

Table 1. Demographics and Genetic Mutations of the Study Cohort

Patient No.	Gender	Race	Diagnosis /Genetic type	Age at initial visit (yrs.)	Age at most recent visit (yrs.)	Visit No.	Follow-up (mo.)	Genetic mutation
1	M	W	RP/AD	57	58	6	14	
2	M	W	RP/AD	39	41	6	16	<i>RP1</i> gene, Arg677stop heterozygous
3	M	W	RP/AD	43	44	3	6	
4	F	W	RP/AD	31	35	13	58	
5	M	W	USH type-II	26	26	3	8	
6	M	W	USH type-II	22	28	4	7	Negative <i>USH2a</i> gene
7	F	W	USH type-I	39	40	6	23	
8	F	W	USH type-II	26	27	4	14	
9	F	AA	RP/ISO	45	49	13	50	
10	F	AA	RP/ISO	46	47	4	12	
11	F	W	RP/AD	22	23	9	22	<i>RHO</i> gene, Pro23His heterozygous
12	M	W	RP/AD	40	42	12	31	
13	F	W	RP/AR	62	67	14	55	Negative for <i>RP1, RHO, and RDS</i> genes
14	F	W	RP/AD	46	50	13	46	
15	F	W	RP/AD	22	25	3	9	
16	M	AA	RP/ISO	39	39	8	31	
17	F	W	RP/AR	25	26	3	9	
18	M	W	RP/ISO	67	68	9	39	
19	F	W	RP/AR	19	22	7	18	Negative for <i>RHO</i> gene
20	M	W	RP/AR	24	24	6	25	
21	F	W	RP/AD	51	55	6	16	<i>RHO</i> gene, Pro347Leu exon 5 heterozygous
22	F	W	RP/AD	21	23	3	8	
23	F	W	RP/ISO	60	61	3	9	
24	F	W	RP/ISO	55	55	3	10	
25	M	W	RP/ISO	49	49	3	6	
26	M	W	RP/ISO	53	53	4	9	
27	M	Asian	USH type-I	20	20	3	8	
28	M	AA	RP/AD	38	38	3	8	
29	F	W	RP/AD	54	54	4	14	
30	M	W	RP/AR	24	24	3	9	
31	F	W	USH type-I	19	20	3	9	<i>MYO7a</i> gene, Arg669stop exon 17 (paternal) & Gln1798stop exon 39 (maternal)
32	F	W	RP/AD	38	39	3	9	

F=female; *M*=male; *W*=white; *AA*=African American; *RP*=retinitis pigmentosa; *USH*=Usher syndrome; *AD*=autosomal dominant; *AR*=autosomal recessive; *ISO*=isolated; yrs. =years; mo. =months.