Sample\_barcode67

Horse

Sample #barcode67

Analyzed: 2025-09-26 Report: 2025-09-26

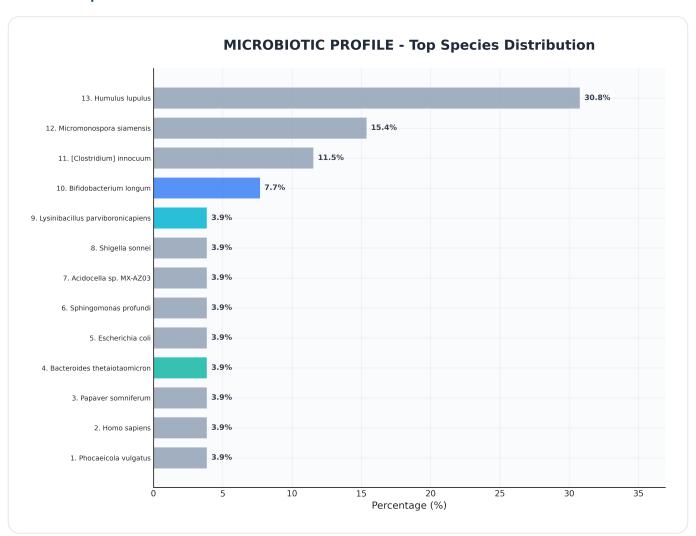
# **Microbiome Sequencing Results**

36.0

Dysbiosis Index

Mild Dysbiosis

## **Bacterial Species Distribution**



## **Dominant Species**

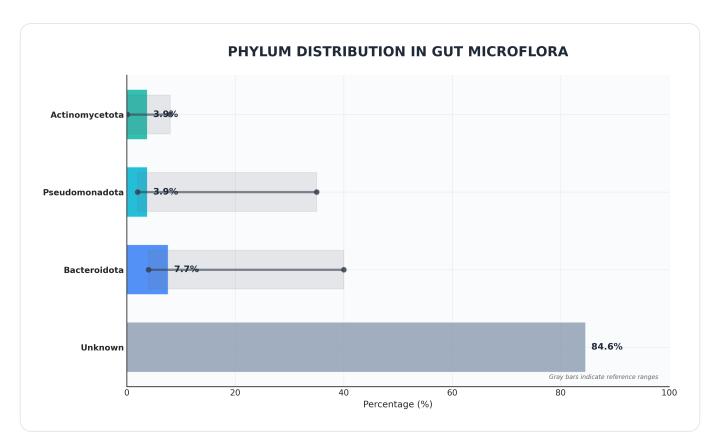
Species	Abundance (%)	Phylum
Phocaeicola vulgatus	30.77%	Unknown
Homo sapiens	15.38%	Unknown
Papaver somniferum	11.54%	Unknown
Bacteroides thetaiotaomicron	7.69%	Bacteroidota
Escherichia coli	3.85%	Pseudomonadota
Sphingomonas profundi	3.85%	Unknown
Acidocella sp. MX-AZ03	3.85%	Unknown
Shigella sonnei	3.85%	Unknown
Lysinibacillus parviboronicapiens	3.85%	Unknown
Bifidobacterium longum MEMT Genetics Laboratory	3.85%	Actinomycetota Page 1 of 4

Sample #barcode67

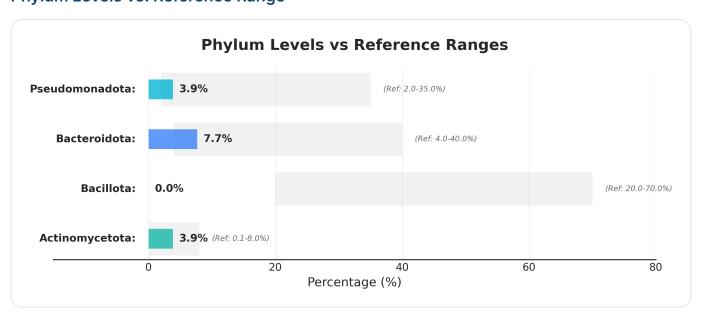
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# **Phylum Distribution Analysis**

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## Phylum Levels vs. Reference Range



Phylum	Patient (%)	Normal Range (%)	Status
Unknown	84.64	0.0 - 100.0	Normal
Bacteroidota	7.69	4.0 - 40.0	Normal
Pseudomonadota	3.85	2.0 - 35.0	Normal
Actinomycetota MEMT Genetics Laboratory	3.85	0.1 - 8.0	Normal Page 2 of 4

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# **Clinical Interpretation**

#### **Clinical Assessment**

The microbiome analysis indicates **mild dysbiosis** with an index of 36.0. This suggests a moderate imbalance in the gut microbial community that may benefit from targeted intervention. While not immediately concerning, this imbalance could impact digestive efficiency and immune function if left unaddressed.

### **Key Findings**

**Low Bacillota:** 0.0% (Normal: 20-70%). Associated with reduced fiber fermentation and butyrate production.

#### **Clinical Recommendations**

- Increase dietary fiber through additional hay supplementation
- → Consider probiotic supplementation (Lactobacillus/Bifidobacterium strains)
- > Reduce grain intake if exceeding 0.5% body weight per feeding

MEMTRe-revaluate microbiome in 8-12 weeks after intervention

Horse

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## **Summary & Management Guidelines**

#### **Report Summary**

Dysbiosis Index: 36.0Total Species Identified: 13Category: MildSample Quality: Adequate

**Dominant Phylum:** Unknown **Analysis Method:** 16S rRNA NGS

#### **Understanding the Dysbiosis Index**

The Dysbiosis Index (DI) quantifies the degree of microbial imbalance in the gut:

- 0-20: Normal, healthy microbiome
- 21-50: Mild dysbiosis requiring dietary adjustment
- >50: Severe dysbiosis requiring intervention

#### **Management Guidelines**

#### **Correcting Mild Dysbiosis**

- Increase forage intake to at least 1.5-2% body weight
- Reduce grain meals to <0.5% body weight per feeding
- Add probiotic supplement (10^9 CFU daily)
- Consider prebiotic fiber sources (beet pulp, psyllium)
- Ensure adequate water intake (30-50L daily)

#### Follow-up Testing

Dysbiosis Category	Re-test Timeline	Monitoring Focus
Normal	12 months	Annual health screening
Mild	2-3 months	Response to dietary changes
Severe	4-6 weeks	Treatment efficacy

#### For questions regarding this report, please contact:

**MEMT Genetics Laboratory** 

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**MEMT Genetics Laboratory**