A Rapid and Sensitive Nucleic Acid Amplification Technique for Mycoplasma Screening of Cell Therapy Products

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INTRODUCTION

Mycoplasma species (spp.) bacteria can infect cell cultures, posing a potential threat to recipients of cell therapy products. Conventional Mycoplasma testing methods are highly sensitive but typically require a minimum of 28 days to produce results. This delay is problematic if rapid results are needed to inform treatment decisions. Nucleic acid amplification technique (NAT) methods have been gaining favor for Mycoplasma testing due to their speed and specificity; however, they must first be qualified as meeting or exceeding the sensitivity of the compendial method. We present herein a NAT method for the detection of Mycoplasma that circumvents the need for live Mycoplasma spp. in the test procedure by instead being qualified using Mycoplasma spp. genomic DNA. We have demonstrated a lower limit of detection that exceeds the regulatory requirements set by Health Canada. This assay is now being used to screen clinical cell therapy products manufactured at our center.

Nucleic acid amplification technique (NAT)-based assays, such as polymerase chain reaction (PCR) techniques, are a potential solution to this issue. NAT-based tests detect the presence of a nucleic acid sequence unique to potentially contaminating microorganisms of interest, and they are highly sensitive and rapidly executable. To meet regulatory requirements, new assays must be qualified in-house to meet or exceed the sensitivity of the compendial methods which, for Mycoplasma spp., is 10 CFU/mL. Qualified PCR-based Mycoplasma detection assays have previously been reported.11–13 These assays used live Mycoplasma spp. for qualification. The use of live Mycoplasma spp. is problematic in facilities that generate cell therapy products because it introduces an unnecessary risk of cell product contamination. Herein, we describe a rapid PCR-based assay that we have qualified for use in testing clinical cell therapy products for Mycoplasma spp. contamination. Briefly, our Mycoplasma detection assay utilizes the commercially available MycoTOOL PCR Mycoplasma detection kit (Roche) with a modified protocol in order to obtain the required 10 CFU/mL sensitivity level. The protocol involves DNA extraction from samples of cell therapy products, amplification of Mycoplasma spp. nucleic acid via highly sensitive staining cell cultures with a fluorescent DNA binding dye followed by visualization by microscopy. The indicator cell culture method is less sensitive than the culture method, with a sensitivity of 100 CFU/mL.10 These conventional methods provide effective Mycoplasma detection; however, they are time-consuming (a minimum of 28 days to perform). This lengthy turnaround time can be problematic in the field of cell therapy, especially for non-cryopreserved cell products that expire quickly (within 24–48 h) and where Mycoplasma test results are needed immediately to inform treatment decisions.

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empirically derived genome copy to CFU (GC/CFU) ratios.14,15 genomic DNA (gDNA) that are converted to CFU/mL values using molecular therapy: methods & clinical development vol. 17 June 2020

usage.10 We qualify to remain unaffected by small but deliberate variations in method parameters.16 Assay robustness.16 was also evaluated as a measure of intermediate precision (measurement of within-laboratory variations)16 was also evaluated as a measure of assay robustness.

RESULTS
To establish the LLOD for the assay, gDNA samples of each Mycoplasma strain were diluted in Elution Buffer (EB) to the indicated concentrations (Figure 1). Positive 400- to 500-bp bands are present in 3 out of 3 reactions for Mycoplasma arginini gDNA diluted to 150 genome copies (GC)/200 μL, and in 10 out of 10 reactions for 15 GC/200 μL. This establishes the LLOD for Mycoplasma arginini at 15 GC/200 μL, which is below the 18 GC/200 μL requirement (equivalent to 10 CFU/mL, as described under “Positive Control gDNA” below and Figure 2). Positive bands are present in 3 out of 3 reactions for Mycoplasma hominis diluted to 1,000 GC/200 μL as well as in 10 out of 10 reactions for 100 GC/200 μL. This establishes the LLOD for Mycoplasma hominis at 100 GC/200 μL, which is also below the 107.2 GC/200 μL requirement.

Specificity Evaluation
The specificity of the assay was analyzed in 17 rounds of testing with CAR-T cell in-process and drug product sample types. All rounds were successful; gels from two representative assays are shown in Figure 3. Both species of Mycoplasma were detected in at least two out of three “spike-test” sample aliquots (Figures 3A and 3B, lanes 3–5), as evidenced by the presence of positive bands in these lanes between 400 and 500 bp. Additional bands (>500 bp) are sporadically observed in Mycoplasma amplification reactions (Figure 3B, M. arginini gel, lane 1). These bands represent non-specific amplification products, and they are not the result of contamination of the PCR reactions. The absence of contamination is confirmed by the lack of positive 400- to 500-bp bands in the Mycoplasma primer PCR negative control reaction (lane 10). The successful amplification of Mycoplasma-specific bands in spike-test samples from both sample types established the specificity of the assay by demonstrating that the assay was capable of detecting the target in the presence of multiple matrices.

Robustness Testing
Four CAR-T cell samples were successfully re-tested by additional operators; the results of two representative assays are shown in Figures 4A and 4B. Gels for each round were analyzed following the acceptance criteria (Materials and Methods) and were found to be in agreement: positive spike-test results as well as β-actin control reactions demonstrate that the assay performs as intended, independent of the operator performing the assay. An additional measure of intermediate precision was executed by the repetition of the assay on the run in-process and drug product samples by the same operator on separate days. The results of run 4 drug product testing are shown in Figures 4C and 4D. Both rounds of testing were successful, and the results are in agreement with each other. The replication of results across different days and by different operators demonstrated that the assay was reliable and established the precision of the assay.

DISCUSSION
We describe herein a protocol for the detection of Mycoplasma spp. that we have qualified for Mycoplasma screening of autologous cell
therapy products. We have demonstrated that the detection levels for both *M. arginini* and *M. hominis* are below the 10 CFU/mL LLOD requirement. We have also demonstrated that the assay detects *Mycoplasma* spp. in the presence of multiple matrices, while repeated rounds of testing established the reliability and robustness of the assay.

The use of the commercial MycoTOOL PCR *Mycoplasma* detection kit in this protocol is advantageous because it has been approved by the US Food and Drug Administration (FDA), Health Canada, and the European Medicines Agency (EMA) for use in *Mycoplasma* spp. testing. Laboratories can leverage data generated by the kit manufacturer in their qualification studies,\(^1\) decreasing the cost of implementation as well as the time necessary to validate their procedures. The universal primers included in the kit target the 16S rRNA sequence conserved across multiple *Mycoplasma* strains,\(^1\) including those representing 90%–95% of *Mycoplasma* cell culture contaminations.\(^1\) This allows for the detection of a wide variety of *Mycoplasma* strains, including fastidious strains that are difficult to detect even by conventional growth-based methods.

The strategy of using gDNA to establish sensitivity enables widespread adoption since the specialized equipment, reagents, and knowledge required for the cultivation of *Mycoplasma* spp. is not necessary, in addition to avoiding cross-contamination of sensitive cultures. This approach has previously been described;\(^1\) however, for qualification of the assay we describe herein, we take the additional step of accounting for the 1/10 sampling at the PCR level in the calculations of LLOD. In our hands, this was necessary for successful qualification of the assay.

This *Mycoplasma* Detection Assay fulfills an important need in the cell therapy field for a rapid assay that can facilitate the prompt decision-making that is critical for the use of live cell therapies. The use of a PCR-based assay avoids the cultivation time necessary for compendial methods while still performing at the required level of sensitivity. The use of gDNA as a positive control in place of live *Mycoplasma* spp. allows the assay to be performed in-house, further decreasing the time required to produce results. We share details of this protocol in the hopes that other laboratories generating biologics that require *Mycoplasma* testing can adopt this approach and expedite *Mycoplasma* testing of their products to ensure patient safety.

**MATERIALS AND METHODS**

**Mycoplasma Detection Assay**

This assay requires standard precautions for PCR setup such as the use of dedicated workstations, filtered tips, and DNA-free materials and reagents. Additionally, the pre-PCR and post-PCR products were spatially segregated, and extreme care was taken to not cross-contaminate samples with positive control *Mycoplasma* gDNA during sample processing and PCR setup steps.

**Positive Control gDNA**

gDNA samples from *M. arginini* strain G230 (ATCC qCRM-23838D) and *M. hominis* strain LBD-4 (ATCC qCRM-27545D) were used to establish LLOD for the assay, as well as to assess matrix interference in cell therapy samples. These strains were chosen as representative of the upper (*M. hominis*, 10 CFU/mL) and lower (*M. arginini*, 0.1 CFU/mL) detection levels of the MycoTOOL PCR *Mycoplasma* detection kit.\(^1\)

*Mycoplasma* gDNA was used to measure assay sensitivity by adding known numbers of genome copies to PCR reactions and assessing amplification success. The GC/CFU ratio for each species was used to convert the desired CFU number to the corresponding genome copies value. We used previously published GC/CFU ratios of 23.82 for *M. hominis* strain LBD-4\(^1\) and 4.0 for *M. arginini* strain G230\(^3\) to calculate the minimum GC that must be detected in the assay in order to satisfy the 10 CFU/mL LLOD requirement. In our assay, a sample of cell product is divided into 450 μL aliquots. DNA is isolated from each aliquot and each sample of isolated DNA has a final volume of 200 μL (Figure 2). The 10 CFU/mL detection requirement means that we need to detect 4.5 CFU in each 450 μL sample aliquot, corresponding to 18 GC of *M. arginini* or 107.2 GC of *M. hominis* gDNA. Each 450 μL sample generates DNA in a final volume of 200 μL, and thus the detection requirement for *M. arginini* is 18 GC/200 μL and 107.2 GC/200 μL for *M. hominis*.

**Primers**

Universal *Mycoplasma* primer A (forward, 5′-GGCGAATGGGTG AGTAACACG-3′) and primer B (reverse, 5′-CGGATAACGC TTGGCAGCTATG-3′) targeting the 16S rRNA gene, originally
described by Wong-Lee and Lovett, are included in the MycoTOOL kit. This primer set has been validated for the detection of *Mycoplasma fermentans*, *Acholeplasma laidlawii*, *Mycoplasma hyorhinis*, *Mycoplasma orale*, *Mycoplasma pneumoniae*, *M. arginini*, *Spiroplasma citri*, *Mycoplasma salivarium*, and *M. hominis* at 10 CFU/mL and it has limited cross-reactivity to phylogenetically similar Gram-positive *Lactobacillus acidophilus*, *Streptococcus bovis*, and *Clostridium sporogenes*.

The MycoTOOL kit includes control primers to amplify Gapdh from the Chinese hamster ovary (CHO) cell line. This primer set is unable to amplify GAPDH in human cells. Consequently, primers targeting human β-actin (hACTB393.f, hACTB642.r; PrimerBank ID: 4501885a1)19 were used to confirm cell lysis and DNA recovery in control PCR reactions. Primers were synthesized by Integrated DNA Technologies at a 25-nmol scale with standard desalting. They were resuspended to 100 μM with TE (100 mM Tris-HCl, 1 mM EDTA) (Molecular Probes) and prepared as a 10 μM mixture (5 μM each primer) with EB buffer (QIAGEN) for use in PCR setup.

**DNA Extraction**

Cell therapy samples were processed using a QC Sample Preparation Kit (Roche). All reagents were included in the kit unless otherwise specified. The DNA extraction procedure is dependent on cell density: samples \( \leq 5 \times 10^6 \) cells/mL are processed following the standard DNA preparation protocol, while samples \( >5 \times 10^6 \) to \( 1 \times 10^8 \) cells/mL are processed following the high cell density DNA preparation protocol.

**Standard DNA Preparation: \( \leq 5 \times 10^6 \) Total Cells/mL**

Cell samples (\( \geq 2 \) mL) were divided into four aliquots of 450 μL each (Figure 2). 50 μL of proteinase K and 450 μL of lysis buffer were added to each vial followed by vortexing three times for 5-s durations. Samples were incubated for 15 min at 56°C/600 rpm in a Thermomixer R with 2.0 mL block (Eppendorf). 630 μL of precipitation reagent and 2 μL of GlycoBlue coprecipitant (Invitrogen) were added to each vial, followed by 20 inversions and vortexing for 5 s. Samples were then centrifuged for 3 min at 16,000 \( \times \) g, and supernatants were removed by pipetting. 1 mL of washing buffer was used to wash each pellet. Vials were inverted five times to mix and DNA was pelleted by centrifugation for 3 min at 16,000 \( \times \) g. Supernatants were completely removed by pipetting. 190 μL of dilution reagent was added to each of the four sample vials, and DNA was resuspended by incubating in the thermomixer at 80°C/900 rpm for 10 min followed by brief vortexing.

**High Cell Density DNA Preparation: \( >5 \times 10^6 \) to \( 1 \times 10^8 \) Total Cells/mL**

Two 950 μL aliquots of cell sample were diluted with 950 μL of DNA-free water (Figure S1). Each of the 1,900 μL diluted aliquots was split further into four 450 μL aliquots, for a total of eight 450 μL samples. 50 μL of proteinase K and 700 μL of lysis buffer were added to each vial, followed by vortexing three times for 5-s durations. Samples were incubated for 30 min at 56°C/600 rpm in a Thermomixer R with 2.0 mL block (Eppendorf). 800 μL of precipitation reagent and 2 μL of GlycoBlue coprecipitant (Invitrogen) were added to each vial followed by 20 inversions and vortexing for 5 s. Samples were then centrifuged for 3 min at 16,000 \( \times \) g, and supernatants were removed by pipetting. 1 mL of washing buffer was used to wash each pellet. Vials were inverted five times to mix and DNA was pelleted by centrifugation for 3 min at 16,000 \( \times \) g. Supernatants were completely removed by pipetting. 95 μL of dilution reagent was added to each of the eight sample vials and DNA was resuspended by incubating in the thermomixer at 80°C/900 rpm for 15 min, followed by...
brief vortexing. Pairs of tubes were pooled to generate a total of four vials containing 190 μL of DNA sample in each.

**Endpoint PCR Assay**

PCR was performed using the MycoTOOL Mycoplasma detection amplification kit (Roche) with two modifications: (1) the CHO-specific Gapdh primer set included with the kit was replaced with a human β-actin primer set; and (2) the PCR reactions were scaled up from 50 μL to 100 μL total volume. This change was necessary to be able to add more input DNA to the reaction, in order to fulfill the 10 CFU/mL sensitivity requirement. We were unable to establish a LOD at or below 10 CFU/mL with the original volumes. All other reagents used are included with the kit. Two master mixes were prepared: one to amplify Mycoplasma spp.-specific templates, with the other targeting β-actin. Each 60 μL of master mix contained 1.4 μL of RM1a, 20 μL of RM1b, 14 μL of MgCl₂ (25 mM), 2 μL of primer mix Mycoplasma or 10 μL of β-actin primer mix (10 μM, 5 μM each primer), 4 μL of detection dye, and 18.6 μL (for Mycoplasma master mix) or 10.6 μL (for β-actin master mix) of PCR-grade H₂O. 60 μL of each master mix was aliquoted to reaction tubes.

Figure 4. Intermediate Precision Testing

The Mycoplasma Detection Assay was run on samples multiple times to demonstrate the reliability of the assay. (A and B) Run 2 in-process samples were tested by operator 1 (A) and operator 2 (B), (C and D) Run 4 drug product samples were tested twice by operator 1 (C [round 1] and D [round 2]). Lane 1, positive control: 18 GC M. arginini or 107.2 GC M. hominis; lane 2, positive control: 1,800 GC M. arginini or 10,720 GC M. hominis; lanes 3–5 and 11, M. arginini or M. hominis spike-test controls; lane 6, blank; lanes 7, 8, and 10, test sample; lanes 9 and 12, negative control; lane L, ladder; lane B, blank. Mycoplasma-specific expected band size was 400-500 bp; β-actin was 250 bp.
Samples were added to reactions in the following order: (1) negative control, (2) test samples, (3) spike-test samples, and (4) PCR-positive controls. 40 µL of EB was added to each of the Mycoplasma and β-actin negative control reactions. 10 µL of EB was added to two of the test samples to bring the final volume to 200 µL. 40 µL of each test sample was run in duplicate in Mycoplasma-specific amplification reactions. To generate the spike-test samples, M. arginini and M. hominis gDNA samples were diluted to the appropriate concentrations and then 10 µL of each diluted sample, respectively, was spiked into the two remaining test sample aliquots (Figure 2). 40 µL of each of these spike-test samples was then added to the Mycoplasma-specific amplification reactions, in triplicate. Each of the test and spike-test control samples (40 µL) was also used as template in β-actin amplification reactions. PCR-positive controls consisted of suitable dilutions of M. arginini and M. hominis gDNA spiked into EB, which were then added to Mycoplasma-specific master mix. All reactions were mixed by pipetting and subjected to PCR cycling.

**PCR Cycling and Amplicon Detection**

Reactions were cycled on a Bio-Rad Dyad thermal cycler in a touchdown PCR program as described in the MycoTOOL Mycoplasma detection amplification kit instructions: samples were incubated at 40°C for 5 min in a carryover prevention step, followed by initial denaturation at 94°C for 10 min. The touchdown portion of the program includes 20 cycles with a denaturation step at 94°C for 30 s, annealing for 30 s with a decreasing temperature profile (2 cycles each at 70°C, 69°C, 68°C, 67°C, 66°C, 65°C, 64°C, 63°C, 62°C, and 61°C), and an elongation step at 72°C for 45 s. This was followed by 25 cycles with a denaturation step at 94°C for 30 s, an annealing step at 60°C for 30 s, and an elongation step at 72°C for 45 s. The final elongation was at 72°C for 4 min, and all samples were then held at 4°C.

Following amplification, 12 µL of PCR product was added to 3 µL of 5× Hi-Density TBE (Tris-borate-EDTA) sample buffer (Invitrogen) and mixed by pipetting up and down. A DNA molecular weight marker (included with kit) was prepared by mixing 16 µL of molecular weight marker with 24 µL of 1× TBE-electrophoresis buffer (Invitrogen), 8 µL of Hi-Density TBE sample buffer, and 1.6 µL of detection dye (included with kit). 10 µL of PCR sample or molecular weight marker was loaded per lane on Novex 6% TBE gels (Invitrogen) and subjected to electrophoresis in XCell SureLock mini-cell (Thermo Fisher Scientific) using a Bio-Rad PowerPac HC at 200 V for 40 min. Gels were visualized on a FLA 9500 (GE Healthcare) using the SYBR Safe (473 nm) settings. Gel results were analyzed based on the acceptance criteria listed below:

1.1.1. The *Mycoplasma* expected band size is ~450 bp; any bands between 400 and 500 bp in size are considered a positive result.11 The expected size for the β-actin band is 250 bp.

1.1.2. The *Mycoplasma* and β-actin negative control lanes do not contain a band of the expected sizes, to rule out contamination of the PCR reactions.

1.1.3. The β-actin control reactions all contain the expected 250-bp band to ensure that the cell lysis and DNA isolation were successful.

1.1.4. At least two of the three spike-test reactions have a positive ~450-bp band to confirm that there was no interference in the PCR reactions by the sample matrix.

1.1.5. If the spike-test controls fail, the results of the positive control reactions can be used to assess whether there was an issue with the PCR reagents/setup/cycling.

1.1.6. If there are 400- to 500-bp bands in any of the four test sample lanes, the sample is considered positive for *Mycoplasma* spp.

**Assay Qualification**

The *Mycoplasma* detection assay qualification process included determination of LLOD, specificity, and intermediate precision testing.

**LLOD Determination**

Regulators require that NAT-based *Mycoplasma* detection assays are validated against a panel of species including *A. laidlawii*, *M. fermentans*, *M. hyorhinis*, *M. orale*, and *M. arginini*.10 The ability of the MycoTOOL PCR Mycoplasma detection kit to detect all of the required species has been previously established11 and was not repeated in our qualification. Instead, we selected two species that were detected with the highest (M. arginini, 0.1 CFU/mL) and lowest (*M. hominis*, 10 CFU/mL) sensitivity using the MycoTOOL kit11 to evaluate the performance of the kit at our center.

*M. arginini* and *M. hominis* gDNA samples were diluted and then spiked into EB at 200 µL total volume to test whether the LLOD was below the 18 GC/200 µL (*M. arginini*) or 107.2 GC/200 µL (*M. hominis*) requirements. *M. arginini* was diluted to 15 and 150 GC/200 µL, and *M. hominis* was diluted to 100 and 1,000 GC/200 µL. 40 µL of the spiked EB dilutions were added to tubes containing 60 µL of *Mycoplasma* master mix and mixed by pipetting. PCR cycling and amplicon detection was performed as described in “PCR Cycling and Amplicon Detection” above.

**Specificity Testing**

Specifity evaluation for NAT-based assays requires confirmation that the test specifically detects a target nucleic acid. As discussed above, the ability of the universal *Mycoplasma* primer set to detect the required panel of *Mycoplasma* species was previously established.11 We were able to leverage these data and did not need to perform this aspect of specificity testing in our qualification process.

Specificity is also a measure of the ability to detect a target in the presence of matrix components, which are any substances present in samples in addition to the target of interest. These substances may interfere with template amplification, and thus each sample type to be tested with the *Mycoplasma* detection assay must be evaluated. We initially developed the *Mycoplasma* detection assay to screen both in-process and final drug product samples generated during CAR-T cell production runs using the CliniMACS Prodigy system.
CAR-T production runs were used in 17 rounds of testing to evaluate (CSL Behring). In-process and drug product samples from eight (Sandoz) and interleukin-7/-15 (Miltenyi Biotec). The drug product medium (Miltenyi Biotec) supplemented with gentamicin sulfate of the CAR-T culture process and consists of cells in TexMACS GMP analysts.16 variation, such as assay performance on different days or by different

Intermediate Precision/Robustness Testing

The robustness of the endpoint MycoTOOL kit was previously established by testing the performance of the assay across different kit lots.11 To qualify the assay for use in our center, we evaluated the intermediate precision as an additional measure of the overall robustness of the assay. Intermediate precision measures within-laboratory variation, such as assay performance on different days or by different analysts.16

Two additional operators re-tested four samples in six separate assays to assess any effect of different analysts on the assay outcome. Two samples were also re-tested by the same operator to evaluate assay performance on separate days (Figure S2).

SUPPLEMENTAL INFORMATION

Supplemental Information can be found online at https://doi.org/10.1016/j.omtm.2020.01.009.

AUTHOR CONTRIBUTIONS

L.D., M.C., R.A.H., and M.B. designed the qualification study plan. L.D., E.Y., and L.L. performed the experiments. J.R.W. and B.H.N. provided samples. L.D. and R.A.H. wrote the manuscript. R.A.H, K.A.H, B.H.N., and N.K. provided supervision. All authors reviewed and edited the manuscript.

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