

Received: 3 October 2017 Accepted: 24 August 2018

Published online: 10 September 2018

OPEN A soy protein Lunasin can ameliorate amyloid-beta 42 mediated neurodegeneration in Drosophila eye

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Alzheimer's disease (AD), a fatal progressive neurodegenerative disorder, also results from accumulation of amyloid-beta 42 (A\Beta42) plaques. These A\Beta42 plaques trigger oxidative stress, abnormal signaling, which results in neuronal death by unknown mechanism(s). We misexpress high levels of human Aβ42 in the differentiating retinal neurons of the Drosophila eye, which results in the Alzheimer's like neuropathology. Using our transgenic model, we tested a soy-derived protein Lunasin (Lun) for a possible role in rescuing neurodegeneration in retinal neurons. Lunasin is known to have anti-cancer effect and reduces stress and inflammation. We show that misexpression of Lunasin by transgenic approach can rescue A\(\beta 42\) mediated neurodegeneration by blocking cell death in retinal neurons, and results in restoration of axonal targeting from retina to brain. Misexpression of Lunasin downregulates the highly conserved cJun-N-terminal Kinase (JNK) signaling pathway. Activation of JNK signaling can prevent neuroprotective role of Lunasin in A β 42 mediated neurodegeneration. This neuroprotective function of Lunasin is not dependent on retinal determination gene cascade in the *Drosophila* eye, and is independent of Wingless (Wg) and Decapentaplegic (Dpp) signaling pathways. Furthermore, Lunasin can significantly reduce mortality rate caused by misexpression of human A β 42 in flies. Our studies identified the novel neuroprotective role of Lunasin peptide, a potential therapeutic agent that can ameliorate A\beta 42 mediated neurodegeneration by downregulating JNK signaling.

Alzheimer's Disease (AD; OMIM: 104300), an irreversible, progressive neurodegenerative disorder, which results in the loss of neurons in the hippocampus and cortex. AD manifests as the loss of memory, cognition functions and eventually results in the death of patient¹⁻⁴. The two major causes for AD are accumulation of amyloid plaques and generation of neurofibrillary tangles (NFTs) due to hyper-phosphorylation of microtubule binding protein Tau^{1,4-7}. The amyloid plaques are generated from improper cleavage of amyloid precursor protein (APP), a transmembrane protein, by β - and then γ -secretase enzymes to form a 42 amino-acid long fragment, which is referred to as amyloid-beta 42 $(A\beta42)^8$. These $A\beta42$ fibrils self-assemble into extracellular $A\beta42$ plaques^{2,9}. Generally, APP cleavage results in a forty amino acid long polypeptide (A β 40). The two extra amino acids in A β 42 polypeptide makes it hydrophobic, which results in accumulation of the amyloid plaques^{4–7,10,11}. These A\(\beta\)42 plaques cause membrane defects, disruption of neural networks, trigger aberrant signaling and disrupt normal cellular processes resulting in neurodegeneration. Thus, the current consensus is that A β 42 conversion and self-assembly into oligomeric forms and plaques is responsible for neuronal death in AD by unknown molecular-genetic mechanism(s)^{2-7,10,12}. Various animal models of AD have been developed to understand the molecular-genetic basis of AD13 as the genetic machinery is highly conserved across organisms including mouse, C. elegans and fruit fly^{4,13-23}.

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Drosophila melanogaster, fruit fly, is a highly versatile organism to model human disease ^{13,14,16,22,23}. The Drosophila eye is used extensively to model human neurodegenerative disorders ^{6,13,15,16,20,24,25} because the important signaling pathways required for development and differentiation of the fly visual system are highly conserved ²⁶. Thus, the information generated from the fly model can help understand molecular-genetic underpinning of the human disease ^{13,14,17,19,27}. Drosophila, a holometabolous insect, has a blue print for its adult organs housed inside the larva referred to as the imaginal discs ^{28,29}. The larval eye-antennal imaginal disc gives rise to the adult compound eye, antenna and head upon differentiation ^{13,26,30-32}. In the developing Drosophila eye, the cell fate specification and differentiation is regulated by a group of genes like twin of eyeless (toy), eyeless (ey), eyegone (eyg), twin of eyegone (toe), Optix (opt); eyes absent (eya), sine oculis (so), dachshund (dac) and optix (opt) ³³⁻³⁹, which are called retinal determination (RD) genes ⁴⁰⁻⁴⁵.

The retinal precursor cells in the eye imaginal disc undergo differentiation to form the photoreceptor neurons in the adult eye^{26,46–48}. Eight photoreceptor neurons (PR1-8) and several support cells form a unit eye called as the ommatidium. The axons from the photoreceptors (retinal neurons) fasciculate together to form an axonal bundle, which traverses through the optic stalk and then innervate the different layers of the *Drosophila* brain^{49,50}. The axons from photoreceptors (PRs) 1–6 terminate in the lamina whereas PR7-PR8 end in a separate layer of medulla after passing through lamina. In the pupal retina, the excessive cells other than the differentiated cells are eliminated by programmed cell death (PCD)⁵¹. However, abnormal extracellular signaling due to inappropriate levels of morphogens may trigger cell death in the larval eye imaginal disc^{52,53}.

Previous work from our lab showed that evolutionarily conserved Wingless (Wg) and Jun-N terminal kinase (JNK) signaling pathway are tightly regulated to allow differentiation to occur and to prevent premature cell death in the developing fields 4. Wg, a member of highly conserved Wnt/Wg, is responsible for regulating early growth, restricting eye fate and later Wg plays a role in triggering programmed cell death (PCD) 55,56 in the pupal retina. Wg also plays a role in developmental cell death during larval eye development 53. Another highly conserved TGF beta (TGF β) signaling pathway, referred to as Decapentaplegic (Dpp) signaling in *Drosophila* 57-59, collaborates with Hedgehog (Hh) signaling to promote retinal differentiation in the developing eye as well as antagonize Wg signaling 60.

Activation of JNK signaling or stress activated kinase proteins of the mitogen-activated protein kinase (MAPK) superfamily trigger cell death $^{27,61-63}$. JNK signaling is activated through a cascade of phosphorylation by MAP Kinases to regulate cell homeostasis $^{61,64-66}$. JNK signaling acts downstream of the Tumor Necrosis Factor (TNF) homolog Eiger (Egr) and its receptor Wengen (Wgn) by Tak1 (TGF- β - activating kinase 1), a JNK kinase kinase (JNKKK), Hemipterous (Hep); a JNK Kinase, Basket (Bsk; Jun kinase) and Jun $^{63-65,67}$. It is known that activation of JNK signaling leads to induction of cell death to eliminate developmentally aberrant cells 63,67 . The functional read out for the activation of JNK signaling is *puckered* (*puc*), which encodes a dual phosphatase, and acts via a negative feedback loop to downregulate the JNK activity 27,63,66 .

We developed a transgenic model system in *Drosophila* eye where we misexpress high levels of human amyloid-beta (A β 42) in the differentiating retinal neurons of the developing fly retina²⁷ using a Glass Multiple Repeat (GMR) Gal4 driver⁶⁸. The transgenic flies with GMR-Gal4 driven UAS-Aβ42 have been abbreviated as GMR > $A\beta 42^{27}$. Targeted misexpression of human $A\beta 42$ (GMR > $A\beta 42$) in the differentiating photoreceptors (retinal neurons) of the developing Drosophila eye69, exhibit progressive neurodegenerative phenotypes that mimic the neuropathology of AD patients²⁷. The frequency of this GMR > A β 42 phenotype is 100%, which makes this Drosophila eye model a highly reliable tool for identifying the genetic modifiers of the GMR > A\(\beta 42\) mediated neurodegeneration 27,70,71 . We used our AD model to test plant-based protein Lunasin for its role in blocking A β 42 mediated neurodegeneration. Lunasin, a soy (glycine max) derived peptide, has multiple roles⁷². Lunasin protein has four functional domains⁷²⁻⁷⁴. It has an N terminal region of unknown function, followed by a chromatin binding helical region, a carboxy terminal RGD cell adhesion motif, and an eight aspartic acid (poly-D) tail. The poly-D tail and RGD motif have been shown to be essential for the bioactivity of Lunasin. Lunasin is known to have anti-cancer effects^{74,75} and reduces stress and inflammation. The odds of manifestation of AD increases with chronic low-grade inflammation like stress, depression, and obesity⁷⁶. Here we present identification of a plant protein Lunasin that can rescue Aβ42 mediated neurodegeneration. This neuroprotective function of Lunasin is achieved by downregulating JNK signaling dependent cell death in the developing retinal neurons of the Drosophila eye. Furthermore, gain-of-function of Lunasin can also reduce the mortality rate of the flies expressing A β 42 in the nervous system.

Materials and Methods

Fly Stocks. All fly stocks used in this study are listed and described in Flybase (http://flybase.bio.indiana.edu). The fly stocks used in this study were Canton-S (Wild-type), GMR-Gal4⁶⁸, elav-Gal4 (BL#485)⁷⁷, UAS-Aβ42²⁷, UAS-puc, puc^{E69}, where lacZ reporter express under the control of puc regulatory element, and acts as the functional read out of JNK signaling pathway⁶⁶. Other stocks used were UAS-Djun^{aspv778}, UAS-hep^{Act}, wg-lacZ⁷⁹, dpp-lacZ⁸⁰, where lacZ reporter⁸¹ express under the control of wg and dpp regulatory element. The UAS- Aβ42 transgenic flies were generated by microinjecting a UAS-construct where two tandem copies of human amyloid -β1-42 (Aβ42) fused to signal peptide for secretion were cloned^{10,25,27}. The rationale of bi-cistronic construct was to mimic APP duplications associated with early onset of familial AD and to express high levels of Aβ42 to induce strong eye phenotype^{25,82}.

Generation of EGFP-Lunasin Transgenic flies. We employed gene tagging approach⁸³ to generate EGFP-Lunasin transgenic flies. The sequence for EGFP (Enhanced Green Fluorescent Protein) was fused to the 5' end of Lunasin sequence⁸⁴. It has been shown that Lunasin tagged with EGFP show no observable differences in bioactivity⁷³. The sequence of EGFP-Lunasin with start, stop codons and the restriction sites was synthesized *in vitro*, sequence verified and cloned into pUAST vector. The GFP reporter can provide spatio-temporal localization

of Lunasin transgene. The clones were sequence verified and microinjected in *Drosophila* embryos and the transgenic flies were generated. These flies were balanced and used for genetic crosses.

Genetic Crosses. We employed a Gal4/UAS system for targeted misexpression studies⁶⁹. All Gal4/UAS crosses were maintained at 18 °C, 25 °C and 29 °C, unless specified, to sample different induction levels. The adult flies were maintained at 25 °C, while the cultures after egg laying (progeny) were transferred to 29 °C for further growth. Misexpression of A β 42 in the differentiating retina (GMR-Gal4 > UAS-A β 42) exhibits a stronger neurodegenerative phenotype at 29 °C with no penetrance^{27,68}. All the targeted misexpression experiments were conducted using the Glass Multiple Repeat driver line (GMR-Gal4)⁶⁸ or embryonic lethal abnormal visual system (elav-GAL4) line⁷⁷. GMR- Gal4 directs expression of transgenes in the differentiating retinal precursor cells of the developing eye imaginal disc and pupal retina⁶⁸. The elav-Gal4 drives expression in the neurons⁷⁷.

Immunohistochemistry. Eye-antennal discs from wandering third instar larvae were dissected, and fixed in 4% paraformaldehyde in Phosphate Buffered Saline (PBS), and stained following the protocol^{85–87}. The primary antibodies used were rabbit anti-Dlg (1:200; a gift from K. Cho), mouse anti-Wg [1:50,Developmental Studies Hybridoma Bank,(DSHB)], rat anti-Elav (1:50; DSHB), mouse anti-Dlg (1:100; DSHB), mouse anti-22C10 (1:100; DSHB), mouse anti-Chaoptin (MAb24B10) (1:100; DSHB⁸⁸), mouse anti-Ey (1:100, DSHB), mouse anti-Eya (1:100, DSHB), mouse anti-Dac (1:100, DSHB), mouse anti-β-galactosidase (1:100; DSHB), rabbit anti-β-galactosidase (1:200) (Cappel), and mouse anti-GFP (1:100, GFP-G1, DSHB). Secondary anti-bodies (Jackson Laboratories) used consisted of donkey anti-rabbit IgG conjugated with FITC (1:200), donkey anti-mouse IgG conjugated with Cy3 (1:250), and goat anti-rat IgG conjugated with Cy5 (1:250). The tissues were mounted in Vectashield (Vector labs) and all immunofluorescence images were captured using the Laser Scanning Confocal Microscopy⁸⁹ (Olympus Fluoview 1000). All images were taken at 20X magnification unless stated otherwise. The final images and figures were prepared using Adobe Photoshop CS6 software.

Detection of Cell Death. Cell death was detected using TUNEL assays^{27,53,54,90,91}. TUNEL assays were used to identify the cells undergoing cell death where the cleavage of double and single stranded DNA is labeled by a Fluorescent tag (TMR Red). The fluorescently labeled nucleotides are added to 3′ OH ends in a template-independent manner by Terminal Deoxynucleotidyl Transferase (TdT). The fluorescent label tagged fragmented DNA within a dying cell can be detected by fluorescence or confocal microscopy⁸⁹. Eye-antennal discs after secondary antibody staining were blocked in 10% normal donkey serum in phosphate buffered saline with 0.2% Triton X-100 (PBT) and labeled for TUNEL assays using a cell death detection kit from Roche Diagnostics (In Situ Cell Death Detection Kit, TMR red,12156792210).

The TUNEL positive nuclei were counted to determine the dying cell population from five sets of imaginal discs and were used for statistical analysis using Microsoft Excel 2013. The P-values were calculated using two-tailed t-test and the error bars represent Standard Deviation from Mean 27,70,71,92 .

Adult Eye Imaging. Adult flies were prepared for imaging by freezing at $-20\,^{\circ}$ C for approximately 2 hours followed by mounting the fly on a dissection needle 54,86 . The needle with the fly was aligned horizontally over a glass slide using molding putty. Images were captured on an MrC5 color camera mounted on an Axioimager.Z1 Zeiss Apotome using Z-sectioning approach. Final images were generated by compiling the individual stacks from the Z-sectioning approach using the extended depth of focus function of Axiovision software version 4.6.3.

Western Blot. Protein sample were prepared from third instar eye imaginal disc from Wild type, GMR > A β 42, GMR > A β 42 + Lun larvae following the standardized protocol^{27,93}. The Phospho SAPK/JNK (Cell Signaling Thr183/Tyr185) (81E11) Rabbit antibody was used at 1:1000 dilution. Signal was detected using Horse Radish Peroxidase (HRP) conjugated goat anti-rabbit IgG using supersignal chemiluminescence substrate (Pierce). Images were captured using the BioSpectrum® 500 Imaging System.

Eclosure Assay. Eclosure assays are desirable assays to screen the effect of genetic backgrounds on eclosion of flies⁷⁷. We collected eggs on a grape plate from elav-Gal4 (control), elav-Gal4 drive UAS- $A\beta42$ (elav-Gal4 $> A\beta42$) and elav-Gal4 drive UAS- $A\beta42$ + UAS-Lunasin (elav-Gal4 > Ab42 + Lun) flies. The eclosion assay was carried out in four sets of 50 larvae each. We seeded first instar larvae (50 in each set) from a synchronous culture in each vial. We counted 200 larvae (4 sets of 50 larvae) for each cross. The larvae were allowed to develop and hatched/eclosed adults were counted. All unhatched pupae were also counted. All four sets of fifty larvae each for every genotype were seeded from the same grape plate to maintain consistency and accuracy in larval staging.

Results

Lunasin can rescue A β 42 mediated neurodegeneration. In comparison to the wild-type eye imaginal discs (Fig. 1A) that develop into adult compound eyes (Fig. 1F), targeted misexpression of A β 42 in the developing *Drosophila* eye using the GMR-Gal4 enhancer results in a strong neurodegenerative phenotype in the eye imaginal disc (Fig. 1D), and the adult eye (Fig. 1I)²⁷. The neurodegenerative phenotype in the GMR > A β 42 eye imaginal disc (Fig. 1D) worsens in the adult eye (Fig. 1I). The adult eyes are highly reduced with glazed appearance due to fusion of individual unit eyes and also exhibits some dark necrotic spots to mark neurodegeneration (Fig. 1I)^{27,70,71,92}. The frequency of GMR > A β 42 phenotype in the eye imaginal discs (Fig. 1D, n = 72) as well as adult eyes (Fig. 1I, n = 151) is 100%. The controls GMR-Gal4 alone (n = 75, 100%, Fig. 1B,G) and the transgene stock UAS-A β 42 alone (n = 75, 100%, Fig. 1C,H) exhibits near wild-type eye phenotype in the eye imaginal discs and the adult eyes. The gene tagging with standardized immune-epitopes or fluorescent tags that permit live imaging and do away with the requirement of generating antibodies against a protein is commonly used approach⁸³. We employed gene-tagging approach to generate UAS- based transgenic flies where the Soy plant

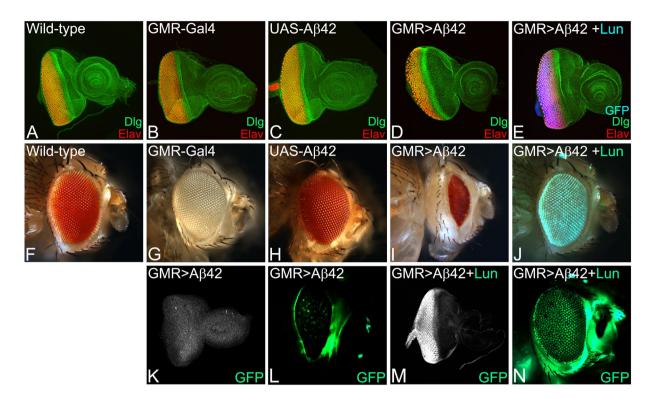


Figure 1. Misexpression of the soy protein Lunasin can rescue A β 42 mediated neurodegeneration. (A,F) Wild-type (A) larval eye imaginal discs, which develop into (F) adult compound eyes comprising of nearly 800 unit eyes. Note that the eye imaginal disc is stained with membrane specific marker Disc large (Dlg: Green) and pan neural marker Embryonic Lethal Abnormal Vision (Elav: Red), which marks nuclei of the retinal neuron. In comparison to the wild-type eyes, (D,I) misexpression of A β 42 (GMR > A β 42) in the differentiating retinal neurons using GMR-Gal4 driver results in the induction of neuronal death as seen in (D) eye imaginal discs and the (I) highly reduced adult eyes. Note that phenotype worsens from (D) larval eye imaginal disc to the (I) the adult eye. The controls used are (B,G) GMR-Gal4 and (C,H) transgene stock UAS-Aβ42. (E,J) Misexpression of soy polypeptide Lunasin (Lun) along with A β 42 (GMR > A β 42 + Lun) results in significant rescue of Aβ42 mediated neurodegeneration as seen in (E) the eye discs and (J) the adult eyes. Since Lunasin is a plant protein, in order to determine if Lunasin (Lun) is actually expressed in our model system it has been tagged with EGFP (as seen in E,J). Note that GFP reporter is detected in GMR domain of (E) eye imaginal discs and (J) adult eyes. (K,M) Expression of Lunasin detected by GFP antibody staining in (K) control GMR > Aβ42 and (M) GMR > $A\beta 42 + Lun$. Note that GFP antibody positively marks the GMR domain only in the (M) $GMR > A\beta 42 + Lun$ eye disc. In adult eyes, GFP expression is detected by GFP reporter expression in control (L) GMR > $A\beta42$ and (N) GMR > $A\beta42$ + Lun. The orientation of all imaginal discs in the figure is posterior to left and dorsal up. Magnification of all eye discs is 20X.

based peptide Lunasin (Lun) is tagged with EGFP (UAS-Lun-EGFP). Misexpression of UAS-Lun-EGFP along with A β 42 (GMR > A β 42 + Lun-EGFP) exhibits a strong rescue (n = 100, 70%) resulting in a near complete wild-type eye (Fig. 1E,J). There were no necrotic spots observed in the adult eyes (Fig. 1J). We confirmed that rescue of GMR > A β 42 neurodegenerative phenotype is due to misexpression of Lunasin based on EGFP transgene expression in the GMR domain of the eye imaginal discs (Fig. 1M) and the adult eyes (Fig. 1N). A strong robust GFP expression was detected by GFP antibody staining in the GMR domain of the GMR > A β 42 + Lun-EGFP eye imaginal discs (Fig. 1M) and GFP reporter expression was seen in the adult eyes (Fig. 1N). The GFP protein was not detected in GMR > A β 42 eye imaginal discs (Fig. 1K), and, no GFP reporter expression was seen in the adult eyes (Fig. 1L). We also verified the data using a UAS-Lun construct which is not tagged with EGFP (data not shown). These data suggest that misexpression of Lunasin can rescue GMR > A β 42 mediated neurodegeneration likely by blocking cell death.

Lunasin can rescue A β 42 mediated cell death in *Drosophila* eye. To test this, we employed the TUNEL staining, which marks the dying cell nuclei by labelling the 5' end of the double and single stranded DNA with a fluorochrome. The fluorochrome tagged TUNEL positive nuclei can be counted to quantify cell death^{27,90,91,94}. We performed TUNEL staining in the third instar eye-imaginal disc in the wild-type (Fig. 2A,A'), GMR > A β 42 (Fig. 2B,B') and GMR > A β 42 + Lun (Fig. 2C,C') backgrounds. The TUNEL staining was performed before the onset of developmentally controlled programmed cell death. The TUNEL positive cells were counted from five sets of imaginal discs from all three backgrounds, and the P values were calculated^{27,70,71,92}. A quantification (n = 5, p < 0.05) of dying cells from these genotypes further confirms that in comparison to the wild-type eye disc, a 3–4 fold increase in cell death was observed in GMR > A β 42 (Fig. 2B,B',D) background. In

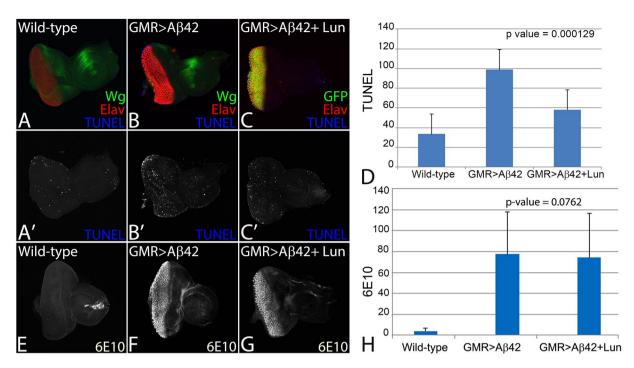


Figure 2. Lunasin can block cell death to rescue A β 42 mediated neurodegeneration. The dying cells nuclei can be marked by TUNEL staining. TUNEL staining was carried out in (A,A') Wild-type, (B,B') GMR > A β 42, (C,C') GMR > A β 42 + Lun eye imaginal discs. The number of dying retinal neurons were counted in these backgrounds (n = 5). Note that the number of dying cells increase nearly 3–4 fold in (B,D) GMR > A β 42 as compared to the (A,D) Wild-type eye discs. (C,D) Misexpression of Lunasin (Lun) along with GMR > A β 42 (GMR > A β 42 + Lun-GFP) results in significant reduction in the dying retinal neurons. The number of TUNEL positive nuclei were counted from five eye imaginal discs for all three backgrounds. (D) A graph comparing the number of dying nuclei of neurons validate that Lun misexpression along with GMR > A β 42 (GMR > A β 42 + Lun-GFP) rescues the GMR > A β 42 neurodegeneration. These numbers are significant based on the calculations of P-values using the two-tailed t- test using Microsoft Excel 2013. (E–H) Accumulation of amyloid plaque was detected using monoclonal antibody 6E10 in (E) Wild-type, (F) GMR > A β 42 and (G) GMR > A β 42 + Lun eye imaginal discs. (H) The signal intensity of 6E10 staining was calculated from five (n = 5) eye discs of each background and plotted on a graph. Note that 6E10 levels are not significantly different between (F) GMR > A β 42 and (G) GMR > A β 42 + Lun background. The levels of amyloid plaques are barely detected in Wild-type background. Magnification of all eye disc is 20X.

comparison to the GMR > A β 42 (Fig. 2B,B',D) background, the number of TUNEL positive dying nuclei were significantly reduced in GMR > A β 42 + Lun (Fig. 2C,C',D). It suggests that expression of Lunasin is capable of inhibiting cell death as seen by the strong rescue phenotype in the adult eyes.

We wanted to test if Lunasin can rescue GMR > $A\beta42$ phenotype by preventing accumulation of amyloid plaques or act downstream of amyloid plaque formation in the GMR > $A\beta42$ + Lun background. Using 6E10 antibody to mark $A\beta42$ plaques, we found that there is a little or no $A\beta42$ present in the wild-type (Fig. 2E,H, n=50, 100%) eye discs. However, there is strong deposition of $A\beta42$ plaques in GMR > $A\beta42$ (Fig. 2F,H, n=52, 100%) and GMR > $A\beta42$ + Lun (Fig. 2G,H, n=46, 100%) eye discs. Furthermore, there is no significant difference between $A\beta42$ plaque formation in GMR > $A\beta42$ and GMR > $A\beta42$ + Lun (Fig. 2H) background. It suggests that Lunasin misexpression does not affect $A\beta42$ plaques formation and acts downstream to GMR > $A\beta42$ plaque formation.

Lunasin can restore axonal targeting defects seen in A β 42 background. The neurodegenerative phenotype in AD encompasses disruption of axonal transport mechanism, which results in impaired axonal targeting (incorrect axonal guidance)^{27,70,71,92}. In order to understand, if these GMR > A β 42 + Lun imaginal discs where neurodegeneration phenotype is rescued have proper connection between retinal neurons and brain, we employed 24B10 (Chaoptin, which marks photoreceptor neurons and their axons)⁸⁸. In the wild-type eye disc, R1-R6 axons of each ommatidium project to the lamina whereas R7 and R8 axons project to the medulla, a separate layer of the optic lobe⁹⁵ (Fig. 3A) in (n = 51)100% eye imaginal discs. In comparison to the wild-type eye imaginal discs, the GMR > A β 42 eye discs show a severe disorganization in axonal targeting (Fig. 3B) in (n = 47)100% larval eye imaginal discs. However, misexpression of Lunasin in GMR > A β 42 (GMR > A β 42 + Lun) background significantly restore the axonal targeting (Fig. 3C) in (n = 50) 70% of larval eye imaginal discs. We also investigated the neurons and their axonal processes using 22C10 marker⁹⁶. In comparison to the wild-type expression of 22C10, that marks the retinal neurons and their processes (Fig. 3D, n = 50,100% eye discs), GMR > A β 42 background show strong neurodegenerative phenotype (reduced neurons

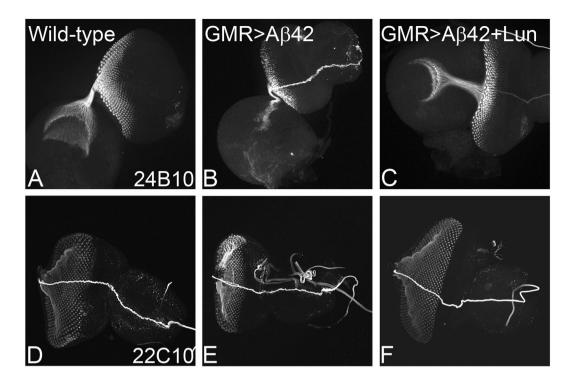


Figure 3. Lunasin (Lun) misexpression can restore A β 42 mediated impairment of axonal targeting from retina to brain. Chaoptin (MAb24B10), a marker for the axonal targeting from retina to the optic lobes of the brain. (A) In wild-type eye imaginal discs, the retinal axons marked by MAb24B10 innervate the lamina and medulla of the brain. (B) Misexpression of GMR > A β 42 in the developing eye imaginal discs result in impaired targeting of retinal axons to the brain. However, (C) misexpression of Lunasin (Lun) along with A β 42 (GMR > A β 42 + Lun) resulted in significant restoration of the axonal targeting to near wild-type. Monoclonal antibody 22C10 marks all axonal sheath of the photoreceptors in the developing eye. 22C10 expression in (D) Wild-type, (E) GMR > A β 42 and (F) GMR > A β 42 + Lun background. Note that impairment of 22C10 expression in GMR > A β 42 background is restored significantly in GMR > A β 42 + Lun background. Magnification of all eye discs is 20X.

and abnormality in their processes) (Fig. 3E, n = 50, 100% eye discs). However, misexpression of Lunasin in GMR > A β 42 (GMR > A β 42 + Lun) background significantly restore the neurodegenerative phenotype (Fig. 3F, n = 50, 80%) as compared to the GMR > A β 42 (Fig. 3E, n = 50,100%) eye discs.

Neuroprotective function of Lunasin is independent of retinal differentiation genes. Since our experimental system is *Drosophila* eye, we need to rule out the possibility of Lunasin affecting the retinal differentiation gene machinery rather than exhibiting the neuroprotective function. We investigated the role of RD genes in neuroprotective function of Lunasin. The Pax-6 homolog *eyeless* (*ey*), eye fate selector gene, is expressed in the early eye and later its expression is restricted anterior to the MF in the eye imaginal disc (Fig. 4A, n = 50, 100%) 30,44,85,97,98 . A tyrosine phosphatase, *eyes absent* (*eya*), is expressed both in the differentiated retinal neurons as well as the retinal precursor cells anterior to MF^{30,31,33,70} (Fig. 4B, n = 50, 100%). Another RD gene *dachshund* (*dac*) is expressed in two different domains one anterior to the MF and another posterior to the MF^{70,99} (Fig. 4C, n = 50, 100%). In GMR > Aβ42 + Lun background the expression of all three RD genes Ey (n = 50, 100%), Eya (n = 50, 100%) and Dac (n = 50, 100%) was not affected (Fig. 4D–F, n = 50, 100%). Our data strongly suggests that the neuroprotective role of Lunasin is independent of RD gene function. In our transgenic model, the Aβ42 expression is triggered at the time of retinal differentiation using a GMR-Gal4 driver^{27,68}, which drives expression of UAS- transgene much later than the event of eye specification, and the onset of retinal determination and differentiation genes expression. Thus, neuroprotective function of Lunasin with respect to Aβ42 mediated neurodegeneration is independent of eye specification function or retinal differentiation machinery.

Lunasin rescues A β 42 mediated neurodegeneration independent of Wingless (Wg) or Decapentaplegic (Dpp) signaling. Previous studies from our lab and others have shown that accumulation of A β 42 plaques (GMR > A β 42) triggers aberrant signaling which results in neurodegeneration^{4,27,70,71,92}. We therefore tested (i) several signaling pathways to discern the mechanism of Lunasin mediated neuroprotective function, and (ii) known genetic modifiers of A β 42 mediated neurodegeneration^{27,70,71,92} (data not shown). The evolutionarily conserved Wg signaling act antagonistically to Dpp during eye development⁶⁰. Wg acts as the negative regulator of eye fate. Gain-of-function of wg can suppress the eye fate and loss of function of wg induces eye enlargements^{44,58,59,85}. In the developing *Drosophila* eye, Wg functions antagonistically to Dpp that promote cell survival. Dpp is also involved in retinal differentiation⁶⁰. During eye development, Wg is expressed

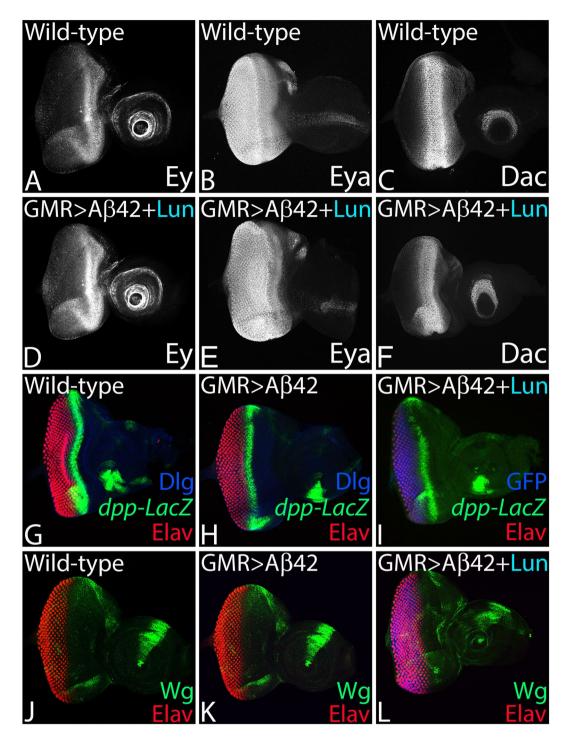


Figure 4. Lunasin neuroprotective function is independent of retinal differentiation gene machinery, Dpp and Wg signaling pathways. (**A**–**C**) Wild-type expression of (**A**) Eyeless (Ey), (**B**) Eyes absent (Eya) and (**C**) Dachshund (Dac), the members of RD gene machinery in developing third instar eye imaginal discs. (**A**) Ey expression is anterior to MF in the third instar eye discs. (**B**) Eya is expressed in the differentiating photoreceptors and anterior to MF. (**C**) Dac is expressed along the MF as well as in antennal region. (**D**) Ey, (**E**) Eya and (**F**) Dac expression is not affected in the GMR > A β 42 + Lun background. (**G**–**I**) In the developing eye imaginal discs, study of dpp expression using dpp-lacZ reporter in the (**G**) Wild-type, (**H**) GMR > A β 42, (**I**) GMR > A β 42 + Lun. Note that dpp-lacZ marks the morphogenetic furrow (MF) in the developing eye. (**J**–**L**) In the developing eye imaginal discs, study of Wg expression using a wg-lacZ reporter in the (**J**) Wild-type, (**K**) GMR > A β 42, (**L**) GMR > A β 42 + Lun. The wg is expressed on the antero-lateral margin of the developing third instar eye imaginal discs. Note that the Lun neuroprotection function is independent of Wg and Dpp signaling. Magnification of all eye discs is 20X.

on the antero-dorso-ventral eye margin (Fig. 4J, n = 50, 100%) where as Dpp is expressed dynamically along with the MF (Fig. 4G, n = 50, 100%). We found that Lunasin misexpression in GMR > A β 42 (GMR > A β 42 + Lun) background does not affect Wg (Fig. 4L, n = 50, 100%) and/or Dpp (Fig. 4I, n = 50, 100%) expression in the eye imaginal discs. Thus, neuroprotective function of Lunasin with respect to A β 42 mediated neurodegeneration is independent of Wg and Dpp Signaling pathways.

Lunasin downregulates JNK signaling to prevent neurodegeneration. We have shown earlier that JNK signaling is involved in A β 42 mediated neurodegeneration²⁷. Activation of JNK signaling triggers a cascade of kinases, which in turn regulates the expression of $puc^{61,63,66,100}$. Puc is a dual phosphatase that negatively regulates JNK signaling by a feedback loop^{27,63} (Fig. 5A). The phospho-Jun kinase, encodes an enzyme which can phosphorylate N-terminal its substrate Jun and can be used to study the activation status of JNK signaling. We tested levels of JNK activation by quantifying levels of phospho-JNK in western blots²⁷. We quantified and compared the amount of phospho-Jun kinase (p-JNK) in wild-type versus GMR > A β 42 and GMR > A β 42 + Lun background. In comparison to the wild-type, p-JNK levels are upregulated in GMR > A β 42 background as seen earlier²⁷. However, in comparison to GMR > A β 42, pJNK levels were significantly reduced in GMR > A β 42 + Lun background (Fig. 5B,B').

We tested if JNK signaling is downregulated by Lunasin (Lun) misexpression. To activate JNK signaling, we misexpressed $Djun^{aspv7}$ and constitutively active hemipterous (hep^{Act}) using the GMR-Gal4 driver and assay its phenotype in the eye imaginal disc and the adult eye (Fig. 5E,F,K,L). In comparison to the wild-type eye (Fig. 5C,D), misexpression of $Djun^{aspv7}$ alone ($GMR > Djun^{aspv7}$) results in the reduced eye phenotype (Fig. 5E,F, n = 50, 100%). The strong neurodegenerative phenotype of highly reduced eye in $GMR > A\beta42 + Djun^{aspv7}$ (Fig. 5G,H, n = 50, 100%) was not rescued in $GMR > A\beta42 + Lun + Djun^{aspv7}$ (Fig. 5I,J, n = 50, 100%) background. Similarly the strong neurodegenerative phenotype due to activation of JNK pathway by using hep^{Act} , as seen in $GMR > hep^{Act}$ (Fig. 5K,L, n = 75, 100%), $GMR > A\beta42 + hep^{Act}$ (Fig. 5M,N, n = 75, 100%) was not rescued by misexpression of Lunasin ($GMR > A\beta42 + Lun + hep^{Act}$, Fig. 5O,P, n = 50, 100%). However, blocking or downregulating JNK signaling by misexpression of puc in GMR > puc exhibits near wild-type eye (Fig. 5Q,R, n = 50, 100%). Misexpression of puc in $GMR > A\beta42 + puc$, exhibits significant rescue of $GMR > A\beta42$ neurodegenerative phenotype (Fig. 5S,T, n = 50, 45%). Furthermore, puc expression in $GMR > A\beta42 + Lun$ background ($GMR > A\beta42 + Lun + puc$) can significantly rescue $A\beta42$ mediated neurodegeneration as seen in the eye imaginal disc and the adult eye (Fig. 5U,V, n = 50, 40%). Thus, Lunasin, which acts upstream of JNK signaling, may downregulate JNK signaling in rescuing $A\beta42$ mediated neurodegeneration in the Drosophila eye.

To test this hypothesis, we analyzed the expression of puc (Fig. 5W,W'), a functional read out of JNK signaling pathway by using a puc-lacZ reporter^{66,101}. There is a robust induction of puc-lacZ in GMR > A β 42 (Fig. 5X,X', n = 30, 100%) as compared to the wild-type puc expression in eye imaginal discs (Fig. 5W,W', n = 30, 100%). However, in GMR > A β 42 + Lun, puc levels are significantly downregulated (Fig. 5Y,Y', n = 30, 70%) as compared to GMR > A β 42 background (Fig. 5X,X'). Our data clearly validate our hypothesis that Lunasin downregulates JNK signaling to rescue A β 42 mediated neurodegeneration.

Lunasin increase the mortality of A\beta42 expressing flies. To rule out the possibility that these studies are not restricted only to the retinal neurons, we employed *elav*-Gal4 that drives expression in the neurons of flies⁷⁷. We misexpressed A β 42 using *elav*-Gal4 (*elav* > A β 42), which resulted in reduced mortality rate as only 50% (n = 200) of the flies could hatch out and survive whereas remaining 50% population were arrested as larvae or pupae. However, all wild-type flies hatched out and did not show any lethality (Fig. 6, n = 200, 100%). We also analyzed mortality rate when Lunasin is misexpressed in *elav* > A β 42 (*elav* > A β 42 + Lun) background. Misexpression of Lun significantly reduced the mortality rate of *elav* > A β 42 background (Fig. 6, n = 200, 80%). Nearly 80% of the flies hatched out and only 20% flies failed to hatch out due to pupal and larval lethality.

Discussion

One of the hallmarks of AD is accumulation of amyloid A β 42 plaques over a period of time, which triggers neuronal death leading to neurodegeneration. This A β 42 mediated neurodegeneration is an outcome of activation of aberrant signaling because of stress in the neurons^{2,4,5,17,27}. Thus, the A β 42 mediated neurodegenerative phenotype observed in AD is not due to a single gene mutation but an outcome of impairment of several signaling pathways^{4,17,19,27,70,71,92}. In order to understand the complexity of this disorder, it is important to identify these signaling pathways.

One of the approaches to discern molecular genetic basis of AD, and to find future cures, it is important to identify the downstream targets of signaling pathways that are triggered as an outcome of the A β 42 accumulation⁴. The efforts have been directed to search for chemical inhibitors that can block/downregulate these downstream targets of aberrant signaling pathways and thereby prevent/delay A β 42 mediated neurodegeneration. In this direction, the repertoire of natural product libraries comprising of plant proteins 102,103 with medicinal properties provide an alternative to the chemical inhibitors which can be screened to identify the ones that can either delay or block the onset of neurodegeneration observed in GMR > A β 42 background.

Several plant products have been identified as therapeutic targets for cancer, inflammation and various other disease^{72,74,75,102,103}. Chronic inflammation has long been implicated in cancer and also plays major role in neurodegenerative disorders like AD. The soy protein Lunasin has multiple interacting domains and may affect different cell biological processes. Lunasin has been reported to have anti-metastatic and chemopreventive activity^{61,72,74,75,104,105}. Lunasin can significantly reduce a melanoma stem cell population¹⁰⁶. It has been suggested that primary anticancer mechanism of Lunasin is based on its activity as a HAT inhibitor^{74,107}. HATs have been known to play role in AD.

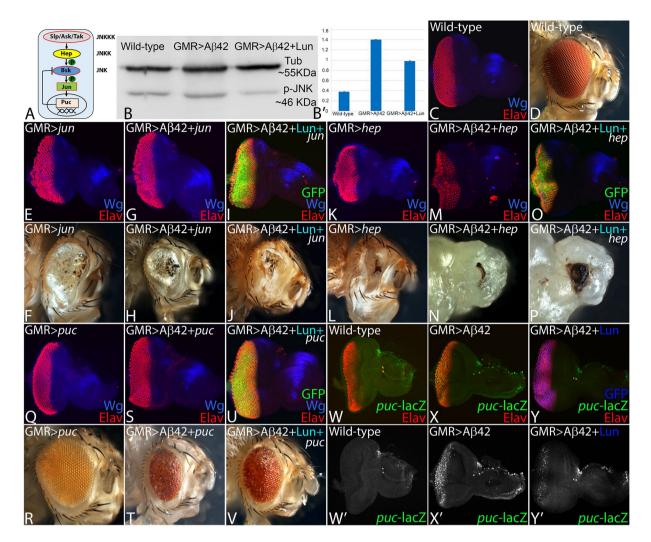


Figure 5. Lunasin downregulates JNK signaling to block Aβ42 mediated neurodegeneration. (A) Schematic presentation of JNK signaling pathway. (B) Levels of phospho-JNK (pJNK) in a semi-quantitative Western Blot can provide the status of JNK signaling. The higher levels of JNK signaling in GMR > A β 42 as compared to the wild-type background were significantly downregulated in GMR > A β 42 + Lun background. The tubulin bands served as controls to normalize the levels of total protein loaded in all three conditions. The p-JNK band staining intensity was calculated by ImageJ. In comparison to the wild-type (C) eye imaginal discs and (**D**) adult eyes, activation of JNK signaling in GMR domain using (**E,F**) $Djun^{aspv7}$ (GMR > jun) and (K,L) constitutively active hep^{Act} (GMR > hep) result in strong neurodegenerative phenotype. Furthermore, activation of JNK signaling in GMR > A β 42 background (**G,H**) GMR > A β 42 + jun (**M,N**) GMR > A β 42 + hep exhibits stronger neurodegenerative phenotype which are not rescued by misexpression of Lunasin (I,J) $GMR > A\beta 42 + Lun + jun$ and (O,P) $GMR > A\beta 42 + Lun + hep.$ (Q,R) Downregulation of JNK signaling by misexpression of puc, a dual phosphatase, results in near wild-type (Q) eye imaginal discs and (R) adult eyes. (S-V) Misexpression of puc in (S,T) GMR > A\(\beta 42\) (GMR > A\(\beta 42\) + puc) and (U,V) GMR > A\(\beta 42\) + Lun $(GMR > A\beta 42 + Lun + puc)$ results in significant rescue as seen in eye imaginal discs and the adult eyes. The puc-lacZ reporter is used as a functional read out of JNK signaling pathway. Expression of puc-lacZ reporter (Green) in (W,W') Wild-type, (X,X') GMR > A\(\beta\)42 and (Y,Y') GMR > A\(\beta\)42 + Lun eye imaginal discs. (W,W') Note that puc has weak expression in the developing photoreceptor neurons in the wild-type eye imaginal discs. However, (X,X') puc expression is dramatically upregulated in GMR > A\(\beta\)42 background. (Y,Y') Misexpression of Lunasin (Lun) along with A β 42 (GMR > A β 42 + Lun) can significantly downregulate *puc* expression in the developing third instar eye disc. Magnification of all eye discs is 20X.

Our studies demonstrated that Lunasin can rescue A β 42 mediated neurodegeneration in the *Drosophila* eye (Fig. 1). Lunasin is known to prevent cancer but its role in neurodegenerative disorders have not been tested to date. We tested the possibility if Lunasin is preventing accumulation of amyloid plaques and thereby preventing A β 42 mediated neurodegeneration. We checked A β 42 plaque accumulation using monoclonal antibody 6E10 in wild-type, GMR > A β 42 and GMR > A β 42 + Lun backgrounds. We found that A β 42 plaque accumulation was comparable between GMR > A β 42 and GMR > A β 42 + Lun backgrounds (Fig. 2). The fact that there is no

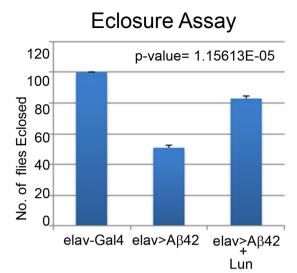


Figure 6. Misexpression of Lunasin can reduce the mortality rate of $elav > A\beta42$ flies. (**A**) A graph comparing the number of flies hatched in Wild-type, $elav > A\beta42$ and $elav > A\beta42 + Lun$ background validates that Lunasin misexpression along with $elav > A\beta42$ ($elav > A\beta42 + LunGFP$) rescues the $elav > A\beta42$ mortality rate. We counted 200 flies in three independent sets from each background and plotted on a graph. These numbers are significant based on the calculations of P-values using the two-tailed t- test using Microsoft Excel 2013.

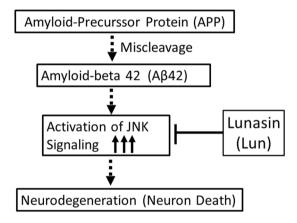


Figure 7. A model to show the mechanism by Lunasin (Lun) blocks A β 42 mediated neurodegeneration. Accumulation of A β 42 plaques trigger a cascade of events, which activates JNK signaling. Activation of JNK signaling in the neuron triggers cell death. Misexpression of Lunasin blocks JNK signaling to promote neuroprotection of retinal neurons in the *Drosophila* eye.

significant difference in the plaque deposition with or without Lunasin in GMR > A β 42 background, proves that Lunasin modulates A β 42 toxicity indirectly, and is downstream of A β 42 plaque accumulation.

In order to identify and characterize the mechanism behind the novel neuroprotective function of Lunasin, we tested various genetic modifiers of A β 42 like C2H2 zinc finger transcription factor, Teashirt (Tsh), CREB binding protein (CBP) and apical basal polarity marker, Crumbs (Crb)^{27,70,71,92}. We found that the neuroprotective function of Lunasin is independent of Tsh, CBP and Crb (data not shown). Interestingly, among various other functions, CBP acts as a histone acetyl transferase (HAT)⁹², and it is known that Lunasin functions as inhibitor of HAT¹⁰⁷. However, we did not see any interaction between Lunasin and CBP in GMR > A β 42 background (data not shown).

In order to discern molecular genetic basis of neuroprotective function of Lunasin, we tested various signaling pathways and found that it is independent of Wg and Dpp signaling (Fig. 4). Finally, we found that neuroprotective function of Lunasin is mediated through downregulation of highly conserved JNK signaling pathway (Figs 5 and 7). Earlier, we have seen that accumulation of A β 42 plaque causes ectopic induction of JNK signaling pathway in the neurons, which in turn triggers neuronal death²⁷. Our data demonstrates that Lunasin misexpression can rescue A β 42 mediated neurodegeneration by downregulating JNK signaling in the *Drosophila* eye (Fig. 7). Thus, our studies provide evidences for the first time that JNK signaling, an important link in onset, manifestation and progression of AD, can be modulated by plant-based protein. Studies in various animal models of AD suggests

the involvement of JNK signaling in AD¹⁰⁸. Our studies open up new avenues where plant proteins expressed by transgenic approach in the neuron can prevent the onset or delay the onset of AD in the animal model of *Drosophila* eye. Since JNK signaling pathway is known to be involved in developmental processes like ageing, development, tissue homeostasis, cell proliferation, cell survival and innate immune response, the modulation of JNK can be of significance in other disease too like Parkinson, stroke etc¹⁰⁹.

The results from our present study support the model of potential therapeutic benefits of Lunasin in Alzheimer's Disease (Fig. 7). Lunasin can prevent progression of A β 42 mediated neurodegeneration by effectively downregulating JNK signaling. The use of plant-based product can be a promising alternative or addition to the use of gene silencing or directly blocking signaling pathways to treat neurodegenerative disorders. Thus, Lunasin, a major bioactive component of the soy-based food has potential to exert a major impact on human health. Further studies are needed to test the efficacy of Lunasin in other vertebrate model organisms to determine if its anti-inflammatory and JNK inhibitory activities show neuroprotective effects. It will be interesting to see if Lunasin can be developed as a potential natural product for the treatment of A β 42 mediated neurodegeneration observed in AD.

Data Availability

The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

References

- Goedert, M. & Spillantini, M. G. A century of Alzheimer's disease. Science 314, 777-781, https://doi.org/10.1126/science.1132814 (2006).
- Hardy, J. The amyloid hypothesis for Alzheimer's disease: a critical reappraisal. J Neurochem 110, 1129–1134, https://doi. org/10.1111/j.1471-4159.2009.06181.x (2009).
- O'Brien, R. J. & Wong, P. C. Amyloid Precursor Protein Processing and Alzheimers Disease. Annu Rev Neurosci, https://doi. org/10.1146/annurev-neuro-061010-113613 (2010).
- 4. Sarkar, A., Irwin, M., Singh, A., Riccetti, M. & Singh, A. Alzheimer's disease: the silver tsunami of the 21(st) century. *Neural Regen Res* 11, 693–697, https://doi.org/10.4103/1673-5374.182680 (2016).
- Crews, L. & Masliah, E. Molecular mechanisms of neurodegeneration in Alzheimer's disease. Hum Mol Genet 19, R12–20, https://doi.org/10.1093/hmg/ddq160 (2010).
- 6. Hirth, F. Drosophila melanogaster in the study of human neurodegeneration. CNS Neurol Disord Drug Targets 9, 504-523 (2010).
- 7. Selkoe, D. J. & Podlisny, M. B. Deciphering the genetic basis of Alzheimer's disease. Annu Rev Genomics Hum Genet 3, 67-99, https://doi.org/10.1146/annurev.genom.3.022502.103022022502.103022 (2002).
- Takami, M. et al. gamma-Secretase: successive tripeptide and tetrapeptide release from the transmembrane domain of betacarboxyl terminal fragment. J Neurosci 29, 13042–13052, https://doi.org/10.1523/JNEUROSCI.2362-09.2009 (2009).
- 9. Hardy, J. & Allsop, D. Amyloid deposition as the central event in the aetiology of Alzheimer's disease. *Trends Pharmacol Sci* 12, 383–388, doi:0165-6147(91)90609-V [pii] (1991).
- 10. Moore, B. D. et al. Short Abeta peptides attenuate Abeta42 toxicity in vivo. J Exp Med 215, 283–301, https://doi.org/10.1084/jem.20170600 (2018).
- Kang, J. et al. The precursor of Alzheimer's disease amyloid A4 protein resembles a cell-surface receptor. Nature 325, 733–736, https://doi.org/10.1038/325733a0 (1987).
- 12. Singh, A. Neurodegeneration a means to an end. Journal of Cell Science and Therapy 3, e107, https://doi.org/10.4172/2157-7013.1000e4107 (2012).
- 13. Singh, A. & Irvine, K. D. Drosophila as a model for understanding development and disease. Developmental dynamics: an official publication of the American Association of Anatomists 241, 1–2, https://doi.org/10.1002/dvdy.23712 (2012).
- 14. Bier, E. *Drosophila*, the golden bug, emerges as a tool for human genetics. *Nat Rev Genet* 6, 9–23, https://doi.org/10.1038/nrg1503
- Cowan, C. M., Shepherd, D. & Mudher, A. Insights from *Drosophila* models of Alzheimer's disease. *Biochem Soc Trans* 38, 988–992, https://doi.org/10.1042/BST0380988 (2010).
- Crowther, D. C., Page, R., Chandraratna, D. & Lomas, D. A. A *Drosophila* model of Alzheimer's disease. *Methods Enzymol* 412, 234–255, https://doi.org/10.1016/S0076-6879(06)12015-7 (2006).
- Pandey, U. B. & Nichols, C. D. Human disease models in *Drosophila* melanogaster and the role of the fly in therapeutic drug discovery. *Pharmacological reviews* 63, 411–436, https://doi.org/10.1124/pr.110.003293 (2011).
- 18. Price, D. L., Tanzi, R. E., Borchelt, D. R. & Sisodia, S. S. Alzheimer's disease: genetic studies and transgenic models. *Annu Rev Genet* 32, 461–493, https://doi.org/10.1146/annurev.genet.32.1.461 (1998).
- 3.2, 461–493, https://doi.org/10.1140/almutev.genet.92.1.461 (1996).
 19. Rincon-Limas, D. E., Jensen, K. & Fernandez-Funez, P. *Drosophila* Models of Proteinopathies: the Little Fly that Coulds. *Curr Pharm Des*, doi:CPD-EPUB-20120130-012 (2012).
- 20. Cauchi, R. J. & van den Heuvel, M. The fly as a model for neurodegenerative diseases: is it worth the jump? *Neurodegener Dis* 3, 338–356, https://doi.org/10.1159/000097303 (2006).
- Iijima-Ando, K. & Iijima, K. Transgenic Drosophila models of Alzheimer's disease and tauopathies. Brain Struct Funct 214, 245–262, https://doi.org/10.1007/s00429-009-0234-4 (2010).
- Wangler, M. F. et al. Model Organisms Facilitate Rare Disease Diagnosis and Therapeutic Research. Genetics 207, 9–27, https://doi. org/10.1534/genetics.117.203067 (2017).
- Yamamoto, S. et al. A Drosophila genetic resource of mutants to study mechanisms underlying human genetic diseases. Cell 159, 200–214, https://doi.org/10.1016/j.cell.2014.09.002 (2014).
- 24. Cao, W. et al. Identification of novel genes that modify phenotypes induced by Alzheimer's beta-amyloid overexpression in *Drosophila. Genetics* 178, 1457–1471, https://doi.org/10.1534/genetics.107.078394 (2008).
- Casas-Tinto, S. et al. The ER stress factor XBP1s prevents amyloid-{beta} neurotoxicity. Hum Mol Genet 20, 2144–2160, https://doi. org/10.1093/hmg/ddr100 (2011).
- 26. Kumar, J. P. The Fly Eye: Through the Looking Glass. Developmental dynamics: an official publication of the American Association of Anatomists. https://doi.org/10.1002/dvdy.24585 (2017).
- 27. Tare, M. et al. Activation of JNK signaling mediates amyloid-ss-dependent cell death. Plos One 6, e24361, https://doi.org/10.1371/journal.pone.0024361PONE-D-11-11860 (2011).
- 28. Cohen, B., Simcox, A. & Cohen, S. Allocation of the thoracic imaginal primordia in the *Drosophila* embryo. *Development* 117, 597–608 (1993).
- 29. Held, L. I. J. In Imaginal Disc (ed. Held, L. I.) 197-236 (Cambridge University Press, 2002).

- Atkins, M. & Mardon, G. Signaling in the third dimension: the peripodial epithelium in eye disc development. Developmental dynamics: an official publication of the American Association of Anatomists 238, 2139–2148, https://doi.org/10.1002/dvdy.22034 (2009).
- Kumar, J. P. Retinal determination the beginning of eye development. Curr Top Dev Biol 93, 1–28, https://doi.org/10.1016/B978-0-12-385044-7.00001-1 (2010).
- 32. Tsachaki, M. & Sprecher, S. G. Genetic and developmental mechanisms underlying the formation of the *Drosophila* compound eye. *Developmental dynamics: an official publication of the American Association of Anatomists* **241**, 40–56, https://doi.org/10.1002/dvdy.22738 (2012).
- Bonini, N. M., Leiserson, W. M. & Benzer, S. The eyes absent gene: genetic control of cell survival and differentiation in the developing *Drosophila* eye. Cell 72, 379–395 (1993).
- 34. Cheyette, B. N. et al. The Drosophila sine oculis locus encodes a homeodomain-containing protein required for the development of the entire visual system. Neuron 12, 977–996 (1994).
- 35. Hanson, I. M. et al. PAX6 mutations in aniridia. Hum Mol Genet 2, 915-920 (1993).
- 36. Jang, C. C. et al. Two Pax genes, eye gone and eyeless, act cooperatively in promoting *Drosophila* eye development. *Development* 130, 2939–2951 (2003).
- 37. Mardon, G., Solomon, N. M. & Rubin, G. M. dachshund encodes a nuclear protein required for normal eye and leg development in *Drosophila*. *Development* 120, 3473–3486 (1994).
- 38. Quiring, R., Walldorf, U., Kloter, U. & Gehring, W. J. Homology of the eyeless gene of *Drosophila* to the Small eye gene in mice and Aniridia in humans. *Science* 265, 785–789 (1994).
- Seimiya, M. & Gehring, W. J. The Drosophila homeobox gene optix is capable of inducing ectopic eyes by an eyeless-independent mechanism. Development 127, 1879–1886 (2000).
- 40. Burgy-Roukala, E., Miellet, S., Mishra, A. K. & Sprecher, S. G. In Molecular Genetics of Axial Patterning, Growth and Disease in the *Drosophila* Eye (eds Amit Singh & Madhuri Kango-Singh) Ch. 1, 1–36 (Springer, 2013).
- 41. Kumar, J. P. My what big eyes you have: How the *Drosophila* retina grows. *Dev Neurobiol*, https://doi.org/10.1002/dneu.20921 (2011)
- Pappu, K. S. & Mardon, G. Genetic control of retinal specification and determination in *Drosophila*. The International journal of developmental biology 48, 913–924, https://doi.org/10.1387/ijdb.041875kp (2004).
- 43. Roignant, J. Y. & Treisman, J. E. Pattern formation in the Drosophila eye disc. The International journal of developmental biology 53, 795–804. https://doi.org/10.1387/jidb.072483jr (2009).
- Singh, A., Tare, M., Puli, O. R. & Kango-Singh, M. A glimpse into dorso-ventral patterning of the Drosophila eye. Developmental dynamics: an official publication of the American Association of Anatomists 241, 69–84, https://doi.org/10.1002/dvdy.22764 (2012).
- 45. Tare, M., Puli, O. R. & Singh, A. In Molecular Genetics of Axial Patterning, Growth and Disease in the *Drosophila* Eye (eds Amit Singh & Madhuri Kango-Singh) Ch. 2, 37–75 (Springer, 2013).
- 46. Kumar, J. P. Building an ommatidium one cell at a time. *Developmental dynamics: an official publication of the American Association of Anatomists* **241**, 136–149, https://doi.org/10.1002/dvdy.23707 (2012).
- Ready, D. F., Hanson, T. E. & Benzer, S. Development of the *Drosophila* retina, a neurocrystalline lattice. *Dev Biol* 53, 217–240, doi:0012-1606(76)90225-6 (1976).
- 48. Wolff, T. & Ready, D. F. In The Development of Drosophila melanogaster. Vol. III (eds Bate, M. & Martinez-Arias, A.) 1277–1325 (Cold-Spring Harbor: Cold Spring Harbor Laboratory Press., 1993).
- Garrity, P. A. et al. Retinal axon target selection in *Drosophila* is regulated by a receptor protein tyrosine phosphatase. *Neuron* 22, 707–717, doi:S0896-6273(00)80730-8 (1999).
- 50. Garrity, P. A. *et al. Drosophila* photoreceptor axon guidance and targeting requires the dreadlocks SH2/SH3 adapter protein. *Cell* **85**, 639–650 (1996).
- Brachmann, C. B. & Cagan, R. L. Patterning the fly eye: the role of apoptosis. *Trends Genet* 19, 91–96, https://doi.org/10.1016/ S0168-9525(02)00041-0 (2003).
- 52. Mehlen, P., Mille, F. & Thibert, C. Morphogens and cell survival during development. J Neurobiol 64, 357–366, https://doi.org/10.1002/neu.20167 (2005).
- 53. Singh, A., Shi, X. & Choi, K. W. Lobe and Serrate are required for cell survival during early eye development in *Drosophila*. *Development* 133, 4771–4781, https://doi.org/10.1242/dev.02686 (2006).
- 54. Tare, M., Sarkar, A., Bedi, S., Kango-Singh, M. & Singh, A. Cullin-4 regulates Wingless and JNK signaling-mediated cell death in the *Drosophila* eye. *Cell Death Dis* 7, e2566, https://doi.org/10.1038/cddis.2016.338 (2016).
- 55. Cordero, J., Jassim, O., Bao, S. & Cagan, R. A role for wingless in an early pupal cell death event that contributes to patterning the *Drosophila* eye. *Mech Dev* 121, 1523–1530, https://doi.org/10.1016/j.mod.2004.07.004 (2004).
- Lin, H. V., Rogulja, A. & Cadigan, K. M. Wingless eliminates ommatidia from the edge of the developing eye through activation of apoptosis. *Development* 131, 2409–2418, https://doi.org/10.1242/dev.01104 (2004).
- 57. Raffery, L. A. & Sutherland, D. J. TGF-beta family signal transduction in *Drosophila* development: from Mad to Smads. *Dev Biol* 210, 251–268, https://doi.org/10.1006/dbio.1999.9282 (1999).
- Ma, C. & Moses, K. Wingless and patched are negative regulators of the morphogenetic furrow and can affect tissue polarity in the developing *Drosophila* compound eye. *Development* 121, 2279–2289 (1995).
- Treisman, J. E. & Rubin, G. M. wingless inhibits morphogenetic furrow movement in the *Drosophila* eye disc. *Development* 121, 3519–3527 (1995).
- 60. Royet, J. & Finkelstein, R. Establishing primordia in the *Drosophila* eye-antennal imaginal disc: the roles of decapentaplegic, wingless and hedgehog. *Development* 124, 4793–4800 (1997).
- 61. Adachi-Yamada, T. & O'Connor, M. B. Morphogenetic apoptosis: a mechanism for correcting discontinuities in morphogen gradients. *Dev Biol* 251, 74–90, doi:S0012160602908213 (2002).
- 62. Dhanasekaran, D. N. & Reddy, E. P. JNK signaling in apoptosis. Oncogene 27, 6245–6251, https://doi.org/10.1038/onc.2008.301 (2008).
- 63. Stronach, B. Dissecting JNK signaling, one KKKinase at a time. *Developmental dynamics: an official publication of the American Association of Anatomists* **232**, 575–584, https://doi.org/10.1002/dvdy.20283 (2005).
- 64. Igaki, T. Correcting developmental errors by apoptosis: lessons from *Drosophila* JNK signaling. *Apoptosis* 14, 1021–1028, https://doi.org/10.1007/s10495-009-0361-7 (2009).
- 65. Igaki, T. et al. Eiger, a TNF superfamily ligand that triggers the *Drosophila* JNK pathway. EMBO J 21, 3009–3018, https://doi.org/10.1093/emboj/cdf306 (2002).
- 66. Martin-Blanco, E. et al. puckered encodes a phosphatase that mediates a feedback loop regulating JNK activity during dorsal closure in *Drosophila*. Genes Dev 12, 557–570 (1998).
- Moreno, E., Yan, M. & Basler, K. Evolution of TNF signaling mechanisms: JNK-dependent apoptosis triggered by Eiger, the *Drosophila* homolog of the TNF superfamily. *Curr Biol* 12, 1263–1268, doi:S0960982202009545 (2002).
- 68. Moses, K. & Rubin, G. M. Glass encodes a site-specific DNA-binding protein that is regulated in response to positional signals in the developing *Drosophila* eye. *Genes Dev* 5, 583–593 (1991).
- 69. Brand, A. H. & Perrimon, N. Targeted gene expression as a means of altering cell fates and generating dominant phenotypes. *Development* 118, 401–415 (1993).

- 70. Moran, M. T., Tare, M., Kango-Singh, M. & Singh, A. Homeotic Gene teashirt (tsh) has a neuroprotective function in amyloid-beta 42 mediated neurodegeneration. *Plos One* 8, e80829, https://doi.org/10.1371/journal.pone.0080829 (2013).
- 71. Steffensmeier, A. M. *et al.* Novel neuroprotective function of apical-basal polarity gene crumbs in amyloid beta 42 (abeta 42) mediated neurodegeneration. *Plos One* 8, e78717, https://doi.org/10.1371/journal.pone.0078717 (2013).
- 72. Jones, G. & Srivastava, A. Understanding Lunasin's biology and potential as a cancer therapeutic by utilizing *Drosophila* genetics. Exp Biol Med (Maywood) 239, 519–528, https://doi.org/10.1177/1535370214522180 (2014).
- 73. Galvez, A. F. & de Lumen, B. O. A soybean cDNA encoding a chromatin-binding peptide inhibits mitosis of mammalian cells. *Nat Biotechnol* 17, 495–500, https://doi.org/10.1038/8676 (1999).
- 74. Shidal, C., Inaba, J. I., Yaddanapudi, K. & Davis, K. R. The soy-derived peptide Lunasin inhibits invasive potential of melanoma initiating cells. *Oncotarget* 8, 25525–25541, https://doi.org/10.18632/oncotarget.16066 (2017).
- 75. Shidal, Č., Al-Rayyan, Ň., Yaddanapudi, K. & Davis, K. R. Lunasin is a novel therapeutic agent for targeting melanoma cancer stem cells. Oncotarget 7, 84128–84141, https://doi.org/10.18632/oncotarget.11554 (2016).
- Businaro, R. et al. Modulation Of Inflammation As A Way Of Delaying Alzheimer's Disease Progression: The Diet's Role. Curr Alzheimer Res, https://doi.org/10.2174/1567205014666170829100100 (2017).
- 77. Yao, K. M. & White, K. Neural specificity of elav expression: defining a *Drosophila* promoter for directing expression to the nervous system. *J Neurochem* **63**, 41–51 (1994).
- 78. Treier, M., Bohmann, D. & Mlodzik, M. JUN cooperates with the ETS domain protein pointed to induce photoreceptor R7 fate in the *Drosophila* eye. *Cell* 83, 753–760, doi:0092-8674(95)90188-4 [pii] (1995).
- Kassis, J. A., Noll, E., VanSickle, E. P., Odenwald, W. F. & Perrimon, N. Altering the insertional specificity of a *Drosophila* transposable element. *Proc Natl Acad Sci USA* 89, 1919–1923 (1992).
- 80. Blackman, R. K., Sanicola, M., Raftery, L. A., Gillevet, T. & Gelbart, W. M. An extensive 3' cis-regulatory region directs the imaginal disk expression of decapentaplegic, a member of the TGF-beta family in *Drosophila*. *Development* 111, 657–666 (1991).
- ausk expression of decapentapiegic, a member of the TGF-beta family in *Drosophila. Development* 111, 657–666 (1991).

 81. Singh, A. Enhacer Trap Technique A novel tool for identification and developmental characterization of *Drosophila* genes. *Current Science* 68, 517–525 (1995).
- 82. Rovelet-Lecrux, A. et al. APP locus duplication causes autosomal dominant early-onset Alzheimer disease with cerebral amyloid
- angiopathy. Nat Genet 38, 24–26, https://doi.org/10.1038/ng1718 (2006).
 83. Kanca, O., Bellen, H. J. & Schnorrer, F. Gene Tagging Strategies To Assess Protein Expression, Localization, and Function in Drosophila. Genetics 207, 389–412, https://doi.org/10.1534/genetics.117.199968 (2017).
- 84. Lam, Y., Galvez, A. & de Lumen, B. O. Lunasin suppresses E1A-mediated transformation of mammalian cells but does not inhibit growth of immortalized and established cancer cell lines. *Nutr Cancer* 47, 88–94, https://doi.org/10.1207/s15327914nc4701_11
- Singh, A., Kango-Singh, M. & Sun, Y. H. Eye suppression, a novel function of teashirt, requires Wingless signaling. *Development* 129, 4271–4280 (2002).
- Wittkorn, E., Sarkar, A., Garcia, K., Kango-Singh, M. & Singh, A. The Hippo pathway effector Yki downregulates Wg signaling to promote retinal differentiation in the *Drosophila* eye. *Development* 142, 2002–2013, https://doi.org/10.1242/dev.117358 (2015).
- 87. Sarkar, A., Gogia, N., Farley, K., Payton, L. & Singh, A. Characterization of a morphogenetic furrow specific Gal4 driver in the developing *Drosophila* eye. *Plos One* 13, e0196365, https://doi.org/10.1371/journal.pone.0196365 (2018).
- 88. Zipursky, S. L., Venkatesh, T. R., Teplow, D. B. & Benzer, S. Neuronal development in the *Drosophila* retina: monoclonal antibodies as molecular probes. *Cell* 36, 15–26, doi:0092-8674(84)90069-2 (1984).
- 89. Singh, A. & Gopinathan, K. P. Confocal microscopy: A powerful tool for biological research. Current Science 74, 841–851 (1998).
- McCall, K. & Peterson, J. S. Detection of apoptosis in *Drosophila*. Methods Mol Biol 282, 191–205, https://doi.org/10.1385/1-59259-812-9:191 (2004).
- 91. White, K. et al. Genetic control of programmed cell death in Drosophila. Science 264, 677-683 (1994).
- Cutler, T. et al. Drosophila Eye Model to Study Neuroprotective Role of CREB Binding Protein (CBP) in Alzheimer's Disease. Plos One 10, e0137691, https://doi.org/10.1371/journal.pone.0137691 (2015).
- 93. Gogia, N., Sarkar, A. & Singh, A. An undergraduate cell Biology lab: Western Blotting to detect proteins from *Drosophila* eye. *Drosophila Information Service* 100, 218–225 (2017).
- 94. McCall, K., Baum, J. S., Cullen, K. & Peterson, J. S. Visualizing apoptosis. Methods Mol Biol 247, 431-442 (2004).
- 95. Gunawardena, S. & Goldstein, L. S. Disruption of axonal transport and neuronal viability by amyloid precursor protein mutations in *Drosophila*. *Neuron* **32**, 389–401 (2001).
- 96. Fujita, S. C., Zipursky, S. L., Benzer, S., Ferrus, A. & Shotwell, S. L. Monoclonal antibodies against the *Drosophila* nervous system. *Proc Natl Acad Sci USA* 79, 7929–7933 (1982).
- Bessa, J., Gebelein, B., Pichaud, F., Casares, F. & Mann, R. S. Combinatorial control of *Drosophila* eye development by eyeless, homothorax, and teashirt. *Genes Dev* 16, 2415–2427, https://doi.org/10.1101/gad.1009002 (2002).
- Kango-Singh, M., Singh, A. & Henry Sun, Y. Eyeless collaborates with Hedgehog and Decapentaplegic signaling in *Drosophila* eye induction. Dev Biol 256, 49–60 (2003).
- Shen, W. & Mardon, G. Ectopic eye development in *Drosophila* induced by directed dachshund expression. *Development* 124, 45–52 (1997).
- Adachi-Yamada, T., Fujimura-Kamada, K., Nishida, Y. & Matsumoto, K. Distortion of proximodistal information causes JNK-dependent apoptosis in *Drosophila* wing. *Nature* 400, 166–169, https://doi.org/10.1038/22112 (1999).
- Agnes, F., Suzanne, M. & Noselli, S. The *Drosophila JNK* pathway controls the morphogenesis of imaginal discs during metamorphosis. *Development* 126, 5453–5462 (1999).
- Arntzen, C. Plant-made pharmaceuticals: from 'Edible Vaccines' to Ebola therapeutics. Plant Biotechnol J 13, 1013–1016, https://doi.org/10.1111/pbi.12460 (2015).
- 103. Yao, J., Weng, Y., Dickey, A. & Wang, K. Y. Plants as Factories for Human Pharmaceuticals: Applications and Challenges. Int J Mol Sci 16, 28549–28565, https://doi.org/10.3390/ijms161226122 (2015).
- 104. Inaba, J., McConnell, E. J. & Davis, K. R. Lunasin sensitivity in non-small cell lung cancer cells is linked to suppression of integrin signaling and changes in histone acetylation. *Int J Mol Sci* 15, 23705–23724, https://doi.org/10.3390/ijms151223705 (2014).
- 105. Jiang, Q. et al. Lunasin suppresses the migration and invasion of breast cancer cells by inhibiting matrix metalloproteinase-2/-9 via the FAK/Akt/ERK and NF-kappaB signaling pathways. Oncol Rep 36, 253–262, https://doi.org/10.3892/or.2016.4798 (2016).
- 106. Dia, V. P. & Gonzalez de Mejia, E. Lunasin potentiates the effect of oxaliplatin preventing outgrowth of colon cancer metastasis, binds to alpha5 beta1 integrin and suppresses FAK/ERK/NF-kappaB signaling. Cancer Lett 313, 167–180, https://doi.org/10.1016/j.canlet.2011.09.002 (2011).
- 107. Hernandez-Ledesma, B., Hsieh, C. C. & de Lumen, B. O. Lunasin, a novel seed peptide for cancer prevention. *Peptides* 30, 426–430, https://doi.org/10.1016/j.peptides.2008.11.002 (2009).
- 108. Morishima, Y. *et al.* Beta-amyloid induces neuronal apoptosis via a mechanism that involves the c-Jun N-terminal kinase pathway and the induction of Fas ligand. *J Neurosci* **21**, 7551–7560 (2001).
- 109. Mehan, S., Meena, H., Sharma, D. & Sankhla, R. JNK: a stress-activated protein kinase therapeutic strategies and involvement in Alzheimer's and various neurodegenerative abnormalities. J Mol Neurosci 43, 376–390, https://doi.org/10.1007/s12031-010-9454-6 (2011).

Acknowledgements

We thank the Bloomington Stock Center for the *Drosophila* strains, K. Cho and the Developmental Studies Hybridoma Bank (DSHB) for the antibodies. We would like to thank Michael Moran for training students and A. Giaquinto for helping with the fly crosses and for maintenance of stocks used in this paper. The authors also thank the members of Singh lab for critical comments on the manuscript. Confocal microscopy was supported by Biology Department central core facility. A.S. and N.G. are supported by Graduate program of Biology. This work is supported by KBRIN-IDEA grant funded through a parent grant 5P20GM103436 to A.S., Start-up support from the University of Dayton, and a subaward on RO1 (CA183991, PI Nakano) to MKS, and National Institute of General Medical Sciences (NIGMS) - 1 R15 GM124654-01, STEM Catalyst Grant from the University of Dayton and start-up support from the University of Dayton to A.S.

Author Contributions

A.S., N.G. and N.G. the three equal first authors performed experiments, analyzed the data and provided comments to the manuscript. A.S. performed earlier experiments about testing the UAS-Lunasin transgene. A.S. screened the genetic modifiers and helped in manuscript writing. G.J., N.P. and A.S. were involved in generating the UAS-Lunasin transgene. A.S. and M.K.S. were involved in developing the concept, designing the experiment, and analyzing the data and writing the manuscript.

Additional Information

Competing Interests: The authors declare no competing interests.

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