Pesticide-induced resurgence in brown planthopper is mediated by

action on a suite of genes that promote juvenile hormone biosynthesis

and female fecundity

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Abstract

Pesticide-induced resurgence, increases in pest insect populations following pesticide application, is a serious threat to the sustainable control of many highly damaging crop pests. Resurgence can result from pesticide-enhanced pest reproduction, however, the molecular mechanisms mediating this process remain unresolved. Here we show that brown planthopper (BPH) resurgence following emamectin benzoate (EB) exposure results from the coordinated action of a diverse suite of actors that regulate juvenile hormone (JH) levels, resulting in increased JH titer in adult females and enhanced fecundity. Following reports of BPH resurgence in rice crops when this species is exposed to EB, we demonstrate that EB treatment results in profound changes in female BPH fitness including enhanced ovarian development and elevated egg production. This enhanced reproductive fitness results from the EB-mediated upregulation of key genes involved in the regulation of JH, including *JHAMT, Met and Kr-h1* and the downregulation of allatostatin (*AstA*) and allatostatin receptor (*AstAR*) expression. The remodulation of gene expression following EB exposure is dependent on the action of this insecticide on its molecular target the glutamate-gated chloride channel (GluCl) receptor. Collectively, these results provide mechanistic insights into the regulation of negative pesticide-induced responses in insects and reveal the key actors involved in the JH-signaling pathway that underpin pesticide resurgence.

Keywords: resurgence; reproduction; pesticide; brown planthopper

Significance

Pesticides remain a key means of controlling many of the world's insect pests, however, in some cases, pesticide applications can result in resurgence of pest populations due to pesticide-induced increases in fecundity. In the current study we show that pesticide resurgence in the brown planthopper (BPH) following exposure to the insecticide emamectin benzoate results from the transcriptional reprogramming of a diverse suite of positive and negative regulators of juvenile hormone (JH), a critical regulator of insect development and reproduction. This in turn leads to profound increases in female BPH reproductive fitness and enhanced fecundity. Our findings unravel the molecular mechanisms mediating pesticide-induced pest resurgence and inform the development of novel strategies to control highly damaging crop pests.

Introduction

Chemical pesticides remain the primary means of controlling many of the world's most damaging arthropod crop pests ([1], [2]). However, pesticide applications can result in pest resurgence, increases in pest insect populations that exceed natural, untreated population sizes, following an initial reduction of the pest population ([3-5]). Two mechanisms have been implicated in pest resurgence - the loss of beneficial insects including natural enemies and pesticide-enhanced pest reproduction ([3]). In the case of the latter, several pesticides, such as the insecticides triazophos, deltamethrin and the fungicide jinggangmycin, have been reported to stimulate pest reproduction ([6-8]). Pesticide-enhanced pest reproduction has been linked to changes in physiology and biochemistry of pest organisms after exposure to pesticides ([3], [4]). However, the molecular mechanisms underlying enhanced reproduction associated with pest resurgence remain poorly resolved.

The brown planthopper (BPH), *Nilaparvata lugens* (Stål), is a notorious pest of rice crops throughout Asia causing annual losses of ~300 million dollars across major rice producing countries ([2], [3]). BPH inhibits the growth of rice plants by feeding, and also transmits highly damaging plant viruses including rice grassy stunt virus and rice ragged stunt virus ([9]). Currently, chemical insecticides play an indispensable role in the control of BPH due to their efficiency, rapid effect, and low cost. However, due to the widespread and intensive use of chemical insecticides, BPH has developed resistance to the majority of compounds used for control ([2], [10]).

Emamectin benzoate (EB) and abamectin are avermectin pesticides, and act as

allosteric modulators of insect glutamate gated chloride channels (GluCls), inhibiting

muscle contractions that lead to the cessation of insect feeding and subsequent death

(11). These insecticides exhibit particularly strong activity against Lepidoptera such as

the rice leaffolder, *Cnaphalocrocis medinalis* Guénee, an important foliage-feeding

insect which attacks rice during the vegetative stage (12). Both BPH and the rice

leaffolder are migratory pests with overlapping migratory paths, however, their

occurrence period in the field differs by approximately one month, with leaffolders

appearing earlier than BPH. Therefore, the use of EB to control rice leaffolder has the

potential to impact BPH arriving later, via exposure to sublethal concentrations of this

compound. In this regard, we have observed that when farmers use EB and abamectin

to control leaffolders on rice crops in China, BPH outbreaks frequently occur in the

same field. While sublethal doses of certain pesticides have been shown to enhance

fecundity in BPH, including the insecticides triazophos and deltamethrin (6, 7, 13, 14)

and the fungicides carbendazim and jinggangmycin (8), whether avermectins trigger

resurgence in BPH via insecticide-enhanced reproduction remains unclear.

Reproduction in insects is influenced by external factors such as light (15),

temperature (16), humidity (17) and nutrition (18), and endogenous factors such as the

juvenile hormone (JH) (19), ecdysone (20), insulin (21) and TOR (22) pathways (23,

24). Of these, JH has been particularly implicated in insecticide-induced enhanced

fecundity, with triazophos and deltamethrin treatments leading to increased circulating

JH III titers in BPH females (3). JH is synthesized and secreted by the corpora allata

in insects (25), and can promote reproduction by regulating the synthesis and secretion

of vitellogenin (vg) in the female fat body, and stimulating the absorption of vg by the

developing oocyte (19). However, the regulation of JH is complex (19) and the key

actors involved in JH-mediated pesticide-enhanced reproduction remains an open

question. Previous finding reported that GluCl receptor is involved in JH biosynthesis

in cockroach (26-28). However, we know little whether insecticide which target on the

GluCl can stimulate reproduction in insect pest and how GluCl receptor regulates JH

production is still unknown.

In this study, we used a diverse range of approaches to investigate the impact of

sublethal doses of avermectins on BPH fecundity, and unravel the molecular

mechanisms mediating enhanced reproduction following exposure to this insecticide

class. We show that avermectin exposure results in profound changes in the

expression of a key suite of genes that in combination regulate JH, resulting in

increased JH titer in adult females, which promotes fecundity. Interestingly, we found

that GluCl channel modulators can inhibit Ast/AstAR expression and then promotes JH

increasing.

RESULTS

GluCl allosteric modulators (emamectin benzoate and abamectin) stimulate

fecundity of female *N. lugens*

To investigate whether GluCl modulators affect fecundity in BPH, we first

determined the sub-lethal and median lethal doses of emamectin benzoate (EB) to

4th instar nymphs, newly emerged males and females of BPH (Table S1). For this we

employed two different bioassay methods, the rice seedling dip bioassay method and

topical application bioassay method (2, 29), in order to assess both the systemic and

contact toxicity of these insecticides (Table S1). We then systemically treated 4th

instar nymphs of BPH with the estimated LC_{15} and LC_{50} concentrations of EB and

examined the fecundity of BPH after these individuals mated with treated or

untreated individuals. We use the term "t" to represent individuals treated with EB

and "ck" to indicate control individuals that were treated with insecticide diluent minus

insecticide. After treatment with the LC_{15} and LC_{50} concentrations of EB, the number

of eggs laid per female of BPH in ♀t × ♂t crosses increased by 1.48 and 1.40 times

compared with control ♀ck × ♂ck crosses (Figure 1A); the number of eggs laid per

female of BPH in ♀t × ♂ck crosses increased by 1.53 and 2.07 times compared with

control crosses (Figure 1B); However, the number of eggs laid by per female of BPH

in ♀ck × ♂t crosses did not increase significantly compared to control ♀ck × ♂ck

crosses (Figure 1C).

Table S1. Determination of the toxicity of emamectin benzoate against BPH in systemic and topical application bioassays.

Treatment method	Developmental stages	Slope ± SE	LC_{15} (95%F.L.) (mg/L) or LD_{15} (95%F.L.) (pg/insect)	LC_{50} (95%F.L.) (mg/L) or LD_{50} (95%F.L.) (pg/insect)	χ^2(df)	*P value*
Systemic route	4th nymph	2.56 ± 0.487	0.87 (0.39-1.32)	2.21 (1.53-2.82)	2.68 (4)	0.61
	Male	2.42 ± 0.35	0.80 (0.40-1.20)	2.13 (1.48-2.78)	2.12 (4)	0.72
	Female	2.47 ± 0.41	2.33 (1.09-3.56)	6.12 (4.18-8.00)	3.56 (4)	0.50
Contact route	4th nymph	2.52 ± 0.35	15.54 (9.46-21.42)	39.90 (30.66-49.56)	2.60 (4)	0.63
	Male	2.77 ± 0.57	6.72 (2.52-10.92)	15.96 (9.66-21.84)	1.23 (3)	0.75

Female	3.31 ± 0.76	18.48 (6.72-28.14)	37.80 (23.10-50.40)	0.79 (3)	0.85

131  Exposure of 4th instar nymphs with the LC$_{15}$ and LC$_{50}$ concentrations of EB in

132 contact bioassays also significantly stimulated fecundity. After treatment with the LC$_{15}$

133 and LC$_{50}$ concentrations of EB, the number of eggs laid by per female of BPH in ♀t ×

134 ♂t crosses increased by 1.18 and 1.26 times compared with the control (♀ck × ♂ck)

135 (Figure 1D); The number of eggs laid per female of BPH in ♀t × ♂ck crosses

136 increased by 1.27 and 1.56 times compared with the control crosses (Figure 1E);

137 However, there was no significant difference in egg-laying number between ♀ck × ♂t

138 crosses and controls (♀ck × ♂ck) (Figure 1F). These results reveal that EB can

139 stimulate the fecundity of females following both the systemic and contact routes of

140 exposure.

141 **Figure 1**

143 **Figure 1.** Fecundity of BPH following exposure to sub-lethal (LC$_{15}$) and median lethal (LC$_{50}$)

144 concentrations of emamectin benzoate following system application bioassays (A: ♀t × ♂t;

145 B: ♀t × ♂ck; C: ♀ck × ♂t) and topical application bioassays (D: ♀t × ♂t; E: ♀t × ♂ck; F:

♀ck × ♂t), respectively. The letter "t" represents treatment with insecticide, while "ck"

indicates controls that was not treated with insecticide. All data are presented as the mean ±

s.e.m. Different lower-case letters above the bars indicate significant differences (One-way

ANOVA with Tukey's Multiple Range Test, $p < 0.05$).

We next examined whether EB treatment of adult BPH also stimulates

reproduction. Indeed, treating newly emerged adults with the LC_{15} and LC_{50}

concentrations of EB significantly stimulated the number of eggs laid per female

(Figure 1-figure supplement 1A). Furthermore, sub-lethal exposure of 4th instar BPH

nymphs to another GluCl allosteric modulator, abamectin (LC_{15} and LC_{50}

concentrations) was also found to significantly enhance reproduction (Figure 1-figure

supplement 1B).

Figure 1-figure supplement 1

Figure 1-figure supplement 1. (A) Fecundity of BPH when newly emerged adults were

treated with sub-lethal (LD_{15}) and median lethal (LD_{50}) concentrations of emamectin benzoate

via topical application. (B) Fecundity of BPH when 4th instar nymphs were treated with sub-

lethal (LC_{15}) and median lethal (LC_{50}) concentrations of abamectin via systemic exposure. All

data are presented as the mean ± s.e.m. Different lower-case letters above the bars indicate

significant differences (One-way ANOVA with Tukey's Multiple Range Test, $p < 0.05$).

To examine if EB also stimulates egg-laying in other insect species we

conducted bioassays on the small brown planthopper, *Laodelphax striatellus*, the

white backed planthopper, *Sogatella furcifera* and fruit flies *Drosophila melanogaster*.

In contrast to our findings on BPH, we found that sub-lethal doses (LC_{15} and LC_{50}) of

EB inhibits fecundity of female *L. striatellus*, (Figure1-figure supplement 2A-C) and

has no impact on the fecundity of *S. furcifera*, (Figure1-figure supplement 2D-F). In

addition, we found that sublethal doses (LC_{15} or LC_{50}) of EB also inhibit fecundity in

D. melanogaster (Figure1-figure supplement 2G and H). These results indicate that

the stimulation of reproduction by EB in BPH is species-specific and does not extend

to even related insect species.

Figure 1-figure supplement 2

Figure 1-figure supplement 2. Fecundity of small brown planthopper, *Laodelphax striatellus*,

(A-C) white backed planthopper, *Sogatella furcifera* (D-F) and fruit fly, *Drosophila*

melanogaster (G and H) when larvae and newly emerged adults were treated with sub-lethal

concentrations of emamectin benzoate. All data are presented as the mean ± s.e.m. Different

lower-case letters above the bars indicate significant differences (One-way ANOVA with

Tukey's Multiple Range Test, $p < 0.05$).

The impact of EB treatment on BPH reproductive fitness

To better understand the effects of EB on the reproductive fitness of BPH, the

preoviposition period, emergence rate, female ratio, female longevity and female

weight were evaluated following exposure using the systemic bioassay. The

preoviposition period of females treated with the LC_{50} of EB decreased significantly

compared with the control (Figure 1-figure supplement 3A). In contrast no significant

effects of EB on emergence rate and female ratio were observed (Figure 1-figure

supplement 3B and C). Female survival analysis showed that exposure of 4th instar

nymphs to the LC_{50} of EB has no impact on female longevity (Figure 1-figure

supplement 3D). Interestingly, brachypterism (long-wing) female ratio and female

weight were significantly increased after EB exposure (Figure 1-figure supplement 3E

and F).

Figure 1-figure supplement 3

Figure 1-figure supplement 3. The impact of emamectin benzoate on the reproductive

fitness of BPH. Fourth instar nymphs were treated with the LC_{50} concentration of emamectin

benzoate in systemic bioassays. All data are presented as the mean ± s.e.m. Different lower-

case letters above the bars indicate significant differences (Student's *t* test, *p* < 0.05).

EB promotes ovarian development in BPH

To investigate the cause of increased female weight following EB exposure we

examined if EB influences ovarian development in BPH. We dissected and compared

the ovaries of females treated with the LC_{50} of EB at 1, 3, 5 and 7 days after eclosion

(DAE) with control females. At 3, 5 and 7 DAE, the number of detained eggs of BPH

in the EB treated group were significantly higher than that of controls (Figure 2A and

B). We also explored whether EB treatment could enhance or impair oogenesis in

BPH. However, dissection of various developmental stages revealed that emamectin

benzoate treatment has no significant effects on oogenesis (Figure 2C and D).

Figure 2

Figure 2. The impact of emamectin benzoate on ovarian development in BPH. Fourth instar

nymphs were treated with the LC_{50} concentration of emamectin benzoate in systemic bioassays.

(A) Ovarian development in EB treated BPH at 1, 3, 5 and 7 days after eclosion (DAE)

compared to untreated controls. Scale bar = 1,000 μm. (B) Number of detained eggs in the

ovaries of EB treated BPH females measured at 1, 3, 5 and 7 DAE compared to controls. All

data are presented as the mean ± s.e.m. Asterisks indicate values significantly different from

the control using student *t* test (ns, no significant; *$p < 0.05$ and **$p < 0.01$). (C) Different

developmental stages of BPH eggs. (D) No impairment of emamectin benzoate on oogenesis

of BPH. Scale bar = 100 μm.

EB exposure enhances the abundances of storage macromolecules and

circulating sugars in BPH

Nutritional status is an important indicator of reproductive fitness. Thus, to

investigate whether EB affects intermediary metabolism and energy storage of BPH,

glycogen, triacylglyceride (TAG), total protein content, cholesterol and four circulating

carbohydrates were quantified in 4th instar BPH nymphs following exposure to the

LC_{50} of EB.

We found that EB exposure has no impact on glycogen levels (Figure 2-figure

supplement 1A). The amounts of TAG in EB-treated BPH were 27% higher (p < 0.05)

than those in controls, but only in BPH of the late fifth instar (5L) stage, with no

significant differences observed in subsequent developmental stages (Figure 2-figure

supplement 1B). The amount of total protein content in EB-treated BPH was higher

than the control groups in the case of all developmental stages from 5L nymph to

7DAE (Figure 2-figure supplement 1C). EB exposure also increased cholesterol

levels at 4 and 5 DAE (Figure 2-figure supplement 1D). Compared with the solvent

control, EB treatment caused significant increases ($p < 0.05$) in the levels of sucrose,

glucose, fructose, and trehalose (Figure 2-figure supplement 1E-H). Thus,

collectively, these data provide evidence that EB exposure leads to energy

mobilization and the metabolism of carbohydrates and lipids in BPH.

Figure 2-figure supplement 1

Figure 2-figure supplement 1. Amounts of Glycogen (A), TAG (B), total protein content (C),

cholesterol (D) and four circulating sugars including sucrose, glucose, fructose and trehalose

(E-H) after BPH exposure to EB. All data are presented as the mean ± s.e.m. The differences

between the EB-treated and solvent-treated BPH were analyzed using unpaired student *t*-test

(*, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$; ****, $p < 0.0001$).

EB stimulates egg-laying that is mediated by the JH signaling pathway

Given the important role of juvenile hormone (JH) in vitellogenesis and egg

development in insects (19, 25, 30-32), we asked whether EB-treatment could

influence the titer of JH in BPH. As measured by ELISA, the juvenile hormone titer of

BPH nymphs treated with the LC_{50} concentration of EB was significantly lower than

that of controls in systemic bioassays during the middle and late stages of the 4th

instar (Figure 3A). However, at 2, 3 and 4 DAE, the JH titer in the EB treated group

was significantly higher than that of the control (Figure 3A). Interestingly, the titer of

another important insect hormone, the steroid ecdysone, was not significantly

different between EB-treated BPH and solvent-treated controls (Figure 3-figure

supplement 1). To independently validate the results of ELISA assays, we employed

HPLC-MS/MS to measure JH titer in BPH following EB exposure (30, 33, 34). The

results showed that JH III titer significantly decreased after EB-treatment at the late

4th instar nymph stage (Figure 3A and B), but significantly increased at the third day

after eclosion (3 DAE) (Figure 3A and C). To further investigate the role of JH in EB-

enhanced fecundity in BPH, we treated BPH with methoprene and pyriproxyfen, JH

analogues or biologically active JH mimics respectively, to determine whether they

can stimulate fecundity in BPH. Both compounds significantly increased egg-laying in

BPH (Figure 3D). Takes together these results reveal that EB stimulates an increase

in JH titer that induces fecundity in BPH. Since we found that EB could induce JH

synthesis in the BPH, we asked whether EB could influence the expression of genes

that are involved in JH synthesis or degradation. For this we treated 4th instar

nymphs of BPH with the LC_{50} concentration of EB using systemic bioassays, and

then collected early (5E), middle (5M) and late (5L) stage of 5th instar nymph and 1-7

DAE female adults (DAE) for analysis. Quantitative PCR was then used to examine

the expression of key genes previously implicated in the regulation of JH (35, 36).

Farnesoic acid O-methyltransferase (FAMeT) (37), is known as an important

enzyme in the JH biosynthetic pathway, catalyzing methylation of farnesoic acid (FA)

to methyl farnesoate (MF) (36). We found that this gene was significantly upregulated

in 5M instar nymph to 4 DAE after EB- treatment (1.0-fold to 3.0-fold) (Figure 3E).

Another gene, juvenile hormone acid methyltransferase (JHAMT) (36, 38), which is

involved in the biosynthesis of JH, was also upregulated in EB-treated BPH at 1 to 5

DAE than controls (1.5-fold to 3.0-fold) (Figure 3F). Methoprene-tolerant (Met),

belongs the basic helix–loop–helix Per/Arnt/Sim (bHLH-PAS) family of transcription

factors and is the intracellular (nuclear) receptor of JH (39, 40). The levels of *met*

mRNA increased in EB-treated BPH at the 5M and 5L instar nymph and 1 to 5 DAE

stages compared to controls (1.7-fold to 2.9-fold) (Figure 3G). Krüppel homolog 1

(*Kr-h1*), a transcriptional target of JH signaling, is reported to be sensitive to levels of

JH and its expression levels are directly correlated with JH titers (35, 41, 42). We

found that *Kr-h1* was significantly upregulated in the adults of EB-treated BPH at the

5M, 5L nymph and 4 to 5 DAE stages (4.7-fold to 27.2-fold) (Figure 3H). Similarly, the

expression levels of vitellogenin (Vg), a key downstream component of JH signaling

triggering ovarian development in insects including BPH (43), was markedly

increased in females at 2–5 DAE by EB (1.7-fold to 5.5-fold) (Figure 3I). Juvenile

hormone esterase (JHE) is the primary JH-specific degradation enzyme that plays a

key role in regulating JH titers (44). Interestingly, we observed a significant

upregulation of JHE mRNA levels in the early and middle 5th instar nymph stage

followed by downregulation in 5L instar nymphs of EB-treated BPH (1.3-fold to 2.6-

fold) (Figure 3J). In combination these results reveal that EB has profound impacts

on the expression of key genes involved in the synthesis of JH or downstream

signaling pathway genes that might promote egg development.

To further understand whether these JH pathway-related genes were involved in

egg-laying behavior in BPH, we performed RNAi experiments to downregulate the

expression of *JHAMT, Met and Kr-h1* (Figure 3-figure supplement 2A-C). We found

that silencing of these three genes downregulated the titer of JH in BPH providing

functional evidence of their role in the regulation of JH (Figure 3-figure supplement

2D). Furthermore, silencing of *Kr-h1* inhibited *FAMeT* and *Vg* gene expression while

increasing *JHE* gene expression (Figure 3-figure supplement 2E-H). Importantly,

silencing of *JHAMT*, *Met* and *Kr-h1* gene in female BPH was also found to suppress

egg-laying (Figure 3K-M). However, this phenotype was rescued by EB treatment,

which restored egg-laying to normal levels in BPH injected with *JHAMT*, *Met* and *Kr-*

h1 dsRNA (Figure 3K-M). Together these results provide a mechanistic

understanding of how EB enhances fecundity in BPH by modulating the expression

of key genes involved in JH synthesis (Figure 3N).

Figure 3

Figure 3. EB induced reproduction in BPH is mediated by components of the JH signaling pathway. (A) The titer of JH III (as measured by ELISA assay) at different developmental stages of BPH when 4th instar nymphs were treated with the median lethal concentrations of EB. (B and C) The titer of JH III (as measured by HPLC-MS/MS) in BPH females at 4L and 3 DAE when treated with median lethal concentrations of EB. (D) Oviposition rate of BPH when 4th instar nymphs were treated with 4 ppm methoprene or 10 ppm

pyriproxyfen. (E-J) Expression of selected JH-related genes (*FAMeT*, *JHAMT*, *Met*, *Kr-h1*, *Vg*,

and *JHE*) in EB-treated BPH. (K) Egg production following silencing of *JHAMT* with or without

EB application. (L) Egg production following silencing of *met* with or without EB application. (M)

Egg production after silencing *Kr-h1* with or without EB application. (N) Schematic illustrating

the proposed impact of EB on the JH signaling pathway leading to enhanced reproduction. The

question mark indicates one or more potential additional signals. All data are presented as

means ± s.e.m. Student's *t* test was used to compare the two samples. One-way ANOVA with

Tukey's multiple comparisons test was used to compare more than two samples. ns, no

significant difference; Asterisks indicate values significantly different from the control (ns, no

significant; *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$ and ****$p < 0.0001$). Different lower-case letters

above the bars indicate significant differences ($p < 0.05$).

Figure 3-figure supplement 1

Figure 3-figure supplement 1. Ecdysone titer at different developmental stages of BPH when

4th instar nymphs were treated with median lethal concentrations of EB.

Figure 3-figure supplement 2

Figure 3-figure supplement 2. (A-C) Expression of *JHAMT* (A), *Met* (B) and *Kr-h1* (C)

following RNAi knockdown. (D) The titer of JHⅢ (as measured by HPLC-MS/MS) at 3 DAE

when female adults were injected with *dsJHAMT, dsMet,* and *dsKr-h1* at 1 DAE. (E-H)

Expression of *FAMeT*, *Vg*, *JHE* and *JHAMT* when *Kr-h1* was silenced in BPH. All data are

presented as means ± s.e.m. *$p < 0.05$; Mann–Whitney test.

EB induces JH biosynthesis through the peptidergic AstA/AstAR signaling pathway

The timing and level of JH biosynthesis can be precisely regulated by the neuropeptides, stimulatory allatotropins (ATs) and inhibitory allatostatins (Asts), in many insects (45-52). Insects can, in a species-specific manner, produce one type of AT and three types of Asts: FGL-amide Ast (AstA) (46, 53); myoinhibitory peptide (MIP or AstB) and PISCF Ast (AstC) (50, 54). In some species, there also exists two paralogue genes of AstCs which are named AstCC and AstCCC (55, 56). Interestingly, the allatostatic activity of these three types of Ast peptides varies between insect species so that in each species only one type of Ast (for example AstA) controls JH production (50, 51, 54, 57).

Analysis of the BPH neural transcriptome sequence data has revealed the presence of one AT, four types of Asts and four corresponding receptors, allatotropin receptor (A16, ATR), AstA receptor (A2, AstAR), AstB (MIP) receptor (A10, AstBR or MIPR) and AstC receptor (A1, AstCR) (58). We cloned the five neuropeptide genes (AT, AstA, AstB/MIP, AstCC and AstCCC) and confirmed the sequence obtained from transcriptome data (Figure 4-figure supplement 1) (58). Interestingly, we found that AstC is missing in the genome of BPH and only AstCC and AstCCC are present (Figure 4-figure supplement 1). Next, we also cloned their corresponding receptors (55) including ATR (A16), AstAR (A2), AstBR (A10) and AstCR (A1) which might be activated by AstCC and/or AstCCC (42, 55, 59). Sequence alignments and phylogenetic analysis are shown in Figure 4-figure supplement 2.

Quantitative PCR was then used to examine if EB treatment could influence the

expression of the genes encoding these neuropeptides and their receptors. Treating

BPH with the LC_{50} concentration of EB significantly increased the expression of *AT*,

ATR and *AstCCC* while resulting in the downregulation of *AstA, AstB/mip, AstCC*,

AstAR and *AstBR/mipr* at the adult stage (Figure 4A and B, Figure 4-figure

supplement 3A-G). Among these, *AstA* and *AstAR* were the most downregulated

genes after EB treatment (Figure 4A and B, Figure 4-figure supplement 3H) and thus

the *AstA/AstAR* signaling system was selected for subsequent functional analysis.

Silencing of *AstAR* in female BPH using RNAi (Figure 4C), resulted in an increased

number of eggs laid per female compared with *dsgfp*-injected controls (Figure 4D).

Interestingly, silencing *AstAR* also resulted in the upregulation of *FAMeT, JHAMT,*

Met and *Kr-h1* which are involved in the JH biosynthesis/signaling (Figure 4E-H).

However, *JHE* was not influenced by AstAR silencing (Figure 4I). We therefore

investigated whether silencing the *AstAR* gene could influence JH titer in BPH. A

significantly increased JH titer was observed in *AstAR* silenced BPH compared with

controls (Figure 4J). Thus, our data strongly suggest that AstA is a key inhibitor of JH

production in BPH.

Finally, we investigated whether injection of mature Ast and AT peptides could

influence the number of eggs laid and JH titer in BPH. We synthesized one AT, six

AstAs (AstA1 to AstA6), one AstCC and one AstCCC peptide according to their

determined sequences (Figure 4-figure supplement 1). Indeed, we found that

injection of AstA1 and AstA6 reduced the number of eggs laid per female in 48 h

(Figure 4K). We also observed that AstA1 injection after 16h and 48h decreased the

JH titer significantly (Figure 4L), and AT injection increased the JH titer after 2h but

levels returned to normal 4h after injection (Figure 4M). Collectively, these results

provide compelling evidence that EB induces reproduction in BPH through the

AstA/AstAR and JH signaling pathways (Figure 4N) and further supports the role of

AstA and AT in regulation of JH titer.

Figure 4

Figure 4. EB induced reproduction in BPH is mediated by the AstA/AstAR and JH signaling pathway.

(A and B) Expression of *AstA* and *AstAR* in different stages of BPH following EB treatment. (C) Downregulation of *AstAR* using RNAi leads to a reduction in mRNA expression level. (D) Egg production in female BPH following silencing of *AstAR* gene. (E-I) Expression of selected JH signaling pathway related genes (*FAMeT*, *JHAMT*, *Met*, *Kr-h1* and *JHE*) in *AstAR* silenced BPH. (J) JHIII titer of BPH females after *AstAR* gene silencing determined by HPLC-MS/MS. (K) Number of eggs laid per female in 48h following injection of six AstA1-AstA6 mature peptides and one AT mature peptide. Fifty nanoliter of PBS (as control) and seven peptides (20 pmol/insect) were injected into female BPH three days after eclosion. (L and M) The JH III titer

of BPH females at different time points following AstA or AT injection. (N) Schematic of the

proposed role of AstA/AstAR in the regulation of JH following EB exposure. The question

mark indicates one or more possible additional signals. All data are presented as means ± s.e.m.

Student's t test was used to compare the two samples. One-way ANOVA with Tukey's multiple

comparisons test was used to compare more than two samples. ns, no significant difference;

Asterisks indicate values significantly different from the control (ns, no significant; *$p < 0.05$, **p

< 0.01, ***$p < 0.001$ and ****$p < 0.0001$). Different lower-case letters above the bars indicate

significant differences ($p < 0.05$).

Figure 4-figure supplement 1

A
AT

```
Nillu-AT : GFKNGISLSTARGF
Locmi-AT : GFKN-VALSTARGF
Scham-AT : GFKN-VALSTARGF
Homvi-AT : GFKN-HALSTARGF
Manse-AT : GFKN-VEMMTARGF
Spofr-AT : GFKN-VEMMXARGF
```

B
AstA

```
Nillu-AstA1 : --------------------MYDFGL
Nillu-AstA2 : ------------DNNRQR---MYAFGL
Nillu-AstA3 : --------------ALK---AYEFGL
Nillu-AstA4 : -----------AAAPQK-LYSFGL
Nillu-AstA5 : ---------------SR----YQFGL
Nillu-AstA6 : GGAFYDGSGERSLRSQR----YNFGL
Drome-AstA1 : ---------------VER----YAFGL
Drome-AstA2 : ---------------LP--VYNFGL
Drome-AstA3 : --------------SRP----YSFGL
Drome-AstA4 : ----------TTRPQP----FNFGL
Spoex-AstA1 : ---------------SP---HYDFGL
Spoex-AstA2 : ---------------LP--VYNFGL
Spoex-AstA3 : --------------SRP----YSFGL
Spoex-AstA4 : --------------ARP----YSFGL
Spoex-AstA5 : --------------AR---AYDFGL
Spoex-AstA6 : ---------------LP---MYNFGL
Spoex-AstA7 : --------------AR---SYNFGL
Spoex-AstA8 : -------------LSSK----FNFGL
Spoex-AstA9 : ----------ERDMHR---FSFGL
Nasvi-AstA1 : ---------------LP---IYQFGL
Nasvi-AstA2 : --------------SQP----FSFGL
Nasvi-AstA3 : --------------TRP----YSFGL
Nasvi-AstA4 : --------------TGG----FNFGL
Nasvi-AstA5 : ---------------DK----YLFGL
```

C
AstB/MIP

```
Nillu-AstB1 : ----AWRDLQS-SW
Nillu-AstB2 : ----GWQDMPSSGW
Nillu-AstB3 : ----GWQDLQG-GW
Nillu-AstB4 : ----AWSSLRG-TW
Drome-AstB1 : ----AWQSLQS-SW
Drome-AstB2 : ----AWKSMNV-AW
Drome-AstB3 : RQAQGWNKFRG-AW
Drome-AstB4 : --EPTWNNLKG-MW
Drome-AstB5 : ---DQWQKLHG-GW
Grybi-AstB1 : ----GWQDLNG-GW
Grybi-AstB2 : ----GWRDLNG-GW
Grybi-AstB3 : ----AWRDLSG-GW
Grybi-AstB4 : ----AWDQLRP-GW
Grybi-AstB5 : ----AWERFHG-SW
Locmi-AstB1 : ----AWQDLNA-GW
Bommo-AstB1 : ----AWQDLNS-AW
```

D
AstCC

```
Nillu-AstCC : RRGN---QKNGRVYWRCYFNAVTCF
Mesma-AstCC : RR------GEGKMFWRCYFNAVSCF
Stear-AstCC : RR------GEGKMYWKCYFNAVSCF
Limpo-AstCC : RR------GERHRFWRCYFNAVSCF
Carma-AstCC : RR----GNGDGRLYWRCYFNAVSCF
Drome-AstCC : RRIQPSGSGGGRAYWRCYFNAVSCF
Strma-AstCC : RR-----RPNGRVYWRCYFNAVSCF
Athro-AstCC : RRG----QAKGRVYWRCYFNAVTCF
Apime-AstCC : RRG----QAKGRVYWRCYFNAVTCF
Locmi-AstCC : RRG----QKTG-QYWRCYFNAVTCF
Dapma-AstCC : RRG----QSSQRVFWRCYFNAVSCF
```

E
AstCCC

```
Nillu-AstCCC : KKRSYWKQCAFNAVSCFGK--
Limpo-AstCCC : RKRSYWKQCAFNAVSCFGRK-
Mesma-AstCCC : RKRSYWKQCAFNAVSCFGRK-
Stear-AstCCC : RKRSYWKQCAFNAVSCFGKK-
Strma-AstCCC : RKRSYWKQCAFNAVSCFGKK-
Carma-AstCCC : RKRSYWKQCAFNAVSCFGKRK
Apime-AstCCC : RKRSYWKQCAFNAVSCFGK--
Athro-AstCCC : RKRSYWKQCAFNAVSCFGK--
Locmi-AstCCC : RKRSYWKQCAFNAVSCFGK--
Dapma-AstCCC : RKRSYWKQCAFNAVSCFGK--
```

Figure 4-figure supplement 1. Alignments of the amino acid sequences of: (A) AT, (B) AstA, (C) AstB/MIP, (D) AstCC and (E) AstCCC peptides from select species. AT, AstA, AstB/MIP and AstCCC are predicted to have a C-terminal amide. The mature peptides belonging to the same species have been highlighted with the same color. Species names are as follows: Nillu (*Nilaparvata lugens*), Locmi (*Locusta migratoria*), Scham (*Schistocerca americana*), Homvi (*Homalodisca vitripennis*), Manse (*Manduca sexta*), Spofr (*Spodoptera frugiperda*), Drome (*Drosophila melanogaster*), Spoex (*Spodoptera exigua*), Nasvi (*Nasonia vitripennis*), Grybi (*Gryllus bimaculatus*), Bommi (*Bombyx mori*); Mesma (*Mesobuthus martensii*), Stear

(*Stegodyphus araneomorph*), Limpo (*Limulus polyphemus*), Carma (*Carcinus maenas*), Strma

(*Strigamia maritima*), Athro (*Athalia rosae*), Apime (*Apis mellifera*), Dapma (*Daphnia magna*).

Black lines under the sequences indicate the locations of the disulfide bridges in the mature

peptides. The accession numbers of the sequences are listed in Figure 4-figure supplement 1

source data.

Figure 4-figure supplement 2

Figure 4-figure supplement 2. Phylogenetic tree of the predicted BPH (*) allatotropin receptor

(A16, ATR), allatostatins A receptor (A2, AstAR), AstB (MIP) receptor (A10, AstBR or MIPR)

and allatostatins C receptor (A1, AstCR) with other insect species. The tree was generated

using the maximum likelihood method. *Drosophila melanogaster* metabotropic glutamate

receptor was included as an outgroup. The accession numbers of the sequences used for this

phylogenetic tree are listed in Figure 4-figure supplement 2 source data.

 Figure 4-figure supplement 3

Figure 4-figure supplement 3. EB induced changes in the expression of *AT*, *AstB*, *AstCC*,

***AstCCC*, *ATR*, *AstBR* and *AstCR* in BPH.** All data are presented as means ± s.e.m. Student's

t test was used to compare the two samples. ns, no significant difference; Asterisks indicate

values significantly different from the control (*$p < 0.05$, **$p < 0.01$, and ***$p < 0.001$).

454 **EB-enhanced fecundity in BPH is dependent on its molecular target protein the**

455 **GluCl**

456   EB and abamectin, are allosteric modulators, which target glutamate-gated

457 chloride channels (GluCls) ([60-62](#)). Hence, we examined whether EB-stimulated

458 fecundity in BPH is influenced by its molecular target GluCl. The full length *GluCl*

459 coding sequence from BPH was cloned and sequenced (Figure 5-figure supplement

460 1) and the impact of EB on *GluCl* gene expression examined using quantitative PCR.

461 Treatment of BPH with the LC_{50} concentration of EB significantly downregulated

462 *GluCl* gene expression at the 5E and 5M nymph stages while upregulating *GluCl*

463 gene expression at 2 DAE and 5 DAE in the adult stage (Figure 5A). To examine the

464 role of *GluCl* gene in BPH fecundity, RNAi was used to knockdown expression of this

465 gene in female BPH (Figure 5B). A significant decrease in the number of eggs laid by

466 per female was observed in *dsGluCl*-injected insects compared with *dsgfp*-injected

467 insects (Figure 5C). However, treatment with EB was found to rescue the decreased

468 egg-laying phenotype induced by *dsGluCl* injection (Figure 5C). To investigate the

469 mechanism by which *GluCl* expression modulates fecundity we examined if silencing

470 *GluCl* influences JH titer and JH-related gene expression. Indeed, we observed that

471 RNAi knockdown of *GluCl* leads to a decrease in JH titer (Figure 5D) and down-

472 regulation of genes including *JHAMT* which is responsible for JH synthesis, and the

473 JH signaling downstream genes *Met and Kr-h1* (Figure 5E-G). In contrast,

474 expression of *FAMeT* and *JHE* were not changed in the *GluCl* silencing insects

475 (Figure 5H and I). We also examined whether silencing *GluCl* impacts the

AstA/*AstAR* signaling pathway. Silencing *GluCl* was found to have no impact on the

expression of *AT*, *AstA*, *AstB*, *AstCC*, *AstAR*, and *AstBR*. However, the expression of

AstCCC and *AstCR* was significantly upregulated in *dsGluCl*-injected insects (Figure

5-figure supplement 2A-H). These results suggest that EB activates *GluCl* which

induces JH biosynthesis and release, which in turn stimulates reproduction in BPH

(Figure 5J).

Figure 5

Figure 5. EB induced reproduction in brown planthoppers through its molecular target protein GluCl.

(A) Expression of *GluCl* in EB-treated and untreated BPH. (B) Expression of *GluCl* following injection of *dsGluCl* in BPH. (C) Egg production after *GluCl* gene knockdown in EB-treated and untreated BPH. (D) The JH III titer of BPH females after *GluCl* gene silencing as quantified using the ELISA method. (E-I) Expression patterns of selected JH-related genes (*JHAMT*, *Met*, *Kr-h1*, *FAMeT* and *JHE*) in *GluCl* silenced BPH. (J) Schematic of the proposed role of GluCl as a molecular target of EB and EB-enhanced reproduction in BPH. The question mark indicates one or more possible additional signals. All data are presented as means ± s.e.m. Student's t test was used to compare the two samples. One-way ANOVA with Tukey's multiple comparisons test was used to compare more than two samples. ns, no significant difference; Asterisks indicate values significantly different from the control (ns, no significant; *p < 0.05 and **p < 0.01). Different lower-case letters above the bars indicate significant differences (p < 0.05).

Figure 5-figure supplement 1

Figure 5-figure supplement 1. Phylogenetic analysis of glutamate-gated chloride channels in

different species. The numbers at the nodes of the branches represent the percentage

bootstrap support (1000 replications) for each branch. The *Sogatella furcifera* GABA-gated

chloride channel and *Nilaparvata lugens* nAchR were used as outgroup. Alignment was

performed with amino acid sequences from TM1-7. The receptor names are listed in the tree.

The accession numbers of the sequences used for this phylogenetic tree are listed in Figure 5-

figure supplement 1 source data.

Figure 5-figure supplement 2

Figure 5-figure supplement 2. The expression of *AT*, *AstA*, *AstB*, *AstCC*, *AstCCC*, *AstAR*,

AstBR and *AstCR* in BPH injected with *dsGluCl or dsgfp*. All data are presented as means ±

s.e.m. Student's *t* test was used to compare the two samples. ns, no significant. Different lower-

case letters above the bars indicate significant differences ($p < 0.05$).

Discussion

Pesticide-induced resurgence of pest insects is a serious problem in rice and several other cropping systems (3). However, the mechanisms underpinning pesticide-enhanced reproduction in insects remain poorly understood. Here we reveal that a suite of molecular actors underlie this trait that, in combination, mediate profound physiological changes in the reproductive fitness of female BPH. Our data provide fundamental insights into the molecular mechanisms by which xenobiotics modify insect reproduction and have applied implications for the control of a highly damaging crop pest. We discuss these topics below.

Sublethal doses of GluCl modulators, EB and abamectin, stimulates fecundity in BPH

We show that in both contact and systemic assays EB and abamectin stimulate reproduction in BPH. Thus, insecticide-enhanced reproduction is likely a key factor in the BPH resurgence observed when farmers use EB and abamectin to control leaffolders in rice crops in China (29). Although this is the first report of sublethal doses of avermectins enhancing insect fecundity, our findings are consistent with previous studies which have shown that certain insecticides, herbicides and fungicides stimulate BPH reproduction (3, 7, 14, 63-68). Intriguingly, we show that EB only induces fecundity in female adults and is specific to BPH, with EB exposure failing to enhance reproduction in two related species, the small brown planthopper, *L. striatellus* and the white backed planthopper, *S. furcifera*, or the model insect *D. melanogaster*. Thus, the

mechanisms underpinning this trait appear to act exclusively on female BPH and may be specific to this species.

Pesticides may stimulate insect reproduction through a variety of physiological and molecular mechanisms. Our data reveal that exposure to sub-lethal concentrations of EB results in profound changes to female BPH fitness, leading to increases in female weight, total protein content, cholesterol content, sugar content and egg production and decreases in duration of the preoviposition period. Some of these findings exhibit parallels with previous studies, which demonstrated that treating third-instar BPH nymphs with either deltamethrin, triazophos, or imidacloprid led to increased soluble sugar levels in the corresponding adults (69). Such metabolites provide the energy that drives BPH reproduction and resurgence (3). Thus, together with prior work, our results suggest that pesticides associated with resurgence stimulate nutritional intake in BPH to fuel enhanced energy-intensive reproduction.

The JH signaling pathway plays an essential role in EB-induced fecundity in BPH

JH is a pleiotropic hormone which plays important roles in development and reproduction in insects (19, 70). Circulating JH titers are regulated by factors that control JH production in the corpora allata including biosynthetic enzymes and catabolic enzymes that regulate JH levels. Our results show that EB increases circulating JH III titers in BPH females over 2–4 days after eclosion (DAE) and promotes ovarian development. Previous studies have reported that triazophos and deltamethrin treatments also lead to increased circulating JH III titers in BPH females

over 1–3 days post emergence. Similarly, jinggangmycin treatments were found to lead to increased JH titers (by approximately 45–50%) in BPH females over two days post emergence (71). Thus our findings, in combination with these previous studies, demonstrate that insecticide treatments can have dramatic effects on the regulation of key insect hormones involved in pest reproduction which can in turn drive pesticide resurgence.

Although increased JH titers following pesticide exposure have been correlated with reduced levels of active JH esterase during the first three days PE (6), the type and number of mechanisms mediating the observed increase in hormone titer has remained an open question. Our data reveal that elevated JH titer in EB-exposed BPH is associated with the upregulation of genes that encode biosynthetic enzymes for JH (*JHAMT*) and downstream signaling genes that can induce *vg* gene expression (*met* and *kr-h1*). Using RNAi we provide functional evidence of the role of these genes in the regulation of JH III and fecundity of female BPH, and demonstrate that EB can restore the reduction in egg production resulting from the knockdown of *JHAMT*, *met* and *kr-h1* expression.

JHMAT is an enzyme that catalyzes the conversion of inactive precursors of JH to active JH in the final stages of JH biosynthesis (36, 38). Interestingly, while it has not been previously implicated in pesticide resurgence, treatment of the stored product pest *Sitotroga cerealella* with diallyl trisulfide, an insecticidal compound in garlic essential oil, was found to increase JHAMT mRNA levels (72). Because JHMAT is the key rate-limiting enzyme in regulation of JH titer our results suggest that its enhanced

expression is a key molecular mechanism of pesticide resurgence in BPH.

Met is a ligand-activated member of the basic helix–loop–helix Per/Arnt/Sim

(bHLH-PAS) transcription factors and is the intracellular receptor for JH ([39], [40]). Kr-h1

is a zinc finger protein that acts downstream of Met and is expressed in response to

JH signaling. Although the genes encoding these proteins have not been previously

linked to pesticide resurgence, our finding that they are upregulated following EB

exposure, and demonstration of their role of in promoting fecundity, is consistent with

previous studies. Specifically, treatment of BPH with JH III or the insecticidal analogues

methoprene or pyriproxifen was found to induce the expression of *Kr-h1* ([73]).

Furthermore, knockdown of *Met* and *Kr-h1* in BPH brachypterous females was found

to result in delayed ovariole development and this was significantly more pronounced

than the response observed in BPH treated separately with ds*NlMet* or ds*NlKr-h1* ([74]).

This finding provides evidence of a possible interaction between *Met* and *Kr-h1* and,

in combination with our data, suggests that *Met* and *Kr-h1* may act in concert to

mediate EB-enhanced fecundity.

EB-induced fecundity in BPH is dependent on the allatostatin signaling pathway

In addition to regulatory proteins that promote JH production, insects have

peptides that inhibit JH biosynthesis. These include the allatostatins: FGLamides

(FGLa; AstA), the W(X)6Wamides (AstB), and the PISCFs (AstC) ([50], [51], [57], [58], [75],

[76]). Interestingly, our results showed that EB exposure results in the marked

downregulation of the expression of the genes encoding allatostatin *AstA* and its

receptor *AstAR*. We provide clear evidence of the functional impact of this on JH

synthesis and BPH fecundity by: i) demonstrating that RNAi knockdown of *AstAR*

expression results in increased JH titer and enhanced female egg production, and, ii)

showing that injection of female BPH with synthetic AStA peptide reduces JH titer and

decreases egg production. Thus, our data provide unequivocal evidence that AstA is a

key inhibitor of JH production in BPH. This finding is consistent with previous work

which has shown that FGLa/ASTs inhibit JH biosynthesis in cockroaches, and termites

(46, 77). To our knowledge, our study is the first report of insecticides inhibiting the

expression of the neuropeptide receptor, *AstAR*, and linking this to increases in JH titer

and enhanced reproduction in insects.

Interestingly knockdown of *AstAR* resulted in significant increases in the

expression of genes involved in JH synthesis/signaling including *FAMeT* and *JHAMT*

Met and *Kr-h1*. Related to this finding, previous work has shown that knockdown of the

AstA receptor gene, *Dar-2*, in *D. melanogaster* results in changes in the expression of

genes encoding Drosophila insulin-like peptides (DILPs) and adipokinetic hormone

(AKH) signaling proteins (78). Together with our findings, this demonstrates that AstA

receptors may modulate the expression of numerous downstream genes involves in

metabolism, energy store and reproduction. In the case of pesticide resurgence our

results imply significant cross-talk in the expression of genes that inhibit JH production

and those that promote it.

The GluCl plays an essential role in EB-induced fecundity in BPH

EB and abamectin are allosteric modulators of GluCls ([60-62](#)). Our data revealed

that EB exposure modifies expression of the GluCl in BPH, and knockdown of GluCl

expression resulted in a reduction in both JH levels and egg production. Interestingly,

the GluCl has been reported to inhibit the biosynthesis of JH in the cockroach,

Diploptera punctata ([28](#)). Recent work has also reported that modulation of

glutamatergic signals may contribute to the photoperiodic control of reproduction in

bean bug, *Riptortus pedestris* ([79](#)). Interestingly, work on *D. punctata* has revealed that

application of the GluCl channel agonist ivermectin, which like EB belongs to the

avermectin family, caused a decline in JH synthesis in corpus allatum glands ([28](#)).

While the inhibitory effect of ivermectin observed in this previous study differs from the

activating effect of EB we observed in our study, it is consistent with our finding of a

role for GluCl channel in the regulation of JH regulation. Interestingly, we found that

knockdown of *GluCl* gene expression results in the down-regulation of *JHAMT*, *Met*

and Kr-h1, further revealing significant convergent relationships between genes

underpinning pesticide resurgence.

Conclusion

Our study has revealed a diverse suite of genes that act in combination to enhance

JH titer and thus fecundity following BPH exposure to EB. A schematic of how these

factors promote ovarian development in the adult stage of *N. lugens* through the JH

signaling pathway is provided in [Figure 6](#). Our findings provide the foundation for

further work to understand exactly how these genes interact and the mechanisms by

which their expression is activated or repressed by EB. Furthermore, our findings

provide fundamental insights into the molecular response in insects to xenobiotic

stress and illustrate that pesticides can have unexpected and negative impacts on pest

populations. In this regard our findings also have applied implications for the control of

a highly damaging crop pest. Previous studies have reported that avermectins such as

abamectin are toxic to the wolf spider *Pardosa pseudoannulata*, which is the main

predator of BPH in rice crops ([80], [81]). Thus, these insecticides both stimulate

reproduction in BPH while killing their natural enemies providing a 'perfect storm' for

damaging BPH outbreaks. Based on these findings, to avoid BPH resurgence, we

suggest that the GluCls, EB and abamectin, should not be (or rarely be) applied to rice

plants at growth stages when BPHs are present. On a more optimistic note, our

findings have identified numerous genes that play key roles in BPH reproduction and

thus represent promising targets for the development of novel controls against this

important pest.

Figure 6

Figure 6. Schematic of the proposed regulatory pathway of EB-enhanced fecundity

in BPH. Emamectin benzoate (EB) exposure results in the upregulation of genes that promote

JH production (*JHAMT, Met and Kr-h1*) and the downregulation of genes that inhibit it

(allatostatin, Ast*A* and allatostatin A receptor, *AstAR*). This transcriptome reprograming is

dependent on the action of EB on its molecular target the glutamate-gated chloride channel

(GluCl) receptor. The resulting increased JH titer promotes vg synthesis and increased

fecundity in EB exposed insects. We observe significant cross-talk in the expression of genes

that inhibit JH production and those that promote it, with *AstAR* inhibiting the expression of

JHAMT, Met and *Kr-h1* and *GluCl* activating the expression of *JHAMT* which is responsible for

JH synthesis, and the JH signalling downstream genes *Met and Kr-h1*.

Materials and methods

Insects

BPH was initially collected from Wanan, JiangXi Province in 2020, reared on 'Taichung Native 1' (TN1) rice seedlings in the laboratory without exposure to any insecticides. The strain was maintained in a climatic chamber at 27 ± 1°C, with relative humidity of 70 ± 10% and a light: dark = 16 h: 8 h photoperiod.

Chemicals

Emamectin benzoate (95.2%) was obtained from Hebei Weiyuan Co., Ltd. (Hebei, China). Abamectin (96.8%) was obtained from Hebei Weiyuan Co., Ltd. (Hebei, China). Pyriproxyfen (98%) was obtained from Shanghai Shengnong Co., Ltd. (Shanghai, China). Methoprene (S)-(+) (mx7457-100mg) was purchased from Shanghai MaoKang Biotechnology Co., Ltd., (Shanghai, China). Juvenile hormone standard sample (J912305-10mg) was purchased from Shanghai Macklin Biotechnology Co., Ltd., (Shanghai, China).

Bioassay

Different life stages of insects were used to perform bioassay to investigate the effects of insecticide on nymphs and adults. To test whether treatment of the nymph stage of insects would promote reproduction in female, we used 4th instar nymphs of BPH or 3rd instar nymph of *Laodelphax striatellus* and *Sogatella furcifera* to perform bioassays. To test whether treatment of the adult stage of insects would promote reproduction in female, we used newly emerged male and female BPH.

Systemic route: The rice-seeding dipping bioassay method was used to evaluate

the susceptibility of BPH, *L. striatellus* and *S. furcifera* to EB. Technical-grade

insecticides were dissolved in acetone as stock solution then diluted in a series of six

concentrations with water containing 0.1% Triton. Selected rice seedlings at the 6-8

cm growth stage were dipped in insecticide solutions for 30 s and then air-dried at

room temperature. The roots of the rice seedlings were wrapped with cotton strips and

placed seedlings placed in a plastic cup 5 cm in diameter. Fifteen insects were

introduced into each plastic cup for each replicate. The top of the cup was then selaed

with gauze to prevent escape. All experiments comprised at least three biological

replicates. Control rice seedings were treated with 0.1% Triton X-100 water solution

only. All treatments were maintained under standard conditions of 27 ± 1 °C and 70–

80% relative humidity with a 16 h light/8 h dark photoperiod. Mortality was assessed

after 4 d for *N. lugens* or 2 d for *L. striatellus* and *S. furcifera* after treatment with

insecticides. The insects were considered dead if they were unable to move after a

gentle prodding with a fine brush.

For *Drosophila* larvae bioassay, we adopted a method described previously in our

lab with minor modifications ([82](#)). Briefly, twenty third instar larvae were placed in fly

vials containing fly food (based on corn powder, brown sugar, yeast and agar)

supplemented with EB of different concentrations. Four concentrations (LC_{10}, LC_{30} and

LC_{50}) were tested together with a negative (no insecticide) control. For *Drosophila* adult

bioassays, we selected virgin females three days after eclosion. Several

concentrations were overlaid onto fly food in standard *Drosophila* vials and allowed to

dry overnight at room temperature. 15 adult flies (three days after eclosion) were then

added to each vial and mortality assessed after 2 d. Four replicates were carried out

for each concentration. Control mortality was assessed using vials containing food with

solvent minus insecticide.

Contact route: For topical bioassays working insecticide solutions were prepared

in acetone. 4th instar nymphs or newly emerged males/females were anesthetized with

carbon dioxide for 5 s, and then 0.04 µl/insect test solution applied topically to the

dorsal plates with a capillary micro-applicator ([29]). Insects were then placed in an

artificial climate incubator with a temperature of 27±1°C, a photoperiod of 16:8 h (L:D),

and a humidity of 70%±10%. Mortality was determined 2 d after treatment. Data with

over 20% mortality in the control treatment were discarded, and the assay was

repeated at least three times.

Fecundity assays

Fourth instar nymphs of BPH and 3rd instar nymph of *L. striatellus* and *S. furcifera*

were treated with the LC_{15} and LC_{50} of EB, and then transferred to fresh rice seedlings.

After eclosion, the adults were used in the following experiment. Newly emerged

treated adults and untreated adults were paired to produce four groups: untreated

males and untreated females (♂ck×♀ck; ck indicates untreated); untreated males and

treated females (♂ck×♀t; t indicates insecticide treated); treated males and untreated

females (♂t×♀ck); treated males and treated females (♂t×♀t). Each group comprised

at least 10 mating pairs. All pairs were transferred to glass cylinders (diameter 5 cm

and height 20 cm) containing rice plants (25 days old almost 20 cm high) as a food

source for eleven days. The number of eggs and nymphs in plants were counted by a

microscope.

For *Drosophila* egg-laying assay, we adopted our previous method ([83]). Briefly,

insecticide-treated virgin females were paired with untreated males for three days and

then the mated females transferred into the *Drosophila* ovipositing apparatus. The

number of eggs were counted after 16 hours.

Fitness analysis

The fitness of EB-treated BPH were analyzed using methods reported previously

([10]). We selected two groups, (♂ck×♀ck) and (♂ck×♀t), to study the effects of the LC_{50}

concentration of EB on BPH fitness. In the case of systemic exposure, 4th instar

nymphs of BPH were treated with the LC_{50} of EB for 4 days and then transferred to

tubes containing untreated rice plants for individual rearing. The rice plants were

replaced every three days with untreated plants. The emergence ratio, female ratio,

preoviposition period, female longevity, brachypterism female ratio and female weight

were calculated.

Examination of ovarian development

Adult females from ♂ck×♀ck control and ♂ck×♀t group on 1, 3, 5, 7 DAE were

dissected for observe the ovarian development. The detained eggs in ovary were

photographed and recorded. Each group has at least fifteen replicates.

To examination whether EB treated impairs egg maturation, we dissected

untreated or EB-treated ovaries and fixed them in 4% paraformaldehyde (PFA) in

phosphate buffered saline (PBS) for 30 min at room temperature. After four washes of

10 min (4 × 10 min) in PAT3 (PBS with 0.5% Triton X-100 and 0.5% bovine serum

albumin), the ovaries were then incubated with DAPI (4',6-diamidino-2-phenylindole,

100 nM) and Actin-stain 670 Fluorescent Phalloidin (200 nM). Imaging was performed

using Zeiss LSM980 confocal laser microscope.

Measurements of glycogen, triglyceride, total protein content, cholesterol and

four sugars

The content of glycogen, triglyceride, cholesterol and total protein was

determined by spectrophotometry at 620 nm, 510 nm, 510 nm and 562 nm

respectively using the glycogen assay kit (A043-1-1), triglyceride reagent kit (A110-1-

1), cholesterol assay kit (A111-1-1) and total protein assay kit (A045-2-2) obtained

from Nanjing Jiancheng Bioengineering Institute following to the manufacturer's

instructions. The determined results were normalized to the protein content in the

sample, which was determined using BCA Protein Assay Reagent Kit (Thermo

Scientific, Waltham, USA). Each sample contained tissue extracts from five adult

female BPH, with three biological replicates per sample.

To assess the content of four sugars (sucrose, glucose, fructose and trehalose)

in the extract of BPH tissue, the same extraction method was used as above. Sugar

content was quantified using the colorimetric method by scanning at 290 nm, 505

nm, 285 nm and 620 nm respectively using the sucrose reagent kit (A099-1-1),

glucose reagent kit (F006-1-1), fructose reagent kit (A085-1-1) and trehalose reagent

kit (A150-1-1) obtained from Nanjing Jiancheng Bioengineering Institute based on the

manufacturer's instructions. Each sample contained tissue extracts from five adult

female *N. lugens*, with three biological replicates per sample.

Determination of Juvenile hormone III and ecdysone titers of BPH by ELISA

The titer of Juvenile hormone III in BPH was measured using the Juvenile

hormone ELISA Kit (Lot Number: HLE92086, Shanghai HaLing Biotechnology Co., Ltd.,

Shanghai, China) which employs the dual-antibody sandwich ELISA method. The titer

of ecdysone in BPH were measured using the ecdysone ELISA Kit (Lot Number:

ZK14705, Shanghai ZhenKe Biotechnology Co., Ltd., Shanghai, China). At least three

biological replicates were employed for each treatment.

Determination of Juvenile hormone III titer in BPH using HPLC-MS/MS

The whole bodies of 5 individuals BPH were mixed with 1 ml of n-hexane, followed

by centrifugation at 10,000×g for 10 min, the upper hexane layer was then dried with

nitrogen, dissolved in methanol and sonicated for 10 min, after centrifugation at

10000×g for 10 min, the supernatant was collected through the organic filter membrane

of 0.22 μm into 2 ml vials for JH Ⅲ determination. JH Ⅲ standard sample (J912305-

10 mg) purchased from (Shanghai McLean Biochemical Technology Co. Ltd),

dissolved in methyl alcohol as stock solution 10,000 mg/L was diluted in a series of six

concentration gradients to serve as a reference. Liquid chromatography-tandem mass

spectrometry (LC-MS/MS) was then carried out using UPLC Xevo TQ-S Micro (Waters

technology), quantitative method according to the external standard, the

chromatographic column was EC-C18 (4.6 mm×150 mm, 2.7 μm), column temperature

was 30°C，injection volume was 20 μl, elution flow rate was 0.3 ml/min, and the mobile

phase was acetonitrile:formic acid water (90:10), detection wavelength was 218 nm,

the peak height was used for quantification.

Cloning, sequence and phylogenetic analysis

The NCBI database and BLAST program were used to carry out sequence

alignment and analysis. Open Reading Frames (ORFs) were predicted with EditSeq.

Primers were designed using the primer design tool in NCBI. Total RNA Extraction was

extracted from 30 adults BPH using TRIzol reagent (Invitrogen, Carlsbad, CA, USA)

according to the manufacturer's instructions. cDNA was synthesized using the Biotech

M-MLV reverse transcription kit. Full-length gene sequences were amplified by PCR

using cDNA as template and Rapid Taq Master Mix (Vazyme Biotech, Cat# P222-02).

The PCR product was purified on a 1% agarose gel, cloned into pClone007 Simple

Vector Kit (Tsingke Biotech, Cat# TSV-007S), and then sequenced using the 3730 XL

DNA analyzer (Applied Biosystems, Carlsbad, CA, USA). Table S2 contains a list of

the primers used in this study.

The exon and intron architectures of AT, AstA, AstB, AstCC and AstACCC were

predicted based on the alignments of putative ORFs against their corresponding

genomic sequences. Sequence similarity/annotations and orthologous gene searches

were performed using BLAST programs available in NCBI. Multiple alignments of the

complete   amino   acid   sequences   were   performed   with   Clustal   Omega

(http://www.ebi.ac.uk/Tools/msa/clustalo).   Phylogeny   was   conducted   using   the

maximum   likelihood   technique   to   create   phylogenetic   trees   and   these   were

bootstrapped with 1000 replications were used using the MEGA 6 software ([84]).

RNA interference

Double-stranded RNA (dsRNA) of *gfp* (green fluorescent protein), *JHAMT*

(juvenile hormone acid O-methyltransferase,), *Met* (methoprene-tolerant), *Kr-h1*

(krüppel homolog 1), *AstAR* (allatostatin-A receptor) and *GluCl* (glutamate-gated

chloride channel) was prepared using Ambion's MEGAscript T7 kit instructions

following the manufacturer's instructions. The primer sequences for double-stranded

RNA synthesis are listed in Table S2. Newly emerged females were injected with 40

nl (5,000 ng/µl) of double-stranded RNA of *gfp* (*dsgfp*) or double-stranded RNA of the

target genes in the conjunctive part between prothorax and mesothorax of insects. In

the RNAi experiments, BPH were then treated with the LC_{50} of EB 24h after dsRNA

injection and the whole body sampled for qRT-PCR analysis.

Quantitative RT-PCR

Fourth instar nymphs of BPH were treated with EB after which total RNA was

extracted from 5th instar nymphs and 1-7 day post-eclosion females of *N. lugens* using

the methods detailed above. The HiScript® II RT SuperMix for qPCR (+gDNA wiper)

kit from Vazyme, Nanjing, China, was used to create first-strand cDNA. Primer3

(http://bioinfo.ut.ee/primer3/) was used to design real-time quantitative PCR (qPCR)

primers listed in Table S2. mRNA levels of candidate genes were detected by qPCR

using the UltraSYBR Mixture (with ROX) Kit (CWBIO, Beijing, China). Each reaction

contained 2 µL of cDNA template (500 ng), 1 µL each forward and reverse qPCR primer

(10 µM), 10 µL of UltraSYBR mixture buffer, and 6 µL of RNase-free water. Q-PCR

was run on an ABI 7500 Real-Time PCR System (Applied Biosystems, Foster City, CA)

under the following conditions: 5 min at 95°C, followed by 40 cycles of 95°C for 10 s

and 60°C for 30 s. Three independent biological replicates and four technical replicates

were used in each qPCR experiment. The housekeeping genes 18S ribosomal RNA

of BPH were selected to normalize the expression of candidate genes. The $2^{-\Delta\Delta Ct}$

method (Ct represents the cycle threshold) was used to measure relative expression

levels (85). Three biological replicates were used for statistical comparison between

samples. Table S2 contains a list of the primers used in this study.

Statistics

PoloPlus v2.0 (LeOra Sofware 2008) was used to calculate the lethal

concentration (LC_{50}) and 95% fiducial limits (95% F.L.). GraphPad Prism 8.0 software

(GraphPad Software Inc., San Diego, USA) was used to generate graphs and perform

statistical analysis of data. Data presented in this study were first verified for normal

distribution using the D'Agostino–Pearson normality test. One-way analysis of

variance (ANOVA) with Duncan's multiple range test was used to test differences

among multiple groups of normally distributed groups of data. Student's t test was used

to test the differences between two groups. If not normally distributed, Mann–Whitney

test was used for pairwise comparisons, and Kruskal–Wallis test was used for

comparisons among multiple groups, followed by Dunn's multiple comparisons. All

data are presented as mean ± s.e.m. The sample sizes and statistical tests used for

each experiment are stated in the figures or figure legends.

 Table S2 Sequences of oligonucleotide primers used in this study.

Primers	Primer sequences (5' - 3')
For cDNA cloning	
Allatotropin：	
NlAT-F	TACGCGGCCAAACACACTTA
NlAT-R	AGGGAAAGAGGGCGAAATTCA
Allatostatins：	
NlAstA-F	TCGGCCGTCACAAGTCAAG
NlAstA-R	CCGAACCCGTACTTCATGCT
NlAstB-F	ACCGGGCTCACAGGAATTTT
NlAstB-R	TGTAGGCGCAGATCTTGAGG
NlAstCC-F	AACACAGCTCTACGAGGCAC
NlAstCC-R	CCAAGCAGGTGACTGCCATA
NlAstCCC-F	TTTGTGTGTGCTTGCAGGTG
NlAstCCC-R	GGATAGAAACGGTAGATTTGGTAGA
Allatostatins receptor：	
NlA16-F	CCTCATTGTGGAACCACCGA
NlA16-R	CGCAGCTGTAAGGTGGAAGA
NlA2-F	GAACGTAATGGGAGTCGGCA
NlA2-R	GTTTTTGTGAGCGCCGACTT
NlA10-F	ATGCAAAACACGGCCAGCCT
NlA10-R	TTAATCGTCTCTGCTCAACTCCAAAGGAAGGT
NlA1-F	CGACCAGACCACTCTACTGC
NlA1-R	ACGTGGACCTCACTATACCAAAAA
For Quantitative RT-PCR	
Q-Nl18S-F	CGCTACTACCGATTGAA
Q-Nl18S-R	GGAAACCTTGTTACGACTT
Q-vitellogenin-F	GTGGCTCGTTCAAGGTTATGG
Q-vitellogenin-R	GCAATCTCTGGGTGCTGTTG
Q-Vitellogenin receptor-F	AGGCAGCCACACAGATAACCGC
Q-Vitellogenin receptor-R	AGCCGCTCGCTCCAGAACATT
Q-JHE-F	GAGCCTCACATCCACAGC
Q-JHE-R	AATGGGAGCCCTACGC
Q-NlMet-F	GGTGGTAAACGGATTGGAAA
Q-NlMet-R	CATCGTCAGCCAACTCGATA
Q-NlFAMet-F	GCAAAGTCAGCAATCCGCAAGAAC
Q-NlFAMet-R	ACACCGTAGTGGGTGACAACGAATG
Q-JHAMT-F	GAACCTGCAGGCCAAACACA
Q-JHAMT-R	ACCACTCGGTTGGGCTGAAT
Q-NlKr-h1-F	TGATGAGGCACACGATGACT
Q-NlKr-h1-R	ATGGAAGGCCACATCAAGAG
Q-NlAT-F	CACGATACGTGGCTTCAAGA
Q-NlAT-R	ACGATCACTTTCGCCAATTC
Q-NlAstA-F	AGGACTTACTGGGCGAGGAT

Q-NlAstA-R	GGTGTCTCGTTTCCTGGTGT
Q-NlAstB-F	AGCGAGCTAGACGAGGACAA
Q-NlAstB-R	TCGTCTCTGCTCAACTCCAA
Q-NlAstCC-F	CTGCTCCCAGTGAAAAGGAG
Q-NlAstCC-R	GCTTCCAGTAACTGCGCTTC
Q-NlAstCCC-F	TGTGCTTGCAGGTGGTAGTC
Q-NlAstCCC-F	AGAAGCATGTGACTGCGTTG
Q-NlA2-F	TCCTGGTGCTGAAGAGTGTG
Q-NlA2-R	CTTTTCGGGCCCATTAATTT
Q-NLA1-F	ATATCGGCACCGAAGATGAG
Q-NLA1-R	GTCTGACCCGACAGGTTCTC
Q-NLA10-F	ACTGGGTGTCGACCAATCTC
Q-NLA10-R	TCGGTAGCGAGGAAGACAGT
Q-NLA16-F	TACCGTTCTGTGGGATGTCA
Q-NLA16-R	CCGGGATATCAAAGACGAGA
Q-NlGluCl-F	CACTGACTGAGGCCAACAGA
Q-NlGluCl-R	GCTGGCCATTCTTAGTGAGC

For double-stranded RNA synthesis

T7-EGFP-F	TAATACGACTCACTATAGGGCGTAAACGGCCACAAGTTCA
T7-EGFP-R	TAATACGACTCACTATAGGGGACTGGGTGCTCAGGTAGTG
T7-Kr-h1-F	TAATACGACTCACTATAGGGCGCCAGTGAAAGTGAGACCT
T7-Kr-h1-R	TAATACGACTCACTATAGGGGAGACCGCAAGTGGTTCTGA
T7-Met-F	TAATACGACTCACTATAGGGCCACCAACCAGCAGATGAACCTGA
T7-Met-R	TAATACGACTCACTATAGGGCCACGCAAAGCCTCGTACTCTTGG
T7-JHAMT-F	TAATACGACTCACTATAGGGCTCCAGGCCATTGTCCCTCA
T7-JHAMT-R	TAATACGACTCACTATAGGGTTGGCCTGCAGGTTCTTTGG
T7-AstA-R-F	TAATACGACTCACTATAGGGTACTGCCGTTCTGGCCTTTT
T7-AstA-R-R	TAATACGACTCACTATAGGGGTGATCTGGAAGAGCGGCTT
T7-Glucl-F	TAATACGACTCACTATAGGGACACATCACCTGCTCACCTG
T7-Glucl-R	TAATACGACTCACTATAGGGGTGTGTTTGCCTGCTGTCTG

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