







CLINICAL GUIDE

Pathogen and Pathobiont Management

Pathogen management guidelines and considerations for pathobiont management to minimise the impact to beneficial gut microbes.

Pathogens defined

| | Pathogen | Opportunistic pathogen | Pathobiont | Commensal |
|------------|---|---|--|---|
| |  |  |  |  |
| Definition | A microbial strain that can cause disease | A microbial strain that can cause disease in susceptible hosts | A microbial species associated with negative health outcomes* | A microbial species associated with positive health outcomes* |
| Management | Medical treatment dependent on pathogen and clinical presentation | Medical treatment may be considered in vulnerable patients | Targeted intervention may be considered if over-abundant | Targeted intervention may be considered if under-abundant |
| Example | <i>E. coli</i> O157 | <i>Klebsiella oxytoca</i> | <i>Bifidobacterium wadsworthia</i> | <i>Faecalibacterium prausnitzii</i> |

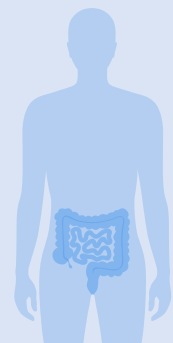
*Often based on cross-sectional studies where causation has not been established

When to consider testing for pathogen identification

Clinical indications for referring for diagnostic pathogen panel include:

- Chronic signs and symptoms of a pathogen infection, including loose stools, frequent defecation, weight loss, bloating, abdominal pain, nausea, vomiting[^]
- History of overseas travel or exposure to environments with reduced sanitation
- Immunocompromised patient
- Suspected or diagnosed post-infective IBS

[^]For acute or severe symptoms referral to the GP for testing is recommended.



Testing to support pathogen identification

Stool testing using Microbiome Explorer Comprehensive can be used for targeted pathogen detection¹ alongside gut microbiome² and gastrointestinal health marker¹ profiles. Microbiome Explorer Comprehensive utilises RT-PCR (real-time polymerase chain reaction) that is routinely used as a highly sensitive method for the detection of target pathogens, species or genera. Microbiome Explorer Comprehensive combines RT-PCR analysis with metagenomics to provide comprehensive insights into the health of the whole gut microbiome to support informed clinical decision-making.



1.The faecal occult blood, real-time polymerase chain reaction (RT-PCR) and enzyme-linked immunosorbent assays (ELISA) used in the Microbiome Explorer™ range are diagnostic and are approved for clinical use. 2.The faeces pH assay used in the Microbiome Explorer™ range is for research use only and not to be used as a basis for diagnosis. The metagenomic assays used in the Microbiome Explorer™ range are to determine the microbiome populations and associated functional pathways in a faecal sample. The application is for research use only and not to be used as a basis for diagnosis.

Pathogen management guidelines

Medical treatment if detected



Entamoeba histolytica

Medical treatment if symptomatic



Clostridium difficile pathogenic strains
C. difficile toxin B,
Hypervirulent *C. difficile*

Giardia lamblia

Medical referral if symptomatic



Escherichia coli pathogenic strains
Enterotoxigenic *E.coli* (ETEC),
Enteroaggregative *E. coli* (EAEC),
E. coli O157,
Shiga toxin
Shigella spp/ enteroinvasive *E.coli* (EIEC),
Enteropathogenic *E. coli* (EPEC)

Not necessarily pathogenic strains
Consideration of clinical
presentation required



Aeromonas spp.
Campylobacter spp.
Cryptosporidium spp.
Cyclospora cayetanensis
Salmonella spp
Vibrio spp.
Yersinia enterocolitica

Metagenomic detected potential pathogens
e.g. *Campylobacter_D upsaliensis*,
Clostridium_P perfringens

Pathogenic role is unclear
Exclude other causes of symptoms
before considering treatment



Dientamoeba fragilis
Blastocystis sub-types (species)*

Clinical context must always be primary consideration.

Pathobiont management checklist

| | | |
|---|--|-----------------------|
| STEP 1 Test the microbiome | Rule out pathogen requiring medical treatment or referral | <input type="radio"/> |
| | Assess microbiome resilience by reviewing diversity and richness | <input type="radio"/> |
| | Identify overabundant pathobionts by reviewing distance from average in species table | <input type="radio"/> |
| | Identify underabundant commensals by reviewing distance from average in species table | <input type="radio"/> |
| | Evaluate functional dysbiosis by reviewing microbial markers | <input type="radio"/> |
| STEP 2 Evaluate the clinical case | Consider clinical context such as clinical symptoms, severity, and duration | <input type="radio"/> |
| | Note the patient's vulnerability, such as immunocompromised status, age, or underlying conditions that may affect treatment decisions | <input type="radio"/> |
| | Ensure any red flag gastrointestinal health markers have been medically investigated (elevated calprotectin or lactoferrin, detected occult blood) | <input type="radio"/> |
| | Assess gut function and environment including inflammation and gut barrier by using gastrointestinal health markers | <input type="radio"/> |
| STEP 3 Consider the cause | Determine gastrointestinal function and environment | <input type="radio"/> |
| | Note functional dysbiosis | <input type="radio"/> |
| | Consider underlying conditions | <input type="radio"/> |
| | Review medication history | <input type="radio"/> |
| | Evaluate dietary factors | <input type="radio"/> |
| | Assess lifestyle and stress factors | <input type="radio"/> |
| STEP 4 Treat the cause | Implement Microbiome Explorer personalised insights to manage microbial markers | <input type="radio"/> |
| | Maximise microbial diversity and richness | <input type="radio"/> |
| | Support gastrointestinal function such as digestive secretions and motility | <input type="radio"/> |
| | Regulate gastrointestinal environment by managing intestinal barrier and intestinal inflammation | <input type="radio"/> |
| | Optimise diet and lifestyle to support microbiome and gut health | <input type="radio"/> |

Considerations for antimicrobial treatment



Consider the use of antimicrobials only when there is sufficient evidence linking the identified pathogen to patient symptoms.



Avoid unnecessary antimicrobial treatment to prevent disruption of the gut microbiome.



If antimicrobials are necessary, carefully weigh the potential benefits against the risks of collateral damage to the microbiome.



Consider using targeted therapies whenever possible to minimise the impact on beneficial microbes.