Supplementary File 1. **List of primers used in the present study.** All primers were designed using the Primer-BLAST tool from the National Center for Biotechnology Information (NCBI). Indicated are the respective primer names, their sequences, and their usage in the experimental protocol.

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| **Primer name** | **Sequence** | **Usage** |
| *fas5*\_dsRNA\_F | GATGCAAAGACCAACAAAGCC | dsRNA synthesis for targeting *fas5* |
| *fas5*\_dsRNA\_R | GCAAAGATTTCGCGATTCCTG |
| *fas5*\_dsRNA\_T7\_F | TAATACGACTCACTATAGGGGATGCAAAGACCAACAAAGCC |
| *fas5*\_dsRNA\_T7\_R | TAATACGACTCACTATAGGGGCAAAGATTTCGCGATTCCTG |
| GFP\_dsRNA\_F | GTGACCACCTTGACCTACGG | dsRNA synthesis for targeting GFP |
| GFP\_dsRNA\_R | TCTCGTTGGGGTCTTTGCTC |
| GFP\_dsRNA\_T7\_F | TAATACGACTCACTATAGGGGTGACCACCTTGACCTACGG |
| GFP\_ dsRNA\_T7\_R | TAATACGACTCACTATAGGGTCTCGTTGGGGTCTTTGCTC |
| *fas5*\_qPCR\_F | CTATGTTTGATGATATCAAGGCTGA | quantitative (q) PCR for ***fas* geneexpression** |
| *fas5*\_qPCR\_R | CATATGTATACTGGTCCCCGTAAG |
| ***fas6*\_qPCR\_F** | **GGGACCACACAAATCAAAATTGTA** |  |
| ***fas6*\_qPCR\_R** | **GTTTTCTCTCCTTCAATCACACTCA** |
| ***fas1*\_qPCR\_F** | **TGAGAAACACAAGCCTACCGA** |
| ***fas1*\_qPCR\_R** | **TTGACTGTCTTGACGTCCTTGA** |
| ***fas2*\_qPCR\_F** | **AATCAAACCTGCCTCGGGTG** |
| ***fas2*\_qPCR\_R** | **GCAGCAAGGCTTCTCCCATT** |
| ***fas3*\_qPCR\_F** | **AGTTCTCCCGTCTCGGTGTA** |
| ***fas3*\_qPCR\_R** | **GTGAACGAAGGTGGCGTAGA** |
| ***fas4*\_qPCR\_F** | **ACTGCTTTGGAGCTTGCGTA** |
| ***fas4*\_qPCR\_R** | **TCCGCAGATAAAACACCTAAGGC** |
| *Elf1a*\_qPCR\_F | GAGCCATCCACCAAAATGCC | quantitative (q) PCR for *Elf1a* **expression ( housekeeping gene)** |
| *Elf1a*\_qPCR\_R | CTTGTAGACGTCCTGGAGGG |