

## Intracellular protozoan parasites of humans: the role of molecular chaperones in development and pathogenesis

This is the Accepted version of the following publication

Shonhai, Addmore, Przyborski, Jude M, Maier, Alexander G and Blatch, Gregory (2011) Intracellular protozoan parasites of humans: the role of molecular chaperones in development and pathogenesis. Protein and Peptide Letters, 18 (2). pp. 143-157. ISSN 0929-8665 (print) 1875-5305 (online)

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Table 1: The diversity and potential role in development and pathology of Hsp100, Hsp90, Hsp70 and Hsp40 chaperones of intracellular protozoan parasites

Chaperone	Leishmania spp.		Trypanosoma cruzi		Plasmodium falciparum		Toxoplasma gondii	
	Number <sup>1</sup>	Dev/Path <sup>2</sup>	Number <sup>1</sup>	Dev/Path <sup>2</sup>	Number <sup>1</sup>	Dev/Path <sup>2</sup>	Number <sup>1</sup>	Dev/Path <sup>2</sup>
Hsp100	1	Hsp100						
Hsp90/HspC:								
Hsp90α/HspC2	17	Hsp90	6	TcHsp83	1	PfHsp90	1	TgHsp90
(cytosolic inducible)							1	
Hsp90β/HspC3							1	
(cytosolic constitutive) Grp94/HspC4 (ER)	1	Grp94	3		1		1	
TRAP1/HspC5	1	G1p34	2		2		1	
(mitochondrial)	1		2		2		1	
Hsp70/HspA:								
Hsp72/HspA1A	2	Hsp70	2		2	PfHsp70-1	1	TgHsp70
(cytosolic inducible)		-				•		
Hsp73/Hsc70/HspA1B								
(cytosolic constitutive)								
BiP/HSPA5 (ER)	1	BiP	1		1		1	
Hsp70/HspA9	5		1		1		1	
(mitochondrial)								
Other isoforms:							1	
Grp170/Hsp110/HspH	2		3		2		1	
Hsp70.a	1		2	TcHsp70.a				
Hsp70.b	1		1					
Hsp70.c	l		$\frac{1}{?^3}$	T II 70 4				
Hsp70.4	1		?	TcHsp70.4				
Hsp40/DnaJ:			_		_		_	
Type I (DnaJA)			3	Tcj2	2	Pfj1	2	
Type II (DnaJB)			1	T D11	9	PFA0660w	3	
Type III (DnaJC)			2	TcDJ1	21	DECA.	26	
Type IV					12	RESA; PF10_0381	2	
Uncategorized	66		61			1110_0001		
(types I-IV)								

The number of chaperones for *L. major* and *T. cruzi* [24], *P. falciparum* [14, 25] and *T. gondii* [14, 131, 144, 148] are listed; where there is no listing indicated for a particular chaperone, these proteins are either not well characterized or have not been discussed in the text

<sup>&</sup>lt;sup>2</sup> Those chaperones potentially involved in development and/or pathogenesis are listed by a common name; for *Leishmania* spp. the species is not designated as a prefix since the information is obtained from more than one species; see text for details

<sup>&</sup>lt;sup>3</sup>LmHsp70.4 has been annotated on the *L. major* genome and a TcHsp70.4 was identified on the basis of amino acid sequence identity to LmHsp70 [65], but has not been annotated on the *T. cruzi* genome