

Ayurgenomics: Bridging Traditional Ayurveda and Modern Genomics – A Narrative Review

¹ Dr Vineetha V ² Dr Deeja C Radhakrishnan

¹Associate Professor, Department of Samhita and Siddhanta, J.S.Ayurveda Mahavidyalaya, Nadiad, Gujarat

²Professor and Head, Department of Samhita and Siddhanta, J.S.Ayurveda Mahavidyalaya, Nadiad, Gujarat

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I. INTRODUCTION

Ayurgenomics is an emerging interdisciplinary field that integrates the principles of Ayurveda—India’s traditional system of medicine—with modern genomics to personalize healthcare. It builds upon the ancient concept of *Prakriti* (individual constitution) and attempts to correlate it with genetic and molecular markers. This integration has the potential to enhance predictive, preventive, and personalized medicine by offering deeper insights into individual susceptibility to diseases, drug responses, and therapeutic strategies.

II. MATERIALS AND METHODS

This narrative review was conducted to synthesize current literature on the convergence of Ayurveda and genomics, particularly the concept and application of Ayurgenomics. A comprehensive literature search was performed using electronic databases including PubMed, Scopus, Google Scholar, and Web of Science. The search covered the period from 2000 to 2024 using keywords such as “Ayurgenomics,” “Prakriti,” “Ayurveda and genomics,” “pharmacogenomics Ayurveda,” “personalized medicine Ayurveda,” and “genetic basis of Prakriti.”

Inclusion criteria consisted of peer-reviewed original articles, reviews, meta-analyses, and high-impact commentaries in English that discussed the genetic, transcriptomic, proteomic, or metabolomic aspects of Prakriti or Ayurgenomics. Additionally, publications correlating Prakriti with disease predisposition, drug response, and biological

markers were included. Articles not addressing both Ayurvedic and genomic components or lacking methodological clarity were excluded.

Reference lists of key articles were also screened to identify additional relevant sources. Selected studies were evaluated for their design, sample size, Prakriti assessment methodology, genomic techniques employed (e.g., SNP analysis, HLA typing, transcriptomics), and relevance to personalized or preventive medicine.

Ethical approval was not required as this review did not involve human or animal subjects directly.

III. DISCUSSION

In Ayurveda, health and disease are determined by the balance of the three Doshas—*Vāta*, *Pitta*, and *Kapha*. The unique combination of these Doshas in an individual, known as Prakriti, governs physical, physiological, and psychological traits. Ayurgenomics proposes that Prakriti types correspond to genotypic variations, thereby serving as a traditional biomarker for personalized healthcare.

Scientific Correlates and Evidence Base

Several studies have attempted to establish genetic and molecular bases for Prakriti types:

- SNP (Single Nucleotide Polymorphism) studies have found significant associations between Prakriti and specific gene variants. For instance, individuals with *Pitta* Prakriti show a higher prevalence of polymorphisms in genes related to metabolism and immunity, such as CYP2C19 and IL6.
- HLA (Human Leukocyte Antigen) profiling has revealed unique patterns among different Prakriti groups, indicating distinct immunogenetic backgrounds.

- A notable study published in *Nature* (2015) highlighted that Pitta-dominant individuals had higher levels of pro-inflammatory markers, while Kapha types had markers indicative of slow metabolism.
- Recent transcriptomic and metabolomic research supports that Prakriti types have differential expression patterns, suggesting biological plausibility behind Ayurvedic classification.

IV. CLINICAL IMPLICATIONS

Ayurgenomics holds promise in several areas:

- **Personalized Medicine:** Integrating Prakriti typing with genomic data can lead to more effective, tailored interventions in chronic diseases such as diabetes, cardiovascular disorders, and autoimmune conditions.
- **Pharmacogenomics:** Ayurgenomics may guide individualized drug therapy by predicting adverse drug reactions and optimizing dosage based on Prakriti-gene interactions.
- **Preventive Health:** Early identification of predispositions based on Prakriti may lead to better lifestyle and dietary interventions aligned with an individual's constitution.
- **Public Health:** Population stratification using Prakriti classification could inform epidemiological research and community health strategies.

V. CHALLENGES AND LIMITATIONS

Despite its promise, Ayurgenomics faces several challenges:

- **Standardization of Prakriti assessment:** Traditional methods vary and lack uniform diagnostic tools, though digital and AI-based models are emerging.
- **Sample diversity:** Most studies are limited in scale and demographic diversity, necessitating large, multicentric, and ethnically diverse research.
- **Interdisciplinary gaps:** Bridging traditional knowledge with rigorous scientific methodology requires collaborative frameworks that respect both epistemologies.

VI. FUTURE DIRECTIONS

To fully realize the potential of Ayurgenomics, the following steps are essential:

1. Development of validated tools for Prakriti assessment integrated with digital phenotyping.
2. Genomic mapping of larger, diverse populations classified by Prakriti to identify robust gene-constitution links.
3. Integration with Artificial Intelligence (AI) and machine learning to analyze complex datasets.
4. Establishment of Ayurgenomics registries and biobanks to support translational research.

VII. CONCLUSION

Ayurgenomics represents a paradigm shift in holistic healthcare by uniting ancient wisdom with cutting-edge genomics. It supports a more individualized approach to health by recognizing constitutional differences at both phenotypic and genotypic levels. While still in its nascent stages, the field promises to reshape preventive, predictive, and personalized medicine, especially in diverse populations where traditional systems like Ayurveda remain culturally relevant. Continued interdisciplinary research, methodological rigor, and cross-cultural collaboration are essential to validate and expand this integrative science.

REFERENCES

- [1] Patwardhan B, Warude D, Pushpangadan P, Bhatt N. Ayurveda and traditional Chinese medicine: a comparative overview. *Evid Based Complement Alternat Med.* 2005;2(4):465–73.
- [2] Prasher B, Gibson G, Mukerji M. Genomic insights into Ayurveda and functional variability of genes. *Curr Sci.* 2016;111(5):901–5.
- [3] Rotti H, Raval R, Anchan S, Kanchan K, Kabekkodu S, Shouche Y, et al. Determinants of prakriti, the human constitution types of Indian traditional medicine and its correlation with contemporary science. *J Ayurveda Integr Med.* 2014;5(3):167–75.
- [4] Govindaraj P, Nizamuddin S, Sharath A, Jyothi V, Rotti H, Raval R, et al. Genome-wide analysis correlates Ayurveda Prakriti. *Sci Rep.* 2015;5:15786.
- [5] Tiwari P, Nayak D, Seth T, Kumar A, Srikanth N, Mehrotra R, et al. Prakriti-based research: Good reporting practices. *J Ayurveda Integr Med.* 2020;11(4):539–42.