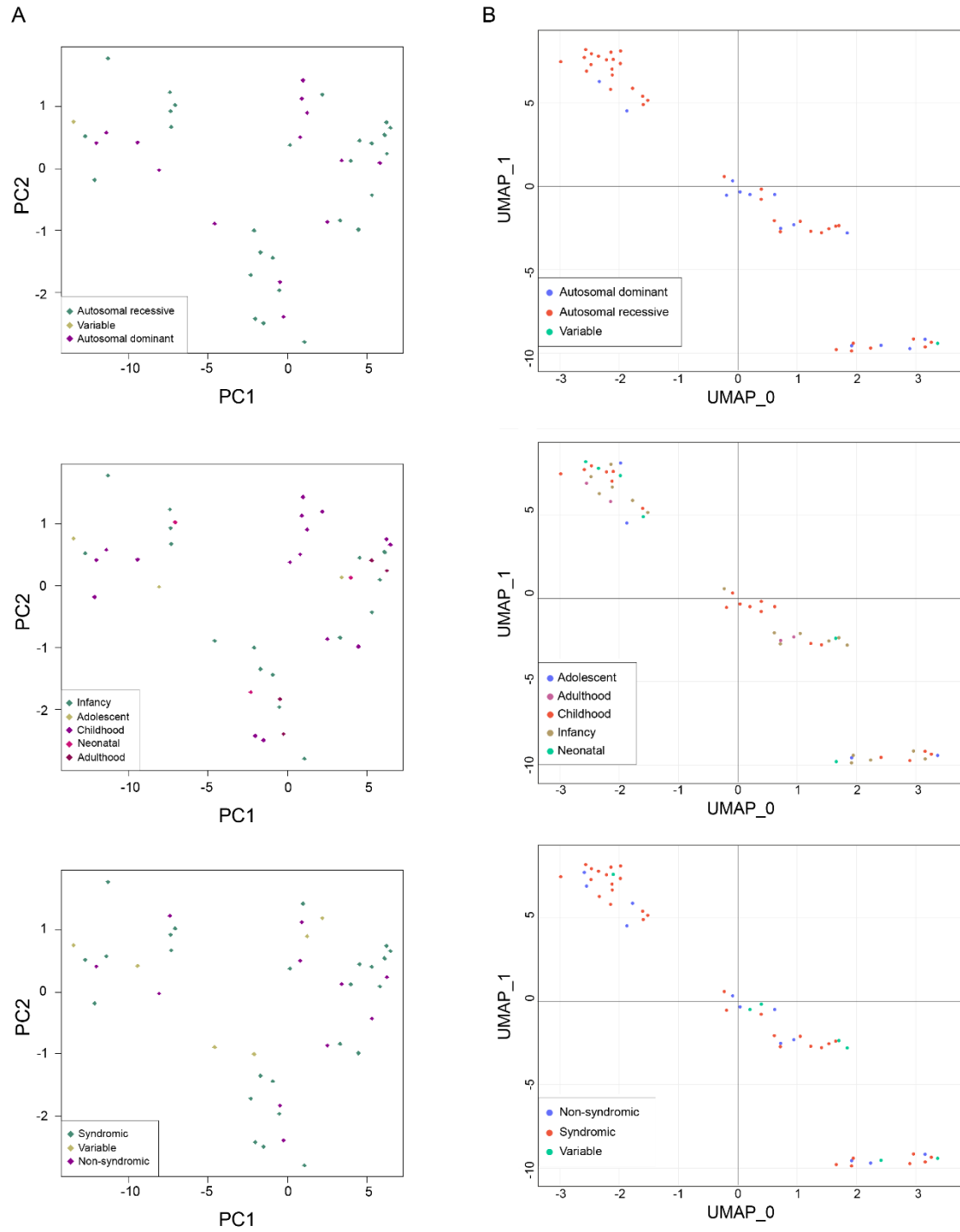
Supplementary Figure S1. Heatmap showing forebrain/hindbrain tissue mRNA expression profiles (TPM) of HSP associated genes in normal human and mouse tissues at each developmental stage.



Supplementary Figure S2. Heatmap visualizing expression patterns of HSP-associated genes across transcriptomic cell types in the human primary motor cortex by sc-RNA-seq.



Supplementary Figure S3. Selection of FEZF2+ pyramidal neuronal populations for further HSP gene expression profile dimensionality reduction analysis.



Supplementary Figure S4. Dimensional reduction analysis of selected mouse isocortical extralencephalic (ET) neuronal populations.

Supplementary Figure S5. Heatmap showing tissue specific mRNA expression levels of HSP associated genes in normal human tissues.

Supplementary Table S1 List of genes associated with HSP

HSP subtype	HGNC approved gene symbol	Reported association to neurological conditions other than HSP
SPG1	<i>L1CAM</i>	Hirschsprung disease, hydrocephalus
SPG2	<i>PLP1</i>	Leukodystrophy
SPG3A	<i>ATL1</i>	Sensory neuropathy
SPG4	<i>SPAST</i>	
SPG5A	<i>CYP7B1</i>	
SPG6	<i>NIPA1</i>	
SPG7	<i>SPG7</i>	
SPG8	<i>WASHC5</i>	Ritscher-Schinzel syndrome
SPG9A/9B	<i>ALDH18A1</i>	
SPG10	<i>KIF5A</i>	Amyotrophic lateral sclerosis, epilepsy
SPG11	<i>SPG11</i>	Amyotrophic lateral sclerosis, Charcot-Marie-Tooth disease, homocarnosinosis
SPG12	<i>RTN2</i>	
SPG13	<i>HSPD1</i>	Leukodystrophy
SPG14	<i>SPG14</i>	
SPG15	<i>ZFYVE26</i>	
SPG16	<i>SPG16</i>	
SPG17	<i>BSCL2</i>	Encephalopathy, distal motor neuropathy
SPG18	<i>ERLIN2</i>	
SPG19	<i>SPG19</i>	
SPG20	<i>SPART</i>	
SPG21	<i>ACP33</i>	
SPG23	<i>DSTYK</i>	
SPG24	<i>SPG24</i>	
SPG25	<i>SPG25</i>	
SPG26	<i>B4GALNT1</i>	
SPG27	<i>SPG27</i>	
SPG28	<i>DDHD1</i>	
SPG29	<i>SPG29</i>	
SPG30	<i>KIF1A</i>	Sensory neuropathy, NESCAV syndrome
SPG31	<i>REEP1</i>	Distal motor neuropathy
SPG32	<i>SPG32</i>	
SPG33	<i>ZFYVE27</i>	
SPG34	<i>SPG34</i>	
SPG35	<i>FA2H</i>	
SPG36	<i>SPG36</i>	
SPG37	<i>SPG37</i>	
SPG38	<i>SPG38</i>	
SPG39	<i>PNPLA6</i>	Boucher-Neuhauser syndrome, Laurence-Moon syndrome, Olver-McFarlane syndrome.

SPG41	<i>SPG41</i>	
SPG42	<i>SLC33A1</i>	
SPG43	<i>C19ORF12</i>	Brain iron accumulation associated neurodegeneration, mitochondrial membrane protein-associated neurodegeneration
SPG44	<i>GJC2</i>	Leukodystrophy
SPG45	<i>NT5C2</i>	
SPG46	<i>GBA2</i>	Cerebellar ataxia
SPG47	<i>AP4B1</i>	
SPG48	<i>AP5Z1</i>	
SPG49	<i>TECPR2</i>	Sensory and autonomic neuropathy
SPG50	<i>AP4M1</i>	
SPG51	<i>AP4E1</i>	
SPG52	<i>AP4S1</i>	
SPG53	<i>VPS37A</i>	
SPG54	<i>DDHD2</i>	
SPG55	<i>MTRFR</i>	
SPG56	<i>CYP2U1</i>	
SPG57	<i>TFG</i>	Charcot-Marie-Tooth disease, motor and sensory neuropathy
SPG61	<i>ARL6IP1</i>	
SPG62	<i>ERLIN1</i>	
SPG63	<i>AMPD2</i>	Pontocerebellar hypoplasia
SPG64	<i>ENTPD1</i>	
SPG73	<i>CPT1C</i>	
SPG74	<i>IBA57</i>	Multiple mitochondrial dysfunction syndrome
SPG75	<i>MAG</i>	
SPG76	<i>CAPN1</i>	
SPG77	<i>FARS2</i>	
SPG78	<i>ATP13A2</i>	Brain iron accumulation associated neurodegeneration, Kufor-Rakeb syndrome
SPG79	<i>UCHL1</i>	
SPG80	<i>UBAP1</i>	
SPG81	<i>SELENOI</i>	
SPG82	<i>PCYT2</i>	
SPG83	<i>HPDL</i>	

Supplementary Table S2 List of HSP genes identified from rhesus monkey brain and average cortical expression by age

	HSP subtype	Symbol	Averaged expression by age		
			16.5 (n =		
			5.75 (n = 4)	2)	24 (n = 1)
Adult.	SPG73	<i>CPT1C</i>	4.989073445	4.989419	4.985567
	SPG33	<i>ZFYVE27</i>	7.109103806	7.10401	7.140942
	SPG76	<i>CAPN1</i>	7.216363539	7.278123	7.120605
Adol.	SPG13	<i>HSPD1</i>	6.069200117	6.074412	6.079972
	SPG30	<i>KIF1A</i>	9.063097963	9.017894	9.059115
Childhood	SPG80	<i>UBAP1</i>	6.248539296	6.195004	6.346933
	SPG39	<i>PNPLA6</i>	6.454761789	6.466153	6.419514
	SPG26	<i>B4GALNT1</i>	4.921820554	4.866581	4.981878
	SPG12	<i>RTN2</i>	4.661147914	4.646705	4.640942
	SPG83	<i>HPDL</i>	1.549666546	1.577375	1.419267
	SPG7	<i>SPG7</i>	6.834505635	6.916359	6.666004
	SPG6	<i>NIPA1</i>	6.724166776	6.715201	6.735772
	SPG5A	<i>CYP7B1</i>	3.715281748	3.705905	3.686841
	SPG3A	<i>ATL1</i>	6.896267671	6.885668	6.883168
	SPG79	<i>UCHL1</i>	8.635203608	8.605775	8.665398
	SPG78	<i>ATP13A2</i>	8.663981318	8.661025	8.661176
	SPG75	<i>MAG</i>	3.491268114	3.454769	3.621977
	SPG48	<i>AP5Z1</i>	6.257186207	6.266809	6.270654
	SPG35	<i>FA2H</i>	4.683975436	4.741617	4.629831
	SPG31	<i>REEP1</i>	3.14573012	3.130801	3.119082
	SPG23	<i>DSTYK</i>	6.384339033	6.394815	6.359493
	SPG17	<i>BSCL2</i>	10.22652729	10.24413	10.17115
	SPG15	<i>ZFYVE26</i>	4.381906137	4.3494	4.441523
	SPG2	<i>PLP1</i>	10.58574904	10.6132	10.5326
Infancy	SPG63	<i>AMPD2</i>	7.407009048	7.39125	7.408616
	SPG28	<i>DDHD1</i>	6.431497958	6.392015	6.476244
	SPG77	<i>FARS2</i>	5.011415642	5.055989	4.9402
	SPG4	<i>SPAST</i>	5.133809311	5.163894	5.08781
	SPG82	<i>PCYT2</i>	6.165597562	6.237175	6.024905
	SPG62	<i>ERLIN1</i>	4.591077338	4.596927	4.58045
	SPG61	<i>ARL6IP1</i>	8.625928848	8.32416	9.236635
	SPG57	<i>TFG</i>	5.695343884	5.701096	5.699016
	SPG54	<i>DDHD2</i>	7.338454584	7.292372	7.429301
	SPG49	<i>TECPR2</i>	4.128680725	4.188984	4.088458
	SPG46	<i>GBA2</i>	6.41666929	6.380121	6.473795
	SPG45	<i>NT5C2</i>	6.44659989	6.456299	6.449627

	SPG18	<i>ERLIN2</i>	6.131058204	6.085902	6.205445
	SPG11	<i>SPG11</i>	6.329148662	6.330945	6.306984
	SPG9A/9B	<i>ALDH18A1</i>	5.918271401	5.882059	5.976919
Neonatal	SPG52	<i>AP4S1</i>	3.56377823	3.643461	3.454884
	SPG51	<i>AP4E1</i>	3.439138519	3.411397	3.521496
	SPG50	<i>AP4M1</i>	3.344894579	3.4122	3.248671
	SPG47	<i>AP4B1</i>	3.832684867	3.811203	3.900783
	SPG1	<i>L1CAM</i>	7.604234024	7.557543	7.64518

Supplementary Table S3 List of HSP genes potentially correlated with clinical presentations of Alzheimer's disease (with reference to www.alzdata.org database)

Gene	eQTL	GWAS	PPI	Earl. DEG	Pathology cor. (ab)	Pathology cor. (tau)	CFG
<i>CYP7B1</i>	-	0	-	Yes	0.37,*	0.601,*	2
<i>KIF5A</i>	0	0	-	-	-0.359,*	-0.82,***	1
<i>RTN2</i>	0	2	-	No	-0.544,***	-0.658,***	2
<i>HSPD1</i>	0	0	APP, APOE	Yes	-0.575,***	Ns	3
<i>REEP1</i>	0	0	-	Yes	-0.621,***	-0.858,***	2
<i>ZFYVE27</i>	0	0	-	No	Ns	-0.608,*	1
<i>GBA2</i>	-	0	-	-	Ns	-0.718,**	1
<i>AP4M1</i>	2	2	-	-	-0.473,**	Ns	3
<i>AP4S1</i>	-	0	-	-	-0.432,**	Ns	1
<i>TFG</i>	0	0	-	-	Ns	-0.622,*	1
<i>AMPD2</i>	1	0	-	-	Ns	-0.649,**	2
<i>MAG</i>	1	0	APP, PSEN2	Yes	0.433,**	Ns	4
<i>CAPN1</i>	2	0	PSEN2	Yes	0.327,*	Ns	4
<i>FARS2</i>	0	0	-	Yes	Ns	0.745,**	2
<i>UCHL1</i>	1	0	-	Yes	-0.338,*	Ns	3
<i>UBAP1</i>	-	0	-	Yes	-0.401,**	Ns	2
<i>PCYT2</i>	1	-	-	Yes	Ns	-0.743,**	3

Supplementary Table S4 Cross-species mean HSP gene expression in human post-mortem, marmoset, and mouse layer 5 extratelencephalic (ET) specimens

Name	Subtype	Earliest reported onset	Motor involvement in mouse	Inheritance	Phenotype	L5 ET mean expr. (human)	L5 ET equivalent mean expr. (marmoset)	L5 ET equivalent population mean expr. (mouse)
<i>L1CAM</i>	SPG1	Neonatal	N.A.	XLR	Syndromic	2.50320506	2.06970191	2.4786849
<i>AP4B1</i>	SPG47	Neonatal	N.A.	AR	Syndromic	1.83428025	1.67597699	0.001410398
<i>AP4M1</i>	SPG50	Neonatal	N.A.	AR	Syndromic	2.30576205	0.488226444	1.67390072
<i>AP4E1</i>	SPG51	Neonatal	N.A.	AR	Syndromic	3.02564096	2.51064634	2.1164546
<i>AP4S1</i>	SPG52	Neonatal	N.A.	AR	Syndromic	2.42715621	1.92109215	3.44838905
<i>CYP2U1</i>	SPG56	Neonatal	N.A.	AR	Syndromic	4.65701485	1.09143281	2.34691644
<i>ALDH18A1</i>	SPG9A/9B	Infancy	N.A.	Var	Syndromic	2.36003065	1.57934618	1.28657353
<i>SPG11</i>	SPG11	Infancy	Yes	AR	Syndromic	4.72942162	4.36034584	1.89516735
<i>ERLIN2</i>	SPG18	Infancy	N.A.	AR	Syndromic	0.739109874	0.351452917	1.36586702
<i>SPG20</i>	SPG20	Infancy	Yes	AR	Syndromic	5.75433397	2.33385515	1.23311675
<i>NT5C2</i>	SPG45	Infancy	No	AR	Syndromic	5.07899904	4.49530315	4.16266584
<i>GBA2</i>	SPG46	Infancy	No	AR	Syndromic	2.74373531	0.962049127	2.8253665
<i>TECPR2</i>	SPG49	Infancy	N.A.	AR	Syndromic	4.1878643	6.16226196	4.9389863
<i>VPS37A</i>	SPG53	Infancy	N.A.	AR	Syndromic	3.30233836	3.68374252	3.65616012
<i>DDHD2</i>	SPG54	Infancy	Yes	AR	Syndromic	4.07473755	3.37199402	5.29261637
<i>TFG</i>	SPG57	Infancy	N.A.	AR	Syndromic	4.55394077	3.77310872	3.26379275
<i>ERLIN1</i>	SPG62	Infancy	N.A.	AR	Syndromic	1.98350084	1.74787068	2.58782506
<i>ENTP</i>	SPG6	Infancy	No	AR	Syndromic	0.756665	2.34869742	0.4370367

<i>D1</i>	4	y			c	23		83
<i>EPT1</i>	SPG8 1	Infanc y	N.A.	AR	Syndromi c	4.665136 81	2.02786827	0
<i>PCYT2</i>	SPG8 2	Infanc y	N.A.	AR	Syndromi c	1.280849 34	1.98453152	4.6629633 9
<i>SPAST</i>	SPG4	Infanc y	No	AD	Variable	3.890479 33	4.24987459	4.3284845 4
<i>FARS2</i>	SPG7 7	Infanc y	N.A.	AR	Variable	7.342730 05	6.18280315	5.2930169 1
<i>DDHD 1</i>	SPG2 8	Infanc y	No	AR	Non- syndromi c	7.729312 42	6.4148922	4.0443997 4
<i>AMPD 2</i>	SPG6 3	Infanc y	No	AR	Non- syndromi c	1.751311 18	1.70578659	0.7999308 71
<i>PLP1</i>	SPG2	Childh ood	N.A.	XLR	Syndromi c	1.181490 42	0.852893293	0.8473243 71
<i>ZFYVE 26</i>	SPG1 5	Childh ood	Yes	AR	Syndromi c	2.581512 21	0.804045618	1.8475801 9
<i>SPG21</i>	SPG2 1	Childh ood	Yes	AR	Syndromi c	2.029647 35	1.27586424	1.0286434 9
<i>DSTYK</i>	SPG2 3	Childh ood	N.A.	AR	Syndromi c	2.772181 03	2.10377002	2.5570519
<i>REEP1</i>	SPG3 1	Childh ood	No	AD	Syndromi c	6.298987 39	5.13639784	7.6637167 9
<i>FA2H</i>	SPG3 5	Childh ood	No	AR	Syndromi c	0.187937 066	0.244301096	0.0973520 43
<i>C19O RF12</i>	SPG4 3	Childh ood	Yes	AR	Syndromi c	2.513184 79	1.31199896	1.4265763 8
<i>AP5Z1</i>	SPG4 8	Childh ood	N.A.	AR	Syndromi c	2.260307 31	1.83817637	1.4368016 7
<i>C12O RF65</i>	SPG5 5	Childh ood	N.A.	AR	Syndromi c	2.111451 15	2.67184377	1.4327778 8
<i>IBA57</i>	SPG7 4	Childh ood	N.A.	AR	Syndromi c	0.927921 057	0.311998993	0.2554549 28
<i>MAG</i>	SPG7 5	Childh ood	Yes	AR	Syndromi c	0.112143 062	0.027179359	0.2325221 3
<i>ATP13 A2</i>	SPG7 8	Childh ood	Yes	AR	Syndromi c	4.252768 04	4.17109203	2.6567893
<i>UCHL1</i>	SPG7 9	Childh ood	Yes	AR	Syndromi c	6.830274 11	2.99317384	6.7002763 7
<i>ATL1</i>	SPG3 A	Childh ood	No	AD	Variable	6.827942 85	6.20265532	5.9586768 2
<i>CYP7B</i>	SPG5	Childh	No	AR	Variable	4.700247	3.40055108	4.0203332

1	A	ood				76		9
<i>NIPA1</i>	SPG6	Childhood	Yes	AD	Variable	3.79010057	2.93336678	4.54365301
<i>SPG7</i>	SPG7	Childhood	Yes	AR	Variable	4.84560728	2.77523804	0
<i>HPDL</i>	SPG83	Childhood	N.A.	AR	Variable	0.013039044	0.013527054	0.065929204
<i>KIF5A</i>	SPG10	Childhood	N.A.	AD	Non-syndromic	3.74235129	4.23039818	6.69980621
<i>RTN2</i>	SPG12	Childhood	No	AD	Non-syndromic	2.6870265	2.89065623	1.49653625
<i>B4GALNT1</i>	SPG26	Childhood	Yes	AR	Non-syndromic	2.16910696	1.19532812	2.04796052
<i>PNPLA6</i>	SPG39	Childhood	N.A.	AR	Non-syndromic	1.86789775	2.37625241	1.81022537
<i>SLC33A1</i>	SPG42	Childhood	Yes	AD	Non-syndromic	2.97526956	1.64698148	2.5734098
<i>UBAP1</i>	SPG80	Childhood	N.A.	AD	Non-syndromic	3.4125874	3.64147043	2.79652238
<i>GJC2</i>	SPG44	Adolescent	Yes	AR	Syndromic	0.216601104	0.013401804	0.060342919
<i>KIF1A</i>	SPG30	Adolescent	No	Variable	Variable	3.75517201	3.90769029	5.89908028
<i>KIAA0196</i>	SPG8	Adolescent	No	AD	Non-syndromic	3.93094397	2.60671353	0
<i>HSPD1</i>	SPG13	Adolescent	N.A.	AD	Non-syndromic	4.02188969	1.36009514	3.55869055
<i>CAPN1</i>	SPG76	Adulthood	Yes	AR	Syndromic	2.95327067	2.37957168	1.49715149
<i>ZFYVE27</i>	SPG33	Adulthood	N.A.	AD	Non-syndromic	3.95735002	4.13739967	2.861449
<i>CPT1C</i>	SPG73	Adulthood	Yes	AD	Non-syndromic	3.3851254	3.05786562	3.45760512

