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## Metabolomics Studies in Silkworm to Enhance Silk Quality: A New Frontier in Sericulture

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Sericulture, the practice of rearing silkworms (*Bombyx mori*) for silk production, is a vital agro-industry with significant economic and cultural importance. Silk quality is influenced by multiple factors, including nutrition, genetics, environmental stress, and disease. Metabolomics, the comprehensive study of small molecules in biological systems, has emerged as a powerful tool to understand the biochemical mechanisms underlying silk protein synthesis and cocoon formation. Recent studies reveal that amino acid metabolism, energy balance, and lipid pathways directly impact fibroin and sericin production, while stress and pathogen exposure alter key metabolic profiles, affecting silk yield and quality. By integrating metabolomics with nutrition management, disease detection, and breeding strategies, researchers can enhance silk quality and productivity. This approach opens new avenues for precision sericulture and the development of high-performance silk for industrial and biomedical applications.

### Introduction

Sericulture, the science of rearing silkworms for silk production, is both an ancient tradition and a modern bio-industry. The quality of silk largely depends on silkworm health, nutrition, genetics, and environmental conditions. With increasing demand for high-quality silk in textiles, biomedical materials, and biomaterials engineering, researchers are now turning to advanced biological tools to understand the factors that influence silk production. One of the most promising modern approaches is metabolomics, a powerful technology that studies small molecules (metabolites) present in biological systems. These metabolites reflect the physiological condition of an organism and provide clues about growth, stress, nutrition, and silk protein synthesis. Through metabolomics, scientists can now identify biochemical pathways associated with silk fibre quality, cocoon weight, and filament strength. Recent metabolomic studies in the silkworm (*Bombyx mori*) are helping researchers develop strategies to improve silk quality through better nutrition, disease management, and breeding approaches. This emerging field is transforming traditional sericulture into a precision-based biotechnology discipline.

### Metabolomics

Metabolomics is the comprehensive study of metabolites such as amino acids, sugars, lipids, nucleotides, and organic acids present in cells, tissues, or organisms. Unlike genomics or proteomics, which study genes and proteins, metabolomics provides a direct snapshot of biochemical activity.

Modern metabolomics uses advanced analytical tools such as:

- Gas chromatography–mass spectrometry (GC–MS)
- Liquid chromatography–mass spectrometry (LC–MS)
- Nuclear magnetic resonance (NMR)
- High-resolution mass spectrometry

These techniques allow researchers to detect hundreds or even thousands of metabolites simultaneously.

### **Role of Metabolites in Silk Formation**

Silk fibre is mainly composed of two proteins:

- Fibroin (structural core protein)
- Sericin (protective gum protein)

The synthesis of these proteins depends on amino acid metabolism, especially glycine, alanine, and serine. These amino acids are essential building blocks of fibroin. Metabolomic studies show that:

- Amino acid metabolism directly influences silk protein synthesis
- Energy metabolism affects larval growth and cocoon formation
- Lipid metabolism influences membrane stability and stress tolerance
- Carbohydrate metabolism affects energy supply for silk gland activity

Recent research using multi-tissue metabolomics revealed that differences in metabolic pathways related to amino acids and energy metabolism can significantly influence cocoon yield and silk productivity. Thus, metabolomics helps identify biochemical markers linked with superior silk traits.

### **Metabolomics and Silkworm Nutrition**

Nutrition is one of the most critical factors affecting silk quality. Traditionally, silkworms feed exclusively on mulberry leaves, but modern sericulture is exploring artificial diets to overcome seasonal limitations.

Metabolomics has helped researchers compare the effects of mulberry leaves and artificial diets. These studies revealed that diet composition affects metabolites related to:

- Protein synthesis
- Immunity
- Growth regulation
- Silk gland metabolism

A metabolomic analysis of silkworm midgut showed hundreds of differential metabolites between artificial and natural diets, many associated with silk quality and disease resistance. Another recent study found that specific amino acids such as valine improved feed efficiency and metabolic performance, suggesting opportunities to optimize artificial diets for better silk production. These findings demonstrate that metabolomics can guide the formulation of nutritionally balanced diets to enhance silk quality.

### **Metabolomics in Understanding Cocoon Quality**

Cocoon quality traits such as shell weight, filament length, and fibre strength are influenced by metabolic efficiency. Metabolomic profiling helps identify biomarkers associated with superior cocoon traits. For example, studies comparing formula feed and mulberry leaves found significant differences in metabolic pathways related to:

- Amino acid biosynthesis
- Carbohydrate metabolism
- Energy production
- Oxidative stress management

These metabolic differences were linked to variations in cocoon yield and silk gland development. Such research helps identify metabolic indicators that breeders can use to select high-performing silkworm breeds.

### **Disease Detection and Stress Management through Metabolomics**

Silkworm health is essential for producing high-quality silk. Diseases and environmental stress can disrupt metabolism and reduce silk yield. Metabolomics allows early detection of metabolic disturbances caused by:

- Viral infections
- Heavy metal exposure

- Environmental pollutants
- Nutritional deficiencies

For example, metabolomic studies of silkworms exposed to chromium showed significant alterations in gut microbiota and metabolic pathways, indicating stress responses that could affect productivity. Similarly, studies on pathogen infection revealed stage-specific metabolic responses that may help identify biomarkers for early disease detection. Such findings can help farmers adopt preventive health management practices to maintain silk quality.

### Metabolomics and Environmental Stress

Environmental factors such as temperature, pollutants, and microplastics can affect silkworm metabolism. Metabolomics helps understand how these stresses influence silk production. Recent studies showed that exposure to environmental microplastics altered gut metabolites and microbial composition in silkworms, which could potentially affect cocoon production. Another multi-omics study demonstrated that environmental contaminants can disrupt metabolic balance and microbiome interactions in silkworms, affecting growth and development. These insights highlight the importance of maintaining clean rearing environments for quality silk production.

### Integration of Metabolomics with Other Omics Technologies

Modern research is moving toward multi-omics approaches combining:

- Genomics
- Transcriptomics
- Proteomics
- Metabolomics

Such integrated approaches provide a complete understanding of silk production biology. For instance, studies combining microbiome analysis with metabolomics have shown that gut microbes contribute to detoxification processes and metabolic regulation in silkworms. This knowledge can help develop probiotic approaches or microbiome-based management strategies to improve silkworm performance.

### Applications of Metabolomics in Silk Quality Improvement

Metabolomics has several practical applications in sericulture:

**Nutritional improvement:** Identification of essential metabolites helps optimize mulberry nutrition and artificial diets.

**Breed selection:** Metabolic biomarkers can help identify high-yielding and stress-tolerant breeds.

**Disease diagnosis:** Early metabolic markers help detect infections before visible symptoms appear.

**Stress tolerance improvement:** Understanding stress metabolism helps develop management practices to reduce environmental impacts.

**Precision sericulture:** Metabolomics supports data-driven decision making in silkworm rearing. These applications demonstrate how metabolomics can modernize sericulture practices.

### Challenges in Metabolomics Research in Sericulture

Despite its potential, metabolomics research faces several challenges:

- High cost of analytical instruments
- Complex data analysis
- Lack of standardized metabolite databases for silkworms
- Need for interdisciplinary expertise
- Limited field-level application

However, as technologies become cheaper and bioinformatics tools improve, these limitations are gradually being overcome.

## Future Prospects

The future of metabolomics in sericulture is promising. Emerging areas include:

- Metabolic engineering for improved silk proteins
- Precision nutrition based on metabolic requirements
- Biomarker-assisted breeding
- AI-based metabolic prediction models
- Climate-resilient silkworm strains

Integration of metabolomics with artificial intelligence and big data analytics may soon enable predictive models for silk quality improvement. In the coming years, metabolomics may also contribute to the development of designer silk with enhanced properties such as:

- Higher tensile strength
- Improved elasticity
- Biomedical compatibility
- Advanced textile applications

Such innovations could expand the role of sericulture beyond textiles into biotechnology and materials science.

## Conclusion

Metabolomics is emerging as a transformative tool in silkworm research. By revealing the biochemical mechanisms underlying silk production, this technology provides new opportunities to improve silk quality through better nutrition, disease management, environmental control, and genetic improvement. Recent studies have demonstrated that metabolic pathways related to amino acids, energy metabolism, and stress responses play critical roles in cocoon quality and silk fibre properties. The integration of metabolomics with other omics technologies is further enhancing our understanding of silkworm biology. As research advances, metabolomics is expected to play a key role in developing precision sericulture systems that combine traditional knowledge with modern biotechnology. This integration will help ensure sustainable silk production while improving the livelihoods of farmers and strengthening the global silk industry. Thus, metabolomics represents not just a research tool but a pathway toward the future of smart sericulture and high-quality silk production.

## References

1. Wu, X., Chen, X., Ye, A., Cao, J., He, R., Pan, M. et al. (2022). Multi-tissue metabolomic profiling reveals potential mechanisms of cocoon yield in silkworms (*Bombyx mori*) fed formula feed versus mulberry leaves. *Frontiers in Molecular Biosciences*, **9**: 977047. DOI: <https://doi.org/10.3389/fmolb.2022.977047>
2. Yuan, S., Sun, Y., Chang, W., Zhang, J., Sang, J., Zhao, J., et al. (2023). The silkworm (*Bombyx mori*) gut microbiota is involved in metabolic detoxification by glucosylation of plant toxins. *Communications Biology*, **6**(1): 790.
3. Chen, Y. Z., Rong, W. T., Qin, Y. C., Lu, L. Y., Liu, J., Li, M. J., et al. (2023). Integrative analysis of microbiota and metabolomics in chromium-exposed silkworm (*Bombyx mori*) midguts based on 16S rDNA sequencing and LC/MS metabolomics. *Frontiers in Microbiology*, **14**: 1278271. DOI: <https://doi.org/10.3389/fmicb.2023.1278271>
4. Wu, J., Li, L., Qin, D., Chen, H., Liu, Y., Shen, G., & Zhao, P. (2024). Silkworm hemolymph and cocoon metabolomics reveals valine improves feed efficiency of silkworm artificial diet. *Insects*, **15**(4): 291. DOI: <https://doi.org/10.3390/insects15040291>
5. Zhang, X., Zheng, W., Shao, W., Yu, W., Yang, Y., Qin, F., et al. (2024). Environmental concentrations of microplastic-induced gut microbiota and metabolite disruption in silkworm, *Bombyx mori*. *Chemosphere*, **358**: 142126.
6. Su, Z., Li, Y., Lin, Z., Huang, Q., Fan, X., Dong, Z., et al. (2025). GC-MS-based metabolomic analysis of silkworm haemolymph reveals four-stage metabolic responses to nucleopolyhedrovirus infection. *Insect Molecular Biology*, **34**(2): 289-301.