## A Systematic Review of Virulence Factors in the Leishmania Genus **Osaru Omoruna,**<sup>1</sup> Avinash N. Mukkala,<sup>2</sup> Ruwandi Kariyawasam,<sup>2</sup> Eric Shao,<sup>3</sup> Priyanka Challa,<sup>4</sup> Michael Klowak,<sup>2</sup> Tianna Chong-Kit,<sup>5</sup> Olamide Egbewumi,<sup>4</sup> Shareese Clarke,<sup>6</sup> Dylan Kain,<sup>7</sup> Jamie Sookhoo<sup>8</sup>, Andrea K. Boggild<sup>2,7,8,9</sup>

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•Leishmaniasis is a neglected tropical disease divided into three major classifications based on clinical presentation: cutaneous (CL), mucocutaneous (MCL) and visceral (VL) •Transmitted by the *Lutzomyia* spp. and *Phlebotomus* spp. sandflies, there are up to 2 million cases of Leishmaniasis globally while 350 million people are at risk •Parasite-determined factors play a complementary role in the pathogenesis of leishmaniasis •Virulence factors (VFs), or pathogen moieties facilitating disease, can potentiate host cell damage by *Leishmania* spp. by increased expression, host cell invasion, stress tolerance, and modulation of the host immune system •Due to large eukaryotic genomes in *Leishmania* spp., there is a wide array of VFs which contribute to different aspects of pathogenesis; we aim to synthesize this knowledge by systematically mapping the literature







TORONTO

 Some common parasite-derived pathogenesis mechanisms in *Leishmania* include: Heat shock adaptation to the host environment

	<ul> <li>stabilization of proteins in stressful host environments</li> <li>Significant expression changes in HSPs as parasite is engulfed in host cells</li> <li>Aid in adapting from poikilothermic insect vector to a homeothermic mammalian host</li> </ul>	
HSP70		
HSP83		
HSP90		
HSP100		
HSP65		
LPG	<ul> <li>Lipophosphoglycan</li> <li>Cell surface anchored molecule</li> <li>Species-specific sugar component</li> </ul>	

• Required to cause infection in the sandfly hindgut

- Evading the immune system
- Increased expression of survival factors
- Preventing innate immunity opsonisation
- Modulation of the host immune system
- Heat shock is mainly directed by heat shock proteins (HSPs):
  - Different HSPs are used preferentially in different species
- HSP23 can protect against thermal, acidic and oxidative stresses
- CyP40 is thought to be a co-chaperone that helps the parasite infect macrophages
- Loss of HSP100 renders L. major and L. donovani non-infective in vitro at physiological temperatures Heat shock and resulting thermotolerance is a crucial

## METHODS

• PubMed (NCBI), MEDLINE (OVID), EMBASE (OVID), Web of Science, and LILACS (VHL) were searched from inception to July 2018 using combinations of the search terms "virulence factor\*", "Leishmania", and "Leishmaniasis\*", while accounting for unique database syntax

 Iterative inclusion and exclusion of search terms was employed to maximize relevant article extraction • Primarily, molecular and mechanistic pathogenesis studies in various model systems, observational studies, review studies, cohort studies, as well as

- Metalloprotease **GP63** 
  - Cleaves C3b complement
  - Halts and hinders innate
    - immunity
  - Protects parasite from cell lysis
- CPB Lowered virulence in
  - macrophages
  - Lowered virulence in mice
  - Required to cause infection
- Elongation factor that is part EF-1alpha of the parasite exosome
  - Blocks Nitric Oxide
    - production
  - Promotes survival

• Exacerbate parasite-derived **A2** immunopathogenesis • Significant in visceral leishmaniasis

method by which *Leishmania* species exert their virulence

## DISCUSSION

• The ability to comprehensively synthesize all the known literature around parasite-determined virulence factors can open new doors into networklevel pathogenesis

- Connecting the dots between virulence factors (if any) to construct a more complete picture of parasite pathogenesis can help better illuminate the underpinnings of different disease manifestations • Once all parasite-determined VFs are mapped, it can elucidate how they may tie into host-determined

clinical trials are included

• Synthesis is done by grouping of similar VFs in similar pathogenesis mechanisms, e.g. heat shock • 760 MEDLINE, 1942 PubMed, 1314 EMBASE, 438 Web of Science, and 8 LILACS records were retrieved for title and abstract screening; after a multi-step deduplication pipeline, 2620 remained • All records undergo double-reviewer screening, with tertiary arbitrators to mitigate any discrepancies

• Catalyze the interconversion MPI of F6P and M6P

- Required for glycoconjugates
- Loss of MPI leads to loss of
  - surface-anchored VF

synthesis, such as

leishmanolysin

immunopathogenesis mechanisms

- Being able to modulate some of these network-level systems can potentially identify novel targets for therapeutics and diagnostics
- This systematic review has implications for painting a more full picture of parasite-determined *Leishmania* pathogenesis and hence help tie the ends between different VFs, and maybe shed light into host environmental factors

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