

UNIVERSAL METHOD FOR PARASITE AND EUKARYOTIC ENDOSYMBIONT IDENTIFICATION

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The Invention

UW-Madison researchers have developed and validated a novel "parasitome" method to accurately characterize parasites in samples. Like published metabarcoding protocols, the new method consists of amplification of a target gene, adding adapter sequences to those amplicons, and deep sequencing them. However, there are three key elements which make the new method unique: 1) a set of newly-designed PCR primers which amplify all parasite groups, 2) a mock community test reagent for standardization ("EukMix"), and 3) a method to reduce interfering host signal using CRISPR-Cas9 digestion with host-specific guide RNAs.

Key Benefits

- · De novo parasite testing
- · Reduces host interference

Additional Information

For More Information About the Inventors

· Tony Goldberg

Tech Fields

- · Animals, Agriculture & Food: Animal health
- <u>Diagnostics & Biomarkers : Diagnostics</u>
- Research Tools: Genomics & proteomics

For current licensing status, please contact Jennifer Gottwald at jennifer@warf.org or 608-960-9854