

**Table S1.** LIAVA haplotype and arrays that cause blue cone monochromacy.

Subject ID	Age (yr)	Genotype	SER OD	SER OS	BCVA
*BCM73-20770	0.25	M <sub>LIAVA</sub>	-8	-9	20/100
*BCM73-20537	3	M <sub>LIAVA</sub>	-4	-4	20/100
S16	10	L <sub>LIAVA</sub> -L <sub>LIAVA</sub> -M <sub>MVVVA</sub>	<-6 D	<-6 D	20/60
*MOL0057 III:3	12	M <sub>LIAVA</sub>	-6	-6	20/160
BCM72-17075	12	L <sub>LIAVA</sub> -M <sub>LIAVA</sub> -M <sub>MVVVA</sub>	0.5	0.5	20/200
*BCM73-17481	14	M <sub>LIAVA</sub>	-14	-14	20/400
BCM93-19164	14	M <sub>LIAVA</sub>	-5	-3	20/67
S17	18	L <sub>LIAVA</sub> -L <sub>LIAVA</sub> -M <sub>MVVVA</sub>	<-6 D	<-6 D	20/40
*MOL0057 III:I	20	M <sub>LIAVA</sub>	-18	-18	20/320
MM_0155	30	L <sub>LIAVA</sub> - M <sub>LIAVA</sub>	ND	ND	ND
*BCM73-16953	51	M <sub>LIAVA</sub>	-12	-12	20/133
S21	73	L <sub>LIAVA</sub>	<-6 D	<-6 D	20/200

\*BCM73-20770, BCM73-20537, BCM73-16953 are relatives, see reference [1] for pedigree. MOL0057 III:3 and III:I are relatives, see reference 15 for pedigree. Subjects with IDs starting with BCM are from reference [1], those starting with S are from reference [2], those starting with MOL are from reference [3]. MM\_0155 is from reference [4].

**Table S2.** LIAVA haplotype and arrays that cause red-green color vision defects.

Subject ID	Age (yr)	Color vision	Genotype	SER OD	SER OS	BCVA OD	BCVA OS	Photopic ERG
*MOL0152 III:2	5	ND	LLIAVA - M <sub>MVVVA</sub>	-5.5	-6.5	20/100	20/200	reduced
*MOL0152 III:3	5	ND	LLIAVA - M <sub>MVVVA</sub>	-8	-8.75	20/67	20/50	reduced
BCM160-23130	6	impaired	LLIAVA - M <sub>MVVVA</sub>	-3.75	-5	20/67	20/40	reduced
*MOL0152 III:1	7	ND	LLIAVA - M <sub>MVVVA</sub>	-9.25	-9.5	20/40	20/50	reduced
ZD314-18057	10	impaired	LLIAVA - M <sub>MVVVA</sub>	-6.5	-6	20/100	20/100	reduced
JC_0609	11	Protanope	LLIAVA - M <sub>MVVVA</sub>	-14	-13.75	20/20	20/20	ND
BCM51-12359	12	impaired	LLIAVA - M <sub>MVVVA</sub>	-14.5	-14.5	20/100	20/100	reduced
MM_0142	13	Protanope	LLIAVA - M <sub>MVVVA</sub>	-2	-2	ND	ND	ND
*MM_0145	15	Protanope	M <sub>LLIAVA</sub> - M <sub>MVVVA</sub>	0	0	20/20	20/30	ND
*MM_0144	19	Protanope	M <sub>LLIAVA</sub> - M <sub>MVVVA</sub>	-8	-8.25	20/70	20/40	ND
JC_0196	28	Protanope	LLIAVA-M <sub>MVVVA</sub> -M	-6	-5	20/40	20/40	ND
JC_0195	29	Protanope	LLIAVA-M <sub>MVVVA</sub> -M	-11.5	-12	20/60	20/60	ND
JC_0084	38	Deutanope	L - M <sub>LLIAVA</sub>	-2.5	-2.5	20/20	20/15	ND

\*MOL0512 III:2, III:3, III:1 are relatives; see reference [3] for pedigree; MM\_0145 and MM\_0144 are brothers, JC\_0195 and JC\_0196 are brothers, the genotype of MM\_0145 not done, presumed to be same as his brother. Subject IDs starting with MOL are from reference [3], those starting with BCM or ZD are from reference [1], those starting with MM or JC are from references [4,5]. ND = not done.

**Table S3.** MIAVA haplotype and arrays that cause red-green color vision defects.

Subject ID	Age (yr)	Genotype	SER OD	SER OS	BCVA
BCM101-19818	3	LLIAVA-MMIAVA	-6	-6	20/200
*ZD379-19195	7	LLIAVA-MMIAVA	1	1	20/125
*S22	9	LLIAVA- LMIAVA-MMIAVA	<-6 D	<-6 D	20/125
*ZD379-19194	12	LLIAVA-MMIAVA	ND	ND	20/125
*S23	68	LLIAVA- LMIAVA-MMIAVA	>-3D	>-3D	20/200
*S24	34	LMIAVA- LMIAVA-MMVVVA	<-6 D	<-6 D	20/80
*S25	51	LMIAVA- LMIAVA-MMVVVA	<-6 D	<-6 D	20/63

Subjects ZD379-19195, -19194 are relatives; see pedigree in reference [1]. Subjects S22 and S23 are relatives, and S24 and S25 are relatives, see pedigrees in reference [2]. Subjects' IDs starting with BCM or ZD are from reference [1], those beginning with S are from reference[2].

**Table S4.** LVAVA haplotype and red-green color vision deficiency.

Subject ID	Age Yrs	Color Vision	Diag.	Genotype	AL OD	AL OS	SER OD	SER OS	BCVA	Pho- topic ERG
BCM194-25474	5	NV	BCM	LLVAVA	ND	ND	-16	-16.25	20/40	re- duced
S18	10	I	XLCD	LLVAVA	ND	ND	<-6D	<-6D	20/50	re- duced
*JC_11445	26	D	XLCD	LLVAVA – LMVVVA – M – M	25.79	25.65	ND	ND	ND	ND
*JC_0758	33	DA	XLCD	LLVAVA – LMVVVA – M – M	29.18	28.53	-12.75	-10.00	20/50	ND
JC_0347	33	D	XLICD	LLVAVA	24.77	24.25	-3.3	-2	20/200	ND
JC_0683	36	DA	XLCD	LLVAVA – LMVVVA – M – M	27.03	26.70	-8.5	-8.5	20/15	ND
BCM66-16407	41	I	BCM	LLVAVA	ND	ND	-13	-14	20/67	absent
S26	80	I	XLCD	LLVAVA-LLVAVA-M <sub>MVVVA</sub>	ND	ND	>-3 D	>-3 D	20/125	re- duced
*BCM112-22852	6	I	BCM/CRD	M <sub>LVAVA</sub>	ND	ND	-9	-9.25	20/67	re- duced
*JC_0451	12	P	XLCD	M <sub>LVAVA</sub> – M – M	28.1	27.6	-9.5	-9	20/20	ND
*JC_0448	14	P	XLCD	M <sub>LVAVA</sub> – M – M	27.75	27.51	-11.5	-10.75	20/25	ND
*BCM112-23518	14	I	BCM/CRD	M <sub>LVAVA</sub>	ND	ND	-24	-23	20/125	re- duced
*JC_0447	17	P	XLCD	M <sub>LVAVA</sub> – M – M	26.54	26.52	-9.75	-9	20/20	ND
*JC_10340	32	P	XLCD	M <sub>LVAVA</sub> – M <sub>MVVVA</sub> – M	28.96	28.08	-17.5	-13	20/20	ND
S19	40	I	XLCD	M <sub>LVAVA</sub>	ND	ND	<-6D	<-6D	20/80	re- duced
JC_0564	45	P	XLICD	M <sub>LVAVA</sub>	27.08	26.58	-5.8	-4.9	20/100	re- duced
S20	49	I	XLCD	M <sub>LVAVA</sub>	ND	ND	<-6D	<-6D	20/63	re- duced

\*JC\_11445 and JC\_11445 are brothers; JC\_0683 and JC\_0758 are cousins; JC\_0451, JC\_0448, and JC\_0447 are brothers and JC\_10340 is their cousin; BCM112-22852 and BCM112-23518 are from the same family. Subject IDs starting with BCM are from reference [1], subject IDs starting with JC\_ are from references [4,5], subject IDs starting with S are from reference [2]. CRD=cone-rod dystrophy. Color vision test results: NV = test not valid; D = deuteranope, DA = deuteranomalous, P = protanope, I = impaired. Yellow shaded rows = subjects with LVAVA haplotype expressed in all non-S cones.

**Table S5.** LVAVA haplotypes and arrays for normal color vision.

Subject ID	Age Yrs	Color Vision	Diag.	Genotype	SER OD	SER OS	BCVA OD	BCVA OS
*BCM133-23364	1	ND	BCM	LLVAVA – M <sub>LVAIA</sub>	-5	-5	ND	ND
*BCM133-20960	10	I	BCM	LLVAVA – M <sub>LVAIA</sub>	-5	-6	20/33	20/40
*BCM133-20961	32	ND	BCM	LLVAVA – M <sub>LVAIA</sub>	-20.75	-19	20/33	20/55
*HM346IV:2	15	Normal	high myopia	LLVAVA – M	-11	-12.25	20/20	20/30
*HM346III:7	27	ND	high myopia	LLVAVA – M	-12.5	-12.5	20/67	20/80
*HM346II:4	47	Normal	high myopia	LLVAVA – M	-13.75	-14	20/200	20/125
*HM346II:2	54	ND	high myopia	LLVAVA – M	-18.25	-16.25	20/200	20/100
*HM295 IV4	6	Normal	high myopia	LLVAVA – M	-10.5	-10.5	ND	ND
*HM295 IV2	12	Normal	high myopia	LLVAVA – M	-8.25	-9	20/30	20/25
*HM295 IV3	12	Normal	high myopia	LLVAVA – M	-9.25	-9.25	20/40	20/40
*HM295 IV7	14	Normal	high myopia	LLVAVA – M	-9.25	-10.25	20/30	20/30
*HM295 IV1	19	Normal	high myopia	LLVAVA – M	-6	-6.25	20/25	20/30
*HM295 IV5	31	Normal	high myopia	LLVAVA – M	-15	-15	20/30	20/40
*HM295 II:5	76	Normal	high myopia	LLVAVA – M	ND	ND	ND	ND
BCM126-20616	41	I	BCM	LLVAVA-M <sub>LVAIA</sub>	ND	ND	20/80	20/100

\*Subject IDs beginning with BCM are from the same reference and BCM133- are from the same family and are from reference [1], subject IDs beginning with HM are from reference [6]. Yellow shaded row = subject expresses LVAVA haplotype in all non-S cones.

**Table S6.** MVAVA haplotype and arrays for normal color vision.

Subject ID	Age Yrs	Color Vision	genotype	AL OD OS	OD SER	OS SER	BCVA OD	BCVA OS	Photopic ERG
Orosz IV:5	40	ND	LLVAVA – MMVAVA	25.16 25.51	-4.25	-5.25	20/20	20/20	ND
Orosz IV:5	57	DA	LLVAVA – MMVAVA	ND	NA	NA	20/25	20/40	ND
Orosz IV:5	62	I	LLVAVA – MMVAVA	ND	NA	NA	20/25	20/40	absent
Orosz IV:7	51	I	LLVAVA – MMVAVA	27.82 27.57	-5.75	-6.5	20/50	20/33	absent
Orosz V:1	32	ND	LLVAVA – MMVAVA	31.44 30.72	ND	ND	20/20	20/20	ND
Orosz V:1	38	I	LLVAVA – MMVAVA	ND	-20.0	-18.0	20/20	20/20	absent
Orosz V:1	46	I	LLVAVA – MMVAVA	33.11 32.28	-21.0	-19.75	20/20	20/25	ND
Orosz VI:3	42	I	LLVAVA – MMVAVA	31.35 32.49	-22.5	-24.5	20/20	20/20	ND
Orosz VI:6	11	N	LLVAVA – MMVAVA	ND	-6.25	-6.5	20/40	20/20	reduced

All subjects are from the same Hungarian family in reference [7]. Color vision: ND = not done; DA = deuteranoma-  
lous, I = impaired, N = normal trichromatic color vision.

**Table S7.** LIAVS haplotype.

Subject ID	Age (Yr)	Genotype	SER OD	SER OS	BCVA OD	BCVA OS
*MOL0250 III:2	50	LLIAVS	-2.25	-1.75	20/67	20/67
*+MOL0250 IV:3	24	LLIAVS	-4.00	-5.50	20/67	20/67
*+MOL0250 IV:3	27	LLIAVS	ND	ND	20/67	20/67
*+MOL0250 IV:3	32	LLIAVS	-2.75	-5.00	20/80	20/100
MOL0267 III:1	34	LLIAVS	ND	ND	20/67	20/67
BCM72- 16874	71	LLIAVS-MLIAVA- MMVVVA	1.75	0.75	20/200	20/200

\*members of the same family; see reference [3] for pedigree. +same individual examined over an 8-year period. BCM72-16874 is from reference [1].

#### References

1. Buena-Atienza, E.; Ruther, K.; Baumann, B.; Bergholz, R.; Birch, D.; De Baere, E.; Dollfus, H.; Greally, M.T.; Gustavsson, P.; Hamel, C.P.; et al. De novo intrachromosomal gene conversion from OPN1MW to OPN1LW in the male germline results in Blue Cone Monochromacy. *Scientific reports* **2016**, *6*, 28253, doi:10.1038/srep28253.
2. Gardner, J.C.; Liew, G.; Quan, Y.H.; Ermetal, B.; Ueyama, H.; Davidson, A.E.; Schwarz, N.; Kanuga, N.; Chana, R.; Maher, E.R.; et al. Three different cone opsin gene array mutational mechanisms with genotype-phenotype correlation and functional investigation of cone opsin variants. *Human mutation* **2014**, *35*, 1354-1362, doi:10.1002/humu.22679.
3. Mizrahi-Meissonnier, L.; Merin, S.; Banin, E.; Sharon, D. Variable retinal phenotypes caused by mutations in the X-linked photopigment gene array. *Investigative Ophthalmology and Visual Science* **2010**, *51*, 3884-3892, doi:doi10.1167/iovs.09-4592
4. Patterson, E.J.; Kalitzeos, A.; Kasilian, M.; Gardner, J.C.; Neitz, J.; Hardcastle, A.J.; Neitz, M.; Carroll, J.; Michaelides, M. Residual Cone Structure in Patients With X-Linked Cone Opsin Mutations. *Invest Ophthalmol Vis Sci* **2018**, *59*, 4238-4248, doi:10.1167/iovs.18-24699.
5. Patterson, E.J.; Wilk, M.; Langlo, C.S.; Kasilian, M.; Ring, M.; Hufnagel, R.B.; Dubis, A.M.; Tee, J.J.; Kalitzeos, A.; Gardner, J.C.; et al. Cone Photoreceptor Structure in Patients With X-Linked Cone Dysfunction and Red-Green Color Vision Deficiency. *Invest Ophthalmol Vis Sci* **2016**, *57*, 3853-3863, doi:10.1167/iovs.16-19608.
6. Li, J.; Gao, B.; Guan, L.; Xiao, X.; Zhang, J.; Li, S.; Jiang, H.; Jia, X.; Yang, J.; Guo, X. Unique variants in OPN1LW cause both syndromic and nonsyndromic X-linked high myopia mapped to MYP1. *Investigative ophthalmology & visual science* **2015**, *56*, 4150-4155.
7. Orosz, O.; Rajta, I.; Vajas, A.; Takacs, L.; Csutak, A.; Fodor, M.; Kolozsvari, B.; Resch, M.; Senyi, K.; Lesch, B.; et al. Myopia and Late-Onset Progressive Cone Dystrophy Associate to LVAVA/MVAVA Exon 3 Interchange Haplotypes of Opsin Genes on Chromosome X. *Invest Ophthalmol Vis Sci* **2017**, *58*, 1834-1842, doi:10.1167/iovs.16-21405.