

Master Biologie-Santé

UE Microbiologie-Pathologies

Rôle des vecteurs dans la
transmission des parasites :
Effet de la salive sur l'infection

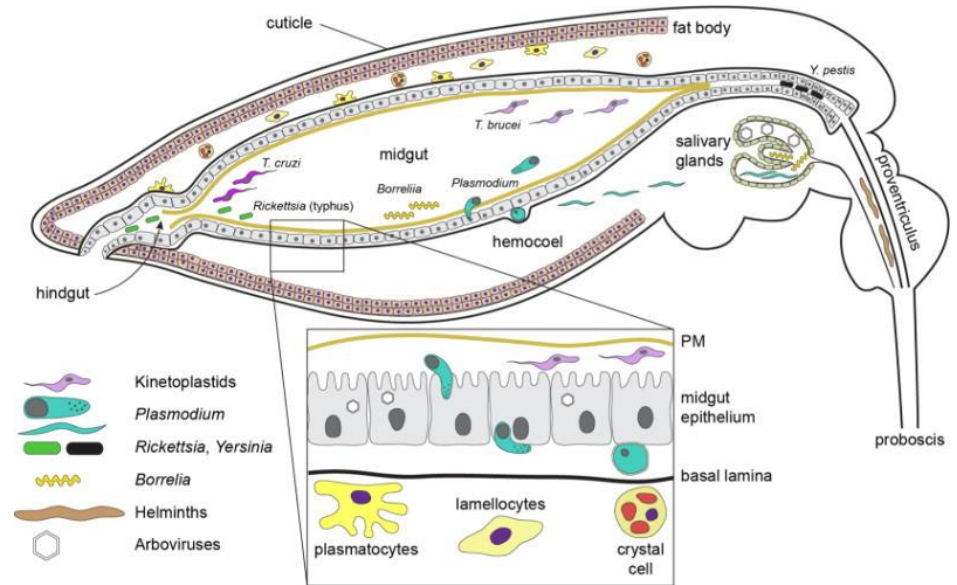
Introduction

Table 1 Taxonomic classification of major vector-borne diseases

Order	Vectors		Diseases
	Family	Genus	
Diptera	Culicidae	<i>Anopheles</i>	Malaria, Lymphatic filariasis
		<i>Culex</i>	West Nile disease Japanese encephalitis
		<i>Aedes</i>	Yellow fever Chikungunya Dengue
	Psychodidae	<i>Phlebotomus</i>	Leishmaniasis
		<i>Lutzomyia</i>	
	Glossinidae	<i>Glossina</i>	Human African Trypanosomiasis
	Simuliidae	<i>Simulium</i>	Onchocerciasis
Tabanidae	<i>Tabanus</i>	Loiasis	
Hemiptera	Reduviidae	<i>Triatoma</i>	Chagas disease
		<i>Rhodnius</i>	
Ixodida	Ixodidae	<i>Amblyomma</i>	Rickettsiosis Tularemia
		<i>Ixodes</i>	Lyme disease Babesiosis
		<i>Haemaphysalis</i>	Tularemia
	Argasidae	<i>Ornithodoros</i>	Tick borne encephalitis Relapsing fever

The taxonomic classification of the major hematophagous arthropod vectors described in the present review is given with their corresponding diseases.

Fontaine et al. *Parasites & Vectors* 2011, 4:187



Baxter et al.

Biochemistry. Author manuscript; available in PMC 2018 February 21.

Telmophage ≠ solénophage

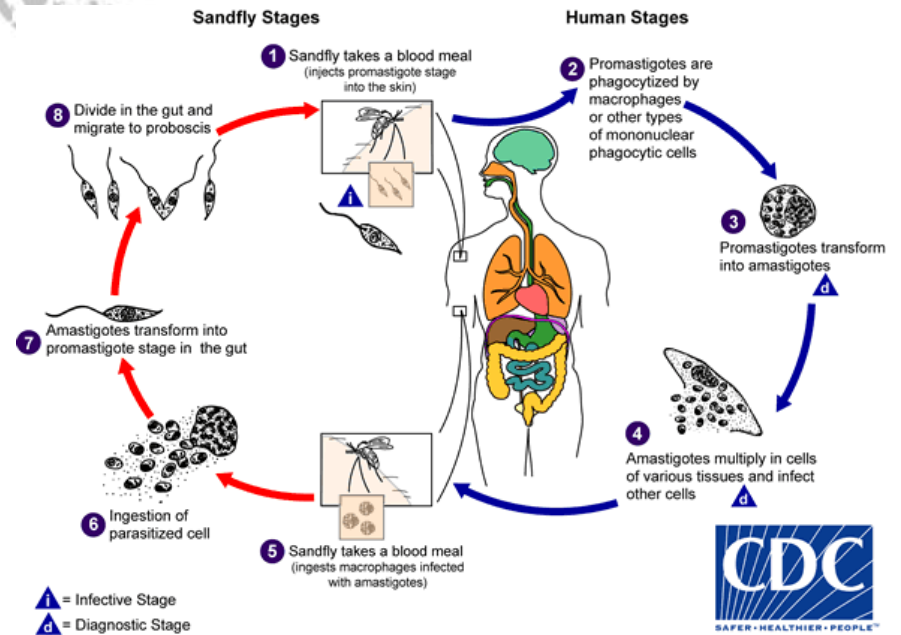
Leishmaniose

Table 1
Medically important species of phlebotomine sand fly and transmission of leishmaniasis

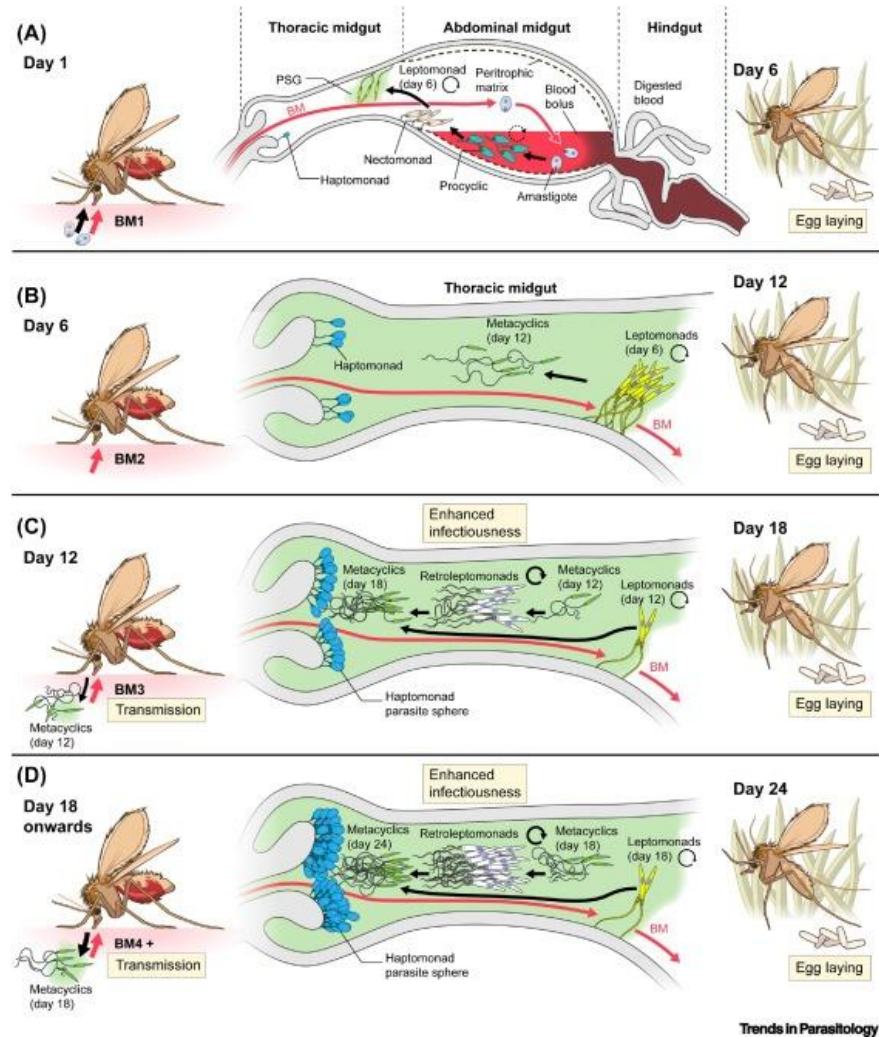
Sand fly species	Geographical distribution	Species of <i>Leishmania</i>	Main disease(s) in humans	Transmission	Important mammalian hosts
<i>Phlebotomus papatasi</i> , <i>Phlebotomus dubosqi</i> , <i>Phlebotomus salehi</i>	Central and West Asia, North Africa, Sahel of Africa, Central and West Africa	<i>Leishmania (Leishmania) major</i>	Cutaneous (oriental sore)	Rural zoonotic	Great gerbil (<i>Rhombomys opimus</i>), fat sand rat (<i>Psammomys obesus</i>)
<i>Phlebotomus sergenti</i>	Central and West Asia, North Africa	<i>Leishmania (Leishmania) tropica</i>	Cutaneous (oriental sore)	Urban anthroponotic	Humans, rock hyraxes
<i>Phlebotomus longipes</i> , <i>Phlebotomus pedifer</i>	Ethiopia, Kenya	<i>Leishmania (Leishmania) aethiopica</i>	Cutaneous diffuse cutaneous	Rural zoonotic	Rock hyraxes (<i>Heterohyrax brucei</i> , <i>Procavia</i> spp.)
<i>Phlebotomus argentipes</i> , <i>Phlebotomus orientalis</i> , <i>Phlebotomus martini</i>	Indian subcontinent, East Africa	<i>Leishmania (Leishmania) donovani</i>	Visceral (kala azar)	Epidemic anthroponotic	Humans
<i>Phlebotomus ariasi</i> , <i>Phlebotomus perniciosus</i>	Mediterranean basin, Central and West Asia	<i>Leishmania (Leishmania) infantum</i>	Infantile visceral	Zoonotic peridomestic	Domestic dog
<i>Lutzomyia longipalpis</i>	Central and South America	<i>L. (L.) infantum (syn. chagasi)</i>	Infantile visceral	Zoonotic peridomestic	Domestic dog, foxes (<i>Lycalopex vetulus</i> , <i>Cerdocyon thous</i>)
<i>Lutzomyia olmeca olmeca</i>	Central America	<i>Leishmania (Leishmania) mexicana</i>	Cutaneous (chiclero's ulcer)	Sylvatic zoonotic	Forest rodents (<i>Ototylomys phyllotis</i> + others)
<i>Lutzomyia flaviscutellata</i>	South America	<i>Leishmania (Leishmania) amazonensis</i>	Cutaneous	Sylvatic zoonotic	Forest rodents (<i>Proechimys</i> spp. + others)
<i>Lutzomyia wellcomei</i> , <i>Lutzomyia complexus</i> , <i>Lutzomyia carrerei</i>	Central and South America	<i>Leishmania (Viannia) braziliensis</i>	Cutaneous mucocutaneous (espundia)	Sylvatic zoonotic	Forest rodents (<i>Akodon</i> spp., <i>Proechimys</i> spp. + others)
<i>Lutzomyia peruensis</i> , <i>Lutzomyia verrucarum</i>	Peru	<i>Leishmania (Viannia) peruviana</i>	Cutaneous (uta)	Upland zoonotic	Reservoir unknown, dog?
<i>Lutzomyia umbratilis</i>	South America	<i>Leishmania (Viannia) guyanensis</i>	Cutaneous, often metastatic (pian-bois)	Sylvatic zoonotic	Sloth (<i>Choloepus didactylus</i>), anteater (<i>Tamandua tetradactyla</i>)
<i>Lutzomyia trapidoi</i>	Central America	<i>Leishmania (Viannia) panamensis</i>	Cutaneous	Sylvatic zoonotic	Sloth (<i>Choloepus hoffmanni</i>)

Various species in the genus *Phlebotomus* are responsible for transmission of leishmaniasis in the Old World and *Lutzomyia* species in the New World. Each sand fly species typically transmits only one species of parasite and each parasite leads to a particular type of disease. Animal reservoirs are important for maintaining the life cycle of many *Leishmania* species and consequently transmission is frequently zoonotic and rural/sylvatic. Important exceptions are *Leishmania tropica* and *Leishmania donovani*, which are transmitted between human beings.

Leishmaniose



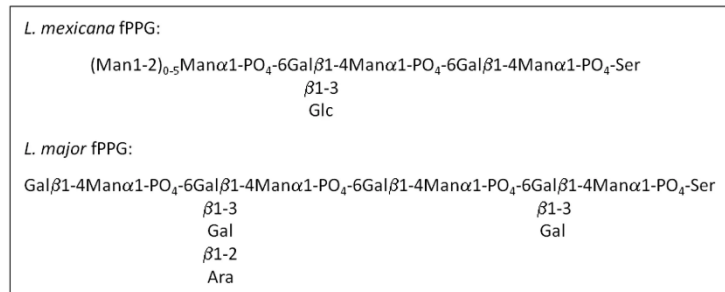
Leishmaniose



Leishmaniose

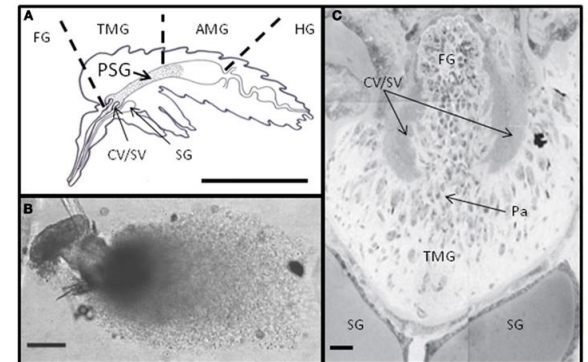
- Transmission promastigotes :
 - Nouveau monde : *Lutzomya*
 - Ancien monde : *Phlebotomus*
- Promastigote Secretory Gel

– fPPG



- Salive

Rogers ME (2012) The Role of Leishmania Proteophosphoglycans in Sand Fly Transmission and Infection of the Mammalian Host. *Front. Microbio.* 3:223. doi: 10.3389/fmicb.2012.00223

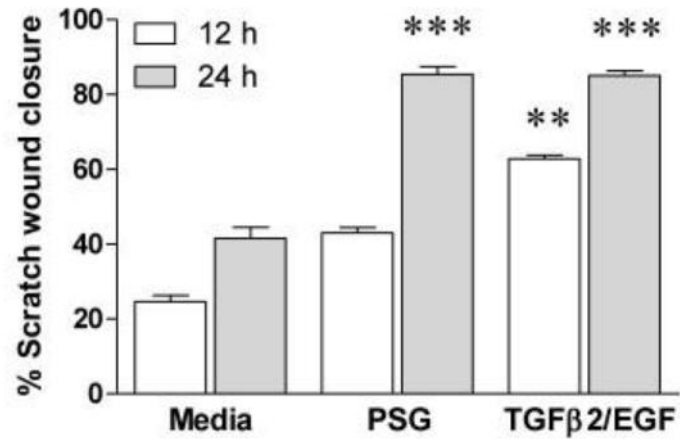


Leishmaniose

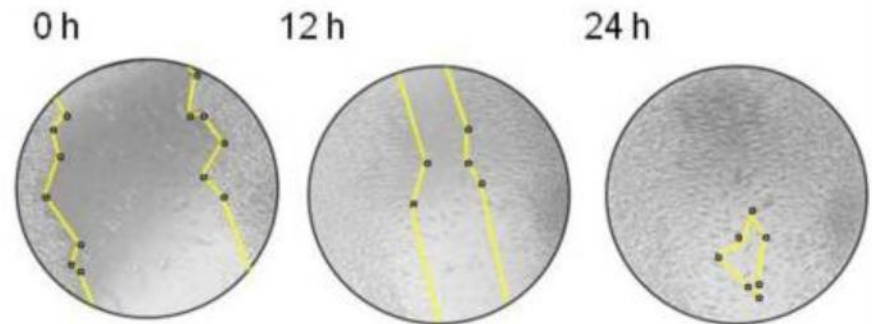
- PSG :
 - Protéophosphoglycane :
 - Cicatrisation accélérée chez la souris
 - Recrutement macrophages
 - Activation des macrophages

Leishmaniose

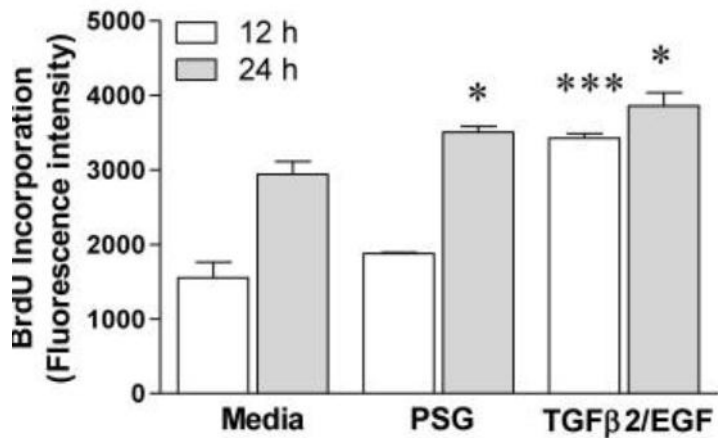
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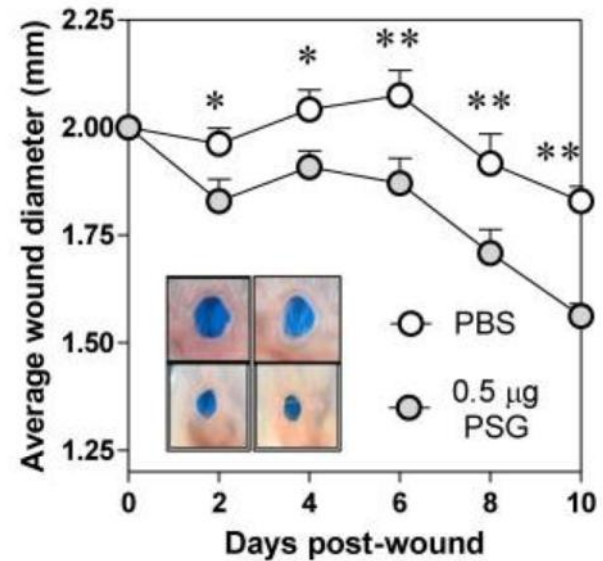
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C



D



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• [PLoS Pathog. 2018 Jan 19;14\(1\):e1006794. doi: 10.1371/journal.ppat.1006794](#)

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Table 1. In skin, PSG induces gene pathways involved in wound healing.

canonical pathways	P Value	Percentage	phase of wound healing
mTOR	7.94E-16	19.4	angiogenesis; fibroplasias and granulation; epithelialisation
hypoxia signalling in the cardiovascular system	5.25E-09	25.0	angiogenesis; fibroplasias and granulation; epithelialisation
IGF-1 Signalling	2.51E-05	15.0	fibroplasias and granulation; epithelialisation
Production of NO and ROS in macrophages	2.63E-05	11.0	inflammation; angiogenesis; fibroplasias and granulation; epithelialisation; remodelling
RhoA signalling	3.63E-05	14.9	inflammation; fibroplasias and granulation; epithelialisation
ERK5 signalling	4.68E-05	18.8	epithelialisation
actin cytoskeleton	5.62E-05	10.5	epithelialisation; contraction
NRF2-mediated oxidative stress response	7.76E-05	11.5	inflammation
ILK signalling	7.76E-05	11.4	fibroplasias and granulation; epithelialisation
integrin signalling	3.16E-04	10.5	fibroplasias and granulation; epithelialisation; contraction; remodelling
p53 signalling	5.89E-04	13.5	inflammation; fibroplasias and granulation; remodelling
glucocorticoid receptor signalling	1.17E-03	8.5	epithelialisation
calcium signalling	1.91E-03	8.7	haemostasis; inflammation; fibroplasias and granulation; epithelialisation; contraction
Cdc42 signalling	2.00E-03	8.3	epithelialisation
NGF signalling	2.19E-03	11.2	inflammation; fibroplasias and granulation; epithelialisation
thrombin signalling	3.39E-03	9.2	haemostasis
androgen signalling	3.63E-03	9.0	inflammation; epithelialisation
HGF signalling	4.07E-03	11.4	epithelialisation
FAK signalling	4.79E-03	10.8	epithelialisation
apoptosis signalling	5.25E-03	11.5	inflammation; fibroplasias and granulation; remodelling
arginine and proline metabolism	8.13E-03	5.1	inflammation; fibroplasias and granulation; epithelialisation; remodelling
EGF signalling	8.51E-03	13.5	epithelialisation

IL-6 signalling	8.71E-03	11.0	inflammation; fibroplasias and granulation; epithelialisation
Gap junction signalling	1.05E-02	8.2	epithelialisation
FGF signalling	1.05E-02	11.1	angiogenesis; fibroplasias and granulation; epithelialisation; contraction; remodelling
VEGF signalling	1.05E-02	10.1	angiogenesis; fibroplasias and granulation; epithelialisation
PDGF signalling	1.07E-02	11.4	inflammation; angiogenesis; fibroplasias and granulation; epithelialisation; contraction; remodelling
Wnt/ β -catenin signalling	1.20E-02	9.2	fibroplasias and granulation; contraction
CCR5 signalling in macrophages	1.29E-02	8.5	inflammation; epithelialisation
IL-4 signalling	1.70E-02	11.0	fibroplasias and granulation; remodelling
IL-1 signalling	1.70E-02	9.4	inflammation
IL-2 signalling	1.82E-02	12.1	inflammation; angiogenesis; fibroplasias and granulation; epithelialisation
regulation of IL-2 in T-lymphocytes	1.91E-02	10.1	inflammation; angiogenesis; fibroplasias and granulation; epithelialisation
PPAR signalling	1.95E-02	9.4	inflammation; fibroplasias and granulation; epithelialisation; remodelling
IL-10 signalling	2.19E-02	10.3	inflammation; fibroplasias and granulation
acute phase response signalling	2.24E-02	8.4	inflammation
IL-8 signalling	2.24E-02	7.8	inflammation; epithelialisation
IL-17 signalling	2.34E-02	10.8	inflammation
chemokine signalling	2.34E-02	11.0	inflammation; angiogenesis; fibroplasias and granulation; epithelialisation
IL-15 signalling	3.72E-02	10.4	inflammation
GM-CSF signalling	3.72E-02	10.4	inflammation; angiogenesis; fibroplasias and granulation; epithelialisation
PTEN signalling	4.68E-02	8.1	inflammation; fibroplasias and granulation; remodelling
angiopoietin signalling	5.00E-02	9.5	angiogenesis

Top canonical pathways involved in innate immunity and one or more phases of wound healing (7.94E-16 < p value < 5.00E-2; FCS, 5% FDR) from BALB/c ear dermis microinjected with 0.5 μ g *L. mexicana* PSG expressed at 6 hours post-inoculation.

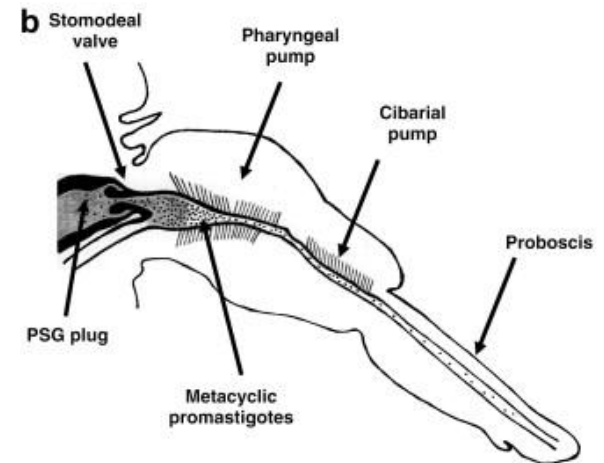
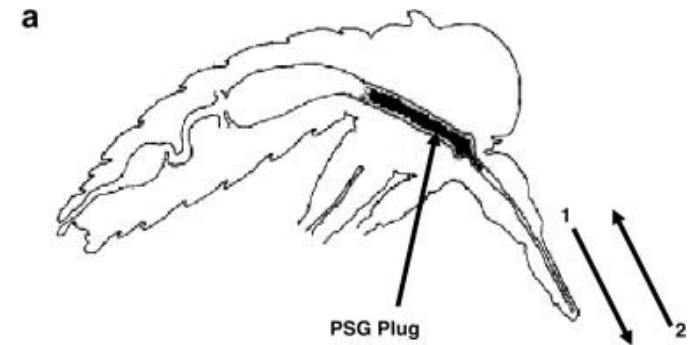
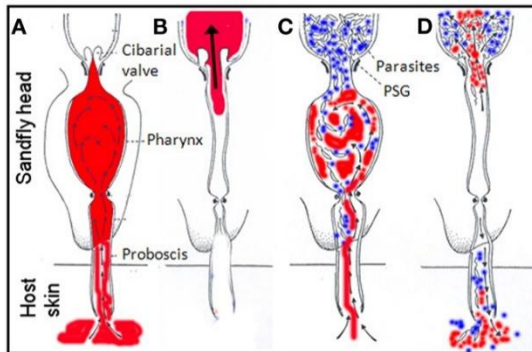
[HHS Vulnerability Disclosure](#)

Leishmaniose

- PSG :
 - Protéophosphoglycanes :
 - Cicatrisation accélérée chez la souris
 - Recrutement macrophages
 - Activation des macrophages
- Échappement parasite
- Amélioration infection

Leishmaniose

- PSG :
- Échappement parasite
- Amélioration infection
 - Modification activité arginase intramacrophage



Leishmaniose

- Salive
 - Composition
 - Rôle :
 - Hémostase
 - Inflammatoire
 - immunologique

Leishmaniose

Table 2
Sand fly salivary proteins with known biological activity.

Sand fly salivary transcriptomes	Biological activities of salivary proteins (molecular weight)										Anti-inflammatory /anti-arthritis
	Inhibitor of contact activation, heparin binding	Biogenic amine binding proteins	Anti-coagulant, inhibitor of factor Xa	Ecto ADPase, inhibitor of platelet aggregation	DNAse activity	Degradation of hyaluronan hydrolysis of chondroitin sulfates	Purine metabolism hydrolysis of adenosine	Vasodilator and inhibitor of platelet aggregation	Nucleotidase	Vasodilator	
	Small odorant binding protein (OBP) – lile (~15 kDa)	Yellow protein (~45 kDa)	Lufaxin/ Lufaxin like (~32 kDa)	Apayrase (~36 kDa)	Endonuclease (~44 kDa)	Hyaluronidase (~42 kDa)	Adenosine deaminase (~56 kDa)	Adenosine	5' Nucleotidase (~61 kDa)	Maxadilan peptide (6 kDa)	LJM111
<i>L. longipalpis</i> (Valenzuela et al., 2004)	LJM04	LJM11 ^a , LJM17 ^a , LJM111 ^a (Xu et al., 2011)	LJL143 (Lufaxin) ^a (Collin et al., 2012)	LuloAPY ^a (Charlab et al., 1999)	LJL138 (Lundep) ^a (Chagas et al., 2014)	LuloHYA ^a (Cerna et al., 2002; Charlab et al., 1999; Rohousova et al., 2012) ^b	ADA ^a (Charlab et al., 1999)		Lulo5NUC ^a (Charlab et al., 1999)	Maxadilan ^a (Lerner and Shoemaker, 1992)	LJM111 ^a (Crespan et al., 2012)
<i>L. intermedia</i> (De Moura et al., 2013)	Linb-7, 8, 28, 59	Linb-21	Linb-17	Linb-35	Linb-46	Linb-54				Linb-147	
<i>L. ayacuchensis</i> (Kato et al., 2013)	LayS32–37, 48–72	LayS22–24, 117, 118	LayS120–132	LayS8–14, 16–21	LayS147						
<i>P. papatasi</i> Tunisia (Abdeladhim et al., 2012)	PPTSP12–15	PPTSP42, 44	PPTSP34 ^b (Collin et al., 2012)	PPTSP36 ^b (Collin et al., 1989b)		^b Cerna et al. (2002))	^b Ribeiro et al. (1999), Charlab et al. (1999), Carregaro et al. (2011)	Adenosine and 5'-AMP ^b (Ribeiro et al., 1999)			
<i>P. dubosqi</i> Mali (Kato et al., 2006)	PduM02–03, 06–07, 12, 31–32, 49–50, 57–58, 60, 62, 99 ^a (Alvarenga et al., 2013)	PduM10, 35	PduM04–05 ^a (Collin et al., 2012)	PduM38–39 ^a (Hamasaki et al., 2009)		^b Cerna et al. (2002)	PduM73				
<i>P. dubosqi</i> Kenya (Kato et al., 2006)	PduK01–03, 40–42, 49, 56–58, 109–110 ^a (Alvarenga et al., 2013)	PduK04–06, 86	PduK70 ^b (Collin et al., 2012)	PduK50 ^a (Hamasaki et al., 2009)		^b Cerna et al. (2002)	PduK60				
<i>P. sergenti</i> (Rohousova et al., 2012)	PsSP9–11, 14–15, 54–55	PsSP18–20, 22, 26	PsSP49	PsSP40–42		^b Cerna et al. (2002), Rohousova et al. (2012)					
<i>P. arabicus</i> (Hostomska et al., 2009)	PabSP2, 45, 63–64, 93,	PabSP26, 53	PabSP34, 32	PabSP39, 40–41	PabSP49	PabSP72 ^b (Rohousova et al., 2012)					
<i>P. argentipes</i> (Anderson et al., 2006)	PagSP01, 02, 07, 12, 13	PagSP04	PagSP09	PagSP03 ^b (Ribeiro et al., 1989b)	PagSP11	^b Rohousova et al. (2012)		Adenosine and 5'-AMP ^b (Ribeiro et al., 1999)			
<i>P. ariasi</i> (Oliveira et al., 2006)	ParSP03, 06, 08	ParSP04, 04B	ParSP09	ParSP01	ParSP10						
<i>P. perniciosus</i> (Anderson et al., 2006)	PpeSP02, 09, 11	PpeSP03, 03B	PpeSP06	PpeSP01, 01B ^b (Ribeiro et al., 1989b)	PpeSP32	^b Rohousova et al. (2012)					
<i>P. perniciosus</i> Madrid Spain (Martin-Martín et al., 2013)	SP02, 09, 11	SP03B	SP06	SP01, 01B ^b (Ribeiro et al., 1989b)		^b Rohousova et al. (2012)					

Leishmaniose

- Salive
 - Composition
 - Rôle :
 - **Hémostase**
 - Inflammatoire
 - immunologique

Leishmaniose

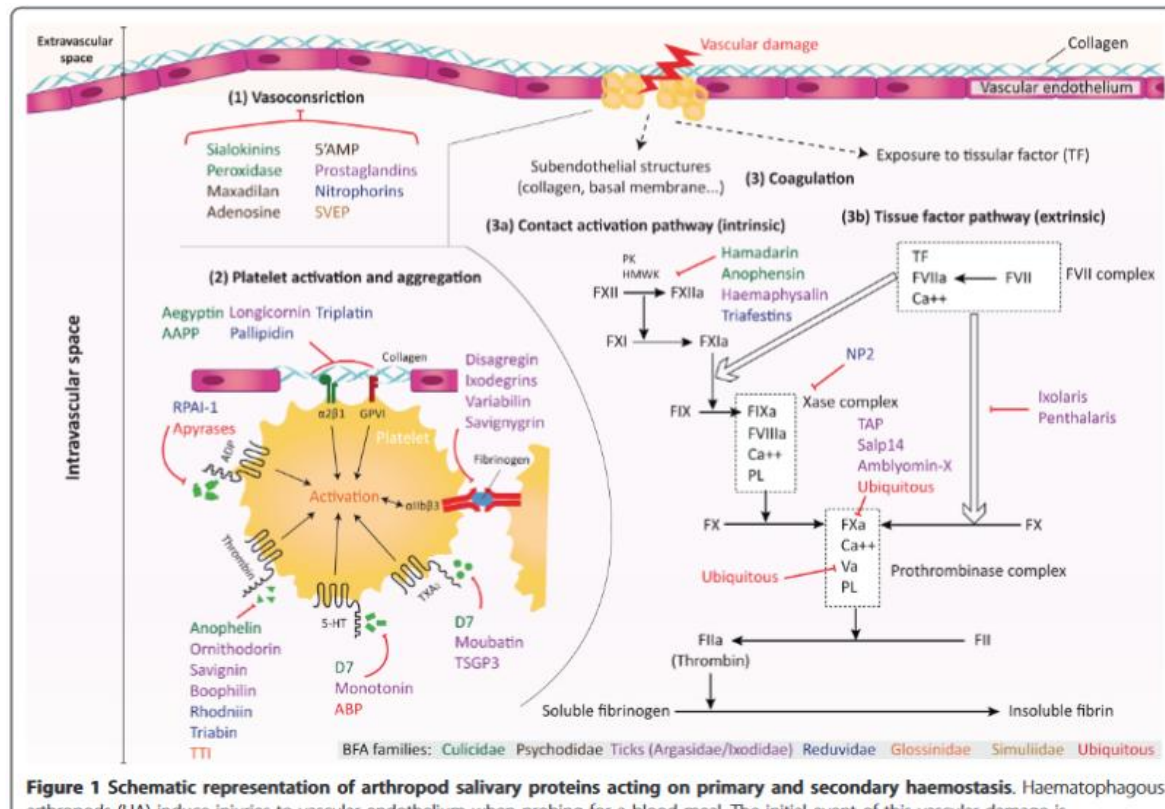


Figure 1 Schematic representation of arthropod salivary proteins acting on primary and secondary haemostasis. Haematophagous arthropods (HA) induce injuries to vascular endothelium when probing for a blood meal. The initial event of this vascular damage is vasoconstriction (1), which retards extravascular blood loss and enhances the adhesion of platelets to exposed subendothelial collagen. This adhesion activates platelets (2) and causes the release of platelet activation agonists (Adenosine diphosphate (ADP), Thrombin, Thromboxane A₂ (TXA₂), serotonin (5-HT)) as well as platelet membrane integrin receptor $\alpha IIb\beta 3$. Fibrinogen binds to this receptor and crosslinks platelets to form a platelet plug. The blood coagulation cascade (3) is then initiated to strengthen the platelet plug with fibrin at the site of injury. The coagulation cascade is separated into two pathways converging into a common pathway. The contact activation pathway (intrinsic) involves high-molecular weight kininogen (HMWK), prekallikrein (PK), factor XII, factor XI and factor IX (3a), and the tissue factor pathway (extrinsic) involves the tissue factor and factor VII complex (3b). Both pathways lead to the activation of factor X. The common pathway leads to the generation of thrombin from prothrombin and the ultimate production of insoluble fibrin from fibrinogen. HA have evolved anti-haemostatic salivary proteins that inhibit specific agonists and factors of platelet aggregation and the blood coagulation cascade. The known actions of some HA salivary proteins listed in Additional file 1 are indicated. (Salivary protein affiliation to HA families is indicated by colour as represented on the bottom right corner legend).

Leishmaniose

- Salive
 - Composition
 - Rôle :
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Leishmaniose

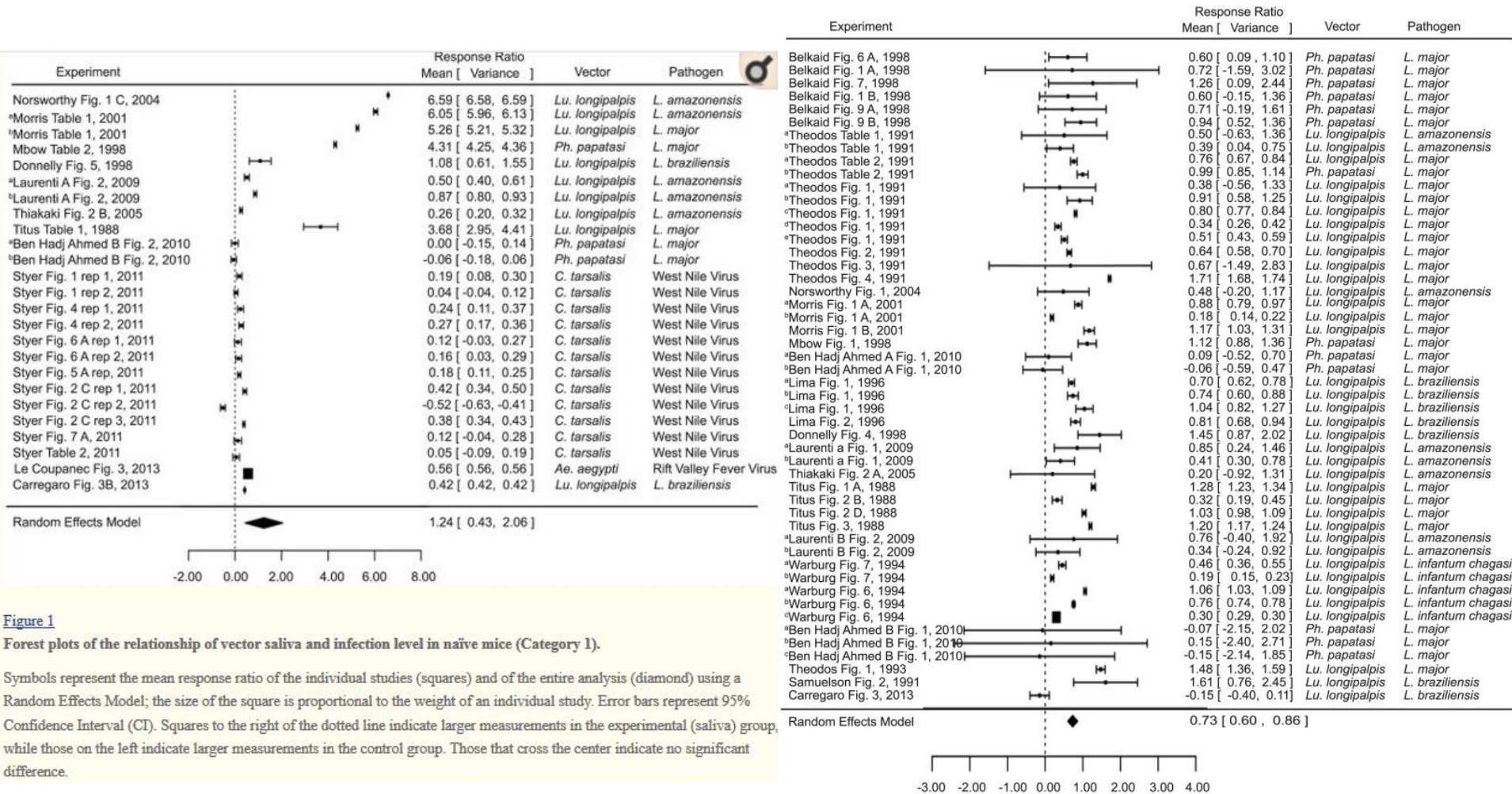


Figure 1
Forest plots of the relationship of vector saliva and infection level in naïve mice (Category 1).

Symbols represent the mean response ratio of the individual studies (squares) and of the entire analysis (diamond) using a Random Effects Model; the size of the square is proportional to the weight of an individual study. Error bars represent 95% Confidence Interval (CI). Squares to the right of the dotted line indicate larger measurements in the experimental (saliva) group, while those on the left indicate larger measurements in the control group. Those that cross the center indicate no significant difference.

Leishmaniose

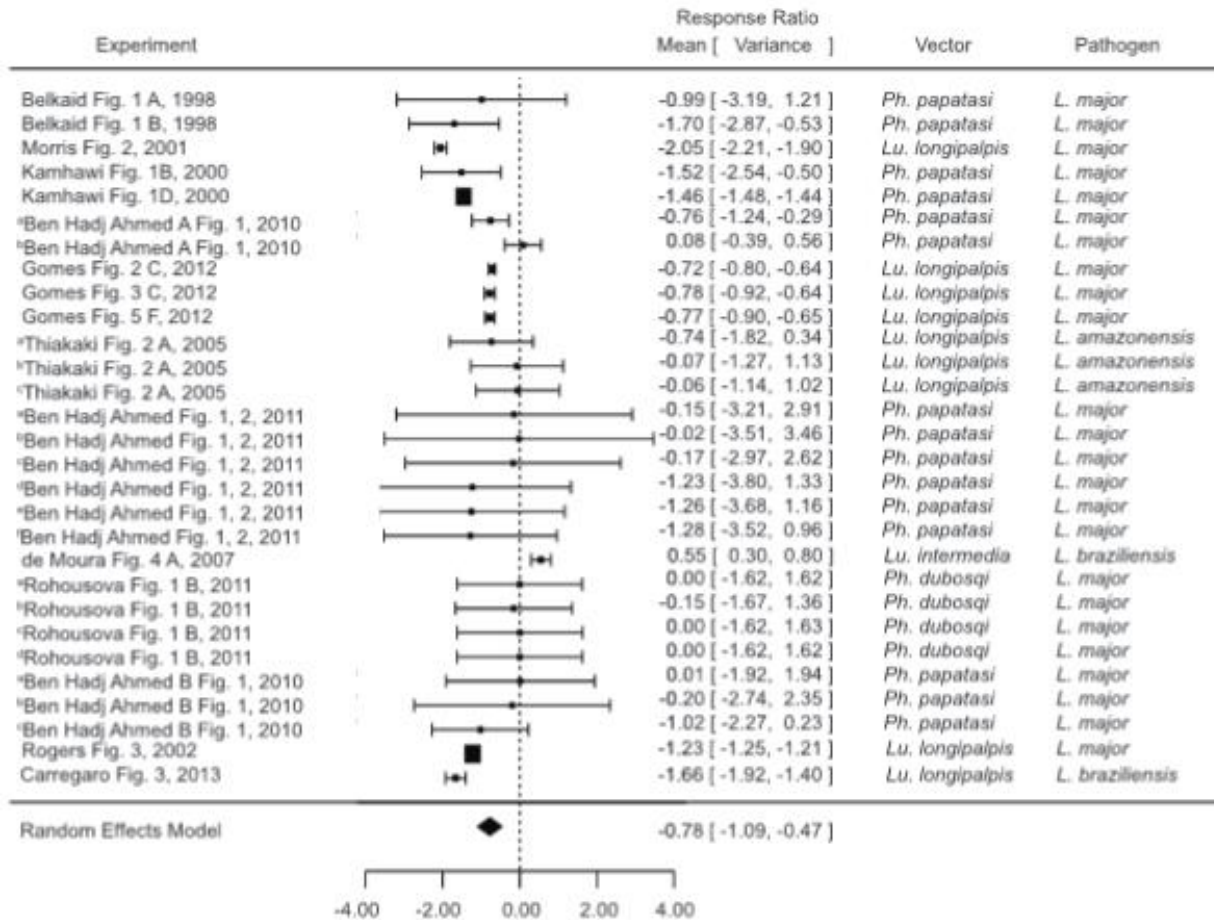


Figure 3. Forest plots of the relationship of exposure to vector saliva before infection and *Leishmania* lesion size (Category 2). Symbols represent the mean response ratio of the individual studies (squares) and of the entire analysis (diamond) using a Random Effects Model; the size of the square is proportional to the weight of an individual study. Error bars represent 95% Confidence Interval (CI). Squares to the right of the dotted line indicate larger measurements in the experimental (pre-exposed) group, while those on the left indicate larger measurements in the control group. Those that cross the center indicate no significant difference.
doi:10.1371/journal.pntd.0003197.g003

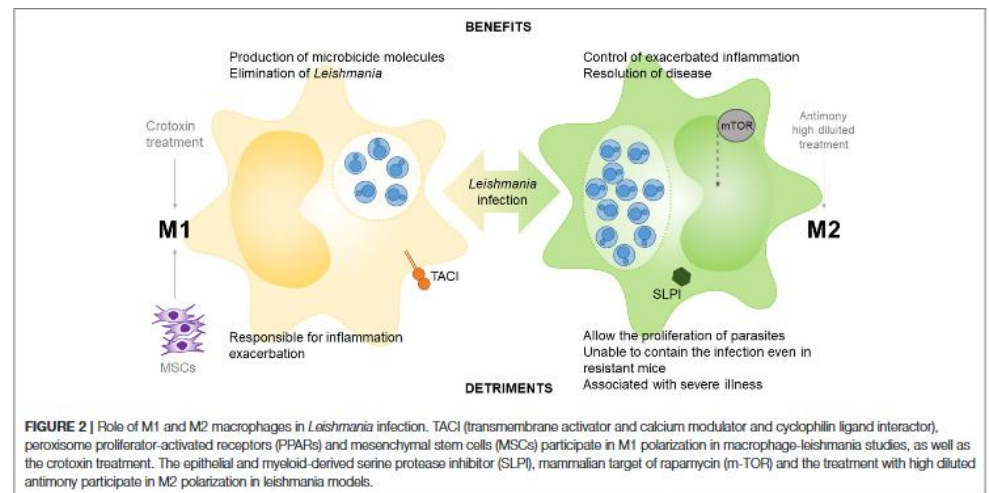
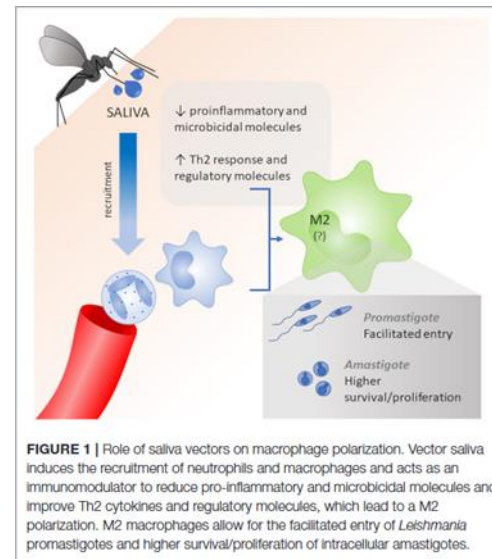
Leishmaniose : salive/réponse immune

- *P. papatasi*/*P. duboscqi* inhibent présentation Ag par CD
- Cellule Dendritique : production PGE2 ; **IL4, IL10**
- Neutrophiles :
 - ↗ nombre
 - Apoptose par *Lu. Longipalpis*
 - Inhibition facteurs chimiotactiques
 - « Trojan rabbit »
- Macrophages : chimiotactisme +
 - Orientation M2
- Th :
 - Orientation Th2: IL4, IL10
 - Inhibition Th1 : IFN, IL12
- Protéines d'intérêt :
 - Maxadilan : Th1, « inactivation » macrophage
 - PpSP44 : ↗ infection
 - LinB-11 : ↘ infection mais salive (*Lu. intermedia*) ↗ infection
 - PpSP15 : IFN=> Th1
 - PdSP15 => rPdSP15 => vaccin
- Anticorps : IgG1, IgE

TABLE 2 | Salivary compounds and their effects on *Leishmania* infection.

Compound	Immunomodulatory effect	References	
Promastigote secretory gel (PSG)	↑ Arg	(77)	
	↑ IL-1β	(78)	
	↑ IL-6		
	↑ IL-10		
	↑ TNF-α		
	↑ CCL2		
	↑ CCL4		
	↑ CCL3		
	↑ CXCL2		
	↑ FGFR2		
	↑ EGF		
	↑ EGFR		
	↑ IGF1		
	Salivary Gland Homogenate (SGH)	↑ MCP-1	↓ INOS (79)
↑ CCR2		↓ NO (80, 81)	
↑ IL-10		↓ IFN-γ (82)	
↑ Eosinophils		↓ IL-13	
↑ Macrophages		↓ IL-5	
↑ IFN-γ			
Salivary Gland Lysate (SGL)	↑ IL-4	↓ IFN-γ (83, 84)	
	↑ IL-6	↓ IL-12 (85–88)	
		↓ INOS	
	Salivary Gland Extracts (SGE)	↑ IL-10	↓ NO (89) (90)
		↑ IL-4	
↑ CD8			
↑ INF-γ			
Salivary Gland Sonicate (SGS)	↑ CD4		
	↑ IL-4	↓ IFN-γ (91, 92)	
	↑ PGE ₂		
Maxadilan (max)	↑ Macrophages		
	↑ LTB ₄		
Adenosine	↑ IL-6	↓ IL-1β (87, 93, 94)	
	↑ IL-10	↓ IL-12p70	
	↑ TGF-β	↓ TNF-α	
	↑ CD86	↓ IFN-γ	
		↓ CD80	
		↓ CCR7	
Adenosine	↑ IL-10	(73)	
	↑ PGE ₂		

CCR, chemokine receptor; CD, cluster of differentiation; IFN, interferon; IL, interleukin; iNOS, inducible nitric oxide synthase; MCP-1, monocyte chemoattractant protein-1; NO, nitric oxide; PG, prostaglandin; TNF, tumor necrosis factor; TGF, transforming growth factor; CCL, chemokines; CXCL, motif chemokine ligand; FGFR, fibroblast growth factor receptor; EGF, epidermal growth factor; EGFR, epidermal growth factor; IGF, insulin-like growth factor.



Tomiotto-Pellissier
(2018) Macrophage Polarization in
Leishmaniasis: Broadening
Horizons. *Front. Immunol.* 9:2529.

Leishmaniose : salive/réponse immune

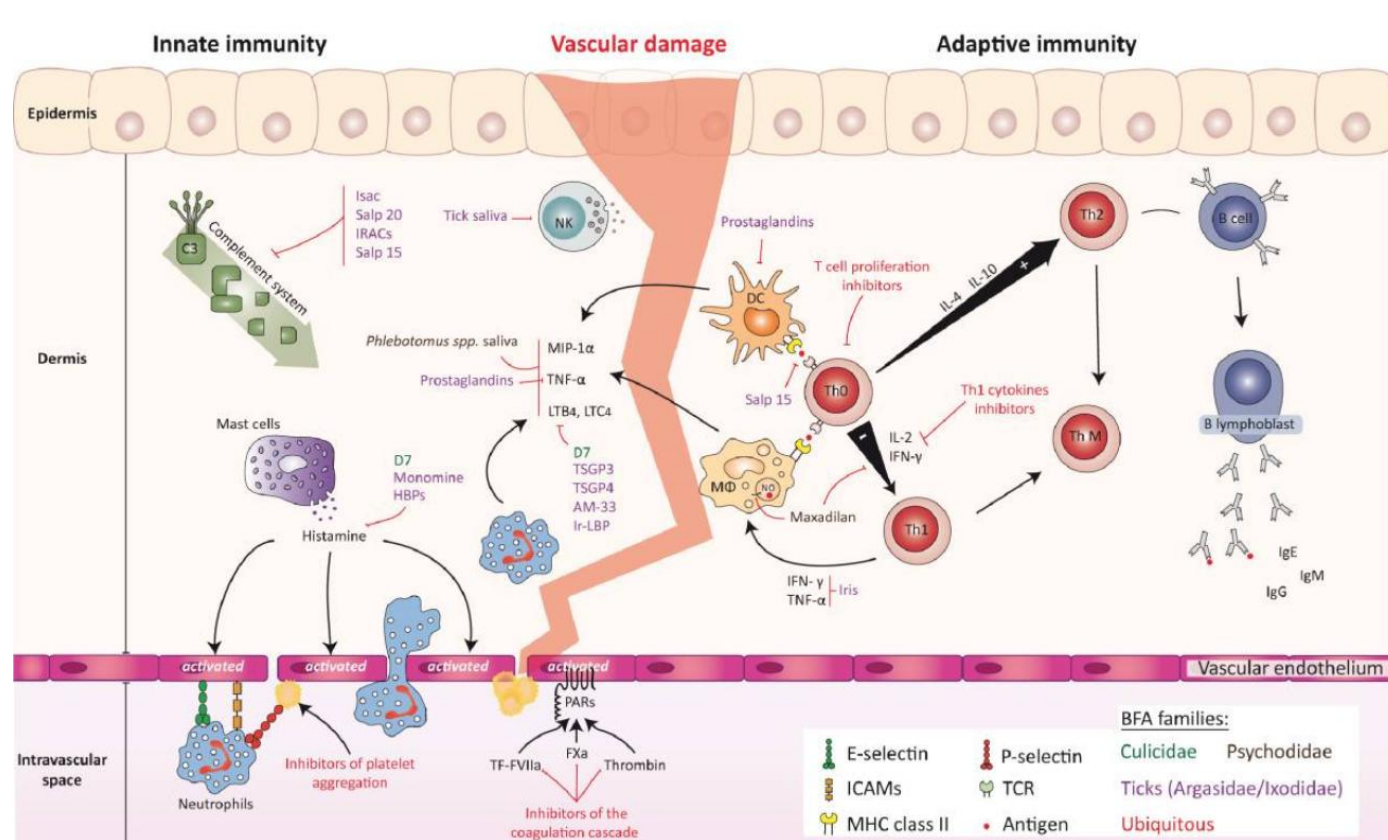
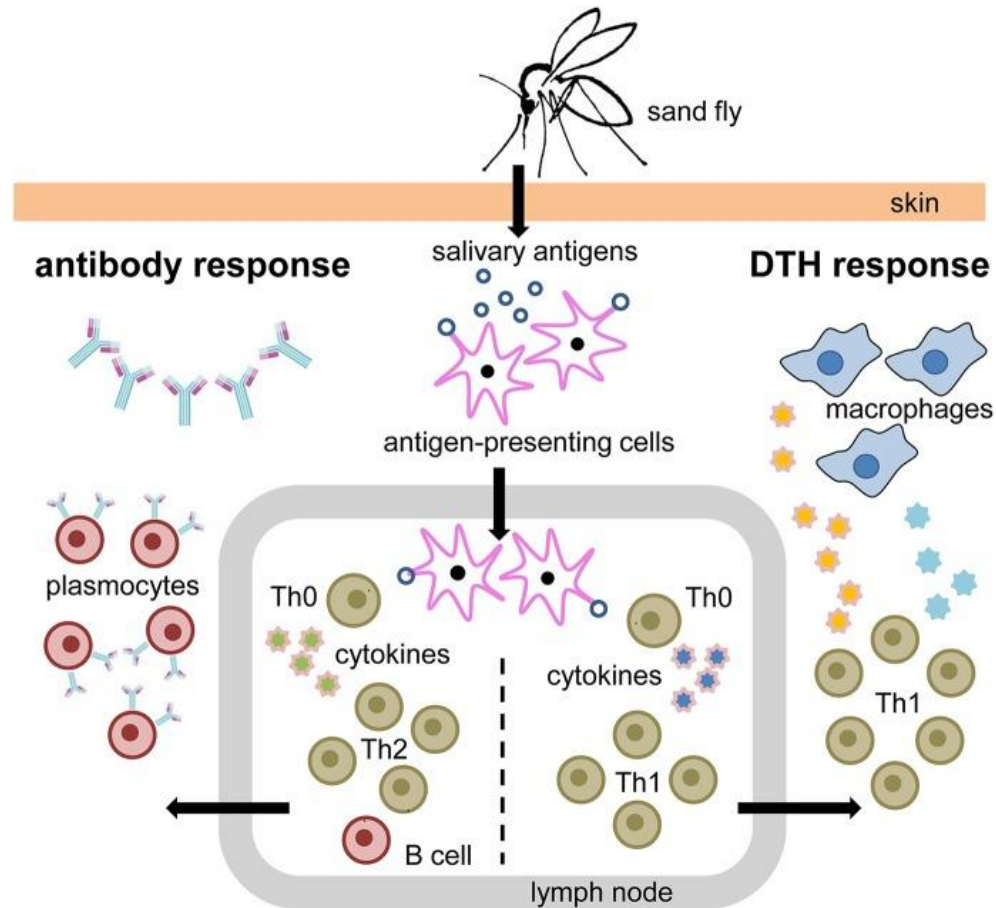


Figure 2 Schematic representation of arthropod salivary proteins involved in the modulation of innate and adaptive immunity.

Protective immunity against haematophagous arthropods (HA) involves both innate and adaptive immunity. Cells involved in the innate response (e.g., neutrophils, natural killers cells (NK), mast cells and macrophages (MΦ)) represent the first line of defence. Once activated, these cells release molecules (e.g., macrophage inflammatory proteins -1 α (MIP-1α), tumour necrosis factor-α (TNF-α) or leukotrienes (LTB₄, LTC₄) that initiate the inflammation process. This local inflammation can further be triggered by the activation of complement, which has chemotactic and inflammatory properties. Endothelial cells and platelets can be activated by the binding of factors of the coagulation cascade to PAR receptors, leading to an over-expression of surface adhesive molecules (ICAMs, E-selectin, P-selectin) that participate in neutrophil migration. Antigen presenting cells, such as dendritic cells (DC) migrate to the lymph nodes where they interact with naïve CD4⁺ helper T lymphocytes (Th0 cells) via the interplay of their T cell receptors (TCR) and major histocompatibility complex (MHC) class II proteins. Th0 cells have the potential to proliferate and to differentiate into two distinct lineages of effectors cells: Th1 and Th2 cells. Memory T helper (Th M) cells, which can improve the quality of the response to a subsequent exposure by developing more efficient memory capacity over time, are also produced. In a general pattern, HA saliva down-regulates the expression of Th1 cytokines (such as IL-2) modulating the adaptive immune response to an antibody mediated Th2 response. The action of saliva or salivary proteins is indicated in the figure as well as their corresponding organism's family. (Salivary protein affiliation to HA families is indicated by colour as represented on the bottom right corner legend).

Leishmaniose : salive/réponse immune



Hypersensibilité retardée => protection

Leishmaniose : salive/applications

- Marqueur d'exposition
- Marqueur du risque de transmission
- Marqueur de réservoir

Leishmaniose : saline/application

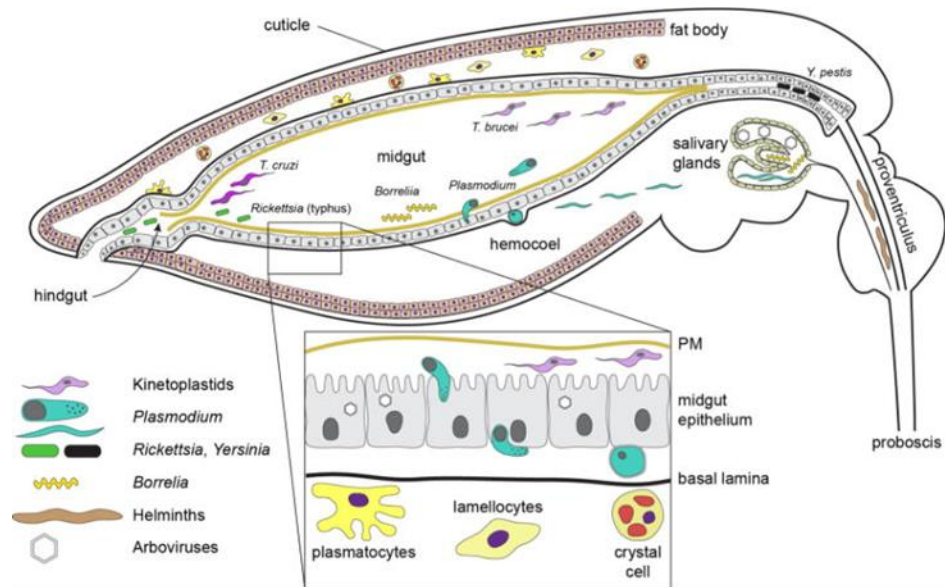
Table 2 Recombinant salivary proteins characterized in hematophagous arthropods and their immunological applications

Protein names	Organisms	Additional informations	MW [kDa]	Application	Ref.
rAed a1	<i>Aedes aegypti</i>	Salivary apyrase	68	Allergy	[151,152]
rAed a2	<i>Aedes aegypti</i>	Belong to the D7 family	37	Allergy	[151,152]
rAed a3	<i>Aedes aegypti</i>	30 kDa salivary gland allergen	30	Allergy	[151,152]
Procalin	<i>Triatoma protracta</i>	Belong to the lipocalin family	20	Allergy	[225]
Arg r 1	<i>Argas reflexus</i>	Belong to the lipocalin family	17	Allergy	[227]
Der-p2	<i>Ixodes ricinus</i>	<i>Dermatophagoides pteronyssinus</i> allergen-like	15.6	Allergy	[226]
TAg5	<i>Glossina m. morsitans</i>	Tsetse Antigen 5	28.9	Allergy	[228]
Maxadilan	<i>Lutzomyia longipalpis</i>	-	9.5	Vaccine candidate	[123]
SP15	<i>Phlebotomus papatasi</i>	-	15	Vaccine candidate	[162]
rLJM19	<i>Lutzomyia longipalpis</i>	-	11	Vaccine candidate	[229]
Salp15	<i>Ixodes scapularis</i>	-	14.7	Vaccine candidate	[163]
gSG6	<i>Anopheles gambiae</i>	-	10	Immunological marker of exposure	[218,219,230,220]
rTC	<i>Amblyomma americanum</i>	Calreticulin	47.5	Immunological marker of exposure	[221]
rLJM11	<i>Lutzomyia longipalpis</i>	Yellow-related protein	43	Immunological marker of exposure	[223,224]
rLJM17	<i>Lutzomyia longipalpis</i>	Yellow-related protein	45	Immunological marker of exposure	[223,224]

Leishmaniose : spécificité/vecteur

- Relation privilégiée vecteur/parasite
- Homologie de la salive/PSG
- *Phlebotomus* vs *Lutzomya*
 - Variation entre genres
 - Conservation intragénre
- Zone endémique/occasionnelle/saisonnalité

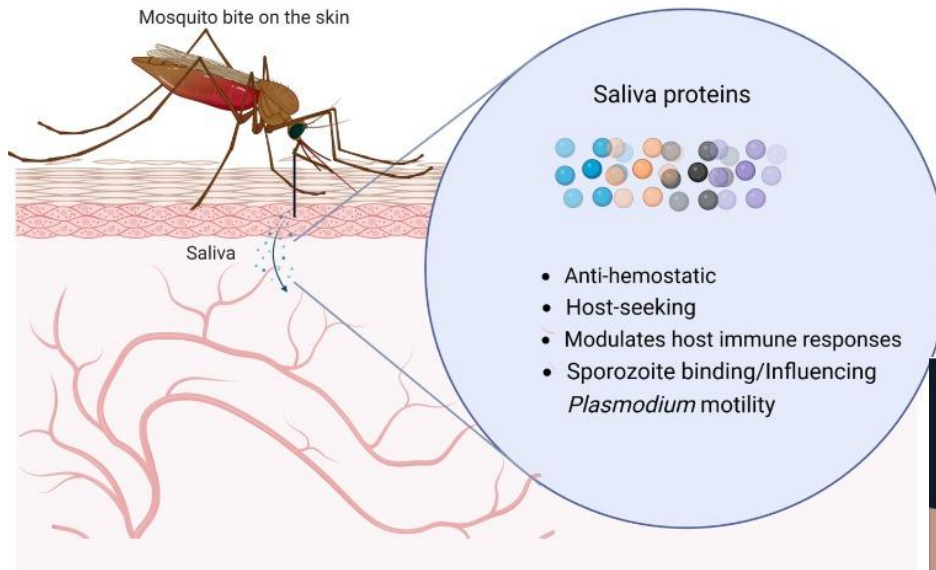
Plasmodium



Baxter et al.
Biochemistry. Author manuscript; available in PMC 2018 February 21.

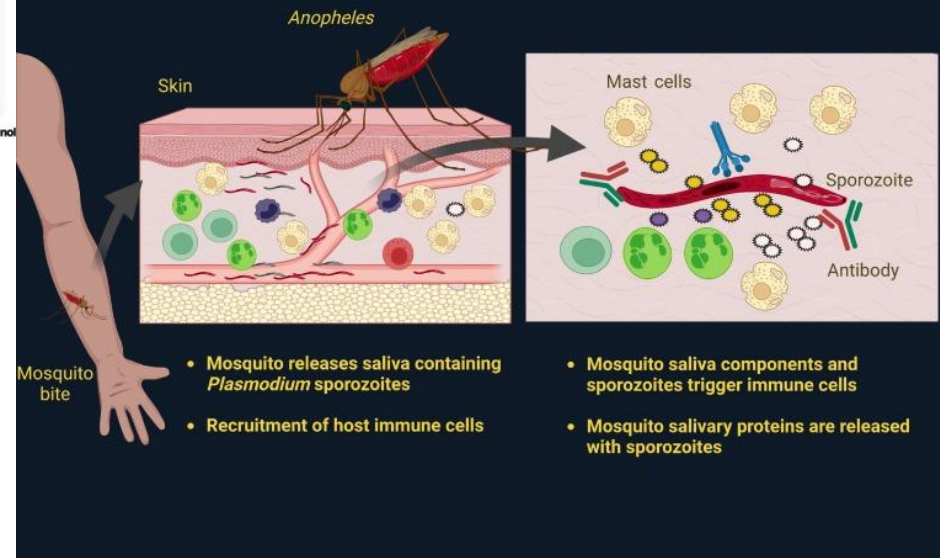
Plasmodium

Mosquito saliva



Trends in Immunol

Immunomodulation by mosquito saliva



Mosquito bite

Trends in Immunology

Plasmodium

- Mastocytes
 - recrutement
 - Dégranulation => afflux sanguin ?
- Neutrophiles :
 - Afflux mais pas d'impact sur l'infection
- Augmentation production INF

Plasmodium

- **Anopheline antiplatelet protein**
 - Baisse coagulation
- **AgTRIO**
 - Augmentée par présence *Plasmodium*
 - Augmente le niveau d'infection
 - Inhibition => baisse infection
- **mosGILT**
 - Baisse motilité sporozoïtes
 - Protection des oocystes
- **SAMSP1**
 - Améliore mobilité sporozoïtes
 - Rôle sur niveau infection
 - Inhibition => baisse infection
- **AgSAP**
 - Baisse immunité locale
 - Rôle sur niveau infection ?
- **D7 protein family**
 - Liaisons à amines biogènes
 - Inhibition vasoconstriction
 - Anticoagulant
 - Peu/pas d'influence sur niveau d'infection

Autres parasitoses vectorisées

Co-infection filaire/*Leishmania* : augmentation risque leishmaniose
... ?

Autres agents infectieux vectorisés

Arboviroses +++

=> vaccins !

Borrelia/Tiques

Conclusion

- Rôle du vecteur
 - Évident pour *Leishmania*... autres parasites ?
- Activité pharmacologique de la salive ++
- Immunisation bénéfique?