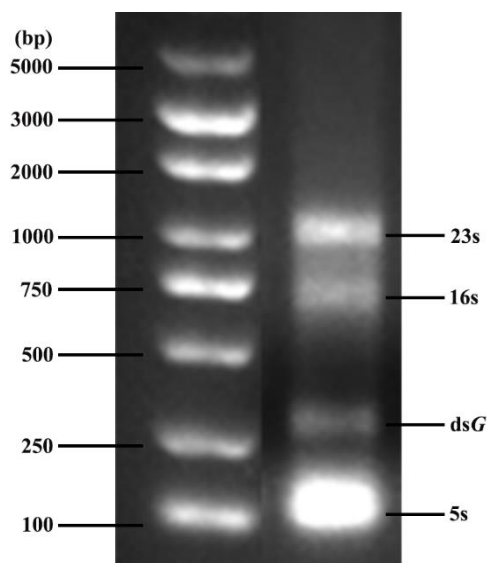
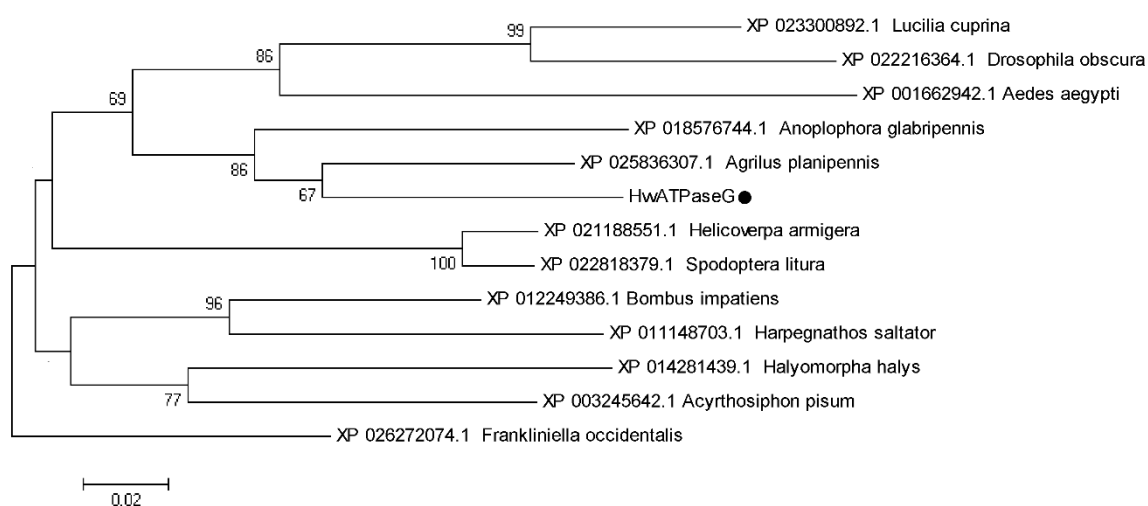


**Table S1.** Primers used in RT-PCR, dsRNA synthesis and qPCR.

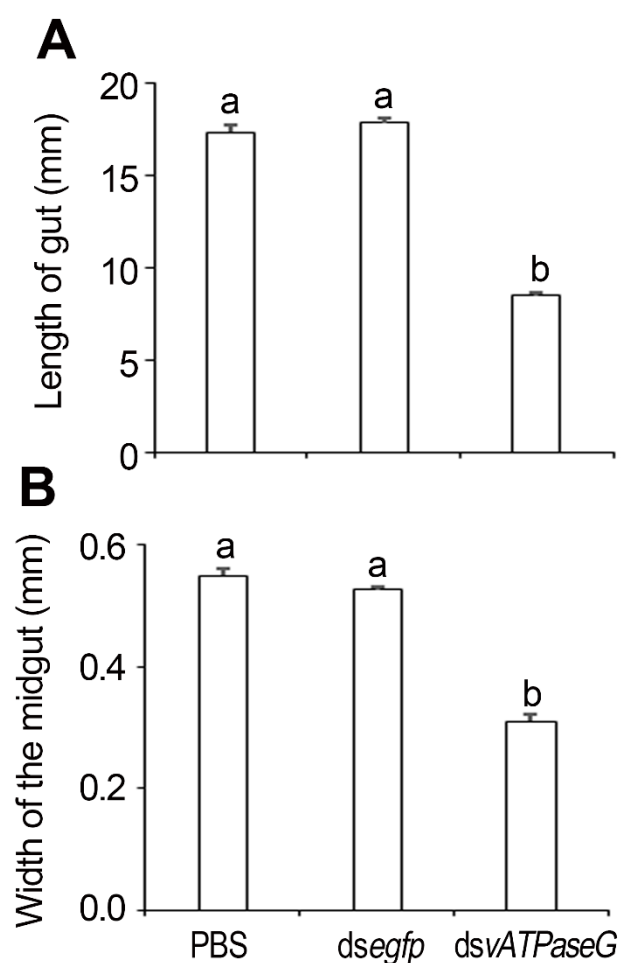
Fragment name	Forward primer	Reverse primer
<b>RT-PCR</b>		
<i>HvATPaseG</i>	GCACATTATGATGTTCAAGCTC	AACTGGGGAGGATTGACTTT
<b>dsRNA synthesis</b>		
dsG	AACTGGCGTTCCCTTTCC	GCAAGTCAAACACAAGGCATT
dsegfp	AAGTTCAGCGTGTCGG	CACCTTGATGCCGTTT
<b>qPCR</b>		
<i>qHvATPaseG</i>	GGCAAGTCAAACACAAGGCA	GGCGTTCCCTTTTCCTTACGA
<i>qHvRPS18</i>	CGCAATCAAAGGTGTTGGAAG	GCCTAGGGTTGGCCATAATAG
<i>qHvRPL13</i>	AGCATCCTTCGCTCGTTTAG	TTCGACAACCTGCCATTAGG



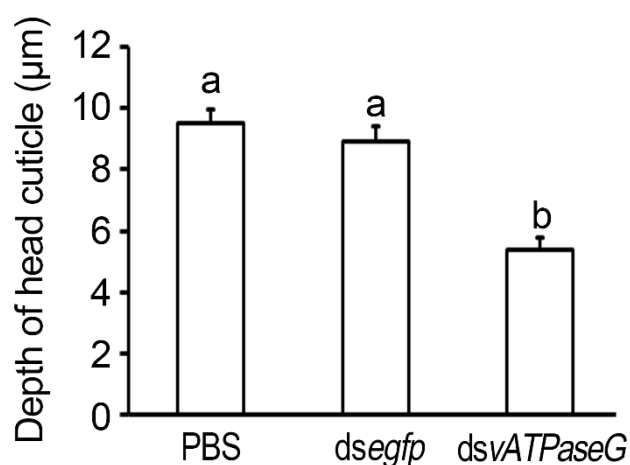
**Figure S1.** Electrophoresis of target dsRNA on 1% agarose gel.



**Figure S2.** Phylogenetic analysis of vacuolar ATPase V1 subunit G proteins (vATPaseGs). The proteins are derived from three Dipteran *Lucilia cuprina*, *Drosophila melanogaster* and *Aedes aegypti*, three Coleopteran *Henosepilachna vigintioctopunctata*, *Anoplophora glabripennis* and *Agrilus planipennis*, two Lepidopteran *Helicoverpa armigera* and *Spodoptera litura*, two Hymenopteran *Bombus impatiens* and *Harpegnathos saltator*, two Hemipteran *Halyomorpha halys* and *Acyrthosiphon pisum*, and a Thysanopteran *Frankliniella occidentalis*. The tree is constructed using the neighbor-joining method based on the full-length protein sequence alignments. Bootstrap analyses of 1000 replications are carried out and bootstrap values > 50% are shown on the tree.



**Figure S3.** The impacts of dsRNA on the lengths (A) and widths (B) in *Henosepilachna vigintioctopunctata*. The newly-ec-dysed fourth instar larvae had ingested PBS-, dsegfp-, and dsvATPaseG-dipped leaves for three days. The guts were dissected 5 days after the initiation of bioassay. The lengths and widths were measured. The bars represent values ( $\pm$  SE). Different letters indicate significant difference at P value  $< 0.05$  using analysis of variance with the Tukey-Kramer test.



**Figure S4.** The impacts of dsRNA on the depth of head capsule in *Henosepilachna vigintioctopunctata*. Table 2. days. The sections of the head capsule were stained using hematoxylin-eosin (HE) method. The depths were measured. The bars represent values ( $\pm$  SE). Different letters indicate significant difference at P value  $< 0.05$  using analysis of variance with the Tukey-Kramer test.