

A Review on the Use of Artificial Intelligence for Human Microbiota Analysis in Clinical Tasks

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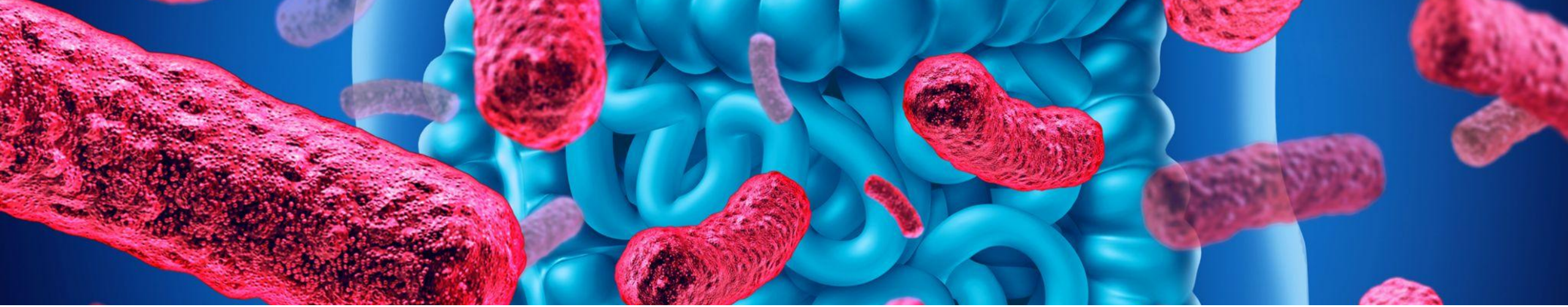
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Agenda

- Background
- Why AI for Microbiota
- Methodology
- Overview of Selected Studies
- Machine Learning Approaches
- Deep Learning Approaches
- Clinical Applications
- Limitations
- Future Directions
- Conclusion



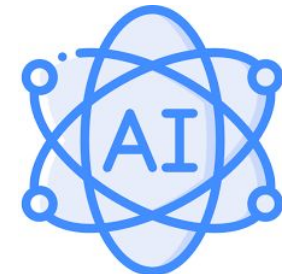


Background : Microbiota

1. Trillions of microorganisms across gut, oral, skin, urogenital sites
2. Key roles: Immunity, Metabolism, Pathogen resistance
3. Dysbiosis linked to major diseases:
 - a. Colorectal cancer
 - b. T2 Diabetes
 - c. IBD
 - d. Liver disease
4. High-dimensional, sparse, noisy data → analytical challenges

Why AI For Microbiota?

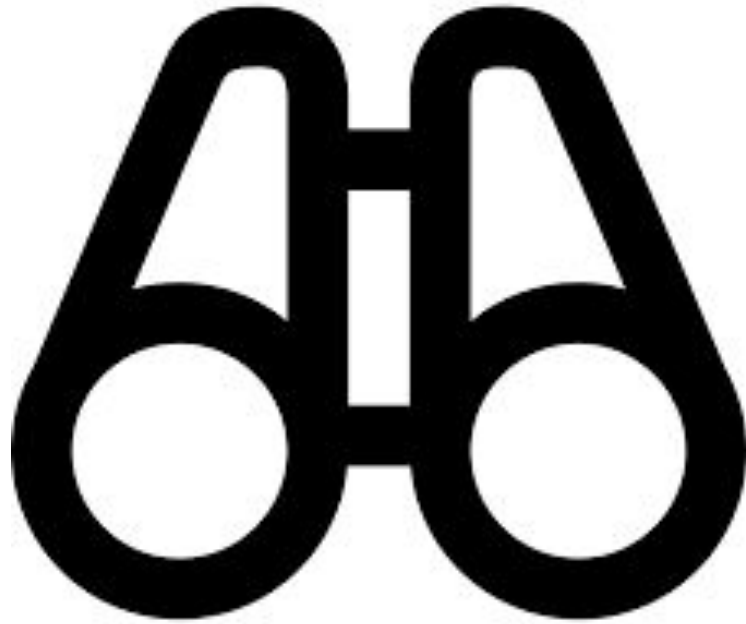
- Learns patterns in complex microbial ecosystems
- Handles high dimensionality and non-linearity
- Supports:
 - Disease prediction
 - Prognosis
 - Patient stratification
 - Biomarker discovery
- Potential pathway to precision medicine



Methodology(PRISMA INSPIRED)



- Databases: PubMed, IEEE Xplore, ScienceDirect, SpringerLink
- Keywords: microbiota · microbiome · ML · DL · clinical
- Inclusion:
 - Human data
 - Peer-reviewed
 - Empirical AI models
- Extracted: model type, data source, clinical goal, performance, limitations



Overview Of Selected Studies

- Majority use classical ML
- Smaller but growing set uses DL, especially CNNs
- Data types:
 - 16S rRNA
 - Whole-metagenome
 - Metabolomics
 - Imaging + microbiota
- Clinical focus: cancer, liver disease, IBD, oral disease, dermatology, drug interactions

Machine Learning Approaches



- Dominant: Random Forest (robust, interpretable)
- Others: SVM · Logistic Regression · XGBoost · Extra Trees
- Applications:
 - Cancer subtype prediction
 - Liver disease prognosis
 - CD vs UC classification
 - Preterm birth prediction
 - CVD screening
 - Drug–microbiota interaction modeling

Deep Learning Approaches

- Mostly CNN-based architectures
- Phylogeny-aware CNNs using taxonomic structure
- Used for:
 - Liver disease staging
 - Dermatology + microbiota multimodal tasks
 - High-resolution ASV identification
- Strength: feature representation
- Weakness: interpretability, data requirements



Clinical Applications of AI

- Diagnosis / Classification
 - Colorectal cancer · Liver disease · IBD · Oral lichen planus · Dermatology
- Prognosis / Risk Prediction
- Survival · CVD · T2D (microbial rhythmicity)
- Therapeutic Support
- Immunotherapy response · Drug metabolism/bioaccumulation
- Multi-disease frameworks emerging



Data Types & Processing

- Mostly gut microbiota (stool samples)
- Growing use of oral & skin microbiota
- Profiling methods:
 - 16S rRNA (common, low resolution)
 - WGS (higher resolution, costly)
- Challenges:
 - Heterogeneous preprocessing
 - OTUs vs ASVs
 - Compositionality (CLR, ALR, TSS)





Key Limitations

- Small, single-center datasets
- Weak external validation
- Batch effects & inter-study variability
- Inconsistent preprocessing pipelines
- Limited interpretability
- Minimal code / pipeline transparency

Challenges: What They Mean



- High AUC \neq reliable model
- Models may learn site-specific artifacts instead of biology
- Poor generalization across regions, devices, populations
- Limited trust due to lack of explainability
- Biological noise + sequencing differences distort patterns

Future Directions



- Standardize microbiome preprocessing workflows
- Multi-center, diverse datasets for validation
- Integrate explainability (SHAP, LIME, attention)
- Expand to oral, skin, vaginal niches
- Multimodal learning (microbiota + metabolomics + imaging + metadata)
- Causal ML & temporal modeling
- Federated learning for privacy-preserving cross-site collaboration

Conclusions

- AI shows strong promise for microbiota-based clinical tasks
- ML dominates; DL excels in multimodal contexts
- Major gaps: generalization, standardization, interpretability
- Clinical translation requires rigorous validation and transparency
- Microbiota + AI → emerging pillar for personalized medicine

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Speaker: Rohith Janwadkar

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