The Development of a Clinically Relevant, High-Throughput High-Content Imaging Assay For Leishmania

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Introduction:

Leishmaniasis is caused by protozoan parasites of the genus Leishmania. The disease is prevalent in over 90 countries and manifests as one of 3 forms: viseral, mucocutaneous or cutaneous. Viseral leishmaniasis is fatal without treatment while mucocutaneous and cutaneous are extremely disfiguring.

The parasite is transmitted by sandfiles and following infection the metacyclic promastigotes are engulfed by macrophages where they transform and multiply as amastigotes causing the clinical symptoms of leishmani-

A efficacious, non-toxic, easily administrable drug is urgently needed. The only drugs available are toxic with long administration schedules and resistance is a growing concern.

not translate to in-vivo efficacy. However, these assays are often not clinically relevant, poorly reproducible and frequently in-vitro activity does Numerous assays have been developed using a variety of life-cycle stages, assay platforms and technologies.

Aim:

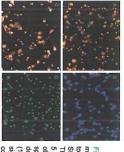
To develop a high-content, high-throughput imaging assay to enable indentification of compounds against Leishmania donovani DD8 intracellular amastigotes.

Methods:

myristate 13-acetate (PMA) cells/mL in presence of 25ng/mL phorbol 12-Plating and transfor-mation of THP-1 (Host) cells at 2.5x10⁶ The leishmania intracellular amastigote assay invovles multiple steps over a 8 day period, summarized below 24 Hours Washing of THP-1 cells remove PMA 48 Hours cells Addition of metacyclic promastigotes to host Day 8: Day 4: 24 Hours Day 8: 72 Hours Addition of drugs internalised parasites. Removal of non-

site nucleic material. Five commercially available nucleic acid stains were evaluated for their ability to selectively stain host cell and para-

2. Selection of Nuclear Acid Stain



to parasite), imaged on a PerkinElmer Opera at 20x magnification. Stained with: A) Acridine Orange (10nM), staining was non-specific B) specific staining of host cell and parasite nuclei observed but difficult to detect parasites within host cell cytoplasm. E) SYBR® Green Thiazole Orange (2.1uM), non-specific staining was observed. C) Draq 5 (5uM), specific staining of host cell and parasite nucleic material was metacyclic promastigotes at a multiplicity of infection of 1:5 (host cells signal to background coupled with a high flurorescent intensity at low (1/10000), specific staining of host cell and parasite nuclei and a large observed but the signal to background was low. D) Hoechst (10uM), igure 2: THP-1 cells, 2.5x105 cells/mL infected with L. donovani DD8

F) SYBR® Green and HCS CellMask™ Deep Red Stain overlay. SYBR® Green when used in combination with HCS CellMask Deep Red easily and clearly allowed the leishmania amastigotes to be visuconcentrations led to this dye being selected

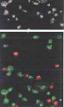
Script Development

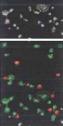
In order to identify lesihmania amastigotes within host cells and thus quantify the number of infected cells an image analysis script was developed using the building blocks within the PerkinElmer Opera Columbus software.











using the find nuclei building block using the SYBR® Green stain image captured on exposure 1, camera 2, Em 540 nm. Size exclusion was used to excluded non-vaible host cells. ©) The find spot tool was utilised to identify amastigotes present within the cytoplasm of the THP-1 host cells using the SYBR® Green stain. D) Host cells with greater than 3 'spots' (amastigotes) in the cytoplasm were classed as infected (Highlighted in green). Figure 3: A)The find cytoplasm building block was used to identify the cytoplasm of the host cells using the HCS-CellMask** Deep Red stain captured on exposure 1, camera 3, Em 690 nm B) Host cell nuclei were identified

Results

PerkinElmer Columbus

Imaging of assay plates on PerkinElmer Opera

Assay plates fixed with 4% paraformaldehyde and stained with SYBR® Green and HCS CellMask™ Deep Red Stain

use in the assay.

Determination of Optimal Host Cells
 The macrophage cell lines; Raw 264.7, J774.1 and THP-1 were extensively profiled in terms of growth rate, infectivity and amenability to high content image (HCI) analysis to determine the most suitable cell line for





Figure 1: Whole well montages (64 fields) of Raw 264.7 (4x10³ cells/mL) (A), THP-1 (2.5x10⁵ cells/mL) (B) and J/74.1 (4x10˚ cells/mL) (C) cells seeded in 394 well CellCarner-384 optically clear bottom tissue outure treated plates (PerkinElmer), stained with HCS CellMask™ Green, imaged on a PerkinElmer Opera at 20X magnification, exposure 1, camera 2 Em 540 nM. The non-contact inhibited growth and morphology of PAW 264.7 (A) and J774.1 (B) cells meant that the cell lines were not compatible with a HCl based assay. THP-1 cells (B) were therefore selected as the host cells in the assay.

4. Establishing Reference Drug Activity

lished L. donovani DD8 intracellular amastigote assay A panel of reference compounds currently used for the treatment of leishmaniasis were screened using the newly estab-

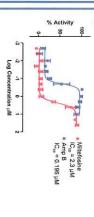








Figure 4: A) The dose-response curves of miltefosine and amphotericin B against intracellular L. donovani DD8 amasligotes. The IC_{0x} values were comparable to previous reported values. THP-cells (2,5xtl0* cells/ml,) infected with L.
donovani DD8 amastigotes at a multiplicity of 1.5 stained with SYBR9 forem and HC8 CellMask™ Deep Red imaged
on a PerkinElmer opera at 20X. Following treatment with 10µM of miltefosine (B) and amphotericin B (C) no amastigotes are detected within the cytoplasm of host cells. In contrast following treatment with 0.1µM of miltefosine (D) and amphotericin B (E) amastigotes could clearly be detected within the cytoplasm of host cells.

