Quantitative Polymerase Chain Reaction - qPCR

Contact us:



We offer a next-generation, real-time quantitative PCR based assay that identifies organisms associated with Onychomycosis

Open Array Format with 18(x3) Assays and 48 Samples



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PREMIER MEDICAL LAB

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• FAST • ACCURATE • CONVENIENT • RELIABLE

We offer a next generation Real-time qPCR



We provide:

- Specimen Collection
- Next Day results
- Easy Access via web portal

Onychomycosis

The most common nail disorder

- 50% of all nail problems
- If neglected can lead to systemic and/or opportunistic infections

* Requires long-term therapy

- Expensive
- Potential side effects of oral antifungals
- New and emerging therapies are genus/species specific
- Precise and timely infection identification
 - Improves clinical outcome
 - Improves quality of life
 - Reduces overall treatment costs
 - Reduces hospitalizations

Organisms Included in PML TFI Assay

- Acremonium strictum
- Alternaria
- Aspergillus fumigatus
- Aspergillus niger
- Aspergillus terreus
- Aspergillus versicolor
- Candida albicans
- Candida parapsilosis
- Curvularia lunata
 - Epidermophyton floccosum
 - Fusarium solani •
 - Geotrichum candidum •
 - Microsporum audouinii
 - Microsporum canis •
 - Microsporum gypseum
 - Microsporum nanum •
 - Neofusicoccum mangiferae
 - Sarcoptes scabiei •
- Scopulariopsis brevicaulis
- Scytalidinum dimidiatum
- Trichophyton interdigitale (T.mentagrophytes)
- Trichophyton rubrum
- Trichophyton soudanense
- Trichophyton violaceum
- Trichophyton tonsurans
- Trichosporon beigelii
- Trichosporon mucoides

Benefits of qPCR vs Traditional Methods

- Full report in 24-48 hours vs up to 6 weeks in traditional fungal culture
- Removes "human factor" in organism identification
- Does not miss slow growing or hard to culture fungi
- Greater precision in identification and characterization of Fungal Infections
- Not limited to the number of pathogens identified per sample
- > Improves clinical outcome
- Clear and easy to interpret reporting including current treatment options for each organism identified

