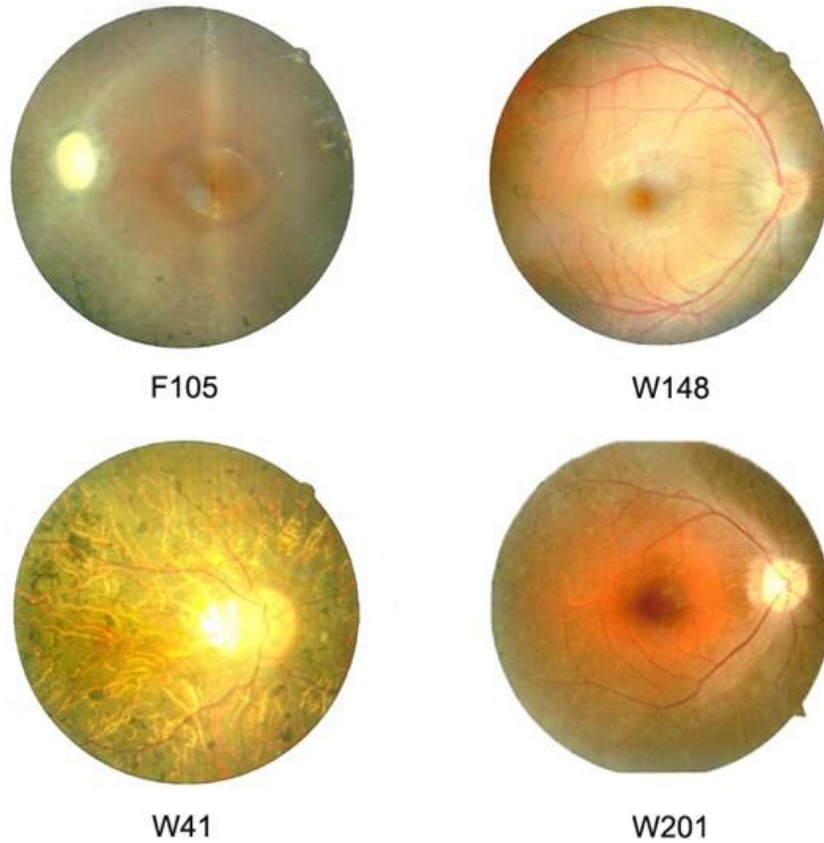


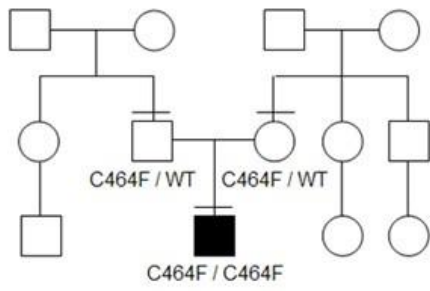
## SUPPLEMENTARY INFORMATION

### Supplementary Figures

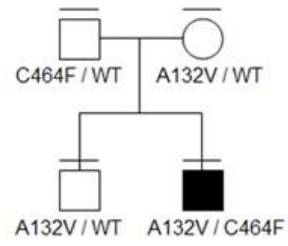


### **Supplementary Fig. 1. Fundoscopy of the four additional patients with distinct SLC7A14 mutations**

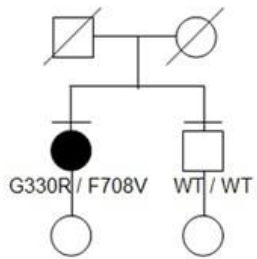
Patients with the s mutation displayed typical manifestations in their fundi including peripheral or pan-retina bone-spicule pigmentation, optic disc pallor and arterial attenuation. F105 had a homozygous mutation; W148, W41 and W201 carried two compound mutations.



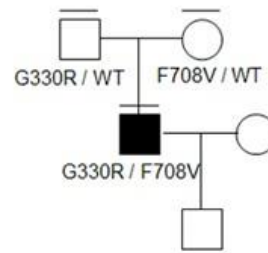
F105



W148

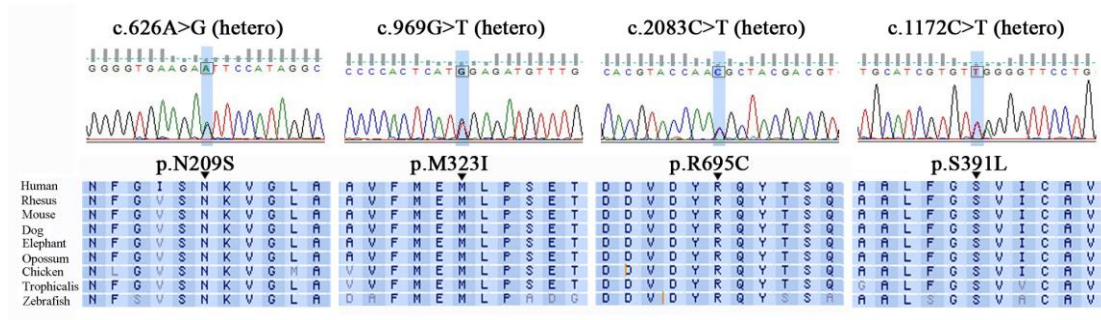


W41



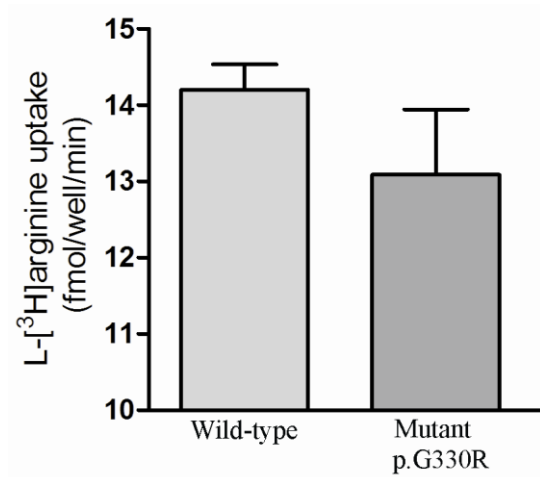
W201

**Supplementary Fig. 2. Pedigrees with SLC7A14 mutations**



**Supplementary Fig. 3. Identified mutations in the SLC7A14 gene.**

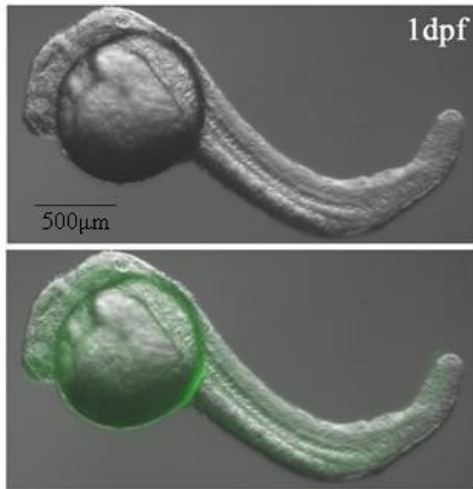
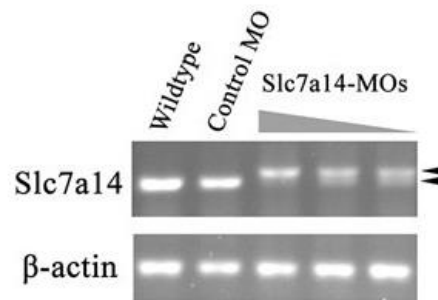
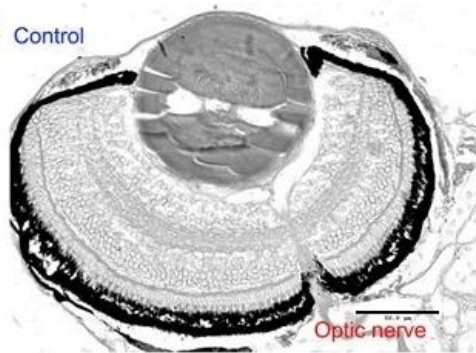
Sequence chromatograms illustrated the causal lesions identified in five unrelated arRP patients. Each mutation leads to alteration of a highly conserved amino acid residue.



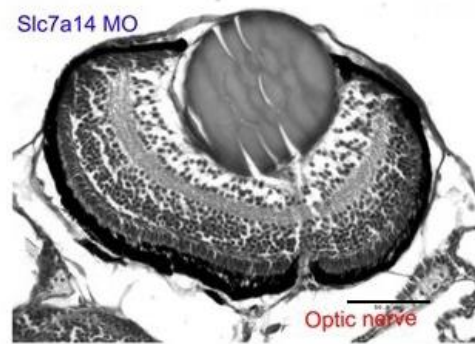
**Supplementary Fig. 4. L-[3H]arginine uptake in photoreceptor cells with either wild-type or mutant *Slc7a14* gene.**

Four independent experiments (N=4) were performed and the results were analyzed using Student's t-test. Bars in the graph represent standard errors of the mean (SEM).

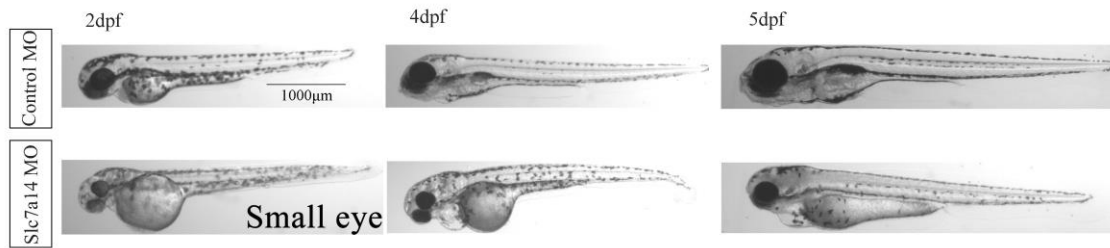
No significance was observed.

**a** Fluoresceinated MO**b** RT-PCR**c** Control

## Slc7a14 MO

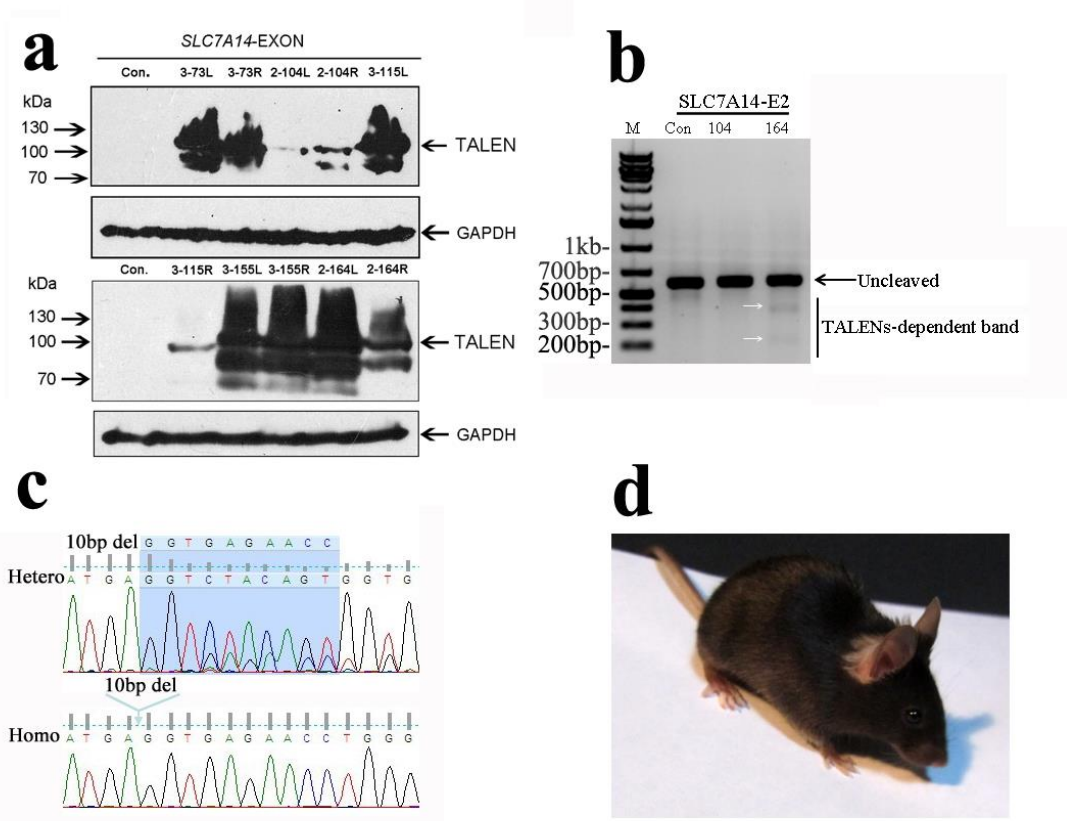
**Supplementary Fig. 5. Confirmation of MO injection in zebrafish Larvae.**

(a) After injection of fluoresceinated standard control MO and *Slc7a14* MO solution, fluoresceins were observed in 1 dpf Larvae. Scale bar, 500 $\mu$ m. (b) RT-PCR confirmed splicing change in *Slc7a14*-MO injected larvae. Three doses (1nL of 0.25nM, 0.15nM, 0.05nM) of MOs were tested. (c) Immunohistology demonstrated no significant changes of retinal structure in the *Slc7a14*-MO zebrafish. Each section was shown with optic nerve. Scale bar, 50 $\mu$ m.



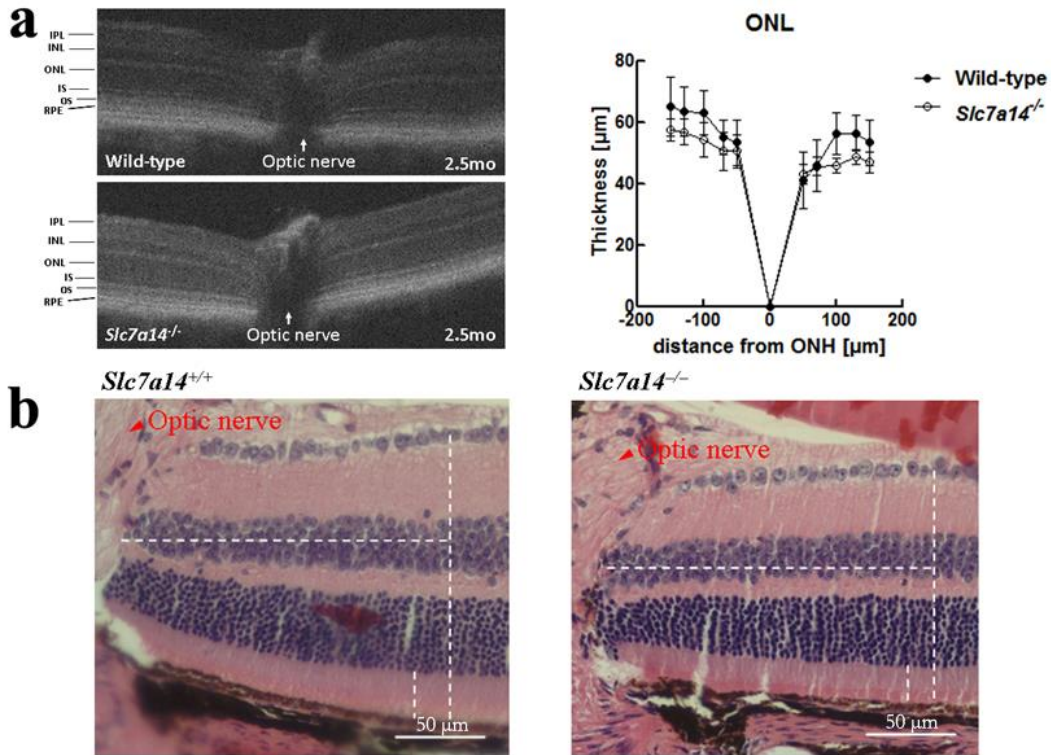
**Supplementary Fig. 6. Representative images of the MO-injected zebrafish.**

Comparing with the control MO-injected zebrafish, Slc7a14-knockdown led to dramatically smaller eye size. Scale bar, 1000µm.



### Supplementary Fig. 7. *Slc7a14*-deficient mice

(a) Western blot confirmed the TALEN expression in the transfected C6 cells. Each lane represents an independent experiment. (b) MSDase assay demonstrated the activity of TALEN pair E2-164. (c) Genotyping of the KO mice revealed a 10bp deletion. Upper image, heterozygous; lower image, homozygous. (d) The *Slc7a14*-deficient mice were viable, fertile and did not show noticeable physical abnormalities.



**Supplementary Fig. 8. OCT results of 2.5-month-old *Slc7a14*-KO mouse**

(a) Compared to age-matched wild-type mice, the 2.5-month-old *Slc7a14*<sup>-/-</sup> mice had a thinner outer nuclear layer (2.5-month age, n=3). Due to the limited number of the mice (n=3), we were unable to perform any statistical analyses. Bars represent SEM. Bonferoni test, no statistical significance. (b) HE staining of a histological section of retina shows a thinner retina in the *Slc7a14*<sup>-/-</sup> mice. Each retinal section shows the optic nerve as an internal reference. Scale bar, 50 $\mu\text{m}$ .



Fig. 3a

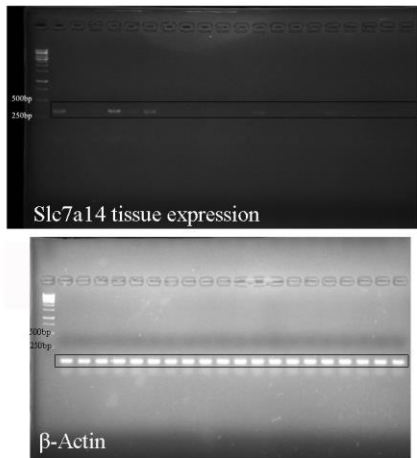


Fig. 3b

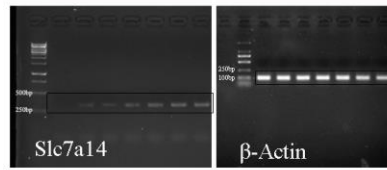
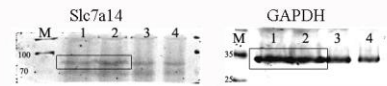
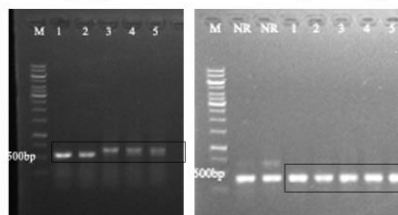


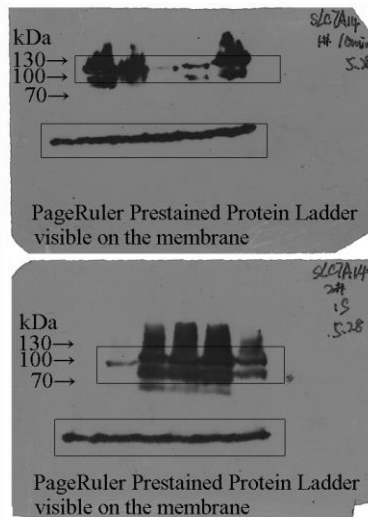
Fig. 3c



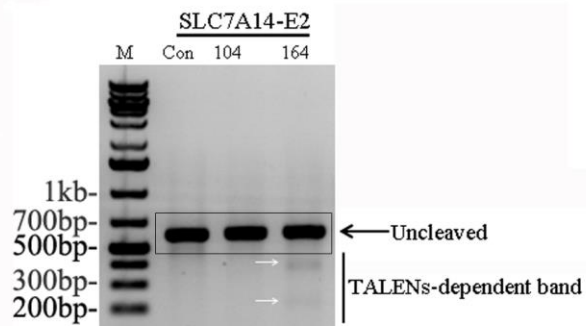
Supplementary fig. 5b



Supplementary fig. 7a



Supplementary fig. 7b



Supplementary Fig. 9. Data of full blots.

Boxes indicate lanes used in the labeled figures.

## Supplementary Tables

**Supplementary Table 1. List of the 15 homozygous variants after adopting filtering**

<i>Chr</i>	<i>Location</i>	<i>Gene symbol</i>	<i>Variant</i>	<i>In homo region&gt;1Mb</i>
1	248813827	<i>OR2T27</i>	c.359A>G,	No
12	53681883	<i>ESPL1</i>	c.4304A>T,	No
19	36349333	<i>KIRREL2</i>	c.235G>A,	No
19	38877754	<i>GGN</i>	c.148T>C,	No
2	219319657	<i>USP37</i>	c.2936T>C,	No
7	87022264	<i>CROT</i>	c.1599C>A,	No
9	125512575	<i>OR1L6</i>	c.449G>A,	No
X	154128188	<i>F8</i>	c.6226G>A,	No
11	47609867	<i>FAM180B</i>	c.433G>A,	No
<b>11</b>	5537592	<b><i>UBQLNL</i></b>	c.80C>G,	<b>Yes</b>
<b>2</b>	21257717	<b><i>APOB</i></b>	c.875C>G,	<b>Yes</b>
<b>6</b>	64422984	<b><i>PHF3</i></b>	c.5500C>T,	<b>Yes</b>
<b>8</b>	27295006	<b><i>PTK2B</i></b>	c.1520A>G,	<b>Yes</b>
<b>X</b>	118586003	<b><i>SLC25A43</i></b>	c.722G>A,	<b>Yes</b>
<b>3</b>	170201230	<b><i>SLC7A14</i></b>	c.988G>A,	<b>Yes</b>

**Supplementary Table 2. Primer sets for *SLC7A14* amplification**

Exon	Forward primer	Reverse primer	Tm	Length
1	GAGAAGGCTGCACTGGGTC	TGATGATCCACTTTTCTGCC	58	565bp
	CAGAAGTGAGGTTGGAAGA	AAGGATGGATAGATTCATAGAG		
2	GG	AAAG	58	576bp
3	CCTCGGCCTCCCAGTTTAC	GGTGATAAAGAGGGTGCTGTG	58	468bp
4	TGCGTGTTTAGCACGAGTTC	GGCTACAGGGGACAAAAGAC	58	413bp
5	GTGAATGGATCCAAACTGCC	GATACATAACAAGGTCCACAGT CC	58	468bp
	TAAATTGGTGGAGCAGGAC	CCATAGAACCAGCCAGATTTTC		1132b
6	C		58	p
7	TGTGGATGCAGTGTTGTCAG	GGTGGTTGCACCTAAGATGG	58	856bp

**Supplementary Table 3. Primers for site-directed mutagenesis**

<b>Primer</b>	<b>Sequence</b>
hSLC7A14-full-F	GGAGCTCGCCACCATGAGTGGCTTCTTCAC CTCGCTG
hSLC7A14-full-R	GGGTACCGCTCTGGAGAGTAATCTAACTCA TC
hSLC7A14-mut-F	GTTTGTGGCTCATAGGTTCTATGCTGC
hSLC7A14-mut-R	GCAGCATAGAACCTATGAGCCACAAAC
hCAT2-NheI-F	GCTAGCATGATTCCTTGCAGAGCCGC
hCAT2-XhoI-R	CTCGAGTTAGAATTCACCTTGTCTTTTCATG
hSLC7A14-fusion-F	GGATCCGCCTACACCTACAGCTATG
hSLC7A14-fusion-R	GGTACCGAAGGAGCAAGACACAGA