Supplementary Information For

Functional characterization of ABCA4 genetic variants related to Stargardt disease

Bo Min Kim^{1,2}, Hyo Sook Song^{1,2}, Jin-Young, Kim¹, Eun Young Kwon¹, Seung Yeon Ha¹, Minsuk Kim¹, Ji Ha Choi^{1,*}

¹Department of Pharmacology, Inflammation-Cancer Microenvironment Research Center, College of Medicine, Ewha Womans University, Seoul, Korea

²B.M.Kim and H.S.Song contributed equally to this study.

*Corresponding author: jihachoi@ewha.ac.kr

Correspondence to

Ji Ha Choi, M.D., Ph.D.

Department of Pharmacology, Inflammation-Cancer Microenvironment Research Center

College of Medicine, Ewha Womans University, Seoul, Korea

25 Magokdong-ro 2-gil, Gangseo-gu, Seoul 07804, Korea

Phone: +82-2-6986-6173; Fax: +82-2-6986-7017; E-mail: jihachoi@ewha.ac.kr

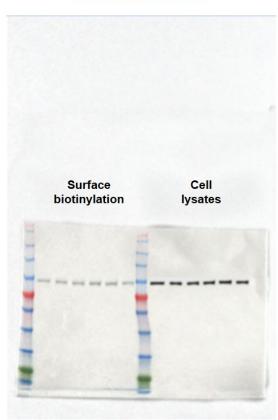
Supplementary Figure 1. Effect of *ABCA4* **variants on** *ABCA4* **surface expression.** HEK-293T cells were transfected with *ABCA4* wild-type or variant plasmids, and a surface biotinylation assay was conducted. Among figures, the figures named 'Repeat 1' were used in the main manuscript.

Repeat 1

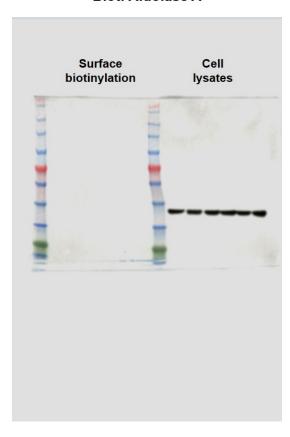
Blot: ABCA4

Surface Cell biotinylation lysates

Blot: Na⁺/K⁺ ATPase

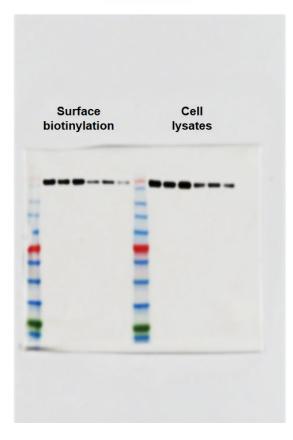


Blot: Aldolase A

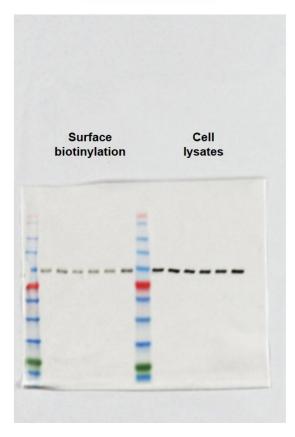


Repeat 2

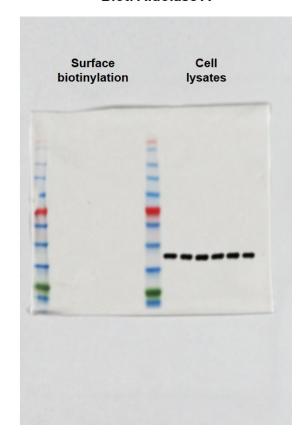
Blot: ABCA4



Blot: Na⁺/K⁺ ATPase

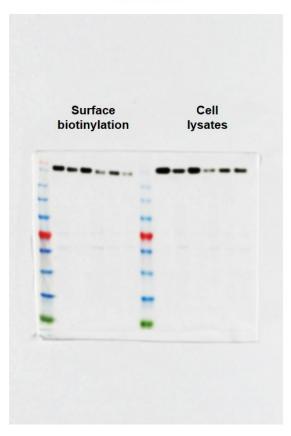


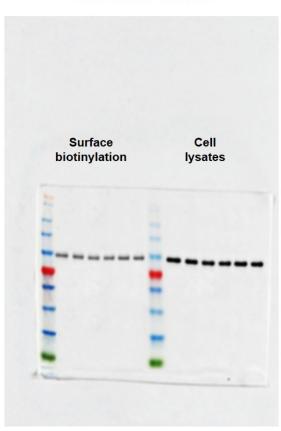
Blot: Aldolase A

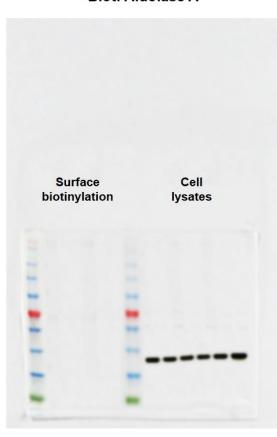


Repeat 3

Blot: ABCA4 Blot: Na⁺/K⁺ ATPase Blot: Aldolase A







Supplementary Figure 2. Effect of MG132 or bafilomycin A₁ on *ABCA4* expression. *ABCA4* expression was examined after transfection with *ABCA4* wild-type or variant plasmids. Immunoblotting was performed after treatment with MG132 (a) or bafilomycin A₁ (b). Among figures, the figures named 'Repeat 1' were used in the main manuscript.

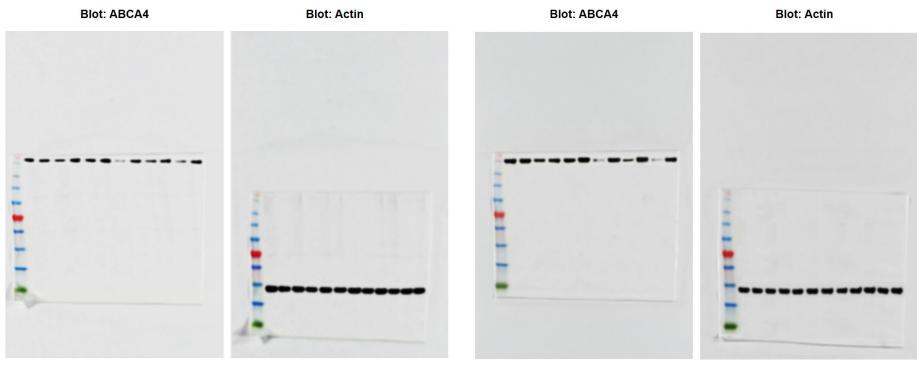
(a)

Repeat 1

Blot: ABCA4

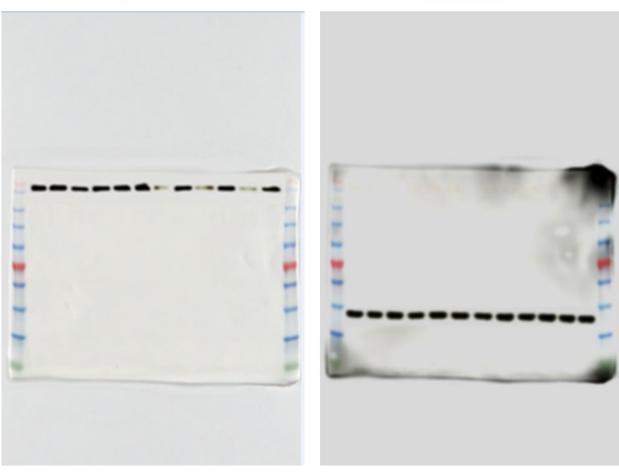
Blot: Actin

Repeat 2 Repeat 3



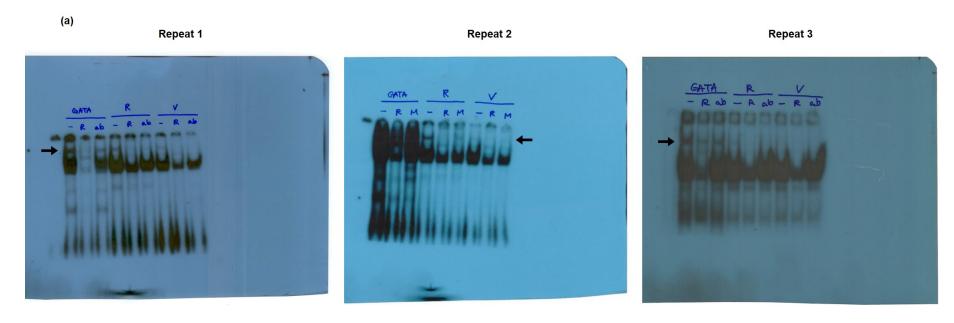
Repeat 1

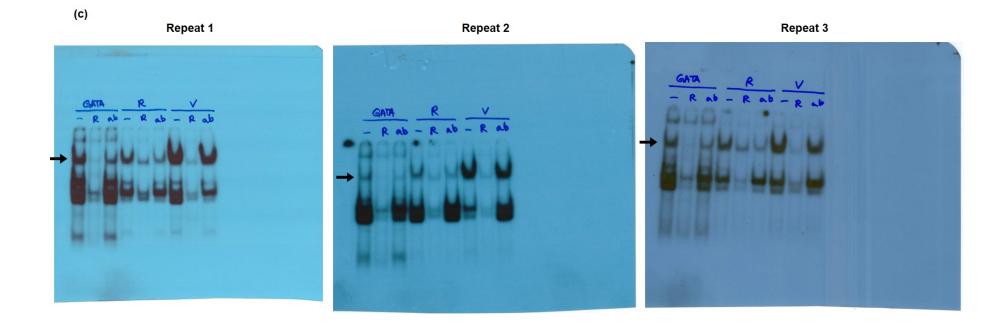
Blot: ABCA4 Blot: Actin



Repeat 2
Blot: ABCA4
Blot: Actin
Blot: ABCA4
Blot: Actin
Blot: ABCA4
Blot: Actin
Blot: ABCA4
Blot: Actin

Supplementary Figure 3. Gel shift assays. ³²P-labeled oligonucleotides [lanes 1-3, GATA-2 consensus; lanes 4-6, wild-type c.-1086A; lanes 7-9, c.-1086C variant, (a), lanes 1-3, HLF consensus; lanes 4-6, wild-type c.-900A; lanes 7-9, c.-900T variant, (b), lanes 1-3, GATA-2 consensus; lanes 4-6, wild-type c.-761C; lanes 7-9, c.-761A variant, (c)] were incubated with nuclear protein extracts. Competition and supershift assays were performed. Each arrow indicates the band of the DNA-protein complex. Among figures, the figures named 'Repeat 1' were used in the main manuscript.





Supplementary Table 1. Oligonucleotides used in the construction of ABCA4 plasmids and or in gel shift assays

Primes for ABCA4 promoter cloning ¹	
Sense (KpnI site)	5'-CGG GGT ACC GGA GCC TTT TTG CCA TCA TAA GC-3'
Antisense (NheI site)	5'-CTA GCT AGC AGG GTC CAG TTC TTC CAG AGC AA-3'
Primers for ABCA4 mutagenesis PCR ²	
p.Arg290Gln (c.869G>A)	5'-TCA AGA GTT TAT CCA TCA GCC GAG TAT GCA GGA CT-3'
p.Asp645Asn (c.1933G>A)	5'-CC CTG CTT CGT GGA CAA TTC TTT CAT GAT CAT C-3'
p.Thr1117Ala (c.3349A>G)	5'-GAA CCA TCA TCA TGT CCG CTC ACC ACA TGG ACG AG-3'
p.Cys1140Trp (c.3420C>G)	5'-GAA GGC TCT ACT GGT CAG GCA CCC CAC-3'
p.Asn1588Tyr (c.4762A>T)	5'-TGG CCG GAT CAT GTA TGT GAG CGG GGG-3'
c1086A>C	5'-GCC TTT TTG CCA TCA TAA GCA GAA ACT CTC TCT CTC TTC TTG-3'
c900A>T	5'-GTC TAG AGT CTT TCA AAG AGT ACA CAT TCT GAG ATT TGA GG-3'
c761C>A	5'-CCA CCC CAT TGC AGG GAT GGA ATG ACA GTA ATG-3'
Oligonucleotides in gel shift assay	
c1086A (wild-type) ²	5'-TAA GCA GAA ACT ATC TCT CTC TTC T-3'
c1086C (variant) ²	5'-TAA GCA GAA ACT CTC TCT CTC TTC T-3'
c900A (wild-type) ²	5'-TCT TTC AAA GAG A AC ACA TTC TGA G-3'
c900T (variant) ²	5'-TCT TTC AAA GAG TAC ACA TTC TGA G-3'
c761C (wild-type) ²	5'-CCC ATT GCA GGG CTG GAA TGA CAG T-3'
c761A (variant) ²	5'-CCC ATT GCA GGG ATG GAA TGA CAG T-3'
GATA-2 consensus ³	5'-CAC TTG ATA ACA GAA AGT GAT AA C TCT-3'
HLF consensus ³	5'-CAG GGT TAC GTA ATC TGC T-3'

¹Restriction endonuclease sites, ²SNP sites, and ³consensus sequences of transcription factors are marked with bold-faced letters.