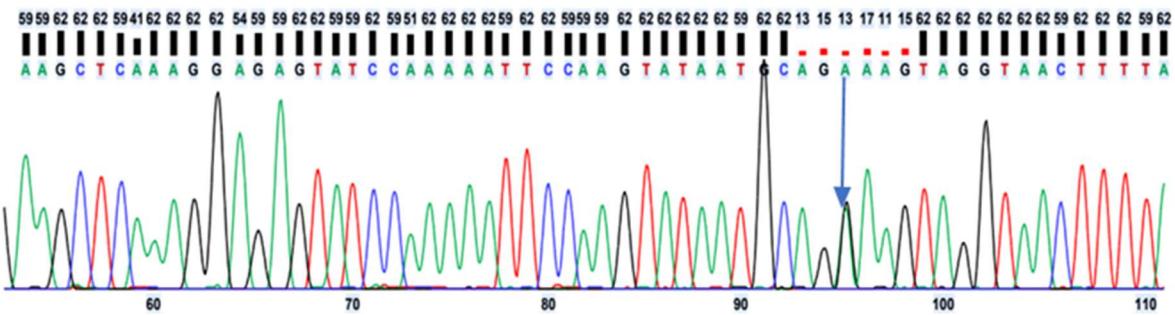


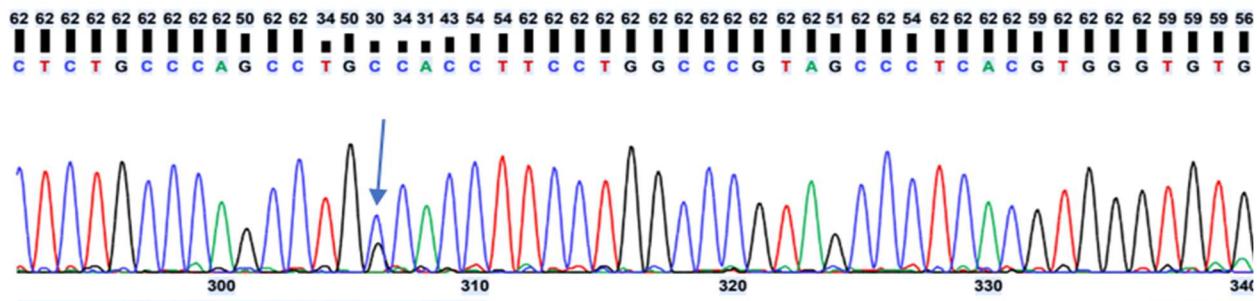
Supplement figure

Supplement figure S1. Sequencing data of the five variants, found the patient

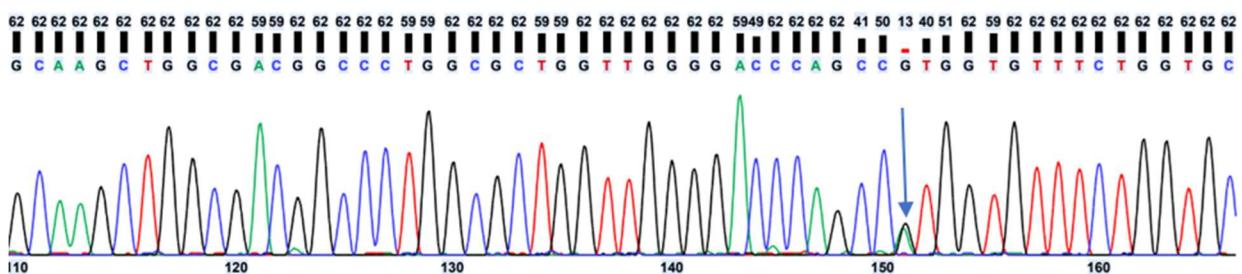
PSEN1 Glu318Gly



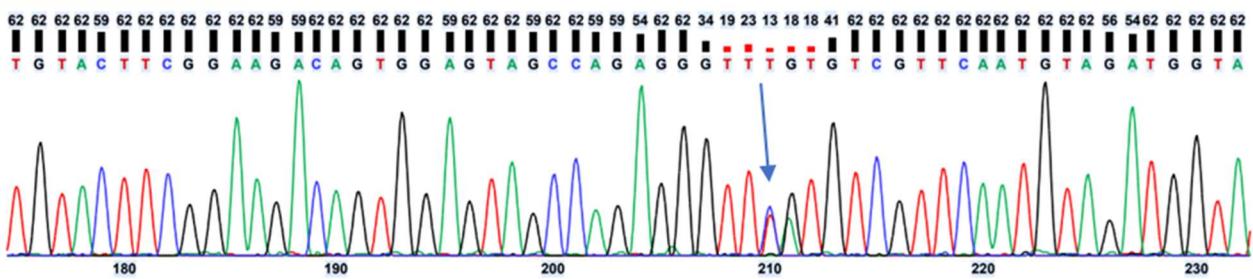
GRN Ala505Gly



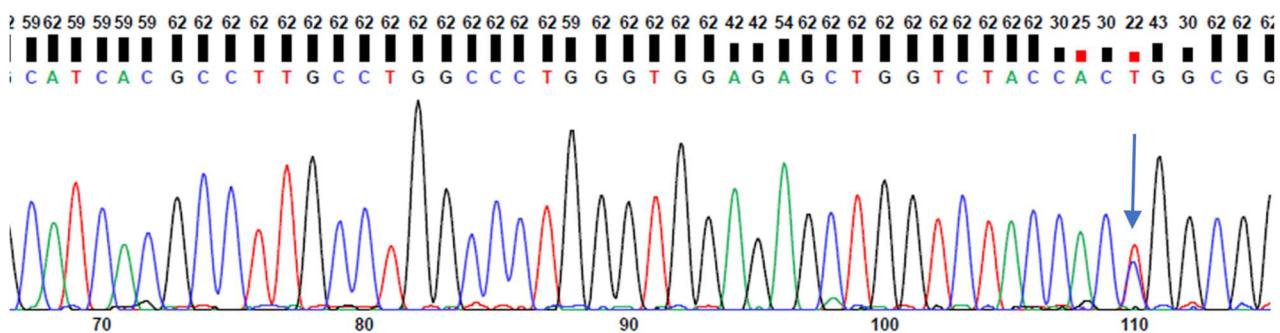
ABCA7 Val1946Met



SORL1 Glu270Lys (3' sequencing). The patient also carried a silent variant H269H



TOMM40 Arg239Trp



Supplementary tables

Supplement table S1. Supplement table S1.: Genes, included in our gene panel

Genes, used in the gene panel
APP', 'PSEN1', 'PSEN2', 'S100A9', 'SACS', 'CR1', 'BIN1', 'TREM2', 'CLU', 'CTNNA3', 'DNMBP', 'SORL1', 'BACE1', 'PICALM', 'GAB2', 'LRP6', 'ADAM10', 'ABCAT7', 'CD33', 'TOMM40', 'MS4A4A', 'MS4A6E', 'TM2D3', 'CD2AP', 'EPHA1', 'CASS4', 'PLD3', 'INPP5D', 'DSG2', 'CDH12', 'CDH18', 'MEF2C', 'NME8', 'PTK2B', 'SLC24A4', 'RIN3', 'ZCWPW1', 'ACE', 'MTHFD1L', 'PINK1', 'PARK7', 'PARK9', 'ATP13A2', 'GBA', 'SNCA', 'PARK2', 'LRRK2', 'ACMSD', 'CD157', 'BST1', 'FBXO7', 'FGF20', 'GAK', 'GIGYF2', 'GPNMB', 'HIP1R', 'PTPRD', 'LAMP3', 'PLA2G6', 'STBD1', 'STK39', 'STX1B', 'SYT11', 'VPS35', 'TDP43', 'TARDBP', 'ATXN2', 'EWSR1', 'HNRNPA2B1', 'PFN1', 'SETX', 'TMEM106B', 'CCNF', 'PPT1', 'TBK1', 'DCTN1', 'NEK1', 'CHMP2B', 'SQSTM1', 'SIGMAR1', 'VCP', 'FUS', 'GRN', 'MAPT', 'TAF15', 'ALS2', 'UBQLN2', 'FIG4', 'OPTN', 'DAO', 'HNRNPA1', 'ANG', 'VAPB', 'SOD1', 'SPAST', 'CYP7B1', 'SPG11', 'CSF1R', 'NOTCH3', 'HTRA1', 'CTSA', 'PRNP', 'STX6', 'GAL3', 'ST1', 'PDIA4', 'BMERB1', 'LPA', 'FGF20', 'TET1', 'ACO1', 'POSTN', 'ABCA13', 'ADAM18', 'ARAP2', 'ATP13A3', 'C1orf51', 'CIART', 'DDX1', 'FAM105A', 'FGF20', ''

FNDC1', 'GBP4', 'GPR63', 'HACL1', 'MYOZ2', 'PDLIM5', 'PKP1', 'POLR1C', 'SFRP4', 'SLAIN2', 'SLC6A5', 'SMC5', 'TWF1', 'VEPH1', 'WDR64', 'ZRANB2', 'ATXN1'
, 'NOTCH4', 'GLA', 'COL4A1', 'COL4A2', 'FOXC1', 'FOXF2', 'TREX1', 'PMF', 'ZCCHC14', 'WDR123', 'ICAL1', 'GCH1', 'S2B3', 'ACOX1'

Supplement Table S2: Variants, found in the patient. The five variants of interest have been highlighted

Gene	HGVS.p	Zygosity	rs ID	p3_1000G	p3_1000G_EAS	gnomAD_exomes	gnomAD_exomes_EAS	SIFT_score	SIFT_predicted
ABCA1	p.Arg555His	HET	rs2361519	0.198083	0.1508	0.145723	0.169234	1	T
ABCA1	p.Ile767Ser	HET	rs17712293	0.194489	0.2153	0.173606	0.224242	0.007	D
ABCA1	p.Phe1540Leu	HET	rs17712299	0.096446	0.1498	0.07085	0.179048	0	D
ABCA1	p.Ala2178Gl	HOM	rs1880736	0.812899	0.9881	0.723538	0.989164	1	T
ABCA1	p.Lys2306Thr	HET	rs78334925	0.097244	0.1508	0.075345	0.186615	0.11	T
ABCA1	p.Arg2674Trp	HOM	rs2222648	0.852236	0.9921	0.803456	0.991853	1	T
ABCA1	p.Thr4550Ala	HET	rs78734445	0.002596	0	0.004008	0	0.112;0.084	T
ABCA7	p.Arg1349Gln	HET	rs3745842	0.390575	0.3482	0.427897	0.335604	0.546;0.546;..	T;T;..
ABCA7	p.Arg1413Arg	HET	rs881768	0.442093	0.3552
ABCA7	p.Gly1527Ala	HET	rs3752246	0.825479	0.6478	0.840229	0.646508	0.877;0.877;..	T;T;..
ABCA7	p.Val1946Met	HET	0.031;0.031;..;	D;D;..;
ABCA7	p.Ala2045Ser	HET	rs4147934	0.605032	0.4325	0.71193	0.446435	0.962;0.962;..;	T;T;..;
ACOX1	p.Ile312Met	HET	rs1135640	0.550719	0.7877	0.653791	0.800441	0.256;0.201	T
ALS2	p.Val368Met	HOM	rs3219156	0.896565	1	0.912953	0.999826	0.191;0.172	T
ARAP2	p.Arg1523Gln	HOM	rs4833069	0.995208	1	0.992128	1	1.0;..	T;..
ARAP2	p.Pro89Leu	HET	rs35822395	0.005591	0	0.008379	0.000116	0.094	T
ATP13	p.Pro1122Leu	HET	rs15786	0.013179	0.001	0.04005	0	0	D
ATP13	p.Ala1072Thr	HET	rs3170740	0.33127	0.246	0.47141	0.295664	0.988;1.0	T
ATP13	p.Leu437Val	HET	rs149372969	0.000599	0	0.00106	0	0.219;0.588;0.236;..0	T;T;T;..T
ATXN1	p.Pro753Ser	HET	rs16885	0.122804	0.0069	0.195486	0.008407	0.015	D
ATXN1	p.Gln225dup	HOM	rs769091061
ATXN2	p.Pro425Leu	HET	0.001;..0.001;..0.002;D;..D;..D; 0.001;0.001;0.003	D;D;D
ATXN2	p.Ser248Asn	HET	rs7969300	0.179513	0.5228	0.076308	0.564288	..0.136;0.128	..T;T
CD33	p.Arg69Gly	HOM	rs2455069	0.335064	0.1042	0.348885	0.080424	0.334;0.335	T
COL4A1	p.Gln1334His	HET	rs3742207	0.288339	0.2629	0.304165	0.251565	0.122	T
COL4A2	p.Arg517Lys	HOM	rs7990383	0.509185	0.2232	0.523827	0.230619	1	T
CR1	p.His1658Arg	HET	rs2274567	0.294329	0.3274	0.250551	0.291633	0.897;0.897;0.897;0.	897;0.309

SNP rs10582052									
Gene	Mutation	Allele	rsID	Frequency	AA	AF	AC	AF	AA
CR1	p.Thr2060Ser	HOM	rs4844609	0.995008	1	0.985426	1	0.804;0.804;0.804;0.804;1.0	T
CR1	p.Ile2065Val	HET	rs6691117	0.493411	0.3433	0.321622	0.3062	1	T
CR1	p.Pro2277Arg	HET	rs3811381	0.26278	0.3224	0.241761	0.288787	0.446;0.446;0.446;0.446;0.394	T
CR1	p.Thr2419Ala	HOM	rs2296160	0.828075	0.6885	0.819882	0.655775	0.987;0.987;0.987;0.987;1.0	T
rs10582052,rs3									
CTSA	p.Leu37del	HOM	080212,rs11468075	0.649361	0.5784
DSG2	p.Arg773Lys	HET	rs2278792	0.240016	0.4643	0.265563	0.466526	0.383	T
EPHA1	p.Met900Val	HOM	rs6967117	0.960264	1	0.940039	0.998493	1	T
EPHA1	p.Arg492Gln	HET	rs11768549	0.00619	0	0.012483	0	0.599	T
EPHA1	p.Val160Ala	HOM	rs4725617	0.94349	0.9921	0.93221	0.991984	0.246	T
FNDC1	p.Ser36Pro	HET	rs295332	0.550919	0.628	0.540541	0	0.981	T
FNDC1	p.Thr438Ala	HET	rs509648	0.508586	0.7232	0.324572	0.751479	1	T
FNDC1	p.Glu463Gln	HOM	rs420137	0.784545	0.621	0.866743	0.637705	1	T
FNDC1	p.Gln1003Glu	HOM	rs370434	0.785743	0.626	0.867677	0.641117	1	T
FNDC1	p.Asp1180Glu	HOM	rs420054	0.782348	0.626	0.850326	0.622257	1	T
FNDC1	p.Leu1261Pro	HOM	rs3003174	0.786142	0.626	0.86882	0.641139	0.345	T
FNDC1	p.Gln1280Arg	HOM	rs2501176	0.786142	0.626	0.869494	0.643025	1	T
FNDC1	p.Thr1479Thr1484del	HOM	rs141435210	0.496406	0.4058
FNDC1	p.Thr1504Lys	HOM	rs369362288	0.784944	0.625	0.872142	0.644028	0.795	T
FOXC1	p.Gly380dup	HET	rs76840944	0.152157	0.2867
rs147426137,rs									
FOXF2	p.Gly306dup	HOM	58230522,rs111257067	0.392772	0.497
GBP4	p.Glu551Gly	HOM	rs561042	0.357428	0.3978	0.442432	0.399049	0.036	D
GBP4	p.Glu551Lys	HOM	rs561037	0.358826	0.3978	0.441993	0.398608	0.505	T
GBP4	p.Leu549Met	HOM	rs608339	0.359425	0.3988	0.43821	0.395317	1	T
GBP4	p.Glu546Lys	HOM	rs1142890	0.359026	0.3988	0.428369	0.38529	0.929	T
GBP4	p.Met545Ile	HOM	rs1142889	0.359026	0.3988	0.427021	0.384029	0.072	T
GBP4	p.Met545Leu	HOM	rs1142888	0.359026	0.3988	0.425834	0.382578	1	T
GBP4	p.Met542Ile	HOM	rs1142886	0.359625	0.3998	0.426034	0.381004	1	T
GBP4	p.Tyr541Asn	HOM	rs655260	0.359625	0.3998	0.430646	0.386015	1	T
GRN	p.Ala505Gly	HET	rs780159686	.	.	7.31E-05	0	0.257;.	T;.
LAMP3	p.Ile318Val	HET	rs482912	0.496406	0.4712	0.651055	0.506262	1	T
LRP6	p.Val1062Ile	HET	rs2302685	0.885583	0.9196	0.850649	0.9335	1	T
LRRK2	p.Arg50His	HOM	rs2256408	0.969249	1	0.993066	1	1	T
LRRK2	p.Met1646Thr	HET	rs35303786	0.004792	0	0.009376	5.81E-05	0.159	T
LRRK2	p.Met2397Thr	HET	rs3761863	0.551717	0.4603	0.618476	0.46794	0.466	T
MAPT	p.Tyr441His	HET	rs2258689	0.312899	0.628	0.281649	0.632031	0.978;0.933;0.978;.	T;T;T;.
MS4A6E	p.Ile6Val	HOM	rs2304935	0.258187	0.254	0.325262	0.270819	1	T
MS4A6E	p.Thr10Ala	HOM	rs2304934	0.258187	0.255	0.325243	0.27093	0.086	T
MS4A6E	p.Val47Phe	HOM	rs2304933	0.257788	0.255	0.325162	0.270275	0.044	D
NME8	p.Cys208Arg	HOM	rs10250905	0.743411	0.5466	0.731553	0.560475	0.046	D
NOTCH3	p.Ala2223Val	HOM	rs1044009	0.629393	0.5724	0.715381	0.584804	0.175	T

PARK2	p.Val380Leu	HET	rs1801582	0.172125	0.0853	0.164749	0.076357	0.377;0.416;0.607;0.	T;T;T;T
								335;0.335;..	..
PARK2	p.Ser167Asn	HET	rs1801474	0.117412	0.3393	0.069934	0.391814	0.219;0.229;0.155	T
PDLIM 5	p.Ser14Phe	HET	rs2452600	0.224641	0.378	0.265912	0.374971	0.024;..	D;..
PDLIM 5	p.Ala374Thr	HOM	rs966845	0.993411	1	0.993729	1	..;1.0;0.702;..;0.709	..;T;T;..;T
PDLIM 5	p.Thr410Ala	HET	rs7690296	0.38758	0.3998	0.44145	0.399779	..;1.0;1.0;..;0.433	..;T;T;..;T
PDLIM 5	p.Ser521Asn	HOM	rs13107595	0.979633	1	0.989681	1	..;0.74;0.571;..;0.91	..;T;T;..;T
PSEN1	p.Glu318Gly	HET	rs17125721	0.005591	0	0.014617	5.80E-05	0.135;0.132;0.135;0..004;0.224	T;T;T;D;T
PTK2B	p.His447His	HOM	rs7005936	0.370208	0.1954
PTPRD	p.Thr781Ala	HET	rs72694737	0.016973	0.001	0.022152	0.000759	1	T
RIN3	p.Thr425Met	HET	rs3742717	0.292532	0.4415	0.241143	0.467308	0.08;..	T;..
RIN3	p.Gly972del	HOM	8153141,rs716 98059
SETX	p.Ile1386Val	HOM	rs543573	0.55611	0.3621	0.736559	0.303514	0.872	T
SETX	p.Gly1252Arg	HOM	rs1183768	0.55611	0.3621	0.736471	0.30334	0.133	T
SETX	p.Asp1192Glu	HOM	rs1185193	0.640575	0.3681	0.769512	0.311333	0.377	T
SFRP4	p.Arg340Lys	HET	rs1802074	0.252995	0.2619	0.197331	0.243691	0.859	T
SFRP4	p.Pro320Thr	HOM	rs1802073	0.542732	0.5635	0.435974	0.550848	0.171	T
SLC6A5	p.Phe124Ser	HOM	rs1443548	0.793131	0.75	0.771566	0.760212	0.44	T
SLC6A5	p.Ala162Gly	HOM	rs1443549	0.99361	1	0.998607	1	1	T
SMC5	p.Val306Ile	HOM	rs1180116	0.855032	0.8532	0.88404	0.855722	0.655	T
SORL1	p.Glu270Lys	HET	rs117260922	0.007788	0	0.015071	0	0.008	D
SORL1	p.Gln1074Glu	HOM	rs1699107	0.984824	1	0.996113	1	0.168;1.0	T
SORL1	p.Val1967Ile	HOM	rs1792120	0.979433	1	0.996169	1	1	T
SYT11	p.Gln48His	HOM	rs822522	0.954673	1	0.990874	0.999942	0.866	T
TET1	p.Asp162Gly	HET	rs10823229	0.258586	0.3919	0.330584	0.383148	0.019	D
TET1	p.Cys832Tyr	HET	rs74925160	0.024161	0	0.005267	0	0	D
TET1	p.Ala985Ser	HET	rs74412312	0.024161	0	0.00526	0	0.013	D
TET1	p.Ile1123Met	HET	rs3998860	0.693291	0.8502	0.777419	0.839626	0.157	T
TM2D3	p.Leu6Arg	HOM	rs2939587	0.993411	1	0.97935	1	0.311;0.318;0.466;0..404	T
TMEM1 06B	p.Thr185Ser	HET	rs3173615	0.595048	0.6508	0.489626	0.657474	0.214	T
TOMM 40	p.Arg239Trp	HET	rs142412517	0.0002	0	0.0005	5.80E-05	0.011;0.011;..;0.011	D;D;..;D
VEPH1	p.Ser522Pro	HET	rs11918974	0.250599	0.3512	0.276265	0.392621	1.0;0.992;0.992	T
WDR64	p.Asn521Asn	HET	rs10802990	0.590855	0.5149
ZCCHC 14	p.Val693Met	HOM	rs3748400	0.461462	0.3115	0.643568	0.307451	0	D