Supplemental Materials Molecular Biology of the Cell

Zhao et al.

Supplemental Figure S1. Strategy for screening recessive mutations on 2L.

(A) The second chromosome of *ey-flp*,*Rh1-GFP*;*FTR40A* flies was isogenized and flies were mutagenized by feeding 25 mM EMS (Sigma) in 2% sucrose for 8 h. They were then mated to *ey-flp Rh1-GFP*;*GMR-hid CL FRT40A/Cyo hs-hid* flies. Flies were heat shocked at 37°C for 1 h to avoid having heterozygous flies among the F1 progeny (the pro-apoptosis gene *hid* was induced by heat-shock (*hs-hid*) in heterozygous flies). The DPP was assessed in F1 flies using a fluorescent stereo microscope day 1 and 5 following eclosion.

Supplemental Figure S2. Alignment of different Drosophila Vha68 proteins.

Amino acid sequences of *Drosophila* Vha68-1, Vha68-2, and Vha68-3, as well as human ATP6V1A are shown. Identical residues, found in at least two proteins, are enclosed in black boxes. The running tally of amino acids is indicated to the right.

Supplemental Figure S3. Decreased light sensitivity and reduced Rh1 levels in *vha68-1*¹ flies.

(A) Light sensitivity assay was performed using ~1-day-old wild-type, *vha68-1*¹, and *vha100-1*² flies. Flies were dark-adapted for 2 min before being exposed to five pulses of 5-sec of white light with increasing intensity. The maximal light intensity was 300 Lux (10⁻¹). (B) Total Rh1 levels were decreased in *vha68-1*¹ flies. Head extracts were prepared from 1-day-old flies of indicated genotypes and labeled using antibodies against Rh1 and INAD. (C) Quantification of Rh1 levels in each genotype. Rh1 levels were normalized to INAD levels, which served as a loading control. *n* = 7, data are presented as mean ± SD, ***p < 0.001 (Student's unpaired t-test).

Supplemental Figure S4. Localization of the scaffold protein INAD and the basolateral membrane protein Na⁺K⁺ATPase were not affected in *vha68-1*¹ flies.

(A) Tangential resin-embedded retina sections from ~1-day-old wild type and *vha68-1*¹ flies were labeled using antibodies against Rh1 (green) and INAD (red). The localization pattern of INAD was indistinguishable between control and *vha68-1*¹ flies. Scale bars are 2 μ m. (B) Adult eyes from wild type and *vha68-1*¹ flies were dissected and stained with antibodies against Na⁺K⁺ATPase (red). GFP fluorescence of Rh1-GFP was directly observed to indicate rhabdomeres. The localization pattern of Na⁺K⁺ATPase was not affected in *vha68-1*¹ flies. Scale bars are 2 μ m.

Supplemental Figure S5. Schematic view and representative images of the Rh1 pulse-chase assay.

(A) Schematic shows the Rh1 pulse-chase assay. Newly hatched flies were exposed to 37°C for 1 h, and then placed in the dark. Flies were collected at indicated time points (6, 18, 24, and 48 h) after heat-shock. Rh1-GFP localized in the rhabdomere, marked by green; hs-Rh1-RFP was synthesized in the ER and transported to the rhabdomere, marked by red. N, nuclear; C, Cell body; R, rhabdomere. (B-E) Cryostat sections of *hs-Rh1-RFP/+* and *vha68-1¹;hs-Rh1-RFP/+* heads at 6 (B), 18 (C), 24 (D), and 48 h (E) after heat shock were labeled with antibodies against RFP. GFP fluorescence of Rh1-GFP was directly observed to indicate rhabdomeres. Scale bars are 5 μm.

Supplemental Figure S6. Vha68-1 functions in Rh1 post-Golgi trafficking to rhabdomere.

(A) Cryostat sections of ninaE-gal4;UAS-shi^{ts1} (ey-flp

rh1-GFP;ninaE-gal4/+;hs-rh1-RFP/UAS-shi^{4s1}), *vha68-1*¹ (ey-flp *rh1-GFP;vha68-1*¹ *FRT40A/GMR-hid CL FRT40A; hs-rh1-RFP /+*) and *vha68-1*¹*ninaE-gal4;UAS-shi*^{4s1} (ey-flp *rh1-GFP; vha68-1*¹ *FRT40A ninaE-gal4/GMR-hid CL FRT40A;hs-rh1-RFP/UAS-shi*^{4s1}) heads 24 h after heat shock were labeled with antibodies against RFP. GFP fluorescence of Rh1-GFP was directly observed to indicate rhabdomeres. Scale bars are 5 µm. (B) Quantification of rhabdomere-localized, newly synthesized Rh1-RFP at 24 h after heat shock. Six different compound eye sections of each sample were quantified. Error bars indicate SD, ***p < 0.001; ns, not significant (Student's unpaired t-test). (C) Adult eyes from wild type, *vha68-1*¹, *ninaE-gal4;UAS-shi*^{ts1}, and *vha68-1*¹*ninaE-gal4;UAS-shi*^{ts1} flies were dissected and stained with antibodies against Rh1 (green). Rh1 vesicles still accumulated in *vha68-1*¹ flies in the *shi*^{ts1} background. Scale bars are 5 µm.

Supplemental Figure S7. Vha100-1 colocalizes with multiple endosome markers in the synaptic terminal in the lamina.

(A) The nSyb-mp reporter and Vha100-1 colocalize in the synaptic terminal in the lamina.
 Cryostat sections of *nSyb-mp/v100-myc* (*ninaE-nSyb-mCherry-pHluorin/ninaE-vha100-myc*) were labeled with antibodies against
 Myc (green) and RFP (red). Scale bars are 5 μm. (B) Cryostat sections of
 rab5-RFP/v100-myc (*ninaE-rab5-RFP/ninaE-vha100-myc*), *rab7-RFP/v100-myc*

(ninaE-rab7-RFP/ninaE-vha100-myc), rab11-RFP/v100-myc

(ninaE-rab11-RFP/ninaE-vha100-myc), and lamp1-RFP/v100-GFP

(*ninaE-lamp1-RFP/ninaE-vha100-GFP*) heads were labeled with antibodies against Myc (green) or GFP (green) and RFP (red). Scale bars are 5 μm.

Supplemental Figure S8. V-ATPase is required to maintain pH in Rh1 secretory vesicles during its trafficking to the rhabdomere and the deglycosylation process.

(A) Tangential resin-embedded retina sections of compound eyes from ~1-day-old wild type, $dmppe^{e02905}$, $vha68-1^{1}$, and $vha68-1^{1} dmppe^{e02905}$ flies were labeled using antibodies against Rh1. Scale bars are 2 µm. (B) Tangential resin-embedded retina sections of compound eyes from ~1-day-old wild type, $vha100-1^{2}$, ninaE-v100-GFP; $vha100-1^{2}$ (ey-flp rh1-GFP; ninaE-vha100-1-GFP/+; $vha100-1^{2}$ FRT82B/GMR-hid CL FRT82B), and $ninaE-v100^{R755A}-GFP$; $vha100-1^{2}$ (ey-flp rh1-GFP; $ninaE-vha100-1^{2}$ (ey-flp rh1-GFP; $ninaE-vha100-1^{R755A}-GFP/+$; $vha100-1^{2}$ FRT82B/GMR-hid CL FRT82B) flies were labeled using antibodies against Rh1. Scale bars are 2 µm. (C) Head extracts were prepared from ~1-day-old wild type, $vha100-1^{2}$, ninaE-v100-GFP; $vha100-1^{2}$, and $ninaE-v100^{R755A}-GFP$; $vha100-1^{2}$, and $ninaE-v100^{R755A}-GFP$; $vha100-1^{2}$ flies and labeled with antibodies against Rh1.

V-ATPase subunit	mutation	phenotype
vha100	vha100-1 ¹	No ERG transient
vha100	vha100-1 ²	No ERG transient
vhaSFD	vhaSFD ^{EY04644}	No homozygous eye
vha68-2	vha68-2 ^{S4214}	No homozygous eye
vha68-2	vha68-2 ^{EP2364}	No homozygous eye
vha68-2	vha68-2 ^{R6}	No homozygous eye
vha44	vha44 ^{KG00915}	No homozygous eye
vhaAC39-1	vhaAC39-1 ^{FY38}	No homozygous eye
vhaAC39-1	vhaAC39-1 ^{FZ29}	No homozygous eye

Table S1: Phenotypes of mutations in V-ATPase genes. All alleles are homozygous

lethal, and eyes ofhomozygous mutationswere generateusing the"Rh1::GFPey-flp/hid"

method.



Vha68-1 1 Vha68-2 1 MEKKSWVAPL TSFTNSNASN EDRARNSGSL NQDGEPRSVP HPTVPAKKCT 50 Vha68-3 ATP6V1A 1 -MSNLRKFKD -MSNLKRFDD DLRRSTD DFSKLPKILD Vha68-1 9 - - - - - - - - - -- - - - - - - - - -. - - - - - - -. 9 Vha68-2 CPTCACQRND HKSNPVPPPP QSRKPVAEPE SPHDTVNDED SLKD 100 Vha68-3 ATP6V1A - - - - - - - - -- M 11 - - - - - - - - - -- - - - - - - -- - - - - - - -- -E E R E S E Y E E R E S K Y E E E E A T M E D K E S T F G R V <mark>Y</mark> A V S G P V G R V F A V S G P V G R T F G V S G P V G <mark>Y V H</mark>G V S G P V Vha68-1 26 26 Vha68-2 OSHKSAHIAL EKNEDSGFVI EOVVDTHKYS SDE 150 Vha68-3 ATP6V1A 28 E L V G E I E L V G E I O L V G E I F L V G E I V T A E A M S G S A V T A E A M S G S A V N A E E M A G A A V T A C D M A G A A L V R V G Y Y L V R V G Y Y L V R V G H S L V R V G H S | R L E | R L E | R L E | R L E G D M A T I Q V Y E G D M A T I Q V Y E G D M A T I Q V Y E G D M A T I Q V Y E M Y M Y Vha68-1 76 шшш E E T S G V T E T S G V T D T S G V S E T S G V S V G D P V G D P V G D P Vha68-2 76 Vha68-3 ΜY 200 78 ATP6V1A ΜΥF V L R T G K P L S V V L R T G K P L S V V Y Q T G K P L S V V L R T G K P L S V L G P G I MG S I L G P G I MG S I L G P G I MG S I L G P G I MG <mark>A</mark> I I G V M T N S I Y I I N E L T E S I Y I I S E L T N S I Y V I S S Q T Q S I Y I Vha68-1 K G V N T <mark>T</mark> A K G V N V P S 126 S P K G V N V P S L S P K G I D T P S L P P R G V N V S A L S Vha68-2 Ε 126 DGIQRPLRS DGIQRPLSD Vha68-3 250 ATP6V1A 128 N -N -K -K N N T L V K - Q R M I N T L V K - H K M I N S M H D H R L I N S L I K - H K I M Vha68-1 174 GDI G G T T T T G G D L Y G L V H E G G D I Y G S V F E G G D I Y G I V S E KVGSHI 174 ν Vha68-2 299 Vha68-3 IDAL K R ATP6V1A VGSHI 177 H T M L Q H T M L Q H T M L Q F T M V Q T E K L P A N H P L T E K L P A N H P L E D K L P S N S P L T E K L P A N H P L Vha68-1 EDIVLE 224 FD D D V V L E T E F D D E V I V E T E F N S D V V L E L E F E GEITKH DEITKH GVKEKF V W P V W P V R Q P R P V V R <mark>R C</mark> R P V V R Q <mark>V</mark> R P V Vha68-2 224 Vha68-3 349 VWPV 227 ATP6V1A G Q R V L D S L G Q R V L D S L G Q R V L D A F G Q R V L D A L 274 Vha68-1 VQGGT VQGGT VQGGT VQGGT G C G C G C T A T A T A T A GCGK s S S S S ΑF s S D٧ Vha68-2 LT С Ρ T T T Õ А Κ Ν 274 G L G K G K S L P P S N S N S 399 Vha68-3 Т G A G A D٧ V QQ Κ Ç G C А ATP6V1A D \ 277 G Vha68-1 R G N E M S E R G N E M S E R G N E M S E R G N E M S E R G N E M S E PELTCE PELSVE POLEVE PELTME 324 DGVTE DGVTE NGTME A S A S A S V A N T S N V A N T S N V A N T S N V A N T S N A A R E A A R E ĞΕ D F D F SSS ΜK Vha68-2 V R ΑL ΜP ۷ 324 Т R D F R D F VAARE G V AL ΜP 449 Vha68-3 Е ۷ ۷ I M K I M K NGTME DGKVE Т ĒV ATP6V1A 327 Vha68-1 374 Y F Y F Y F TLS TLS TLS DMGYNVSMM DMGYHVSMM DMGYHVSMM DS DS DS T S T S T S R W A E A R W A E A R W A E A R E R E G R L A E G R L A E G R L A E MPADSG MPADAG MPADSG 374 Vha68-2 Α Ρ E R G G S S S A A A Vha68-3 R А Y Ρ 499 Т E RE Î Α 377 ATP6V1A G Vha68-1 L G A R L A <mark>T</mark> F L G A R L A S F L G A R L A S F L G A R L A S F L G A R L A S F Y E Y E Y E Y E R A G R V K C L G N R A G R V K C L G N R A G L V K C L G S R A G R V K C L G N P E R E G S P E R E G S P D R E G S P E R E G S P P G G D F P P G G D F P P G G D F P P G G D F 424 D P V T S A D P V T S A D P V T S A D P V T S A D P V T S A G G s S s S S S V V V G A V S G A V S G A V S G A V S s s s Vha68-2 V V V 424 Т İ L S L G 549 Vha68-3 T T ATP6V1A 427 474 Vha68-1 Q V F WG L D K PS PS PS PS A Q R K H F A Q R K H F MRALDEYYDK MRALD<mark>DF</mark>YDK K Y K Y K Y K Y 15 s Y Y Y ΚL NWL Vha68-2 474 A Q R K H F A Q R K H F s MRTLDTYYEE MRALDEYYDK S N P E F T H L R A H F T E F V P L R T V N W L OS V N W L IS Vha68-3 599 L 477 ATP6V1A V K E I L Q E E E V K E I L Q E E E A K <mark>K V</mark> L Q E E D A K E I L Q E E E | V Q L V G | V Q L V G | V Q L V G | V Q L V G K A S L A E T D K V K A S L A E T D K I K S S L N E E D K I K A S L A E T D K I T L E V A K L L K D T L E V A K L L K D T L E V A K M L K D T L E V A K M L K D
 x
 x
 x
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
 y
D L S E D L S E D L A E D L A E Vha68-1 524 K V K A K A Vha68-2 524 Vha68-3 649 527 ATP6V1A N T I R E S M G G I N V I R E A M G N I G L I R I K A A N I S I I R E H M G D I Vha68-1 574 D R V C D R F C т VES V G M L R N I I D F T G M L K N M M T F V G M L S N M I A F Ρ F ΥK Y D M A R H S V E S Y D A A I L S V O N Y D M A R R A V E T TAQSENKITW 574 Vha68-2 T T T F F K F Y K P T A <mark>D N E A R V</mark> T W T A Q S D N K I T W Vha68-3 699 DAYC YDRFC ΡF ATP6V1A 577 Vha68-1 V K D G E Q K I V K D G E A K I - K L G E A T V L K D G E A K I K A D Y D Q L Y E D K A D F E Q L H E D L E S M N K L H D N L Q Q A F R N L E D L Q Q A F R N L E D I L S T F R E I E D MQ N A F R S L E D 614 Q L S S M K F K Q L S S M K F K E L S T M K F I K L S S M K F K DPV DP-DPL ΜY Vha68-2 614 NVEDY 743 Vha68-3 KSDYAQLLED ATP6V1A 617

Α wild type vha68-11 vha100-12 5 mv 5s <u>10</u>-3 10-2 10-5 10-4 **10**⁻¹ ſ С В 1.5 -*** Normalized Rh1 level *** wild type vha68-11 vha100-12 INAD 70 - 35 Rh1 0.0 wild type vha68-11 vha100-12 Α

В



Rh1-GFP Na*K*ATPase Merge



Α

24h after heat-shock





С

В





Α

В

С



2µm

2µm

