Activated AKT/PKB signaling in C. elegans uncouples temporally distinct outputs of DAF-2/insulin-like signaling
Minaxi S Gami, Wendy B Iser, Keaton B Hanselman and Catherine A Wolkow*

Address: Laboratory of Neurosciences, NIA, NIH, Baltimore, MD, USA
Email: Minaxi S Gami - gamimi@grc.nia.nih.gov; Wendy B Iser - iserwe@grc.nia.nih.gov; Keaton B Hanselman - keaton.hanselman@duke.edu; Catherine A Wolkow* - wolkowca@grc.nia.nih.gov
* Corresponding author

Abstract

Background: In the nematode, Caenorhabditis elegans, a conserved insulin-like signaling pathway controls larval development, stress resistance and adult lifespan. AGE-1, a homolog of the p110 catalytic subunit of phosphoinositide 3-kinases (PI3K) comprises the major known effector pathway downstream of the insulin receptor, DAF-2. Phospholipid products of AGE-1/PI3K activate AKT/PKB kinase signaling via PDK-1. AKT/PKB signaling antagonizes nuclear translocation of the DAF-16/FOXO transcription factor. Reduced AGE-1/PI3K signaling permits DAF-16 to direct dauer larval arrest and promote long lifespan in adult animals. In order to study the downstream effectors of AGE-1/PI3K signaling in C. elegans, we conducted a genetic screen for mutations that suppress the constitutive dauer arrest phenotype of age-1(mg109) animals.

Results: This report describes mutations recovered in a screen for suppressors of the constitutive dauer arrest (daf-C) phenotype of age-1(mg109). Two mutations corresponded to alleles of daf-16. Two mutations were gain-of-function alleles in the genes, akt-1 and pdk-1, encoding phosphoinositide-dependent serine/threonine kinases. A fifth mutation, mg227, located on chromosome X, did not correspond to any known dauer genes, suggesting that mg227 may represent a new component of the insulin pathway. Genetic epistasis analysis by RNAi showed that reproductive development in age-1(mg109);akt-1(mg247) animals was dependent on the presence of pdk-1. Similarly, reproductive development in age-1(mg109);pdk-1(mg261) animals was dependent on akt-1. However, reproductive development in age-1(mg109); mg227 animals required only akt-1, and pdk-1 activity was dispensable in this background. Interestingly, while mg227 suppressed dauer arrest in age-1(mg109) animals, it enhanced the long lifespan phenotype. In contrast, akt-1(mg247) and pdk-1(mg261) did not affect lifespan or stress resistance, while both daf-16 alleles fully suppressed these phenotypes.

Conclusion: A screen for suppressors of PI3K mutant phenotypes identified activating mutations in two known pathway components, providing insights into their regulation. In particular, the interdependence of akt-1 and pdk-1, even in activated forms, supports the existence of AGE-1-independent pathways for these phospholipid-dependent kinases. Phenotypic analysis of these alleles shows that the larval and adult outputs of AGE-1/PI3K are fully separable in these mutants.
Background
In the nematode *C. elegans*, development, lifespan and stress resistance are regulated by a conserved insulin-like signaling pathway [1]. Under optimal conditions, *C. elegans* nematodes develop through four larval stages (L1-L4) to become reproductive adults. In adverse environmental conditions, such as low food abundance or high population density, larvae can enter diapause and arrest development as dauer larvae. Dauer larvae are non-feeding, long-lived, sexually immature and stress resistant animals. Morphologically they are thinner and darker than reproductive adults [2,3]. Once favorable conditions resume, dauer larvae can re-enter the reproductive developmental pathway to become fertile adults.

In *C. elegans*, three genetic pathways influence the decision to arrest development at the dauer larval stage [3,4]. One is a neuronal daf-11 guanylyl cyclase pathway [4,5]. The second is a TGF-β pathway that regulates dauer arrest through expression of the DAF-7/TGF-β ligand [6]. Ultimately, the daf-11 and daf-7 pathways act on a nuclear hormone receptor, encoded by daf-12 [4,7]. The third pathway regulating dauer arrest is an insulin-like signaling pathway that includes the genes daf-2, encoding an insulin/IGF-1 receptor-like protein, *age-1*, encoding the p110 catalytic subunit of PI3K, as well as a collection of protein kinases, encoded by the genes akt-1, -2, pdk-1 and sgg-1, which mediate phospholipid signaling downstream from AGE-1 [8-12]. Phosphoinositide bound PDK-1 activates the protein kinases which, in turn, phosphorylate and inhibit nuclear translocation of the FOXO transcription factor DAF-16 [13-16]. AGE-1/PI3K-generated phospholipids are believed to bind to both PDK-1 and AKT-1. Phosphoinositide bound AKT-1 undergoes a conformational change that promotes phosphorylation by PDK-1 and this phosphorylation is greatly enhanced if PDK-1 is also bound to phosphoinositides. However, kinase activity can also be detected in the absence of phosphoinositides suggesting that phospholipid-binding may not be absolutely required for PDK-1 catalytic activity [17,18].

In addition to regulating dauer arrest in larvae, DAF-2/insulin signaling also regulates lifespan and stress resistance in adult animals [9,19-23]. As with dauer larval arrest, *daf-16* activity is required for the longevity and stress resistance phenotypes in *daf-2* pathway mutants [20,21]. DAF-2/insulin signaling regulates the expression of many genes involved in metabolism and stress resistance, including some genes that appear to be direct transcriptional targets of DAF-16, such as *daf-15/RAPTOR* and *sod-3/Mn-SOD* [24-26].

Most of the known *daf-2* pathway components were identified in genetic screens on the basis of dauer arrest phenotype. In particular, a screen for suppressors of dauer larval arrest in animals with the null *age-1(mg44)* mutation uncovered hypermorphic alleles of *akt-1* and *pdk-1* [11,12]. In order to determine whether additional effectors of *AGE-1/PI3K* signaling could be recovered by this approach, we repeated this screen, using the missense *age-1(mg109)* allele [10].

Here, we describe five mutations identified in this screen, comprising two *daf-16* alleles and new hypermorphic alleles of *akt-1* and *pdk-1* that appear to promote activation of AKT/PKB signaling in the absence of *AGE-1/PI3K*. The *akt-1* and *pdk-1* alleles also uncoupled the larval dauer arrest, adult longevity and stress resistance phenotypes of *age-1(mg109)* animals. We also describe an uncloned allele that has the unusual phenotype of enhancing the longevity of *age-1(mg109)* adult animals. These findings suggest that the larval and adult phenotypes of insulin signaling in *C. elegans* may be differentially affected by AKT/PKB signaling.

Results and discussion
Genetic identification of *age-1(mg109)* suppressors
In *C. elegans*, both lifespan and dauer formation are regulated by *age-1* activity [10,27]. Animals with maternal, but not zygotic, *age-1* activity do not arrest as dauer larvae, but grow into reproductive adults that are stress resistant and long-lived. In the absence of both maternal and zygotic *age-1* activity, animals arrest development as dauer larvae. In order to identify genetic modifiers of *age-1*, we conducted a screen for mutations that could suppress the dauer-constitutive (*daf-C*) phenotype of *age-1(mg109)*, which carries a missense mutation in the lipid kinase domain of *AGE-1/PI3K* [10]. Previously, a similar genetic screen identified the AKT-1 and PDK-1 effectors of *AGE-1/PI3K* [11,12]. We repeated this screen in an effort to identify additional mutations capable of activating signaling in the absence of *AGE-1/PI3K*. From a screen of approximately 20,000 haploid genomes, we identified 40 alleles that could suppress the dauer-constitutive phenotype of *age-1(mg109)* animals (see Methods). Five of these alleles were chosen for further study, based on high penetrance and strong suppression of *age-1(mg109)* constitutive dauer arrest. In particular, four of these alleles, *mg242*, *mg255*, *mg261* and *mg227*, strongly suppressed the *age-1(mg109)* dauer-constitutive phenotype (Table 1). The *mg247* allele partially suppressed this larval phenotype, as *age-1(mg109);mg247* animals bypassed dauer arrest but then developed into sterile adults (Table 1). Previous studies have shown that adult sterility is correlated with modest decrements in *AGE-1/PI3K* signaling [27,28]. In addition to the adult sterility phenotype, *mg247* animals also exhibited a slight developmental delay, which has also been previously observed in insulin pathway mutants [19,27,28]. Thus, we conclude that the *mg247* allele only
partially restores phospholipid signaling in *age-1(mg109)* animals.

Using standard SNP snip-mapping protocols, we mapped four of these alleles to genetic regions that contained known components of the *age-1* pathway [29]. Molecular lesions in candidate genes were identified by direct DNA sequencing (see Methods). Two alleles, *mg242* and *mg255*, corresponded to nonsense mutations in *daf-16* (Figure 1A). *daf-16(mg242)* is a Trp220Amb nonsense mutation in *daf-16* and truncates both major *daf-16* splice forms, *daf-16a* and *daf-16b*, resulting in a predicted null phenotype. *daf-16(mg255)* carries a Trp144Amb nonsense mutation in the forkhead domain of the *daf-16a* splice form only (Figure 1A). Two other suppressor alleles corresponded to mutations in the serine/threonine kinases that are regulated by phosphoinositides (PIP) products of AGE-1/PI3K. The *mg247* allele corresponded to a missense mutation in *akt-1* (Ala149Thr). This mutation changes a nonconserved residue in the linker region between the pleckstrin homology (PH) and kinase domains of AKT-1 and is predicted to affect both *akt-1a* and *akt-1b* splice forms (Figure 1B). The *mg261* allele is the result of two mutations in different regions of PDK-1 of which both are missense mutations. The first corresponds to a nonconserved residue in the linker region between the kinase domain and the PH domain (Ala447Val) and the second in a conserved residue in the PH domain (Glu530Trp) (Figure 1C). A fifth mutation, *mg227*, was mapped to a region of chromosome X that does not contain any known components of the *daf-2* pathway. Two known dauer pathway genes, *sgk-1* (X 23.70) and *daf-12* (X 2.39), are also on chromosome X but are on the outside of the candidate region and thus do not correspond to the gene mutated by *mg227* (see Methods) [7,8].

Both *akt-1* and *pdk-1* encode proteins that are effectors of signaling downstream of AGE-1/PI3K [11,12]. The observation that the *akt-1(mg247)* and *pdk-1(mg261)* alleles suppressed the daf-C phenotype of *age-1(mg109)* animals suggested these alleles may activate AKT-1 and PDK-1 function in the absence of the phospholipid products of AGE-1 (Table 1). Consistent with this hypothesis, both *akt-1(mg247)* and *pdk-1(mg261)* segregated in a dominant fashion (progeny of heterozygous mothers were scored, *akt-1(mg247)*, 72.2 % bypassers, 27.8 % dauer, *n* = 467 animals: *pdk-1(mg261)*, 75.8 % bypassers, 24 % dauer, *n* = 153 animals). To confirm that *mg247* and *mg261* were activating mutations, we tested whether RNA interference (RNAi)-mediated expression of these genes could revert their suppression of *age-1(mg109)* dauer arrest. Since *age-1(mg109);akt-1(mg247)* adults were mostly sterile at 25°C, we conducted these experiments at 20°C using the feeding RNAi protocol [30]. Consistent with *mg247* being a hypermorphic *akt-1* allele, RNAi of *akt-1* in *age-1(mg109);akt-1(mg247)* adults fully reversed the suppression of dauer arrest and animals arrested as dauer larvae (Figure 2). Similarly, *pdk-1* RNAi reversed the dauer suppression phenotype in *age-1(mg109);pdk-1(mg261)* animals.

### AKT-1 and PDK-1 are interdependent when activated by mutations

We next examined the epistatic relationships of *akt-1(mg247)* and *pdk-1(mg261)* with other components of the AGE-1/PI3K effector pathway using RNAi. First, we found that *pdk-1(RNAi)* caused constitutive dauer larval arrest in *age-1(mg109);akt-1(mg247)* animals, suggesting that *mg247*-activated AKT-1 required PDK-1-mediated phosphorylation for activity (Figure 3). In contrast, *akt-2(RNAi)* did not have a significant effect on the development of *age-1(mg109);akt-1(mg247)* animals. PDK-1 is

### Table 1: Suppression of the *age-1(mg109)* dauer arrest phenotype.

| Genotype                          | 20°C (% of population) | 25°C (% of population) | (n)  |
|-----------------------------------|-------------------------|-------------------------|------|
|                                   | Dauer larvae            | Adults                  |      |
|                                   | Sterile                | Fertile                      |      |
| Wildtype                          | 0                       | 0                        | 100  |
| *age-1(mg109)*§                   | 100                     | 0                        | 0    |
| *age-1(mg109);mg242*              | 0                       | 0                        | 100  |
| *age-1(mg109);mg255*              | 0                       | 0                        | 100  |
| *age-1(mg109);mg247*              | 0                       | 100                      | 64.4 |
| *age-1(mg109);mg261*              | 0                       | 100                      | 64.4 |
| *age-1(mg109);mg227*              | 0                       | 100                      | 64.4 |

Some animals exhibited Egl (egg retention) phenotype: * 35 %, ** 4.7 %, *** 32 %, # 100 %.

§ These are homozygote progeny of *age-1(mg109)* homozygous hermaphrodites that have been maternally rescued for dauer arrest [27].
believed to be constitutively active under basal conditions [17,18]. Thus, in age-1(mg109) animals carrying the akt-1(mg247) allele, PDK-1 may be active at low levels.

Next, we observed that reproductive development of age-1(mg109);pdk-1(mg261) animals was suppressed by akt-1(RNAi), and this effect was enhanced in combination with akt-2(RNAi), showing that the activated form of PDK-1 remained coupled to the AKT/PKB outputs (Figure 3). Bypass of dauer larval arrest in age-1(mg109);akt-1(mg247) and age-1(mg109); pdk-1(mg261) animals was unaffected by RNAi-mediated knockdown of sgk-1, which encodes another effector of AGE-1/PI3K signaling (data not shown) [8].

**Figure 1**

**Mutations identified as suppressors of constitutive dauer arrest in age-1(mg109) animals.** Suppressors were mapped using standard SNP-snip-mapping protocols with the Hawaiian strain CB4856 [29]. Mutations were confirmed using PCR and direct DNA sequencing. (A) daf-16(mg242) is a Trp220Amber substitution in exon 5, affecting both daf-16a and daf-16b splice forms. daf-16(mg255) is a Trp144Amber substitution in exon 3 that only affects the daf-16a splice form. Both mutations fall within the Forkhead domain (indicated by the grey area) involved in DNA interaction. (B) akt-1(mg247) is an Ala149Thr substitution that falls in the linker region, between the pleckstrin homology domain (PH, indicated by the grey boxes) and the protein kinase C domain (KD, indicated by the black boxes). (C) pdk-1(mg261) is a Ala447Val substitution in the linker region and a Glu530Trp substitution in the pleckstrin homology domain (PH, grey boxes) of PDK-1.
We also analyzed the epistatic interactions of \( mg227 \) with \( akt-1 \), \( akt-2 \), \( sgk-1 \) and \( pdk-1 \). In these experiments, \( akt-1 \text{ (RNAi)} \) partially suppressed bypass of dauer larval arrest in \( age-1(mg109) \); \( mg227 \) animals, indicating that \( mg227 \) may act upstream of \( akt-1 \). However, development in \( age-1(mg109);mg227 \) animals was not affected by \( akt-2 \), \( sgk-1 \) or \( pdk-1 \) RNAi (Figure 3).

age-1(mg109) dauer arrest suppressors differentially affect lifespan

Although all 5 of these alleles could suppress dauer arrest in \( age-1(mg109) \) animals, there were significant differences in their effects on the \( age-1(mg109) \) adult longevity and stress resistance phenotypes. Only the \( daf-16 \) alleles, \( mg242 \) and \( mg255 \), fully suppressed \( age-1(mg109) \) adult longevity (Log-Rank test, \( P \leq 0.0001 \) vs. \( age-1(mg109) \), Table 2; Figure 4A). In addition, \( age-1(mg109) \) animals carrying either the \( mg242 \) or \( mg255 \) alleles also lived significantly shorter than wildtype animals, consistent with previous reports of shortened lifespan by \( daf-16 \) mutations [16,31]. The \( daf-16(mg242) \) allele, which disrupts both the \( daf-16a \) and \( daf-16b \) isoforms, caused a slightly greater lifespan reduction in \( age-1(mg109) \) animals, than the \( daf-16a \)-specific \( mg255 \) allele (Log-Rank test \( P \leq 0.0001 \) \( mg242 \) vs. \( mg255 \)). However, this difference was minor, indicating that the \( daf-16a \) isoform provides the majority of the DAF-16 function required for longevity in \( age-1(mg109) \) animals.

Neither \( akt-1(mg247) \) nor \( pdk-1(mg261) \) suppressed \( age-1(mg109) \) adult longevity (Table 2; Figure 4B). Interestingly, the \( mg227 \) allele enhanced longevity of age-

| Genotype             | % of population, 20°C |
|----------------------|-----------------------|
|                      | L4/Adult              |
|                      | Dauer                 |
| wildtype             |                       |
| (-) control          | 206 126 131           |
| akt-1(RNAi)          | 131 102 286           |
| akt-2(RNAi)          |                       |
| act-1 + 2 (RNAi)     |                       |
| pdk-1(RNAi)          |                       |
| age-1(mg109)         |                       |
| (-) control          | 249                   |
| akt-1(RNAi)          | 740 129               |
| age-1(mg109); akt-1(mg247) |                   |
| (-) control          | 1493 97               |
| pdk-1(RNAi)          |                       |

Figure 2
RNAi against components of the insulin signaling pathway reverts \( age-1(mg109) \) dauer arrest suppression by \( akt-1(mg247) \) and \( pdk-1(mg261) \) mutant alleles. Animals were fed bacteria containing either control vector, L4440 or vectors producing gene-specific dsRNA. Numbers on the side of the bars represent the total number of animals examined. The bars show the percentage of animals that developed into L4 larvae or adults (grey) and those that arrested as dauer (black). RNAi experiments were diluted with control L4440 RNAi to match the mixed \( akt-1 \) and \( akt-2 \) RNAi dose. \( akt-1 \), \( akt-2 \) and \( pdk-1 \) RNAi experiments were conducted on wildtype, \( age-1(mg109) \), \( age-1(mg109);akt-1(mg247) \) and \( age-1(mg109);pdk-1(mg261) \) animals.
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RNAi epistasis analysis of age-1(mg109) suppressors reveals interdependence of AKT-1 and PDK-1. To establish whether akt-1(mg247), pdk-1(mg261) and mg227 mutations suppressed dauer arrest through the AGE-1/PI3 kinase pathway we carried out a series of genetic epistasis experiments using RNAi. Bars show the percentage of animals that bypassed dauer arrest, developing into L4 larvae or adults (grey), and those that arrested as dauers (black). Numbers on the side of the bars represent the total number of animals examined. RNAi experiments were diluted with control L4440 RNAi to match the mixed akt-1 and akt-2 RNAi dose. (A) age-1(mg109);akt-1(mg247) animals arrested as dauer larvae on pdk-1 RNAi. Similarly, age-1(mg109);pdk-1(mg261) animals arrested as dauer larvae on akt RNAi. (B) Epistasis analysis on age-1(mg109);mg227 animals suggests mg227 requires akt-1 but not akt-2, sgk-1 or pdk-1 to bypass dauer larval arrest.

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Dauer arrest suppressors differentially affect stress resistance and FIRE response

Reductions in AGE-1/PI3K signaling are associated with increased resistance to oxidative and thermal stresses [22,23]. We therefore examined the effects of dauer arrest suppressors on survival of age-1(mg109) animals under conditions of oxidative and thermal stress. Both daf-16 alleles, mg242 and mg255, completely suppressed oxidative stress resistance of age-1(mg109) animals tested by treatment with 10 mM paraquat, an intracellular free radical generator (Figure 5a). However, neither akt-1(mg247), pdk-1(mg261) or mg227 suppressed oxidative stress resistance of age-1(mg109) animals to paraquat. Double mutant age-1(mg109) animals carrying any of these suppressor alleles appeared to have increased survival in the presence of 10 mM paraquat, compared to parental age-1(mg109) animals (Figure 5a). Similar results were observed for suppression of thermotolerance of age-1(mg109) adults, as tested by survival at the stressful temperature of 35°C (Figure 5b).

We also analyzed the intestinal response to fasting in these strains by monitoring the FIRE response (Fasting-Induced Redistribution of Esterase activity) [32]. After a 6-hour fast, intestinal esterase activity in wildtype adults redistributes from cytoplasmic to nuclear localization
In *C. elegans*, signaling by the DAF-2/insulin receptor, via AGE-1/PI3K, regulates developmental arrest at the dauer larval stage, as well as longevity and stress resistance in adult animals. Here, we report the characterization of new mutations identified through a genetic screen designed to identify components of the AGE-1/PI3K signaling pathway. This screen identified 2 mutations inactivating one or two isoforms of *daf-16*, which encodes the FOXO transcription factor that is the ultimate output of AGE-1/PI3K signaling. Both of the *daf-16* alleles suppressed all the *age-1(mg109)* dauer and adult phenotypes examined, consistent with the conclusion that *daf-16* activity is necessary for both the larval and adult phenotypes of *age-1* mutant animals. Additionally, we identified 3 mutations that uncoupled the larval and adult phenotypes of AGE-1/PI3K signaling. These mutations appear to activate the AKT/PKB effector pathway of AGE-1/PI3K signaling, as evidenced by the dependence of these 3 alleles on wildtype *akt-1* and/or *pdk-1* activity. Previous reports also identified activating mutations in *akt-1* and *pdk-1*, yet ours is the first report to probe in depth the effects of this class of mutations on adult AGE-1/PI3K functions [11, 12]. In addition, we report the phenotypic characterization of a new mutation, *mg227*, which has the unusual phenotype of enhancing adult longevity in *age-1(mg109)* animals in addition to suppressing larval dauer arrest.

In this work, we identified an activating mutation in the AKT-1 linker region between the PH and kinase domains. Interestingly, the previously identified *mg144* allele also contained a linker region mutation (Ala183Thr) [12]. The linker region in mammalian AKT/PKB proteins is considerably shorter than in the *C. elegans* protein, so it is not clear whether linker region function is conserved in the *C. elegans* and human proteins or whether the *C. elegans* linker region possesses additional functions. The structure of the PH domain from human AKT has been determined [33]. This AKT fragment contained part of the linker region which appeared to face away from the residues that mediate phospholipid binding and probably does not play a significant role in determining lipid binding specificity. Instead, linker region mutations have been postulated to relax autoinhibition of the kinase and PH domains in the inactive AKT protein [12]. Interestingly, our epistasis analysis showed that the gain-of-function phenotype of *akt-1(mg247)* required the presence of wild type *pdk-1*. This suggests that the mutant protein encoded by *akt-1(mg247)* may be more easily phosphorylated by PDK-1 in the absence of AGE-1/PI3K-generated phospholipids.

We also identified an activating allele of *pdk-1*, encoding the kinase that phosphorlates and activates phospholipid-bound AKT/PKB. The *mg261* mutant contains mutations of both a nonconserved residue in the linker region

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Table 2: Adult lifespan of *age-1(mg109)* dauer arrest suppressors.

| Genotype                      | Adult Lifespan | Median | Maximum | SE  | n   | Log-Rank |
|-------------------------------|----------------|--------|---------|-----|-----|----------|
| Wildtype                      | 14.9 ± 0.33    | 255    | -       | -   | -   | -        |
| *age-1(mg109)*                | 21 ± 0.66     | 142    | -       | -   | -   | -        |
| *daf-16(mg242)*/age-1(mg109)* | 9.0 ± 0.11    | 225    | <0.001  | -   | -   | -        |
| *daf-16(mg255)*/age-1(mg109)* | 9.8 ± 0.11    | 283    | <0.001  | -   | -   | -        |
| *age-1(mg109)/akt-1(mg247)*   | 25.5 ± 0.42   | 205    | <0.0001 | -   | -   | -        |
| *age-1(mg109)/pdk-1(mg261)*   | 20.2 ± 0.69   | 166    | <0.6941 | -   | -   | -        |
| *age-1(mg109)/mg227*          | 30.5 ± 0.38   | 279    | <0.0001 | -   | -   | -        |

a Lifespan assays performed at 25°C.

b These are homozygote progeny of *age-1(mg109)* homozygous hermaphrodites that have been maternally rescued for dauer arrest [27].

*Although the akt-1(mg247) allele increased mean lifespan compared to *age-1(mg109)* control animals, the median lifespan and maximum lifespan was not significantly affected (age-1(mg109) ± b, 25 days median, 32 days maximum; age-1(mg109)/akt-1(mg247) ±, 24 days median, 33 days maximum, Figure 4)

d Wilcoxon P value = 0.2370.

e Lifespan assays performed at 20°C.

f P vs *age-1(mg109) ± b, SE, Standard error.

n indicates the number of animals scored in at least three independent trials.

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(Figure 6A, B). This response is altered in *age-1(mg109)* adults, which continue to exhibit the cytoplasmic pattern of esterase activity after a 6-hour fast (Figure 6C). Both the *daf-16(mg242)* and *daf-16(mg255)* mutations suppressed the altered FIRE response in *age-1(mg109)* animals, while *akt-1(mg247)*, *pdk-1(mg261)* and *mg227* had no effect on the altered FIRE response of *age-1(mg109)* animals (Figure 6D–F).

Finally, we investigated the effects of these mutations on DAF-16::GFP localization. In wildtype animals, 35°C heat stress could induce the nuclear localization DAF-16::GFP translational fusion (Figure 7A, B). In unstressed wildtype animals DAF-16::GFP was detected in both the nucleus and cytoplasm. Defects in the insulin signaling pathway should cause nuclear accumulation of DAF-16::GFP as previously reported [13-15]. We observed that DAF-16::GFP fluorescence was predominantly nuclear in *age-1(mg109)* adults containing the *daf-16(mg255)* mutation (Figure 7C). In contrast, DAF-16::GFP showed both nuclear and some cytoplasmic localization in *age-1(mg109)* animals carrying the *akt-1(mg247)*, *pdk-1(mg261)* or *mg227* alleles (Figure 7D–F). Thus we conclude that the mutations activating AKT signaling may partially rescue DAF-16::GFP nuclear localization by the *age-1(mg109)* allele.
and a conserved residue in the PH domain which may mediate binding to the phospholipid products of AGE-1/PI3K [34,35]. A previously identified activated \textit{pdk-1} allele, \textit{mg142}, corresponded to an Ala303Val substitution in the PDK-1 kinase domain and was shown to increase the basal level of PDK-1 kinase activity [11]. The regulation of PDK-1 signaling is complex and depends on the downstream target in question [36]. In \textit{C. elegans}, the major targets of PDK-1 are believed to be AKT-1, -2 and SGK-1 [8,11,12]. One possibility is that the \textit{mg261} mutation in the PH domain could alter the specificity of PDK-1 phospholipid binding, enhancing PDK-1 activity in the absence of AGE-1/PI3K phospholipid products. However, PDK-1 is active in \textit{age-1(mg109)} as evidenced by \textit{age-1(mg109);akt-1(mg247)} mutants. Therefore it is not entirely clear how the \textit{pdk-1(mg261)} allele is activating AKT-1.

daf-16 activity is required for both the larval dauer arrest and adult longevity and stress resistance phenotypes that result from genetic deficiencies in the \textit{age-1} pathway [21,27]. Consistent with previous findings, the \textit{daf-16} alleles described here suppressed both larval and adult phenotypes of \textit{age-1(mg109)} animals. In contrast, the \textit{akt-1(mg247)}, \textit{pdk-1(mg261)} and \textit{mg227} alleles did not affect the \textit{age-1(mg109)} adult phenotypes. Thus, the larval and adult outputs of AGE-1/PI3K signaling are differentially affected by \textit{akt-1(mg247)}, \textit{pdk-1(mg261)} and \textit{mg227} alleles. It was previously shown that \textit{pdk-1} activity is necessary for wildtype adult lifespan, demonstrating that PDK-1 mediates the adult outputs of AGE-1/PI3K signaling [11]. Thus, we propose that the differential suppression by \textit{pdk-1(mg261)} of the \textit{age-1(mg109)} larval versus adult phenotypes results from insufficient activation of PDK-1 activity for adult, but not larval, phenotypes.

A similar explanation may account for the inability of \textit{akt-1(mg247)} to suppress adult phenotypes of \textit{age-1(mg109)} animals. However, a previous report suggested that SGK-1, and not AKT-1 or -2, may be the primary mediator of DAF-2 and AGE-1 signaling in adult animals [8]. Thus, it is possible that \textit{akt-1} itself does not affect adult-specific functions, but only the dauer-specific functions of insulin signaling. However, we have observed that the DAF-16::GFP reporter is distributed in the cytoplasm in adult \textit{age-1(mg109); akt-1(mg247)} animals, suggesting that \textit{mg247} at least partially restores wildtype regulation of DAF-16 in adult animals. Thus, there may be additional...
pathways that converge with AGE-1/PI3K signaling to regulate DAF-16 activity and, ultimately, lifespan in adult animals.

One possible explanation for the different effects of these mutations on the larval versus adult phenotypes of age-1(mg109) is that they partially restore insulin pathway signaling and negative regulation of DAF-16. According to this model, wildtype levels of insulin pathway signaling may strongly inhibit DAF-16 and promote reproductive development and adult lifespan. Whereas, low insulin signaling promotes both dauer larval arrest and adult longevity [19-21]. The mutations activating AKT signaling appear to promote intermediate levels of insulin pathway signaling, allowing reproductive development, but prolonging adult lifespan. This phenotypic gradient could reflect a gradient of DAF-16 activity in these strains, in which the highest levels of DAF-16 activity causes dauer larval arrest (Figure 8). Thus the mutations activating AKT signaling may allow an intermediate level of DAF-16 activity, which is insufficient to promote dauer larval arrest, but does promote adult longevity. This is similar to a DAF-16 gradient model that has been previously proposed [8].

**Conclusion**

In this study we identified mutations capable of activating signaling in the absence of AGE-1/PI3K. We describe five mutations including two mutations inactivating one or two isoforms daf-16 and two new hypermorphic alleles of akt-1 and pdk-1. The daf-16 alleles described in this work suppress age-1(mg109) dauer and adult phenotypes examined consistent with previous reports. The akt-1 and pdk-1 alleles uncouple the dauer arrest, adult longevity and stress resistance phenotypes of age-1(mg109) mutants. Furthermore, genetic epistasis analysis suggests these hypermorphic alleles are interdependent. These findings suggest larval and adult phenotypes of insulin signaling are fully separable in these mutants.

**Methods**

**Strains and general methods**

The C. elegans strains used in this work were N2 Bristol (wildtype), sqt-1(sc13) age-1(mg109)/MM1; and CB4856 Hawaiian. Strains were provided by the Caenorhabditis
Genetics Center at the University of Minnesota. All strains were cultivated at 15°C on NGM agar plates seeded with the *E. coli* strain OP50 following standard protocols [37].

**Suppressor isolation and mapping**

Suppressors were isolated as previously described on a parent strain carrying the *sqt-1(sc13) age-1(mg109)* mutation balanced with *mnC1* and F2 progeny were screened for suppression of the *age-1(mg109)* constitutive dauer arrest phenotype [12]. From a screen of 20,000 haploid genomes, 40 suppressors were identified of which 5 were crossed with Hawaiian strain males (CB4856). DNA from these suppressor recombinants were collected and analyzed using SNP-snip mapping for each chromosome [29]. For candidate gene sequencing, genomic DNA from the mutant strains (*sqt-1(sc13) age-1(mg109); m/m*) was PCR amplified with exon-specific primers and sequenced to identify molecular changes. *mg227* SNP-snip mapping showed strong linkage to the right arm of chromosome X between 10.5 – 13.5 m. u. Candidate genes were PCR amplified and sequenced. Genes sequenced included known dauer forming genes, *daf-12*, (F11A1.3, X2.39), *lin-14* (T25C12.1, X4.28), *bra-1* (F54B11.6, X13.84), *akt-2* (F28H6.1, X16.56) and *sgk-1* (W10G6.2, X2.70). Furthermore, a cosmid carrying the *daf-12* gene did not rescue the *mg227* daf-C suppression phenotype in *age-1(mg109);mg227* animals (5 transgenic lines examined, 94% of animals bypassed dauer arrest, 6% arrested as dead eggs).

**RNA interference**

A 500 bp region spanning the largest exon in both *akt-2* and *pdk-1* was amplified from N2 DNA and cloned into the RNai feeding vector L4440 [38]. *akt-1 (RNAi) and daf-16 (RNAi)* was obtained from the RNAi feeding library [30]. LB broth supplemented with 100 μg/mL of ampicillin was inoculated with RNAi clones and grown overnight at 37°C without shaking. RNAi experiments were typically carried out on 6-well plate NGM media supplemented with 50 μg/mL ampicillin and 1 mM IPTG. Overnight bacterial cultures were seeded on duplicate wells and allowed to grow at room temperature for a further 12 – 16 hr. *G1* and *G2* progeny of fertile hermaphrodites were collected in a 5-hour egg lay on RNAi bacteria. The *G2* progeny were allowed to grow at 20°C for 3 days after which any phenotypes were scored. Plates were re-examined 24 hr later for changes in phenotypes if any. Each experiment was done in triplicate.

**Figure 6**

**Suppression of altered FIRE response in *age-1(mg109)* animals.** Distribution of intestinal esterase activity in (A) well-fed or (B) fasted *age-1(+)* young adult animals exhibited the fasting-induced redistribution (arrows) of esterase from the intestinal cytoplasm (A) to a nuclear pattern (B). (C) *age-1(mg109)* adults were resistant to the FIRE response. Altered FIRE response of *age-1(mg109)* animals was suppressed by (D) *daf-16(mg242)* and (E) *daf-16(mg255)* alleles, but not by (F) *akt-1(mg247)*, (G) *pdk-1(mg261)* or (H) *mg227* mutations. Scale bars, 50 μm.
**Figure 7**

*age-1(mg109) suppressor mutants differentially redistributes DAF-16:GFP localization.* Intestinal DAF-16:GFP localization redistributed to the nucleus of wildtype animals after a period of heat stress (35°C) (A, B). DAF-16:GFP localization is primarily in the nucleus of daf-16(mg255);age-1(mg109) animals (C) reflecting localization of DAF-16 in a *age-1(mg109)* animal. In contrast, DAF-16:GFP localization was seen in both the cytoplasm and nucleus of age-1(mg109);akt-1(mg247) animals (D), age-1(mg109);pdk-1(mg261) animals (E) and age-1(mg109);mg227 (F). Scale bar 50 μm.
Phenotypic assays
To assay dauer larval arrest, gravid adult hermaphrodites were allowed to lay eggs at room temperature for 4 hours to produce a synchronized population. Animals were then shifted to 25°C or 20°C and left to develop for a further 72 or 96 hours after which the number of dauer and adults (sterile or fertile) was counted. Sterility was scored as the absence of embryos in the uterus in young adults (day 1–2).

For lifespan analysis, larvae were raised at 15°C until the first day of adulthood (Day 0) prior to being shifted to 25°C or 20°C on standard NGM plates supplemented with FUDR (50 μg/mL). Survival analyses were always performed in the presence of FUDR, since some of these strains have an egg-laying defect that causes eggs to accumulate within the uterus and hatch, thereupon killing the parent hermaphrodite. FUDR treatment helps to remedy this problem as it is an inhibitor of DNA synthesis and prevents embryonic development. Animals were scored every two days for death by failure to respond to gentle prodding on the head and tail. All aging experiments were repeated in at least three separate trials and cumulative results were analyzed. Lifespan statistical analysis was performed using the JMP software package (version 5.1).

For paraquat and thermotolerance survival, young adults were obtained as for lifespan assays. For paraquat treatments, day 0 adults were shifted to standard NGM plates supplemented with 10 mM paraquat at 20°C. Survival of
animals was scored at 2 or 3 day intervals by failure to respond to gentle prodding. For thermotolerance assays, synchronized young gravid adults were placed on bacteria-free NGM agar with 100 μg/mL ampicillin and incubating at 25°C for 6 hours. Fasted adults were fixed with methanol (-20°C) for 10 minutes and incubated overnight at room temperature in staining solution described previously [32]. Stained specimens were mounted on 2% agarose pads and viewed on a Nikon E800 microscope with Nomarski optics. Images were collected using a Hamamatsu ORCA-ER CCD camera using OpenLab software (Improvision, Lexington, MA).

**DAF-16:GFP localization**

A gfpdaf-16 cDNA fusion was constructed using an intestinal gly-19 promoter as previously described [32]. In brief, the DAF-16:GFP plasmid was constructed by inserting a gfp coding sequence in-frame with the daf-16a full-length cDNA. The intestinal promoter was cloned upstream of gfpdaf-16 using unique SphI and KpnI sites. Plasmid were injected into adult hermaphrodites at a final concentration of 25–100 μg/mL [32].

**Authors’ contributions**

MSG carried out genetic mapping of daf-16(mg242), daf-16(mg255), abt-1(mg247) and mg227, lifespan analysis, RNAi experiments, oxidative stress and heat stress assays on all mutants, and wrote the manuscript. WBI performed the FIRE assays and DAF-16:GFP localization analysis. KH carried out mapping of pdk-1(mg261). CAW carried out the genetic screen as a postdoctoral fellow in Gary Ruvkun’s laboratory and wrote the manuscript.

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