Insights into role of Non coding RNAs in Insects

Rupali J S¹, Impana², Sagar D^{2*}, Vidya Madhuri E¹, Pranita Roy² and Ratna Prabha^{3,4}

¹Division of Entomology, ICAR-Indian Agricultural Research Institute, New Delhi, India

²ICAR-National Bureau of Agricultural Insect Resources, Bengaluru, India

³Agricultural Knowledge Management Unit (AKMU), ICAR-Indian Agricultural Research Institute, New Delhi, India

ISSN: 3049-3374

Corresponding Author: garuda344@gmail.com

Introduction

Ribonucleic acid (RNA) is an essential biological macromolecule that is involved in numerous cellular processes, such as gene expression, regulation, and catalysis. RNA is usually a single-stranded molecule and possesses a wide range of structural and functional characteristics. RNA participates in all levels of genome organization, cell structure and gene expression, through RNA-RNA, RNA-DNA and RNA-protein interactions, often involving repeat elements including small interspersed nuclear elements in 3' untranslated regions. These interactions are involved in the regulation of chromatin architecture and transcription splicing (especially by antisense lncRNAs), protein translation and localization and other forms of RNA processing, editing, localization and stability. The central dogma of molecular biology, which posited that genetic information flows from DNA to RNA to protein, has been significantly revised with the discovery of functional noncoding RNA (ncRNAs). These RNA molecules do not encode proteins but contribute to numerous cellular functions.

RNA is classified into two categories: coding RNA and non-coding RNA. Coding RNA is mRNA that carries genetic information from DNA to ribosomes for protein synthesis and non-coding RNA (ncRNA) do not encode proteins but perform regulatory and structural functions. Housekeeping ncRNA (constitutive), which are ribosomal RNA (rRNA) it is major component of ribosomes, facilitating protein synthesis, transfer RNA (tRNA) adapts amino acids to codons during translation, small nuclear RNA (snRNA) plays a role in mRNA splicing within the spliceosome and small nucleolar RNA (snoRNA) involved in chemical modifications of rRNA, tRNA, and snRNA.

Regulatory ncRNAs, including microRNAs (miRNAs), are small RNA molecules (approximately 22 nucleotides) that control gene expression by binding to complementary mRNA sequences, resulting in mRNA degradation or translational repression. Small interfering RNA (siRNA) are short double-stranded RNA molecules that play a role in RNA interference (RNAi) and gene silencing. Long non-coding RNA (lncRNA) are RNA molecules that are longer than 200 nucleotides and play a role in chromatin remodeling, transcriptional regulation, posttranscriptional processes (Lee, 2012). PiRNA plays a crucial role in transposon silencing and maintaining genome stability, especially in germline cells. Circular rna (circRNA) a covalently closed RNA that regulates gene expression and acts as a miRNA sponge. Circular RNA (circRNA) a covalently closed rna that regulates gene expression and acts as a miRNA sponge.

An overview of non-coding RNA molecules: regulators that extend beyond the protein-coding genome.

Non-coding RNAs (ncRNAs) have gained significant attention as regulators of gene expression, with ncRNAs now acknowledged for their role in cellular differentiation, development, and defense mechanisms. ncRNAs are RNA molecules that do not code for proteins but play essential roles in gene expression and regulation, ncRNAs such as microRNAs (miRNAs) and small interfering RNAs(siRNAs) in the transcriptional and post-transcriptional regulation of gene expression. MiRNAs and siRNAs are referred as small RNAs (sRNAs) based on their small length ranging between 18 to 24 nucleotides, longer than 200 nucleotides defined as long non-coding RNAs (lncRNAs). LncRNAs are similar to mRNAs in their structure and the way they are produced, they are mainly transcribed by RNA polymerase II and polyadenylated. Non-coding RNAs without protein-coding potential have regulatory functions in downstream mRNA expression through epigenetic, transcriptional, and posttranscriptional mechanism.

The functions of non-coding RNA/ mediators of expression: gaining insight into non-coding RNA functions

ncRNAs regulate gene activity at the level of transcription and translation. Although ncRNAs do not possess the ability to encode proteins, they still play crucial regulatory functions in various biological processes. miRNAs play a role in controlling genes that are important for the development of organisms and their ability to respond to stress (ambros, 2004). Gene regulation, miRNAs and siRNAs control gene expression by interacting with mRNA. In genome defense piRNAs help silence transposable elements, protecting genome integrity. RNA processing snRNAs and snoRNAs are critical for splicing and RNA modification. One of the most studied classes, miRNAs regulate gene expression at the post-transcriptional level by binding to complementary sequences in target mRNAs, leading to



translational repression or degradation of the mRNA (Bartel, 2009). siRNAs perform a similar function, often involved in pathways that interfere with RNA and responses to viruses. The lncRNA Xist plays a crucial role in the process of xchromosome inactivation during female development (Lee, 2012). ncRNAs can also modulate epigenetic states by guiding chromatin-modifying complexes to specific genomic loci, thereby affecting the accessibility and transcription of target genes (Rinn & Chang, 2012). Some ncRNAs act as molecular sponges that bind and sequester miRNAs, preventing them from interacting with their target mRNAs, thus indirectly upregulating gene expression (salmena et al., 2011). There is growing evidence that extracellular miRNAs, often enclosed within exosomes, can function in cell-to-cell communication and may serve as biomarkers for disease (Valadi et al., 2007). The multifaceted roles of non-coding RNAs highlight their importance in controlling cellular processes and shaping the growth and development of organisms. Recognizing their roles provides valuable insights into intricate biological processes and potential therapeutic applications.

Overview of synthesis of non-coding RNAs

ncRNAs are synthesized through transcription by RNA polymerases, primarily RNA polymerase II and RNA polymerase III. miRNAs, lncRNAs and some snRNAs are transcribed by RNA polymerase II, often undergoing capping, splicing, and polyadenylation similar to mRNAs (Cech & Steitz, 2014). The biogenesis of miRNAs begins with the transcription of a primary miRNA (pri-miRNA), which is processed in the nucleus by the Drosha-DGCR8 complex into a precursor miRNA (pre-miRNA). This pre-miRNA is exported to the cytoplasm and further processed by Dicer, an RNase III enzyme, into a mature miRNA duplex. One strand (the guide strand) is incorporated into the RNA-induced silencing complex (RISC) to carry out gene silencing (Bartel, 2009).

In insects ncRNA synthesis follows similar molecular pathways, but with species-specific regulatory proteins and developmental timing influencing expression. In *Drosophila melanogaster*, the Drosha homolog is known as Pasha, and miRNA processing involves insect-specific cofactors that guide biogenesis during metamorphosis and tissue differentiation (Carthew & Sontheimer, 2009). Insect miRNAs play vital roles in regulating hormonal pathways, especially ecdysone signaling, which controls molting and metamorphosis (Sempere *et al.*, 2003). Piwi-interacting RNAs (piRNAs), another important class in insects, are especially prominent in germline cells, where they protect the genome from transposable elements.

Non-Coding RNAs in Insects: Emerging Players in Gene Regulation and Evolution

ISSN: 3049-3374

Biological processes are regulated not only by protein coding genes but also by ncRNA, miRNA, lncRNA, and circRNA (Beermann *et al.*, 2016). In insects, ncRNAs play vital roles in growth, development, immunity and recognized as crucial regulators of gene expression and evolutionary adaptation. miRNAs regulate metamorphosis by way of controlling hormone signaling pathways. siRNAs make contributions to antiviral defense by means of degrading viral RNA, that is critical for insect survival (Keene *et al.*, 2004). Insects which represent the maximum diverse organization of animals, depend on ncRNAs for essential biological procedures. Understanding those molecules gives insights into evolutionary biology and gives novel avenues for pest control.

The defining trait of non-coding RNA (ncRNA)

- Non-protein coding hence no protein synthesis, ncRNAs do not encode proteins. They perform various regulatory functions directly through their RNA structure
- Diversity of types as small ncRNAs consists of miRNAs, siRNAs, piRNAs. they are concerned in RNA silencing and regulation of gene expression. Long ncRNAs are extra versatile, regularly concerned in transcriptional regulation and splicing control.
- Formation of Complexes: many ncRNAs function by means of interacting with proteins to form ribonucleoprotein complexes, that can then adjust processes like transcription, splicing or translation.
- Conserved Sequences and Evolutionary Conservation: ncRNAs are notably conserved in evolution throughout species, indicating their crucial roles in fundamental cellular processes.
- Gene Regulation: ncRNAs regulate gene expression in diverse ways. This includes regulating transcription, influencing RNA splicing, and affecting mRNA stability and translation.
- RNA processing and cell signaling: some ncRNAs can act as signaling molecules, influencing cellular behavior.

Function of non-coding RNA(ncRNA) in insect development: lncRNAs and miRNAs play key roles in insect development, miR-34 regulates ecdysone signaling during metamorphosis in *Drosophila* (Sempere *et al.*, 2002). Law of Immune device and Immune Responses as siRNAs offer immune protection towards viruses in insects. In mosquitoes, RNAi pathways are activated upon viral infection, mainly for the degradation of viral RNA. In reproductive features



piRNAs are vital for maintaining genome integrity in insect germ cells. They suppress transposable factors, ensuring proper reproductive characteristic. The jobs of ncRNAs in insects are numerous and especially integrated, starting from regulating fundamental developmental tactics to complicated protection mechanisms and orchestrating massive-scale changes inside the genome, making sure proper organismal development, immune responses and protection mechanisms. Their potential to alter gene expression and coordinate diverse cellular techniques is crucial for the survival and flexibility of insects of their ever-converting environments.

As insects face several ecological pressures, including pathogens, environmental stressors, and competition, the useful diversity of ncRNAs gives a dynamic approach of regulating gene expression, allowing them to adapt quick to their environment. The roles of ncRNAs in insects opens up new avenues for exploring their biology and may provide insights into pest manipulate, evolutionary biology, and the improvement of novel techniques for coping with insectborne sicknesses. Metamorphosis and life cycle well ncRNAs regulate key transitions during an organism's life cycle. In insects, miRNAs are involved in regulating the hormonal changes that occur during these transitions, ensuring that the insect progresses through various life stages correctly, miRNA-14 in Drosophila regulates the timing of metamorphosis, coordinating the transition from the larval to pupal stage and lncRNAs participate in the segmentation of embryos and tissue-specific gene expression (Xu et al., 2003).

Insights into the function of non-coding RNAs

Insect Immunity as siRNAs and piRNAs are concerned in antiviral responses in insects. studies on *Aedes aegypti* have verified the function of RNAi pathways in controlling viral infections (Campbell *et al.*, 2008).

Insecticide Resistance as miRNAs adjust cleansing enzymes, thereby influencing insecticide resistance in agricultural pests. In *Helicoverpa armigera*, miR-998 modulates cytochrome P450 genes, affecting resistance to pesticides (Gong *et al.*, 2013). RNAi-based totally Pest manipulate, RNA interference has emerged as a promising approach for pest management. dsRNA targeting vital genes in pests like Leptinotarsa Decemlineata (Colorado potato beetle) has proven efficacy in decreasing pest populations (Baum *et al.*, 2007).

• gambling numerous roles across duplicate, and environmental response.

 ncRNAs are tightly regulated at some point of key developmental tiers consisting of embryogenesis, metamorphosis, and molting.

Conclusion

ISSN: 3049-3374

Non-coding RNAs (ncRNAs) are imperative players within the molecular device, orchestrating essential processes that allow for proper development, effective immune responses, and genomic balance. ncRNAs are critical for regulating gene expression and keeping cellular homeostasis. Their multifunctional roles make them vital within the life cycle and evolution of insects, highlighting the significance of ncRNAs beyond simple gene expression regulation and ncRNAs are worried in essential methods that impact organismal health and development. Information of ncRNA mechanisms can offer insights into insect body structure and provide ability targets for pest control techniques the discovery of lncRNAs has accelerated the expertise of genomic manipulate beyond the protein-coding genes (Wu et al., 2010). In insects it holds colossal capability for biotechnology and pest control. Advances in CRISPR-based totally gene editing can further enhance RNAi techniques, paving the way for more sustainable agricultural practices. Their study not most effective enriches our understanding of molecular biology but additionally presents novel equipment for pest manage and offers capacity programs in pest manipulate via the use of RNA interference (RNAi). As studies continues to uncover the complexities of these molecules, new strategies for manipulating gene expression in insects may additionally emerge, supplying valuable gear for biotechnology and agriculture.

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