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Use of mutants to dissect the role of ethylene signalling in organ senescence and the regulation of yield in *Arabidopsis thaliana*

Running title: Ethylene signalling and senescence in Arabidopsis

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Abstract

The role of ethylene in regulating organ senescence in *Arabidopsis* has been investigated by studying the development of mutants that have an attenuated capacity to perceive the gas. The onset of leaf senescence and floral organ abscission was delayed in the ethylene insensitive mutant *etr1*. The photosynthetic life span of rosette leaves was similarly extended in the gain of function mutant *ers2* and this mutant also exhibited a delay in the timing of pod dehiscence primarily as a consequence of an extension in the final stages of senescence. A detailed analysis of yield revealed that whilst thousand grain weight was increased, by as much as 20%, in *etr1*, *ein4* and the loss of function mutant *etr2*, only the latter showed a significant increase in total weight of seeds produced per plant. The other mutants studied exhibited a reduction in total seed yield of almost 40%. These observations are discussed in the context of the possible role of ethylene in regulating organ senescence and their significance in the breeding of crop plants with enhanced phenotypic characteristics.

Keywords:

Ethylene signalling, mutant, senescence, abscission, *Arabidopsis*, yield, seed, pod development, dehiscence

Introduction

Since its serendipitous discovery over one hundred years ago ethylene has been known to regulate plant growth and development (Neljubov 1901), however, it was not until 1934 that it was officially classified as a plant hormone. This gaseous hydrocarbon is synthesized by most, if not all, plant tissues and affects a wide variety of processes including: germination, seedling growth, organ senescence, abscission, fruit ripening, grain filling, and responses to biotic and abiotic stresses (Abeles and others 1992; Gao and others 2003; Yang and others 2006; Robert and others 2008; Carbonell-Bejerano and others 2011).

Ethylene is only one of many factors affecting organ senescence, a process which also requires the plant to be at an appropriate stage of developmental competency to respond to the gas. For instance, young leaves fail to undergo senescence even when exposed to concentrations of ethylene that are known to promote the process in mature tissues (Hensel and others 1993; Grbić and Bleecker 1995; Buchanan-Wollaston 1997; Jing and others 2005) and these observations have given rise to the concept of a 'senescence window' in plant development (Jing and others 2002). The presence of reproductive structures, which constitute a resource sink in the plant, has also been shown to affect both the onset and/or rate of leaf senescence (Bennett and others 2012).

In *Arabidopsis* ethylene is perceived by a suite of receptors (ETR1, ETR2, ERS1, ERS2 and EIN4 - Chang and others 1993; Hua and others 1995; Sakai and others 1998) which are homologous to bacteria two-component histidine kinase (HK) sensors (Binder and others 2012) and predominantly located on the endoplasmic reticulum (ER). Based on their phylogeny and structure the ethylene receptors can be split into two subfamilies with ETR1 and ERS1 constituting subfamily I, whilst ETR2, EIN4 and ERS2 form subfamily II. The major difference between these groups is that the His kinase domain of subfamily II lacks any catalytic activity (Guo and Ecker 2004). Although it has been concluded that HK activity is not necessary for receptor signalling and therefore the subfamily II receptors are still able to inhibit ethylene responses (Hall and others 2012). The receptors act as negative regulators of ethylene signalling and in their unbound state they activate the Ser/Thr kinase CTR1, which leads to a suppression of the downstream ethylene signalling component EIN2 (Bleecker and Kende 2000). When ethylene binds to the receptors, with the aid of a copper cofactor provided by the copper transporter RAN1 (Rodriguez and others 1999; Binder and others 2010), this prevents the receptors from signalling, thereby releasing the repression of the

EIN2 pathway and enabling ethylene responses to occur. Consequentially mutations affecting the ability of a receptor to bind ethylene can lead to a dominant insensitivity to the gas (Chang and others 1993; Hua and others 1995; Sakai and others 1998).

In recent years, advances in ethylene research have rendered the concept of a linear signalling pathway somewhat oversimplified and despite the ethylene receptors being genetically similar and all contributing to the ethylene signalling pathway, it is emerging that they are not functionally redundant and are not able to fully compensate for one another (O'Malley and others 2005). For example ETR1 has a distinctly different role compared to the other receptor isoforms in ethylene-stimulated nutational bending of *Arabidopsis* hypocotyls (Binder and others 2006). Instead they act in a synergistic manner to enhance receptor output and are capable of interacting with either each other, or CTR1, to form homodimers and larger complexes (Gamble and others 2002; Gao and others 2003; Gao and others 2008; Grefen and others 2008; Liu and Wen 2012). Within this system it is believed that the subfamily I receptors play a more central role in the ethylene signalling pathway (Qu and others 2007, Gao and others 2008). This is possibly due to subfamily I having a stronger association with CTR1 and it has been proposed that a degree of subfamily II receptor signalling relies upon the presence of subfamily I receptors (Cancel and Larsen 2002; Qu and others 2007). Interestingly the receptors are also capable of directly interacting with EIN2 (Bisson and Groth 2010). The emerging number of potential receptor interactions could help explain why plants are able to respond differently to such a vast range of ethylene concentrations across diverse tissue types (Liu and Wen 2012, Shakeel and others 2013). Equally it is also possible that they are able to signal down CTR1 independent pathways (Desikan and others 2005; Ju and Chang 2012; Shakeel and others 2013) to bring about a diverse array of responses. The need for multiple receptors is highlighted by the fact that in all plants studied to date more than one ethylene receptor has been identified (Binder and others 2012). In addition the ethylene signalling system in *Arabidopsis* shares a great deal of similarity with crop plants such as tomato, rice and maize (Gallie and Young 2004; Klee 2004; Wuriyangan and others 2009), making it easier to translate research from a model plant into commercially important crops.

As the structure and function of the ethylene receptors begins to be unravelled it is becoming increasingly clear that their role in plant development, and specifically organ senescence, has yet to be fully elucidated. It has been predicted that delaying the start of senescence by just 2 days could lead to an 11% increase in the carbon contributed to the plant from a leaf (Thomas and Howarth 2000), which if harnessed correctly could lead to

yield increases. For example decreased expression of the ethylene receptor *ETR2* in rice increased thousand grain weight by up to 4% due to an alteration of starch and sugar accumulation in the filling grain (Wuriyanghan and others 2009). In addition postponing the onset of senescence is also crucial for delaying postharvest degradation and extending shelf life. One way of achieving this is to block the ethylene receptors so they are no longer capable of sensing ethylene and the pathway remains repressed. Commercially this has been achieved by the application of 1-methylcyclopropene (1-MCP) (Sisler and Serek 1997), which is highly effective in bagged leaves such as basil not only in terms of delaying senescence but also extending shelf life and retaining quality (Hassan and Mahfouz 2010). Whilst successful, 1-MCP application only provides a short term solution and requires a direct intervention to apply the chemical, therefore a more permanent solution could involve reducing the receptors sensitivity to ethylene perception (Agarwal and others 2012). By controlling receptor expression Gallie (2010) demonstrated that it was possible to control temporally the extent of the ethylene response. To date most research has focused upon *ETR1* as it was the first receptor to be isolated and characterised (Bleecker and others 1988) and the dominant ethylene insensitive mutant *etr1-1* has been shown to delay the onset of leaf senescence and consequentially extend leaf lifespan by approximately 30% (Grbić and Bleecker 1995). An ethylene insensitive phenotype of coriander has been generated by transforming the plant with a mutated version of *ERS1* and this was found to delay significantly leaf and flower senescence (Wang and Kumar 2004). Similarity wheat *W-er1* may be involved in leaf senescence (Ma and Wang 2003).

Whilst the much research effort has focused on the role of ethylene in flower development and climacteric fruit ripening (Serek and others 2006; Agarwal and others 2012) less attention has been given to its impact on crop quality and yield. Given that siliques share similar characteristics with climacteric fruit (Kou and others 2012) and that ethylene is believed to have a central role in pod senescence (Wagstaff and others 2009) the work described in this paper focuses on enhancing our understanding of the role of ethylene perception and signalling in *Arabidopsis* leaf, flower and fruit senescence and the impact of this on seed yield. To help us dissect the role of ethylene signalling we have utilised receptor mutants that exhibit either a gain (*etr1*, *ein4*, *ers2*) or a loss (*etr2*, *ers1*) of function to the gas.

Materials and Methods

Plant material and growth conditions

Arabidopsis (*Arabidopsis thaliana*) ethylene receptor mutants *ein4* (NASC ID: N8053), *ers1* (NASC ID: N3978), *ers2* (NASC ID: N8854), *etr1* (NASC ID: N237) and *etr2* (NASC ID: N657634) plus the background ecotypes they originated from, Columbia (Col-0), Noosen (No) (NASC ID: N3081) and Wassilewskija (Ws) (NASC ID: N22659) were obtained from The European *Arabidopsis* Stock Centre (NASC; Nottingham, UK; <http://arabidopsis.info/>). All plants underwent a 4°C stratification cold treatment for 2-3 days before being transferred into a Fitotron plant growth chamber (Weiss Gallenkamp) set to 20°C, 55% relative humidity with a photon flux density of 200 $\mu\text{mol cm}^{-2} \text{s}^{-1}$ and 16h photoperiod. Plants were grown in individual 90mm diameter pots containing horticultural potting medium (William Sinclair).

Leaf senescence measurements

Plants were harvested at set intervals after the date leaf 6 was tagged and aerial images taken, subsequently leaf 6 was removed from the plant and immediately frozen in liquid nitrogen for chlorophyll *a* and *b* analysis. Chlorophylls were extracted from frozen rosette leaf material, leaf 6, using N,N-dimethylformamide and left for 7 days in the dark at 4°C. The sample absorbance was measured at A_{664} , A_{647} and A_{480} nm using a plate reader (Spectra max 340 PC) and the chlorophyll *a* and *b* concentrations were calculated using the coefficients of Wellburn (1994).

Time course measurements

The time taken for pods on the primary inflorescence to develop was visually scored based upon the following scale (Wagstaff and others 2009), flower anthesis: flowers fully open, petal abscission: petals begun to abscise, pod stage 1: pods are green and fully mature, pod stage 2: pods are 50% yellow, pod stage 3: pods are 100% yellow, pod stage 4: pods are brown and just about to dehisce, pod shatter: pod opened to release some seeds. The total plant lifespan is a measure of the time taken for the plant to go from bolting to the end of flowering.

Physiological measurements

Seed physiological measurements were taken from pods at different stages of development; the numbers of seeds per pod and seed weight per pod were obtained from stage 4 pods

which were fully senescent and about to dehisce. Whereas pod length and area were determined from stage 2 pods which were visually scored as being 50% yellow. Pod area and the number of seeds per pod were established using Image J (NIH Image). For the seed weight per pod and thousand grain weight (TGW) all seeds were oven dried (GENLAB) at 70°C until they reached a constant weight. Total seed yield was determined by weighing the seeds collected from plants grown in aracons which had become fully senescent and reached the end of their life span. The TGW was calculated by taking 100 seeds from the total seed yield experiment, drying them until they reached a constant weight and multiplying this figure by 10 to represent 1000 seeds. The number of pods per plant, height of the primary inflorescence, number of rosette leaves and number of axillary stems were all measured when pod shatter on the primary inflorescence began.

Statistics

Statistical analysis was performed using MINITAB version 14 (Minitab Inc., State College, PA, USA). Means were compared between an ethylene receptor mutant and their wildtype ecotype (Col, No or Ws) using a two-sample t test with a significance level of $P = 0.05$. Data are presented as % relative to the parent, the results for the wildtype ecotype were set at 100% and the percentage difference between the ethylene receptor mutants and their corresponding wildtype ecotype is presented.

Results

Leaf development in ethylene signalling mutants

It is well documented that exposure of leaf material to ethylene accelerates senescence. In this programme of research the time course of leaf senescence was studied in ethylene perception and signalling mutants in comparison to their wild type (WT) ecotypes. The genotypes studied were three gain of function mutants (*etr1*, *ers2* and *ein4*) that exhibit insensitivity to the gas as revealed by their inability to exhibit a triple response when exposed to the gas (Table 1), and two loss of function mutants where expression of the gene encoding the ethylene receptor *ETR2* or *ERS1* (*etr2* and *ers1*) are substantially down-regulated (Alonso and others 2003). Leaf senescence was measured in an individual organ by tagging leaf six from the time of emergence through to end of life and measuring chlorophyll content every six days over a 48 day period (Fig. 1A). To determine a more global senescence profile whole

rosettes were assessed over the same timeframe (Fig. 1B). Individual leaf 6 of *etr1* plants exhibited a delayed onset of senescence (Fig. 1A), with chlorophyll sustained in the middle part of leaf development (12-30 days after emergence) and a similar observation was seen in the rosette (Fig. 1B) which remained fully green at least six days longer than the WT and senescence, when it did occur, was primarily noticeable at the leaf tips. Global leaf lifespan was prolonged in the *ers2* mutant (Fig. 1A & B) and this was conferred by maintenance of chlorophyll in the early stages of leaf life (2-12 days after emergence) although once senescence had been induced this took place faster than the WT (Fig. 1A). The whole rosette of *etr2* plants appeared to retain chlorophyll for longer than WT (Fig. 1B), although leaf 6 did not exhibit delayed senescence (Fig. 1A). Individual leaves of the *ein4* mutant lost chlorophyll more rapidly than WT plants at the very end of senescence (48 days in Fig. 1B) although the rate of global senescence of the rosette may have been delayed at the earlier stages in relation to the parental ecotype. Clearly the plants were under stress since they appeared to accumulate anthocyanins overlying the remaining chlorophyll at days 36-42 more than the other mutant genotypes, in line with the anthocyanin accumulation apparent in the WT in the same time period.

Flower and pod development in ethylene signalling mutants

The time course of development of reproductive structures in *Arabidopsis* plants was investigated in the different mutants by tagging individual flowers on the main inflorescence and then following them through from petal abscission through to pod dehiscence (Fig. 2). Petals of *etr1* flowers were retained for almost twice as long as WT flowers and senescence of the pods in this mutant was also delayed and as a consequence the time from flowering to pod shatter of an individual *etr1* pod was approximately 40% longer than in WT plants. *ers2* pods also displayed a significantly extended life span primarily as a consequence of a delay in the final stages of pod senescence although the timing of petal abscission was also somewhat delayed compared to WT. *ein4* exhibited similar features to *etr2* exhibiting a 10% extension in pod life span compared to WT primarily as a consequence of a more prolonged period of time spent at S3 when pod senescence was taking place (Fig. 2).

Whole plant development and seed yield in ethylene signalling mutants

The height of the primary inflorescence is significantly reduced in *etr1* plants with the stature of the mutant attaining only 60% of WT plants (Fig. 3A). Furthermore, the lifespan of *etr1* plants is also substantially curtailed (Fig. 3B). In contrast, *ers1* plants significantly exceed the

height of their parental WT lines by approximately 10% (Fig. 3A) but their lifespan was unaffected (Fig. 3B). The height and lifespan of the other mutants studied was indistinguishable from their ecotype controls.

Although it is evident that mutations in ethylene receptor signalling can attenuate the development of leaves, flowers and siliques in *Arabidopsis* what is even more critical is the impact that these changes have on plant yield. We therefore carried out a detailed analysis of the total seed yield per plant and the individual components that contributes to this parameter. Overall, the mutants *etr1*, *ein4*, *ers1*, and *ers2* have a significantly deleterious impact on total weight of seed produced per plant compared to WT plants (Fig. 4D). For all 4 mutants the reduction was approximately 40%. Only *etr2* showed a significant increase, of approaching 20% in total yield of seed (Fig. 4D). Although the majority of mutants exhibited reduced yield, thousand grain weight (TGW) was significantly higher, by as much as 20%, in *ein4*, *etr1* and *etr2* (Fig. 4C). These three mutants retained significantly fewer seeds per pod as did both *ers1* and *ers2* albeit to a smaller degree compared to their parental WT lines (Fig. 4A). In general, the mutants produced a similar total number of pods per plant with the exception of *ers1* that generated approaching twice as many pods as WT (Fig. 4B).

Discussion

Although it is well documented that exposure of plants to exogenous ethylene can influence both the timing and rate of organ senescence the role of endogenous levels of the gas in regulating the process in vivo is less well defined (Graham and others 2012). One strategy that has been adopted to explore this phenomenon is to block ethylene action using inhibitors such as 1-MCP (Sisler and Serek 1997) or silver ions (McDaniel and Binder 2012). Whilst these approaches have generated valuable data the impact of inhibitors on other cellular events is always open to question and in this study we have explored the use of ethylene receptor mutants to dissect the role of the gas in regulating *Arabidopsis* development. In particular we have studied the progression of leaf, flower and pod senescence and correlated these events with seed yield. We have previously shown that senescence in *Arabidopsis* is regulated in unique ways in these three organs (Wagstaff and others 2009) and that ethylene biosynthesis and binding appeared to be more important in silique and petal senescence than in leaves, although some elements were conserved in all three tissue types examined. The family of *Arabidopsis* ethylene receptors comprises five members and because of the nature

of the signalling process mutations in the ethylene binding domain can generate dominant mutants such as *etr1*, *ein4* and *ers2* that are blind to the gas (Binder and others 2012). We have studied these three gain of function mutants in detail and also the two loss of function mutants, *ers1* and *etr2* where, due to a T-DNA insertion in the respective genes, we have been able to probe the specific contributions of *ERS1* and *ETR2* to plant senescence and the results are summarised in Fig 5.

Reports of delayed senescence in ethylene mutants and transgenics generally relate to flower longevity (Jones, 2008) or individual leaf longevity (Grbic and Bleecker, 1995; Wang and Kumar, 2004). Our data show that the senescence pattern of an individual leaf (leaf 6) was changed in *etr1*, *etr2* and *ers2* mutants. In *etr1* and *ers2* this also resulted in delayed senescence of the entire rosette, but this was not the case for *etr2* which began the onset of senescence at a similar time to the WT plants but the rate of senescence was reduced. There is a question of how functional the observed leaf 'stay-green' phenotypes are since extended leaf longevity is not necessarily related to plant longevity suggesting that photosynthates are not available for remobilization, supporting evidence from Grbic and Bleecker (1995) that leaves functionally senesced even though the apparent life span of the leaf was prolonged. Re-allocation of resources around the plant did not always occur in proportion to the photosynthetic period; *ers2* maintained leaf chlorophyll at a time when senescence in the WT had started, but senescence occurred extremely rapidly in the mutant once it was triggered.

There is a substantial body of evidence that ethylene can accelerate the onset of flower abscission although the gas is not critical for organ shedding to take place (Roberts and others 2002). In accord with these observations and reports by others (Patterson and Bleecker 2004) whilst the timing of petal abscission was delayed in *etr1* plants shedding ultimately took place. It has previously been shown that the introduction of the mutated version of *ETR1* into other species can delay both flower senescence and abscission (Yang and others 2008). Interestingly the timing of shedding in *ein4*, that has also been classified as ethylene insensitive on the basis of its inability to undergo the triple response, was indistinguishable from WT. This observation provides clear evidence that the different receptor family members have different signaling capacities and that conclusions about the roles of ethylene in a particular developmental process based on the phenotypes of individual mutants should be interpreted with caution.

Pods from *ers2* plants underwent shatter significantly later than WT primarily as a consequence of the final stages of senescence being prolonged. Pod dehiscence is an important event in the regulation of yield in species such as *Brassica napus* (Roberts and

others 2002) and this observation may be of value in identifying strategies to reduce premature shatter. A slower rate of pod senescence was observed in both *etr2* and *ein4* plants which spent longer at stage 3, mimicking the pattern observed in individual leaves, but the total reproductive period was not extended in either genotype. In *etr1* the longevity of individual flowers and pods was also increased, but the reproductive period was significantly shorter than WT. In contrast to leaves, it would appear that in *etr1*, *etr2* and *ein4* the extended pod green period is potentially functional, or that resource partitioning was affected by the mutations, since there was a significant increase in TGW, but overall seed yield was reduced due to fewer seeds per pod. These data support an important role for local photosynthesis of the pod in seed filling. The RNAi-driven down-regulation of *ETR2* in rice has also been reported to increase TGW but overall plant seed yield or seeds per panicle was not recorded (Wuriyangan and others 2009). The same authors reported an early onset of flowering in the rice *etr2* mutants, suggesting that the life cycle was accelerated; in our hands the only *Arabidopsis* mutant with an accelerated life cycle was *etr1*. The curtailed life span in *etr1* plants correlates with both a significantly reduced plant height and a small reduction in total number of pods. This indicates that ethylene signaling through ETR1 is important to maintain the 'normal' developmental sequence in *Arabidopsis* and a failure to achieve this results in a substantial decline in seed yield.

The *ers1* mutant produces more pods and is taller, thereby making more reproductive positions available during its life cycle compared to the WT. Although the mutant produces a similar number of seeds per pod as WT both TGW and total yield is reduced indicating that the seed content is reduced. Previous studies have shown that ethylene-insensitive mutants of *Arabidopsis* exhibit poor germination in relation to the WT (Bleecker and others 1988), perhaps indicating a reduction of available assimilates, and conversely the over-producing mutant *eto1* has accelerated germination (Cheng and others 2009). Global reproductive lifespan is unaffected in *ein4*, but TGW is increased compared to WT which was shown to be achieved through investing in fewer, heavier seeds per pod although the absolute mass of seeds per plant was reduced. Like *etr1*, *ers2* has a foreshortened period where the pod is fully photosynthetic but this is compensated for by spending longer in the late stages of pod development. Plant overall seed yield in *ers2* is also reduced but there is no change in TGW, indicating that the late produced seeds are less well filled. The *etr2* mutant maintains seed filling throughout the reproductive lifespan, resulting in an increased TGW and overall plant seed yield of nearly 20% above WT values in both cases.

A possible mechanism to explain the organ-specific differences observed between the different mutants was first proposed by Klee (2004) who suggested that the concentration of receptors varies between tissues, enabling the plant to have differential temporal and spatial sensitivity to ethylene. Receptors are also known to form dimer complexes (Liu and Wen, 2012); thus mutating one half of a dimer could impact on the function of the other partner in the complex and make teasing apart the functions of individual genes rather more difficult, but also provides a tool by which ethylene responses can be repressed to different extents. The results presented in this paper clearly support a role for individual family members to contribute to the perception of ethylene at different developmental stages and supports the hypothesis that responses may be tissue specific as asserted by Shakeel and others (2013). Others have noted unique roles for particular isoforms; *ETR1* has long been associated with delayed leaf senescence and larger leaves (Grbić and Bleeker, 1995), but more recently has been shown to be required for nutation of seedlings (Binder and others, 2006). *ETR2* is essential for correct trichome branching, which is controlled by microtubule assembly (Plett and others, 2009). *ETR1*, *ETR2* and *EIN4* are all important for growth recovery following exposure to ethylene (Binder and others, 2004), however the *ers1* mutant appears to cause growth inhibition in low ethylene environments (Liu and others, 2010); in the present study we showed that this is not necessarily correlated with low yield.

The controlled induction of mutant ethylene signalling phenotypes could have implications for crop production in the future. Maintenance of green leaf colour in transgenic coriander using the mutated version of the *ERS1* gene has already been shown to be beneficial for the production of leafy crops (Wang and Kumar 2004) and transformation of tomatoes with an inducible version of the ethylene insensitive *ETR1* gene provides a strategy for regulating fruit ripening (Gallie 2010). Our observations indicate that the timing and duration of organ senescence can be manipulated by attenuating ethylene perception and signaling and that this could enhance leaf, flower and pod longevity and have consequences for seed quantity and possibly quality. Further work will be necessary to develop a strategy in individual crop species that might lead to beneficial phenotypes for both the producer and consumer.

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Figure legends

Figure 1

Leaf chlorophyll content during lifespan. (A) Chlorophyll content of leaf 6 from time of emergence until full senescence (n=3). Wild type lines are shown next to the relevant mutants. (B) shows rosette leaf senescence in the mutants compared to the wild type plants. (G) indicates a gain of function mutant; (L) indicates a loss-of-function mutant. Parental ecotypes are indicated as follows: Columbia-0 (Col), Wassilewskija (Ws) and Nossen (No).

Figure 2

Ethylene mutant reproductive development time course. Time spent at each stage of development is shown with anthesis taken as time zero. Pod stage 1 = Pods are green and fully mature, pod stage 2 = pods are 50% yellow, pod stage 3 = pods are 100% yellow, pods stage 4 = pods are brown and just about to dehisce, n=16. Bars represent the mean \pm SE and those which are statistically significant from their parent line (Col, No or Ws) are indicated by an asterisk, * P < 0.05, ** P < 0.01, *** P < 0.001. (G) indicates a gain of function mutant; (L) indicates a loss-of-function mutant.

Figure 3

Whole plant development (A) Height of the primary inflorescence compared to the relevant WT plant (B) Reproductive lifespan; time from bolt emergence to the end of pod senescence. Bars indicate SEM from n=5. Asterisks indicate lines which are significantly different from their parent line (Col, No or Ws), * P < 0.05, ** P < 0.01, *** P < 0.001. (G) indicates a gain of function mutant; (L) indicates a loss-of-function mutant.

Figure 4

Yield parameters from ethylene mutants.(A) Number of seeds per pod (B) Number of pods per plant (C) Thousand Grain Weight (D) Total mass of seed yield per plant. All values are expressed as a % of the relevant WT line with the original units of measurement indicated on the y axis. Bars indicate SEM from n=5. Asterisks indicate lines which are significantly different from their parent line (Col, No or Ws), * P < 0.05, ** P < 0.01, *** P < 0.001. (G) indicates a gain of function mutant; (L) indicates a loss-of-function mutant.

Figure 5

Interactions between developmental and yield parameters. Key to figure as follows: (ND) No sig difference from parent; (---)*** sig difference from parent in minus direction; (--) ** sig difference from parent in minus direction; (-) * sig difference from parent in minus direction; (+++) *** sig difference from parent in positive direction; (++) ** sig difference

from parent in positive direction; (+) * sig difference from parent in positive direction. Blue shading indicates a negative interaction, red shading indicates a positive interaction.

Table 1

Ethylene receptor mutants used in the study along with their corresponding backgrounds and location of the mutation.