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## Conditional Gene Expression and RNAi Using MEC-8-Dependent Splicing in *C. elegans*

Andrea Calixto<sup>1</sup>, Charles Ma<sup>1,2</sup>, and Martin Chalfie<sup>1,\*</sup>

<sup>1</sup>Department of Biological Sciences, Columbia University New York, NY 10027

### Summary

We describe a method for conditional regulation of gene expression based on the processing of an intron cassette by a splicing factor. The RNA processing factor MEC-8 is necessary for the function of the *Caenorhabditis elegans* touch receptor neurons; *mec-8* mutants are touch insensitive. We show here that this insensitivity involves the loss of MEC-8-dependent splicing of *mec-2*, which encodes a component of the mechanosensory transduction complex. MEC-8 is needed to remove the ninth intron in *mec-2* pre-mRNA to form the longest of three mRNAs, *mec-2a*. Without MEC-8, splicing causes the termination of the transcript. Inclusion of *mec-2* intron 9 is sufficient to convey *mec-8*-dependent regulation on other genes and, in *mec-8(u218ts)* mutants, resulted in their temperature-dependent expression. Because *mec-8* is expressed ubiquitously in embryos and extensively in larvae, this system should produce temperature-sensitive expression for most genes. We report a strain that exhibits temperature-dependent RNA interference.

### Introduction

Conditional expression of gene activity provides a means to examine gene function that is restricted in time or cell type, and is particularly useful when complete elimination of gene activity leads to lethality. Several methods produce conditional gene expression. These methods include the Cre-*lox* system for inducible gene targeting in mice<sup>1</sup>, a heat-induced signal to target protein degradation in yeast<sup>2</sup>, a tetracycline-responsive promoter in mammalian cells<sup>3</sup>, and three heat-shock based systems for regulated gene expression in *Caenorhabditis elegans*<sup>4, 5, 6</sup>.

One of the first means of obtaining conditional gene expression was the use of temperature-sensitive mutations. Classic examples include studies of essential genes in *Neurospora*<sup>7</sup> and of morphogenesis in the bacteriophage T4 (ref. 8). The usefulness of such mutations is limited, however, by their relative rarity; temperature-sensitive alleles do not exist for most

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\*To whom correspondence should be sent at: Department of Biological Sciences, 1012 Fairchild, MC#2446, Columbia University, 1212 Amsterdam Avenue, New York, NY 10027, Phone: 212-854-8870, Fax: 212-865-8246, mc21@columbia.edu.

<sup>2</sup>Present address: Phalanx Biotech Group, 1400 Page Mill Road, Building B, Palo Alto, CA 94304-1124

### Author contributions

A.C., C.M. and M.C. designed experiments; A.C. performed the experiments; A.C. and M.C. wrote the manuscript.

genes. For example, in our own work identifying mutations affecting touch sensitivity in *C. elegans*, only seven of eighteen mapped genes (affected by 418 mutations) had temperature-sensitive alleles<sup>9</sup>.

In this paper we describe a method to generate splicing-dependent and temperature-sensitive expression of virtually any *C. elegans* gene for which a loss-of-function allele is available, using temperature sensitive MEC-8-dependent splicing. Unlike previous *C. elegans* methods<sup>4, 5, 6</sup>, which require heat shock to allow gene expression, our method employs a temperature shift at normal growth temperatures. The *mec-8* gene is needed for touch sensitivity in *C. elegans*<sup>10</sup>. In addition, the MEC-8 protein, which has two RNA recognition motifs (RRMs), is required for the processing of the *unc-52* transcript in the hypodermis<sup>11</sup>. We sought potential targets of *mec-8* regulation among the genes needed for touch sensitivity. We found that MEC-8 is needed for the removal of the ninth intron of *mec-2*, which encodes a subunit of the channel complex needed for mechanosensory transduction<sup>12, 13</sup>. This discovery enabled us to generate a system that results in the *mec-8*-dependent expression of many *C. elegans* genes. Moreover, using the *mec-8(u218ts)* allele, which produces a temperature-sensitive phenotype<sup>9</sup>, we can express virtually any gene containing *mec-2* intron 9 in a temperature-dependent manner.

## Results

### MEC-8 regulates alternative splicing of *mec-2*

The expression of *mec-8* begins in the 50-cell embryo and is ubiquitous. As the animals develop, *mec-8* expression persists in the hypodermis, intestine, the six touch receptor neurons (TRNs), chemosensory neurons of the head and tail, and in vulval nuclei (ref. 14; and our unpublished observations).

To identify *mec-8* targets in the TRNs, where it is known to act cell-autonomously, we looked for differences in RT-PCR products of genes needed for touch sensitivity (listed in Methods) using mRNA from wild-type and *mec-8* mutants. The only cDNAs that were differentially amplified were those of *mec-2* (Fig. 1). The *mec-2* gene produces three mRNAs, as identified by RT-PCR ([www.wormbase.org](http://www.wormbase.org)). The largest (*mec-2a*: 1.4 kb) is formed from 13 exons<sup>12</sup> (Fig. 2a). We could amplify *mec-2a* cDNA from wild type, but not from worms homozygous for the *mec-8* null mutations *u314* or *e398* (Fig. 1a). The *mec-2a* mRNA was made, however, in *mec-8* mutants rescued with *mec-8(+)* expressed in the TRNs (Fig. 1b). The *mec-2b* mRNA has exons 1 – 9 of *mec-2a* but lacks exons 10 – 13, containing instead a small exon (exon 9') at its 3' end. The inclusion of exon 9' adds an in-frame stop codon and a consensus polyadenylation site to the mRNA. The *mec-2c* mRNA begins with an exon within intron 2 (exon 2') and continues with exons 3 – 9' as in *mec-2b*. Using cDNA-specific primers, we amplified *mec-2b* and *mec-2c* cDNAs in wild type and the two *mec-8* null strains.

The three *mec-2* mRNAs encode polypeptides with a prohibitin homology (PHB) domain preceded by a hydrophobic region that inserts the protein into the cytoplasmic leaflet of the plasma membrane. N- and C-termini surround the hydrophobic and PHB domains in

MEC-2A (Fig. 2a). MEC-2B has the N-terminal but not the C-terminal domain, and MEC-2C lacks both domains (Fig. 2a).

We had previously generated antibodies against the N terminus of MEC-2A (M2N) and against the PHB domain through the C-terminus of MEC-2A (M2C). We stained wild type and *mec-8* mutants with the M2N and M2C antibodies and, as expected from the RT-PCR results, the M2N antibody labeled TRNs in wild-type and *mec-8* worms; M2C antibody labeled only wild type TRNs (Fig. 2b). Since the antibodies were purified by preabsorption to extracts of *mec-2* null mutants and since *C. elegans* has several PHB domain proteins<sup>15</sup>, the M2C antibody is likely to recognize only the C-terminus of MEC-2A, which is missing in *mec-8* mutants.

### ***mec-2* intron 9 is sufficient for *mec-8*-dependent splicing**

To test whether *mec-2* intron 9 contains all the necessary elements for conferring *mec-8* dependency, we inserted *mec-2* intron 9 before the coding sequence for Yellow Fluorescent Protein (YFP). We expressed *intron 9::yfp* in TRNs using the *mec-18* promoter (Fig. 3a). No YFP fluorescence was observed in *mec-8(e398)* or *mec-8(u314)* worms transformed with this construct (*P<sub>mec-18</sub>intron 9::yfp*). In contrast, fluorescence was seen in all six TRNs in the *mec-8/+* progeny of transgenic worms when they were crossed with wild-type males (Fig. 3b). These results indicate that YFP expression was dependent on MEC-8.

The *mec-8*-dependent splicing of *mec-2* intron 9 resulted in the temperature-dependent production of YFP fluorescence in worms with the temperature-sensitive *mec-8* allele *u218* (ref. 10; Fig. 3c). The *u218* mutation produces an Ala278Thr change in the second RRM16. *mec-8(u218ts)* mutants are wild type at 15° C and touch insensitive (Mec) at 25° C. *u218* mutants transformed with *P<sub>mec-18</sub>intron 9::yfp* have fluorescent TRNs at 15° C, but not at 25° C (Fig. 3c). We also assessed the production of *yfp*-containing messages by RT-PCR of mRNA from *u218* mutants expressing *P<sub>mec-18</sub>intron 9::yfp* grown at different temperatures. Amplifiable material representing the splice from exon 9 to the *yfp* coding region was detected in worms grown at 15° C but not at 25° C (Fig. 3d). These results demonstrate that the control of MEC-8 over intron 9 splicing is very tight at the non-permissive temperature.

To show that *mec-8* and intron 9 of *mec-2* can be used to regulate gene expression in other tissues, we placed *gfp* in the intron 9 cassette driven by promoters specific for neurons (*sng-1*), hypodermis (*dpy-5*) and intestine (*nhx-2*) and injected the constructs into worms of the *mec-8(u218ts)* genotype. Worms expressing *P<sub>sng-1</sub>intron9::gfp* grown at 15° C had fluorescence in the neurons in the head and tail (including the TRNs) where *mec-8* is expressed. Even though *sng-1* is pan-neuronal, *gfp* was not seen in the ventral cord, where *mec-8* is not expressed (ref. 14; and our unpublished observations). Worms grown at 25° C did not show fluorescent neurons (Fig. 4a). *mec-8(u218)* expressing *P<sub>dpy-5</sub>intron9::gfp* and *P<sub>nhx-2</sub>intron9::gfp* showed fluorescence in the hypodermis and intestine, respectively, at 15° C but not at 25° C (Fig. 4a). Consistent with these results, the expression of the mRNA for *gfp* under the *sng-1*, *dpy-5* and *nhx-2* was seen only at 15° C and not at 25° C (Fig. 4b). These experiments suggest that gene expression can be made conditional using intron 9 and *mec-8(u218ts)* for genes in many tissues.

## Temperature-sensitive gene expression

The finding that *mec-2 intron 9* can convey *mec-8* dependence suggests that *intron 9* can be used to generate temperature-dependent constructs of potentially any *C. elegans* gene; these constructs can then be transformed into appropriate null backgrounds. Certain characteristics of the *mec-8* gene and the *mec-8(u218ts)* allele suggest that *intron 9* constructs of a gene of interest would mimic the expression of the endogenous gene product. First, suppression studies and temperature-shift experiments suggest that a relatively small amount of the wild-type MEC-8 may be needed for function, i.e., MEC-8 is not rate-limiting. Specifically, a single dose of a tRNA suppressor mutation can suppress an amber allele of *mec-8*, but not amber mutations in other *mec* genes<sup>9, 10</sup>. Second, shifting *mec-8(u218ts)* worms from the permissive to the restrictive temperature at hatching (so that only embryonic MEC-8 is available) results in touch sensitive adults (ref. 9; Fig. 5a) that detectably express YFP (Fig. 5b); thus, *mec-8* displays considerable perdurance. Third, worms shifted from the restrictive to the permissive temperature as late larvae (Fig. 5a and 5b) produce sufficient *mec-8* activity for touch sensitivity and YFP production in adulthood. Indeed, when *mec-8(u218ts)* worms were shifted to the permissive temperature as adults (Fig. 5a and 5b, dotted lines) and scored 48 hours later, they were both touch sensitive and YFP positive, an indication that MEC-8 is produced even after adulthood in the TRNs.

To test whether the use of *intron 9* and *mec-8(u218ts)* could mimic the results of an endogenous temperature-sensitive mutation, we compared *mec-8(u218ts); mec-4(u253)* worms expressing  $P_{mec-4intron9}::mec-4$ , with *mec-4(u45ts)* worms. Although the temperature shift curves of touch sensitivity for *mec-8(u218ts)* and *mec-4(u45ts)* are quite different (Fig. 5a and 5c; ref. 9), worms with the *intron 9::mec-4* construct showed temperature-shift touch-sensitivity curves that were essentially the same as those from *mec-4(u45ts)* worms (compare Fig. 5c and 5d). These results suggest that the timing of *mec-4* expression, but not of *mec-8*, is critical for *mec-4* function in touch sensitivity in adults.

## Temperature-sensitive RNAi

We have used the *mec-2 intron 9* to produce temperature-dependent RNA interference (RNAi). We took advantage of the fact that RDE-1, a *C. elegans* Argonau protein, is required for RNAi<sup>17</sup>. Transformation of *mec-8(u218ts); rde-1(ne219)* worms (*ne219* is a null allele) with  $P_{rde-1intron9}::rde-1(+)$  resulted in worms that were responsive to RNAi at 15° C but not at 25° C (Table 1). We observed the relevant temperature-sensitive RNAi phenotype using bacteria making dsRNA for *unc-22* (expressed in muscle), *unc-52* (expressed in the hypodermis) and *rpl-3* (needed for embryonic viability), indicating that the RNAi effects can be detected in many tissues. To test how quickly the RNAi phenotype could be detected, we fed newly hatched *intron9::rde-1* worms with bacteria making dsRNA for *unc-22* at 25° C for 24 hours and then shifted them to 15° C. The *unc-22* twitcher phenotype was seen within 3 hours of the shift (Fig. 6). In comparison, worms placed on the bacteria at the time of the shift did not show a phenotype for at least 24 hours. Since RDE-1 is thought to act as part of the RISC complex<sup>18</sup>, worms presumably load with dsRNA at the restrictive temperature but cannot execute RNAi. Switching to the permissive temperature

allows RNAi to proceed, thus making this method particularly useful for the study of late effects of genes whose loss is lethal.

## Discussion

We have developed a method to produce conditional mutants that depends on the correct processing of their transcripts by MEC-8. Taking advantage of the existence of a temperature-sensitive allele of *mec-8*, researchers can now produce tightly controlled and temperature-sensitive gene expression for virtually any gene in *C. elegans*. A further elaboration of this method was the generation of a strain that allows temperature-sensitive expression of RNAi.

Previous work by Lundquist *et al.*<sup>11</sup> found that *mec-8* facilitated exon skipping in *unc-52* mRNAs, allowing splicing from exon 15 to exon 19 and exon 16 to exon 19. Similarly, our data show that *mec-8* is required for the skipping of exon 9' in *mec-2*, allowing the production of *mec-2a* mRNA. Since the C terminus of MEC-2A is needed for enhancing MEC-4 channel activity<sup>19</sup>, the loss of *mec-2a* mRNA underlies, at least in part, the touch-insensitive phenotype of *mec-8* mutants. At this time we do not know whether MEC-8 regulates *mec-2* processing by enhancing the exon 9–10 splice or inhibiting the exon 9–9' splice. Based on examples of alternative splicing regulated by other RRM-containing proteins<sup>20, 21</sup>, however, we suggest that MEC-8 prevents the splice to exon 9'.

We have not tried to determine the minimal sequences needed for *mec-8*-dependent splicing of *intron 9*. The smallest number of exonic nucleotides in our constructs was 18 on each side (see Methods, TU#839). Before publication we distributed our constructs to Poon *et al.*<sup>22</sup> who truncated the construct further (with 13 nucleotides at the 5' end and 15 at the 5' end). Their construct appeared to show a modest cold-sensitive expression (instead of heat-sensitive expression) in ventral cord neurons (which do not detectably express *mec-8*). Their modification appears to modify the splicing of *intron 9*.

Several characteristics of *mec-8* contribute to the usefulness of this conditional system. First, many *mec-8* mutants are viable at all growth temperatures. Second, a small amount of MEC-8 is sufficient for its function. Third, *mec-8* is expressed in almost every cell and from the early embryo into adulthood. This last property allows conditional expression of the wild-type product from *mec-8*-dependent constructs at a wide variety of times using temperature shifts. Fourth, the use of *mec-8* mutants and different promoters to drive expression of both *mec-8(+)* and *intron 9*-containing genes, provides a means of producing both spatial and temporal regulation of expression. For example, the use of different promoters with minimally overlapping expression patterns can permit cell-specific splicing. We have used such combinatorial expression to label specific cells with recGFP<sup>23</sup> and to kill them with recCaspase<sup>24</sup>. *Intron 9* and *mec-8(+)* would allow a similar recSplicing system to be used in *C. elegans*. For example, the use of a heat-shock promoter to drive the expression of an *Intron 9*-containing gene and a cell-specific promoter to drive the expression of *mec-8(+)* in a *mec-8* mutant would result in temporal regulation in selective cells.

Since little MEC-8 is needed for touch sensitivity, a potential problem with using *intron 9* and *mec-8(u218ts)* is that perdurance of MEC-8 may obscure the results of shifting from the permissive to the restrictive temperature. Nonetheless, the effect appears minimal, and could be potentially corrected by using a rapidly-degraded MEC-8, such as one with an attached RING domain from an E3 ubiquitin ligase<sup>25</sup>. Another potential problem is that comparisons are made between animals with and without *mec-8* activity. For this reason, care must be taken to show that the *mec-8* state does not affect the activity being studied.

Two other systems have used alternative splicing to control gene expression. The first relied on the temperature-dependent release from a block in splicing of an artificial intron to allow gene expression in yeast<sup>26</sup>. This method required transitions from 16° C to 36° C and resulted in 15–20% of the wild-type activity. The second method utilized sex-specific splicing of *transformer* in *Drosophila* to express genes only in females<sup>27</sup>. The method we describe here differs from these previous methods in that it does not require temperature extremes, appears to be highly efficient and tightly controlled, and is not restricted to one sex. In addition, the *mec-8* system can be rapidly activated (within three hours at 15° C), whereas the artificial yeast intron system required growth for 3 days at 36° C.

We have not tested whether this system can be used in organisms other than *C. elegans*. Other splicing regulators that have identified targets, however, could be used to develop similar regulatory control in other animals and in cells in culture. Such components include *Drosophila* ELAV, for which temperature-sensitive alleles exist<sup>28</sup>, and either *ewg* intron 6 (ref. 29) or the *nrg* terminal neuronal intron<sup>20</sup>; and three sets of mammalian components, FOX-1/2 and the alternative neuronal intron of the Calcitonin/CGRP gene<sup>21</sup>, Nova-1 and clusters of the consensus binding sequence UCAY<sup>30</sup>, and ETR-3 and cardiac troponin T exon 5 (ref. 31).

## Methods

### *C. elegans* growth and strains

Wild-type *C. elegans* (N2) and strains with mutations in *mec-8(u314, e398, or u218ts)I*, *rde-1(ne219)V*, or *mec-4(u45ts or u253)X9, 10, 17*, were usually grown at 20° C as previously described<sup>32</sup>. For temperature sensitivity experiments, worms were tested after growth for several generations at either 15° C or 25° C.

### Detection of *mec-2* transcripts

RT-PCR was used to detect transcript differences between wild type and *mec-8* mutants for the following genes: *mec-1, mec-2, mec-3, mec-4, mec-5, mec-6, mec-7, mec-8, mec-9, mec-10, mec-12, mec-17, and mec-18*. The presence of *mec-2a, mec-2b, and mec-2c* was tested using RT-PCR and RACE using the SMART RACE Amplification Kit (Clontech). Total RNA was isolated with TRIreagent (Sigma) and poly-A RNA with the Oligotex mRNA Maxi Kit (Qiagen). All the primers used in our experiments are listed in Supplementary Table 1. RACE products were TA cloned, sequenced and analyzed using Sequencher 4.5 (Gene Codes Corporation). We were not able to see *mec-2c* in our RACE experiments, probably because it is underrepresented compared with *mec-2a* and *mec-2b*.

## Expression constructs and transformation

Three DNA fragments containing the 1.8 kb sequence of *mec-2 intron 9* and different amounts of flanking exon codons were amplified by PCR from the plasmid DT#94 (ref. 12), which contains the full genomic sequence of *mec-2*, using primers that introduced 5' and 3' *Bam*HI sites. The resulting PCR products were cut with *Bam*HI and cloned between the *mec-18* promoter and the *yfp* coding sequence in TU#739 (ref. 24) to generate TU#821 [*P<sub>mec-18</sub>intron 9::yfp(fragment I)*], TU#838 [*P<sub>mec-18</sub>intron 9::yfp(fragment II)*], and TU#839 [*P<sub>mec-18</sub>intron 9::yfp(fragment III)*] or placed between the promoter and the genomic coding sequence of *rde-1* [TU#820, *P<sub>rde-1</sub>intron 9(fragment II)::rde-1*] or between promoter and the genomic coding sequence of *mec-4* [TU#837, *P<sub>mec-4</sub>intron 9(fragment II)::mec-4*] in Fire vector pPD95.75 ([www.ciwemb.edu/pages/firelab.html](http://www.ciwemb.edu/pages/firelab.html)). *rde-1* was amplified from genomic DNA with primers including *Pml*I/ *Bam*HI sites for 2.1 kb of promoter region and *Bam*HI/ *Apa*I sites for the coding sequence.

The *mec-4* gene was amplified in two parts from genomic DNA TU#12 (ref. 33) using primers including *Hind*III/ *Xba*I sites for the 823 bp promoter region and *Bam*HI/ *Eag*I sites for the coding sequence. Tissue-specific constructs were made by inserting the *intron 9* cassette (fragment III, cut *Bam*HI from TU#839) into pPD95.75 creating TU#871. 2 kb of promoter sequences of *sng-1*, *dpy-5* and *nhx-2* were cloned into plasmid TU#871 using primers that created *Pst*I sites for *P<sub>sng-1</sub>*, *Sph*I/ *Xba*I sites for *P<sub>dpy-5</sub>*; and *Sph*I/ *Xba*I sites for *P<sub>nhx-2</sub>*. The insertion of the *mec-2* sequences before *yfp* (fragments I and III) or *gfp* (fragment III) introduced several amino acids to YFP and GFP (52 for fragment I and 6 for fragment III). Fragment II, which was inserted before the genomic sequences for *rde-1* and *mec-4*, did not introduce a new start codon, so their translation start was unaltered.

We generated transgenic worms by microinjection<sup>34</sup>. The injection mix contained 10– 40 ng/μl of one of the intron 9 constructs; combined with either 40 ng/μl of the dominant Roller marker plasmid<sup>34</sup> (for the YFP vector), 20 ng/μl of pCW2.1 (a *ceh-22::gfp* plasmid; ref. 35) for the *rde-1* and *yfp* plasmid, 20 ng/μl of *P<sub>myo-3gfp</sub>* for the *mec-4* plasmid or 10 ng/μl of *P<sub>mec-3dsred</sub>* (a gift from I. Topalidou) for *sng-1*, *dpy-5* and *nhx-2* plasmids. pBSK (Stratagene) plasmid was used as filling DNA to a final concentration of 100 ng/μl for all injection mixes. At least 5 stable lines were generated for each construct and all behaved similarly.

## Detection of MEC-8-dependent splicing

Production of *mec-2 exon-9::yfp* or *mec-2 exon-9::gfp* mRNA at 15° C and 25° C was assessed by RT-PCR using primers for the last 18 nucleotides of *mec-2* exon 9 and for 863 nucleotides of *yfp* or *gfp*. The reverse primer amplifies both *yfp* and *gfp* messages, since it does not target the regions of divergence.

## Immunocytochemistry

Whole-mount immunocytochemistry was carried out as in Ruvkun *et al.*<sup>36</sup>. We used acetone powder-purified MEC-2 antibody (ref. 37; 1:1000) and rhodamine-conjugated goat anti-rabbit IgG (H+L) antibody (Jackson ImmunoResearch Laboratories, Inc., West Grove, PA;

1:1000). Both MEC-2 antibodies were used in parallel and gave the same results in wild type. MEC-2 antibody-staining pattern was scored in blind tests.

### Phenotype characterization

Temperature shifts and YFP expression: Worms synchronized at hatching were grown at the appropriate temperatures for at least two generations. They were shifted from 15° C to 25° C or 25° C to 15° C at the indicated times, and tested for touch sensitivity and/or YFP expression as egg-laying adults. YFP fluorescence was observed using a Zeiss Axiophot II or a Leica stereodissecting microscope equipped for fluorescence. DIC optics were used to observe the TRNs when YFP was absent.

RNAi sensitivity: RNAi responses were tested by growth on bacteria making dsRNA for *unc-22*, *unc-52*, or *rpl-3* according to Timmons and Fire38. All RNAi experiments were performed in triplicates in at least 3 independent experiments on different days. Thirty to fifty worms were used and scored in each plate. Temperature-sensitive mutants were grown at 25° C or 15° C for more than two generations prior to the experiments. The scoring was performed blindly.

The time course of RNAi induction was determined using worms grown at 25° C in the presence of *unc-22* dsRNA expressing bacteria or OP50 (for three generations) and were synchronized and re-fed as L1 larvae in the desired condition. After 24 hrs, worms were moved to 15° C and scored for the *unc-22* twitcher phenotype at intervals of 3 hrs for the first day, 12 hrs for the second day, and 24 hrs for subsequent days.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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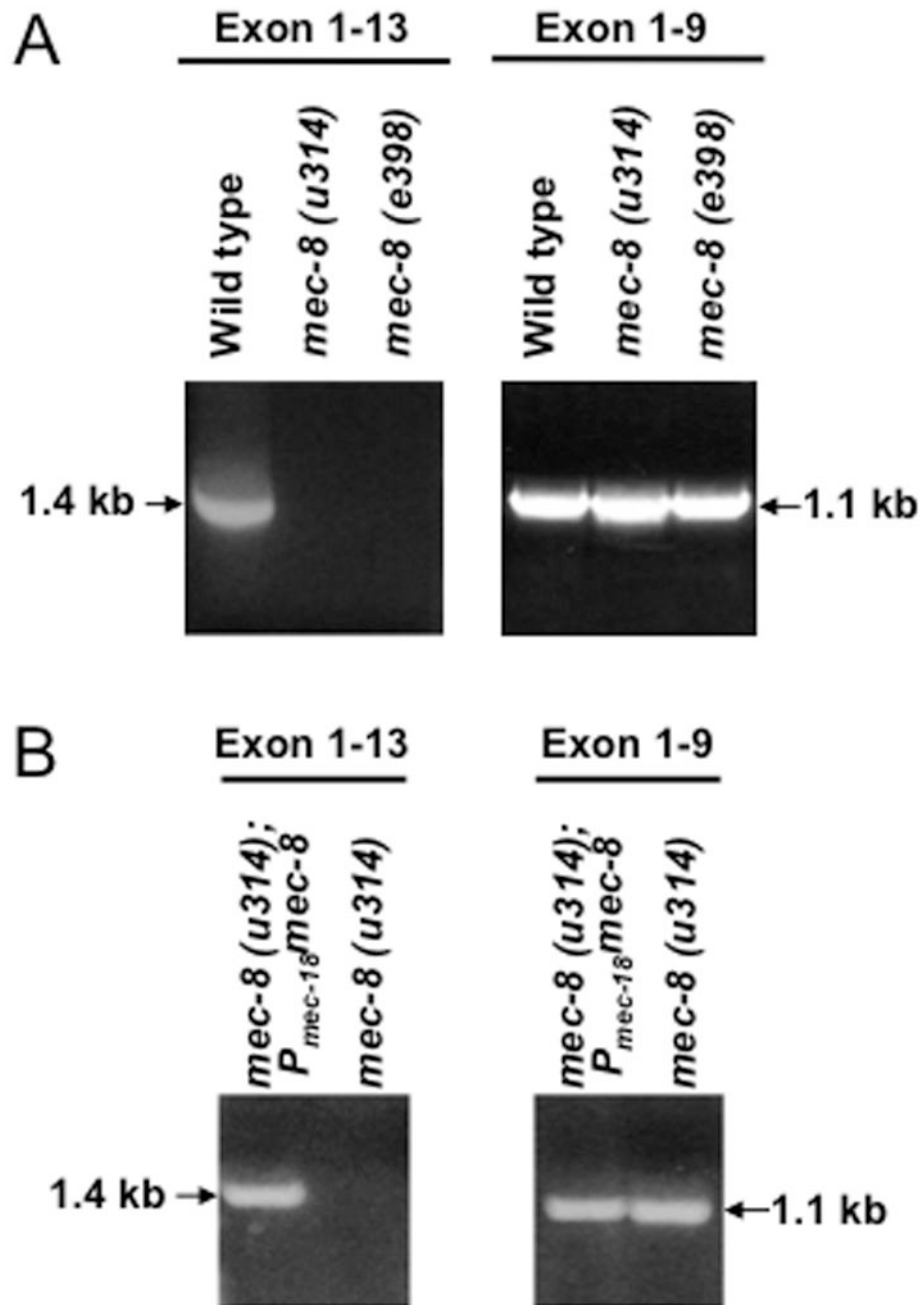
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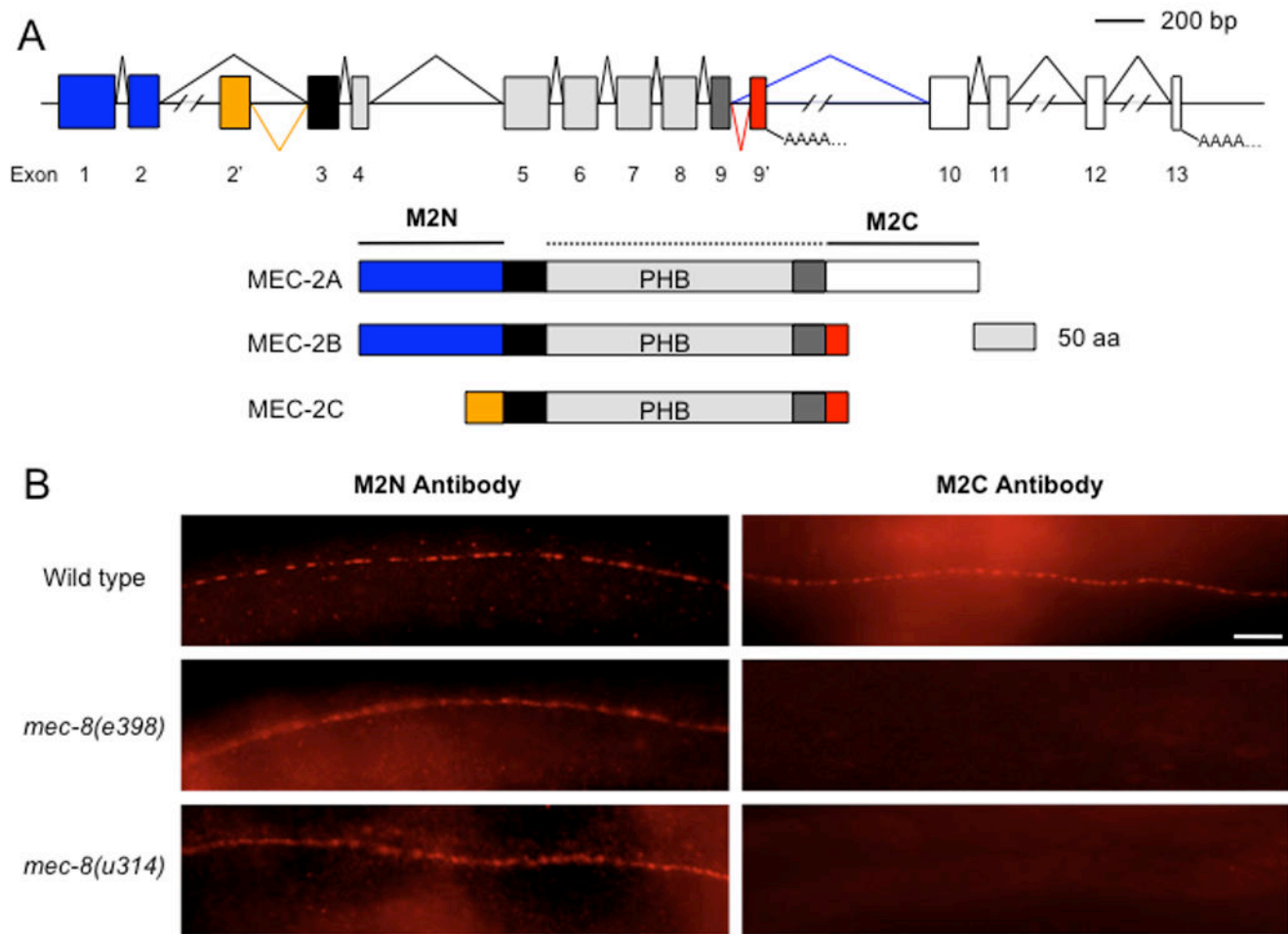
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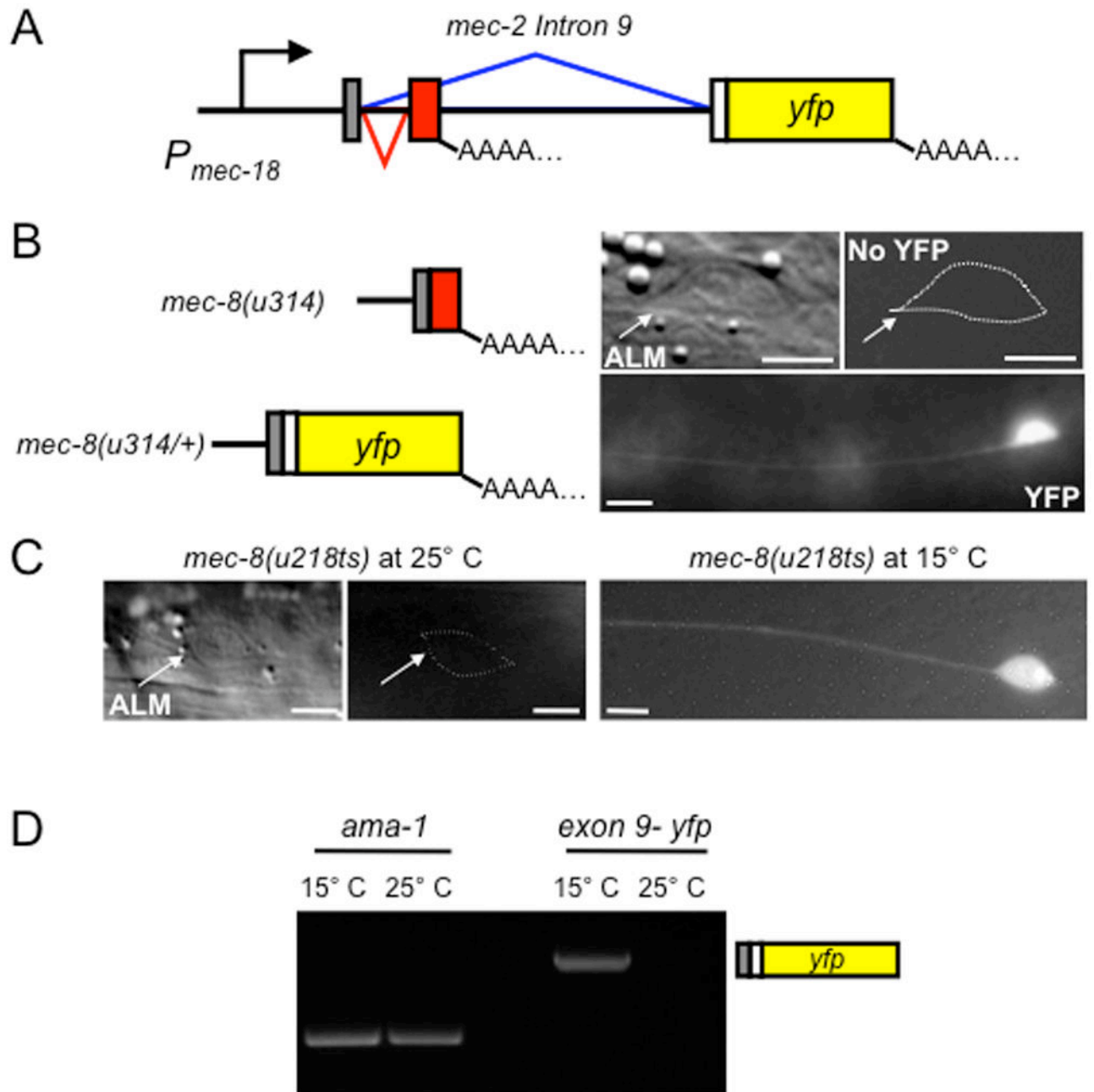
**Figure 1. *mec-2a* mRNA is not produced in *mec-8* mutants**

RT-PCR using primers that amplify exons 1–13 or 1–9 of *mec-2* in wild type and the indicated *mec-8* mutant strains (a) and in *mec-8* mutant strains in which the expression of *mec-8* is restored in TRNs (b). The 1.4 kb band corresponds to *mec-2a* and the 1.1 kb band to *mec-2b*.



**Figure 2. MEC-8 regulates the alternative splicing of *mec-2***

(a) *mec-2* splicing pattern and MEC-2 proteins. Colors denote corresponding regions of the proteins and the transcripts (blue: MEC-2 N terminus; orange: exon 2'-encoded polypeptide; black: hydrophobic region; gray: PHB domain; dark gray: the portion of the PHB domain contributed by exon 9; red: exon 9'-encoded polypeptide; and white: MEC-2 C terminus). The *mec-8*-dependent splice is shown in green; the *mec-2b*- and *mec-2c*-specific splices are shown in red and orange, respectively. Distances between exons 2 and 2', exons 9' and 10, exons 11 and 12, and exons 12 and 13 are 6327, 1592, 1164 and 2678 bp, respectively. Regions of the proteins recognized by the M2N and M2C antibodies are indicated. (b) Immunofluorescent staining of worms of the indicated genotypes with M2N and M2C antibodies. Scale bar 10  $\mu$ m



**Figure 3. *mec-2* intron 9 conveys MEC-8 dependence**

(a) Diagram of the *intron 9::yfp* construct under the control of the *mec-18* promoter. Colors here and in (b) are the same as in Fig. 2; in addition, the YFP coding region and polypeptide are colored yellow. (b) YFP fluorescence in ALM TRNs in the indicated strains. The predicted mRNA is also depicted and for homozygous *mec-8(u314)* worms, a DIC image is shown. (c) Temperature-dependent expression of *intron 9::yfp* in a *mec-8(218ts)* mutant. Scale bars, 5  $\mu$ m. (d) RT-PCR using primers for the last 18 nucleotides of *mec-2* exon 9 and

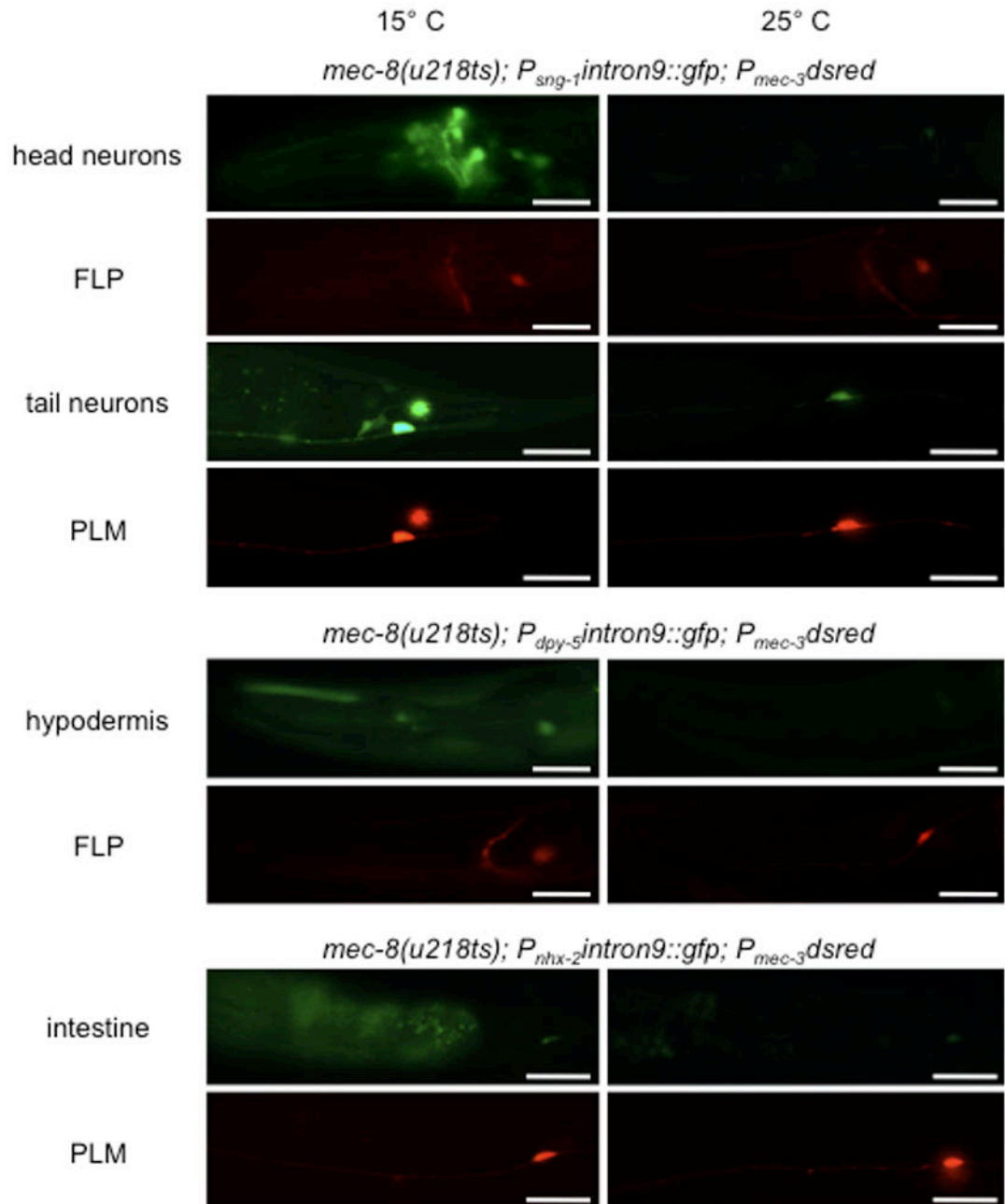
863 nucleotides of *yfp* of worms grown at 15° C and 25° C. *ama-1*, encoding the large subunit of RNA polymerase II, is used as control.

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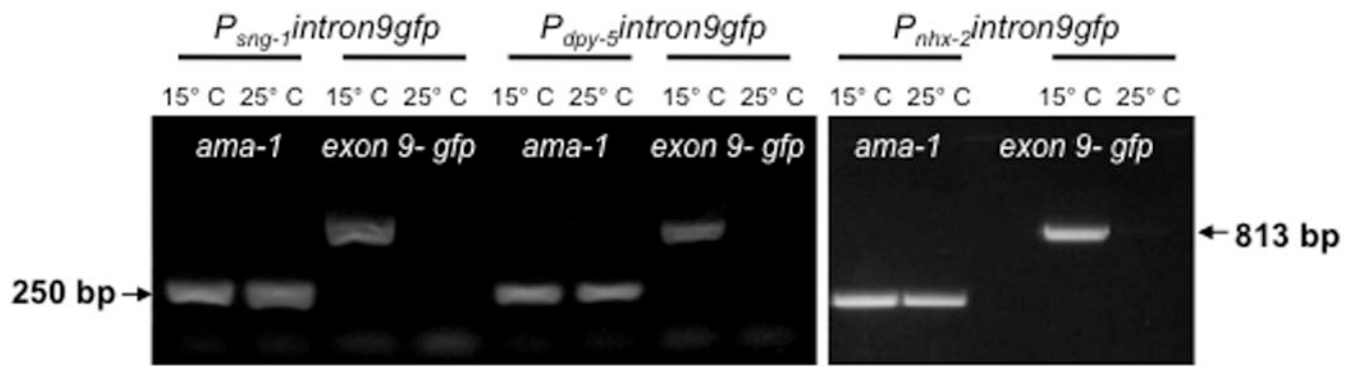
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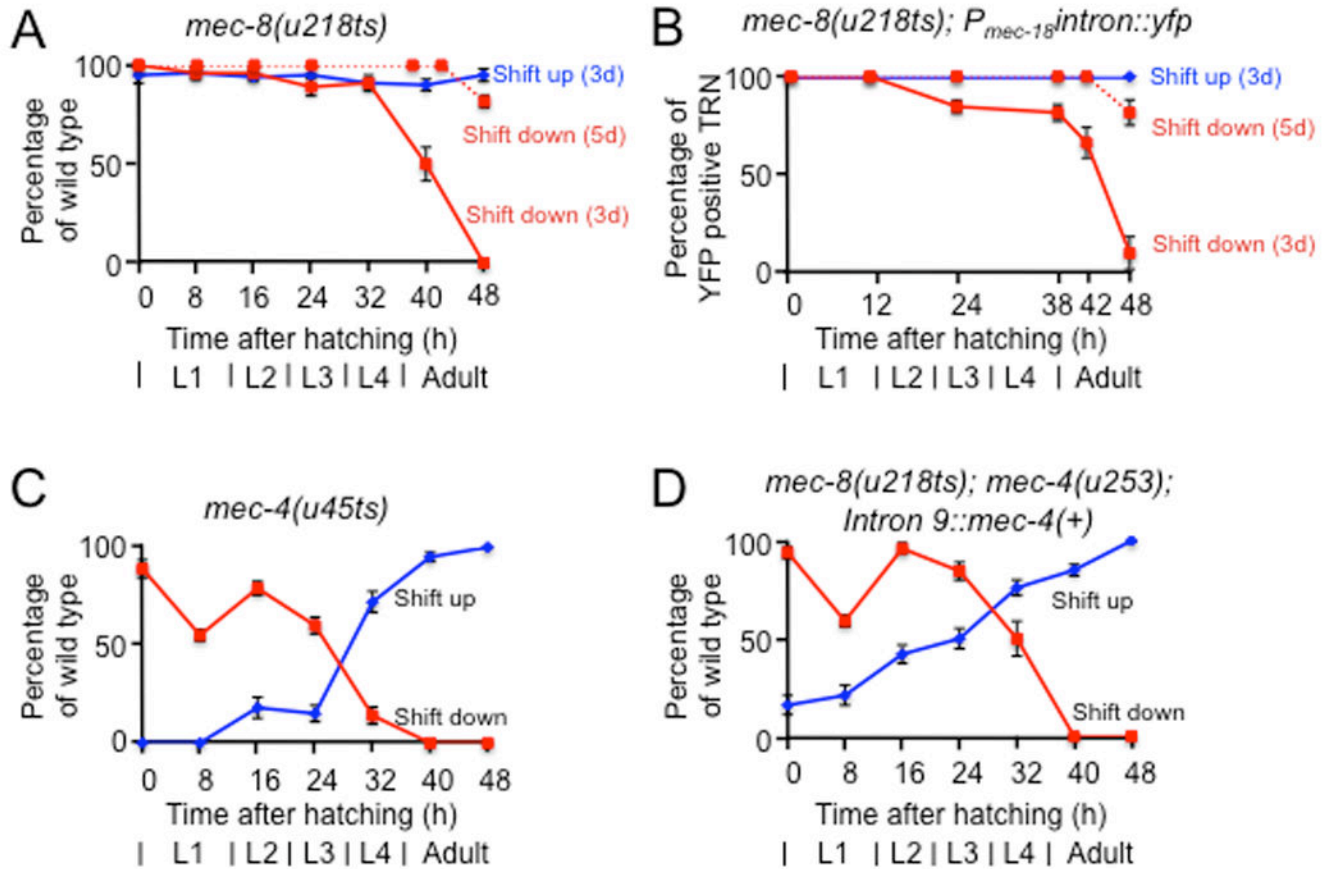
**Figure 4. MEC-8-dependent expression of *mec-2 intron 9::gfp* occurs in many tissues**  
**(a)** Fluorescence micrographs of single worms showing temperature-dependent expression of *intron 9::gfp* (green) in neurons (using the *sng-1* promoter), hypodermis (*dpy-5* promoter), and the intestine (*nhx-2* promoter). Expression of dsRed (red) in the FLP and PLM neurons (using the *mec-3* promoter) is seen at both temperatures. Scale bars 25 μm. **(b)** RT-PCR on the indicated strains grown at 15° C or 25° C, using primers for the last 18 nucleotides of *mec-2* exon 9 and for 863 nucleotides of *gfp*. The ubiquitously expressed *ama-1* gene was used as a control.



**Figure 5. Intron 9-containing strains and strains with missense temperature-sensitive mutations show similar responses to temperature**

Panels **a-d** show the touch response of adults with the indicated genotypes that have been shifted up (blue lines and rhomboids) or down (red lines and squares) at the indicated times after hatching. Touch sensitivity and YFP fluorescence were scored at 72 hours (solid lines) and 96 hours (dotted lines) after hatching ( $n = 3$  sets of 30–50 worms). Error bars represent the standard error of the mean.





### Figure 6. Conditional RNAi is potent and rapidly triggered

Conditional *intron 9::rde-1(+)* expression allows for rapid RNAi.

Newly hatched *mec-8(u218ts); rde-1(ne219); P<sub>rde-1</sub>intron 9::rde-1(+)* worms were grown for 24 hrs at 25°C (or for the equivalent developmental period – 48 hrs – at 15°C; Initial) and then placed at indicated temperature (Final). Worms were fed either bacteria with dsRNA for *unc-22* (+) or control bacteria (OP50)(-). Worms were tested for the *unc-22* twitcher phenotype (%Unc) at various times after the shift (times are given in 15°C equivalents). Error bars represent the standard error of the mean.

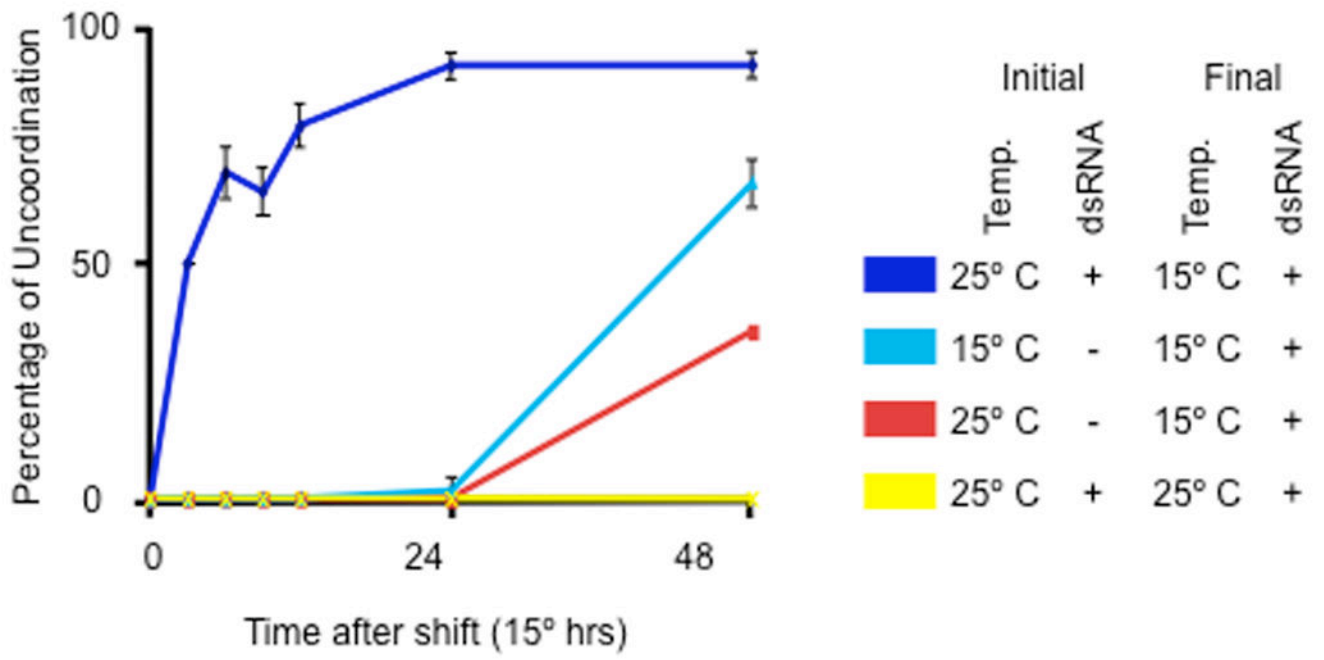


Figure 7.

RNAi responses in the presence or absence of *rde-1* and *mec-8*.

<i>rde-1</i> allele	<i>Intron9::rde-1</i>	<i>mec-8</i> allele	RNAi (%) <sup>a</sup>		
			<i>unc-22</i>	<i>rpl-3</i>	<i>unc-52</i>
+	-	+	98 ± 5	97 ± 3	83 ± 9
+	-	<i>e398</i>	99 ± 4	98 ± 3	86 ± 6
<i>ne219</i>	-	+	0	0	0
<i>ne219</i>	+	+	92 ± 7 <i>b</i>	90 ± 7 <i>b</i>	78 ± 8 <i>b</i>
<i>ne219</i>	+	<i>e398</i>	0	0	0
<i>ne219</i>	+	<i>u218 25°C</i>	0	0	0
<i>ne219</i>	+	<i>u218 15°C</i>	90 ± 5 <i>b</i>	87 ± 7 <i>b</i>	79 ± 7 <i>b</i>

<sup>a</sup> Values indicate the percent of animals that show a phenotype. Values are the mean ± std. dev. (n = 3 sets of 30–50 worms). The phenotypes produced by the three dsRNAs were 1) *unc-22* Twitcher by L4 and Paralyzed by adulthood, 2) *unc-52* Paralyzed or Uncoordinated, *rpl-3* Sterile or Growth defective (arrest as young larvae).

<sup>b</sup> These worms responded in a less severe fashion than wild type and *mec-8* mutants did at 15°C. The number of worms affected was not significantly different from wild type, but the phenotypes were less severe though still clear (for *unc-22*, the worms twitched throughout adulthood, for *unc-52*, worms were uncoordinated and paralyzed as later adults, and for *rpl-3*, worms were mostly sterile adults but some were arrested as late (L3 and L4) larvae).

**Table 1**