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**DANUZA LEITE LEÃO**

**NOVOS PARÂMETROS PARA AVALIAÇÃO ANDROLÓGICA EM MACACO-  
DE-CHEIRO (*Saimiri collinsi* Osgood, 1916): DISTÂNCIA ANOGENITAL,  
MICROMORFOLOGIA E PERFIL PROTEICO ESPERMÁTICO**

**BELÉM  
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Tese apresentada a Universidade Federal Rural da  
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na Amazônia, para a obtenção do título de Doutor.  
Área de Concentração: Produção Animal.  
Orientador (a): Prof<sup>ª</sup>. Dra. Sheyla Farhayldes  
Souza Domingues.  
Coorientador: Prof. Dr. Arlindo de Alencar  
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**BANCA EXAMINADORA**

---

**Prof<sup>ª</sup> Dr. Ednaldo da Silva Filho**  
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**Universidade Federal do Pará - UFPA**

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**Prof. Dr. Thiago Velasco Guimarães Silva**  
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**Prof<sup>ª</sup>. Dr<sup>ª</sup>. Priscila Di Paula Bessa Santana**  
**Universidade Federal Rural da Amazônia - UFRA**

---

**Prof. Dr. Bruno Moura Monteiro**  
**Universidade Federal Rural da Amazônia - UFRA**

*Aos meus pais, Jerceu e Ione, minha  
irmã Danuta, minha avó Benedita e a Pérola  
que sempre me apoiaram nessa jornada ♥*

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“Em todas as páginas de um livro  
A celulose de um ser vivo  
Alerta da preservação”

(Papel - Falcão, P., 2018)



## RESUMO

O macaco-de-cheiro (*Saimiri collinsi*), um primata neotropical endêmico da Amazônia brasileira, apresenta uma sazonalidade reprodutiva em vida livre marcante, contudo pouco estudada. Recentemente essa espécie foi adotada como modelo experimental biológico para pesquisas reprodutivas sobre o gênero *Saimiri*, contribuindo para conservação desses animais. Para melhor compreensão da fisiologia reprodutiva do gênero *Saimiri*, esse estudo teve como objetivos: (i) avaliar a correlação entre a distância anogenital com os parâmetros espermáticos e hormônios reprodutivos (estradiol e testosterona) em macacos-de-cheiro (*S. collinsi*); (ii) descrever as características micromorfológicas e ultraestruturais dos espermatozoides de macacos-de-cheiro (*S. collinsi*) por meio da microscopia eletrônica de varredura e transmissão; (iii) caracterizar o perfil proteico dos espermatozoides de macaco-de-cheiro (*S. collinsi*); (iv) associar as expressões de proteínas espermáticas e as características espermáticas de macaco-de-cheiro (*S. collinsi*). Quanto a análise da qualidade seminal, além do pH, não houve diferença na qualidade seminal entre as estações seca e chuvosa. No tocante as proteínas espermáticas, foram identificadas 2.343 proteínas presentes nas amostras de espermatozoides ao longo dessas duas estações. Das 79 proteínas expressas diferencialmente entre as duas estações do ano, 39 proteínas relacionadas à espermatogênese, motilidade espermática, capacitação, fecundação e sistemas de defesa contra o estresse oxidativo foram aumentadas na estação seca. A micromorfologia analisada por microscopia de varredura revelou que os espermatozóides normais de *Saimiri collinsi* medem  $71,73 \pm 0,71 \mu\text{m}$  com inserção lateral da cauda, cabeça achatada em forma de remo e acrossoma ocupando a maior parte da cabeça, e complementarmente, pela análise da ultraestrutura por microscopia de transmissão foi observado que a peça intermediária é composta por um axonema central com 9 pares de  $9 + 2$  microtúbulos rodeado por nove fibras densas, e as mitocôndrias justapostas formando a bainha mitocondrial. A distância anogenital foi correlacionada com o volume seminal, motilidade, vigor e integridade da membrana plasmática, mas não com o peso corporal, hormônios reprodutivos e volume testicular. Assim, nossos resultados ajudam a avançar na compreensão da fisiologia reprodutiva de *S. collinsi*, fornecendo informações valiosas para a melhoria dos protocolos utilizados nas técnicas de reprodução assistida para a conservação de espécies ameaçadas de extinção no gênero *Saimiri*.

**Palavras-chaves:** Primatas neotropicais, espermatozoides, proteômica, microscopia eletrônica, distância anogenital.

## ABSTRACT

The squirrel monkey (*Saimiri collinsi*), a Neotropical primate endemic to the Amazon in Brazil, with a remarkable but little studied free-living reproductive seasonality. Recently this species was adopted as an experimental biological model for reproductive research on the genus *Saimiri*, contributing to the conservation of these animals. To better understand the reproductive physiology of the genus *Saimiri*, this study aimed to (i) to evaluate the correlation between anogenital distance with sperm parameters and reproductive hormones (estradiol and testosterone) in squirrel monkeys (*S. collinsi*); (ii) describe the micromorphological and ultrastructural characteristics of spermatozoa from squirrel monkeys (*S. collinsi*) through scanning and transmission electron microscopy; (iii) characterize the protein profile of squirrel monkeys sperm (*S. collinsi*); (iv) associate the expression of sperm proteins and the sperm characteristics of squirrel monkey (*S. collinsi*). The study approach based on bottom-up proteomics (Shotgun proteomics) allowed the identification of 2343 proteins present in the sperm samples throughout these two seasons. Of the 79 proteins that were differentially expressed between the two seasons, 39 proteins that were related to spermatogenesis, sperm motility, capacitation, fecundation, and defense systems against oxidative stress were upregulated in the dry season. The SEM electromyographies revealed a normal *Saimiri collinsi*'s sperm measure  $71.73 \pm 0.71$   $\mu\text{m}$  with lateral tail insertion, a paddle-shaped flattened head and a acrosome occupied most of the head, and completely, the TEM also showed that the middle piece is characterized by a central 9 + 2 microtubule axoneme surrounded by nine dense fibers, and the mitochondria were juxtaposed forming the mitochondrial sheath. Anogenital distance was correlated with seminal volume, sperm motility, vigour, and plasma membrane integrity, but not with body weight, reproductive hormones, and testicular volume. Thus, our results help to advance our understanding of the reproductive physiology of *S. collinsi*, providing valuable information for the improvement of protocols used in assisted reproduction techniques for the conservation of endangered *Saimiri* species.

**Keywords:** Neotropical primates, sperm, proteomics, electron microscopy, anogenital distance.

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## LISTA DE ABREVIATURAS E SIGLAS

<b>cm</b>	Centímetro
<b>dpi</b>	Pontos por polegada
<b>µg</b>	Micrograma
<b>µL</b>	Microlitro
<b>g</b>	Gravidade
<b>g</b>	Gramma
<b>h</b>	Hora
<b>kDa</b>	Quilodalton
<b>kg</b>	Quilograma
<b>L</b>	Litro
<b>M</b>	Molar
<b>m</b>	Metro
<b>mg</b>	Miligrama
<b>mA</b>	Miliamperagem
<b>min</b>	Minuto
<b>mm</b>	Milímetro
<b>mM</b>	Milimolar
<b>mL</b>	Mililitro
<b>mOsm</b>	Miliosmol
<b>NaCl</b>	Cloreto de sódio
<b>nm</b>	Nanômetro
<b>°C</b>	Celsius
<b>p</b>	Probabilidade
<b>pH</b>	Potencial hidrogeniônico
<b>TRIS</b>	Tris (hidroximetil) aminometano
<b>V</b>	Volts
<b>W</b>	Watts

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## 1 CONTEXTUALIZAÇÃO

De acordo com o relatório divulgado pela União Internacional para a Conservação da Natureza – IUCN, duas espécies de primatas não-humanos (PNH) brasileiras: o macaco-caiarara (*Cebus kaapori*) e o bugio-marrom (*Alouatta guariba guariba*), estão entre os 25 primatas mais ameaçados do mundo (SCHWITZER et al., 2017). Diante dessa realidade, faz-se prioritária a aplicação de alternativas para a conservação *ex-situ* de PNH, sobretudo o desenvolvimento de biotécnicas reprodutivas (DOMINGUES et al., 2011; SAMPAIO et al., 2017b).

Pensando nesse contexto, a espécie neotropical *Saimiri collinsi*, popularmente conhecida como macaco-de-cheiro, tem sido proposta como modelo experimental para o emprego de biotécnicas da reprodução em outras espécies do mesmo gênero (OLIVEIRA et al., 2015; 2016ab), como é o caso das espécies *S. oerstedii*, *S. vanzollini* (vulneráveis) e *S. ustus* (quase ameaçada), que já constam na lista vermelha de espécies ameaçadas de extinção (BOUBLI; RYLANDS, 2018).

Contudo, em *S. collinsi*, assim como em outras espécies de PNH, a ocorrência de sêmen coagulado é fator limitante para aplicação de biotécnicas da reprodução animal (VALLE et al., 2004), visto que após sua dissolução *in vitro*, os parâmetros espermáticos apresentam-se com valores diminuídos (LEÃO et al., 2015; LIMA et al., 2017; OLIVEIRA et al., 2011, 2015, 2016a, 2016b), com gametas imóveis (NAGLE; DENARI, 1983), e até mesmo sêmen azoospérmico (BENNETT, 1967), o que pode inviabilizar o uso deste material biológico em programas de reprodução assistida para esses indivíduos. Dessa forma, se faz necessário pesquisas mais detalhadas acerca das características reprodutivas de *S. collinsi*, tendo em vista sua representatividade como referência para outras espécies de primatas neotropicais.

Um outro fator peculiar às espécies do gênero *Saimiri*, é a ocorrência de sazonalidade reprodutiva (LINDBURG et al., 1987; STONE, 2014; WOLF et al., 1975). Desse modo, é possível que durante e após o período de sazonalidade reprodutiva, o sêmen desses animais apresente peculiaridades quanto a micromorfologia e ultraestrutura, assim nos componentes biológicos espermáticos, principalmente os proteicos. Porém, até o presente momento não existem relatos nesse primata Neotropical que demonstrem uma correlação entre tais componentes e a qualidade espermática.

Nesse contexto, apesar da espécie *S. collinsi*, ser utilizado em pesquisas voltadas a reprodução e conservação de espécies do gênero *Saimiri*, ainda existem limitações na seleção de machos para programas de reprodução assistida, principalmente quando se envolve técnicas de colheita de sêmen, em que os animais passam por um procedimento de sedação, o que pode ocasionar estresse durante captura e refletir na baixa qualidade seminal desses animais. Pensando nisso, esse trabalho também trouxe como proposta um novo método não invasivo, de fácil execução e barato que pode ser realizado tanto no criadouro quanto a campo ser requerer uma estrutura física complexa e nem equipamentos, para avaliação e seleção de machos aptos a reprodução ou para serem utilizados em programas de reprodução assistida. E para isso foi utilizada a distância anogenital, o qual foi proposto como um biomarcador para a fertilidade no homem, o qual é baseado na mensuração da distância do ânus à base posterior do escroto (AGD<sub>AS</sub>) e do ânus à inserção cefálica do pênis (AGD<sub>AP</sub>) (MENDIOLA et al., 2011).

Sabe-se que a distância do ânus aos órgãos genitais (distância anogenital; AGD) é considerado uma característica sexualmente dimórfica em algumas espécies de mamíferos, incluindo ratos e humanos. No homem, as concentrações aumentadas de andrógeno levam ao aumento da AGD, resultando em uma AGD quase duas vezes maior que as mulheres (MACLEOD et al., 2010; THANKAMONY et al., 2016; WELSH et al., 2008; WOLF et al., 1999). A AGD é determinada durante o desenvolvimento inicial, pela exposição fetal ao meio androgênico, o útero, durante a janela de programação da masculinização (EISENBERG et al., 2012a, 2012b; MENDIOLA et al., 2011; WELSH et al., 2008;).

Estudos em humanos também mostram que um tamanho menor de AGD foi associado a um tamanho menor de testículo (DEAN; SHARPE, 2013), como baixos níveis de testosterona e síndrome de disgenesia testicular, e, conseqüentemente, menor concentração espermática (LÓPEZ-ESPÍN et al., 2017; MENDIOLA et al., 2011, 2015). O *Saimiri collinsi* possui testículos simétricos e existe uma correlação positiva entre o volume testicular e a fração líquida do ejaculado (OLIVEIRA et al., 2016), no entanto, não sabemos se existe uma relação entre o volume testicular e os parâmetros espermáticos com o tamanho da AGD. Desta forma, a medição do AGD pode ser usada como parâmetro não invasivo auxiliar no exame andrológico para a seleção de primatas



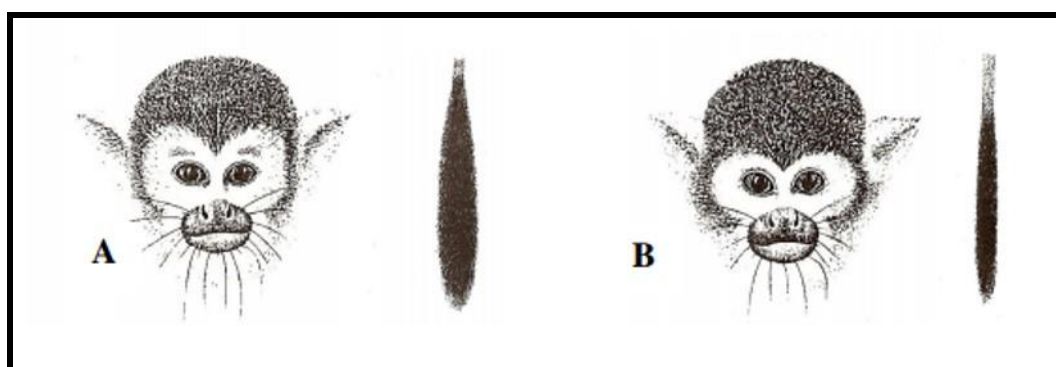
não-humanos em programas de biotecnologia para a conservação de espécies ameaçadas de extinção, bem como estudos em fisiologia reprodutiva.

### 1.1 Taxonomia e aspectos gerais do gênero *Saimiri*

Atualmente são reconhecidos no gênero *Saimiri* a existência de oito táxons: *S. sciureus*, *S. collinsi*, *S. oerstedii*, *S. cassiquiarensis*, *S. macrodon*, *S. ustus*, *S. boliviensis* e *S. vanzolinii* (MERCÊS et al., 2015; PAIM et al., 2013; RYLANDS et al., 2013). As espécies que compõem o gênero têm como características: pequeno porte, com coloração preta esverdeada que varia conforme a espécie e tons que vão do preto ao amarelo-alaranjado (GROVES, 2005). Ao redor dos olhos, possuem uma máscara de pelos branco, que formam um arco, cujo o padrão os divide em dois grupos: o gótico e o romano (Figura 1) (GROVES, 2005; INGBERMAN; STONE; CHEIDA, 2008).

São animais de hábitos diurnos (INGBERMANN; STONE; CHEIDA, 2008), arborícolas, que se locomovem predominantemente de forma quadrúpede (BOINSKI, 1989), e se alimentam de frutas, pequenos vertebrados, flores, goma e insetos (STONE, 2007).

Figura 1- Diferenças entre os arcos superciliares e o pincel caudal entre os grupos gótico (A) e romano (B).



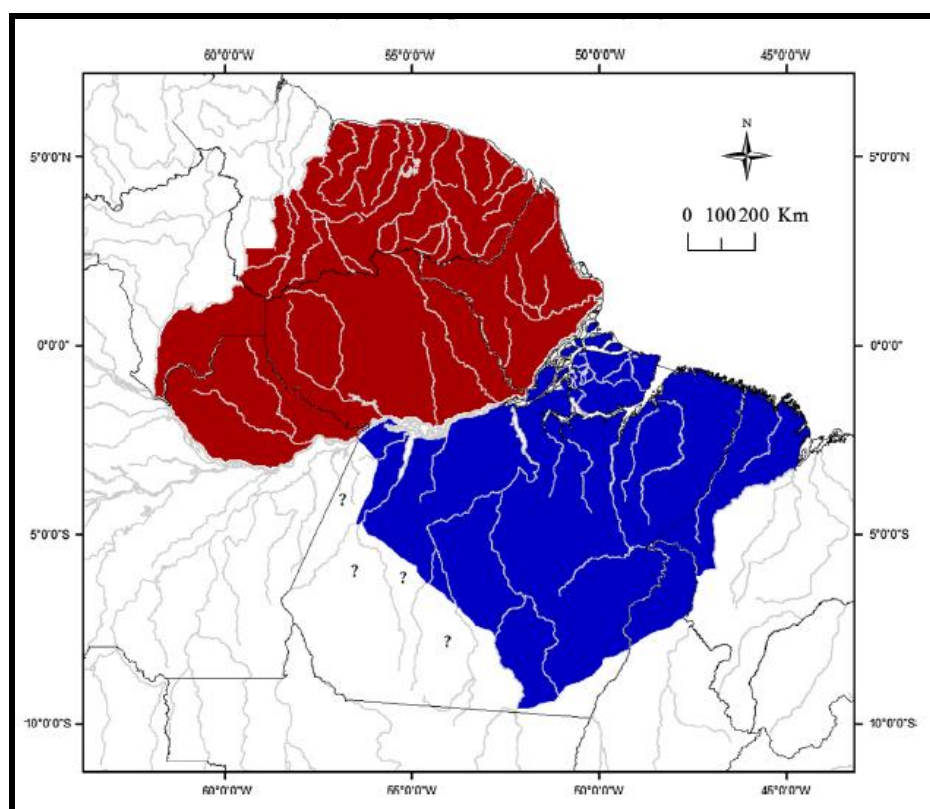
Fonte: HERSHKOVITZ (1984).

Vivem em grupos sociais compostos por vários machos e várias fêmeas (GOODALL; MITTERMEIER, 1999) cujo tamanho varia de dezenas a centenas de indivíduos (BALDWIN; BALDWIN, 1981). A longevidade conhecida em torno de 20 anos (MENDOZA, 1999), mas estima-se que possam chegar aos 30 anos (WILLIMS, 2008).

## 1.2 A espécie *Saimiri collinsi*

A espécie *Saimiri collinsi* (Osgood, 1916), é uma espécie endêmica brasileira que ocorre no arquipélago do Marajó (Estado do Pará), e também na região situada ao sul do rio Amazonas, entendendo-se até o Estado do Maranhão (Figura 2) nas matas de transição entre os biomas Amazônia e Cerrado. Contudo, o limite sul ainda é desconhecido e a porção oeste da distribuição de *S. collinsi* se estende até a margem esquerda do rio Tapajós (MERCÊS et al., 2015).

Figura 2 - Distribuição geográfica de *S. collinsi* (azul).



Fonte: MERCÊS et al. (2015).

Em vida livre vivem em grupos de múltiplos machos e múltiplas fêmeas (cerca de 45-50 indivíduos) (STONE et al., 2015). Em cativeiro, os machos adultos pesam cerca de 620–1.115g (OLIVEIRA et al., 2015, 2016<sup>a</sup>, 2016b) (Figura 3). No tocante a alimentação, ainda não está bem descrita a composição alimentar do *S. collinsi* em seu habitat natural. Entretanto, em cativeiro, são ofertadas frutas, legumes, leite, ração

industrializada, e larvas de besouro tenébrio gigante (*Zophobas morio*) (OLIVEIRA et al., 2015, 2016<sup>a</sup>, 2016b).

Figura 3 - Exemplar macho da espécie *S. collinsi* oriundo do Centro Nacional de Primatas (Ananindeua/Pará/Brasil).



Fonte: A autora.

### 1.3 Aspectos reprodutivos de Saimiri sp.

Em *S. sciureus*, a maturidade sexual ocorre entre 2,5 a 3,5 anos de idade (RICHTER; LEHNER; HENDRICKSON, 1984). São animais poligâmicos (sistema multimachos-multifêmeas) (DIXON; ANDERSON, 2002; FORTMAN et al., 2002; e, como já mencionado, são sazonais com estações reprodutivas bem definidas (BALDWIN; BALDWIN, 1981; TAUB; ADAMS; AUERBACH, 1978).

A reprodução sazonal é a distribuição temporal de cópulas durante um período do ano para ser seguido por nascimentos em outro período. Em PNH há três tipos de reprodução: a sazonalidade discreta, estrita e a ausência de sazonalidade (LINDBURG, 1987).

As espécies do gênero *Saimiri* são consideradas sazonais estritas, visto que a cópulas só ocorrem em um determinado período do ano (época de acasalamento), logo, os nascimentos são agrupados no período seguinte. Na região amazônica, *S. sciureus*

reproduzem-se de julho a setembro, com os nascimentos e lactação acontecendo entre dezembro e fevereiro (DUKELOW, 1983). Stone (2007) define que para essa espécie a estação chuvosa corresponde ao nascimento e aleitamento dos filhotes, enquanto a estação seca corresponde ao acasalamento e gestação.

Durante a estação reprodutiva ocorrem alterações morfofisiológicas e comportamentais tanto nos machos quanto nas fêmeas do gênero *Saimiri* (CHEN et al., 1981). Dentre as alterações morfológicas, nos machos, ocorre um fenômeno conhecido como “*fatted*”, que é o aumento da massa corporal (85 a 222g) por meio do acúmulo de gordura e água (DuMOND; HUTCHISON, 1967; STONE, 2004), principalmente na parte superior do corpo, que produz um aspecto de “gordo” (MENDONZA et al., 1978; STONE, 2004), essa condição de “*fatted*” tem início cerca de dois a três meses antes do início do acasalamento e é restrita à estação reprodutiva (BALDWIN, 1985). Contudo, em indivíduos *S. sciureusi* mantidos em cativeiro, a condição de “*fatted*” não foi necessária para manutenção da capacidade fecundante espermática *in vitro*, visto que ao longo de todo o ano (estação seca e chuvosa) foi possível obter sêmen com espermatozoides viáveis para a fecundação *in vitro*, com obtenção de zigoto (KUEHL; DUKELOW, 1979).

Não se sabe ao certo quais fatores ambientais teriam a maior influência sobre a sazonalidade reprodutiva dos macacos-de-cheiro. Acredita-se que este padrão temporal reprodutivo seja influenciado pelo índice pluviométrico, temperatura e fotoperíodo, entretanto o papel de cada uma dessas variáveis ainda não é completamente estabelecido (DUMOND, 1968; HARRISON; DUKELOW, 1973, HEARN, 1983).

Estudos realizados por Granados et al. (2014) relacionaram a variação da concentração espermática com as porções do ejaculado (líquido seminal, coágulo seminal e *plug* copulatório). Esses estudos ainda pautaram a emissão de coágulo seminal ao longo do ano em *Macaca arctoides*, demonstrando que a concentração espermática foi significativamente maior no coágulo seminal, a qual variou sazonalmente em relação ao fotoperíodo, visto que a concentração espermática encontrada foi maior quando o período de dia e noite apresentaram horas semelhantes, e diminuiu em que o período do dia foi maior.

Já em *Saimiri* sp. mantidos em semi-cativeiro, DuMond e Hutchinson (1967) revelaram uma espermatogênese sazonal em machos mantidos em grupos sociais compostos por ambos os sexos. Nessa pesquisa, foram realizadas biopsias testiculares

durante 13 meses, os quais revelaram que na estação de acasalamento em quase todos os túbulos seminíferos havia a presença de células germinativas em todas as fases, e fora da estação de acasalamento os túbulos regrediram, com uma parede reduzida à camada basal e com pouca indicação de atividade mitótica.

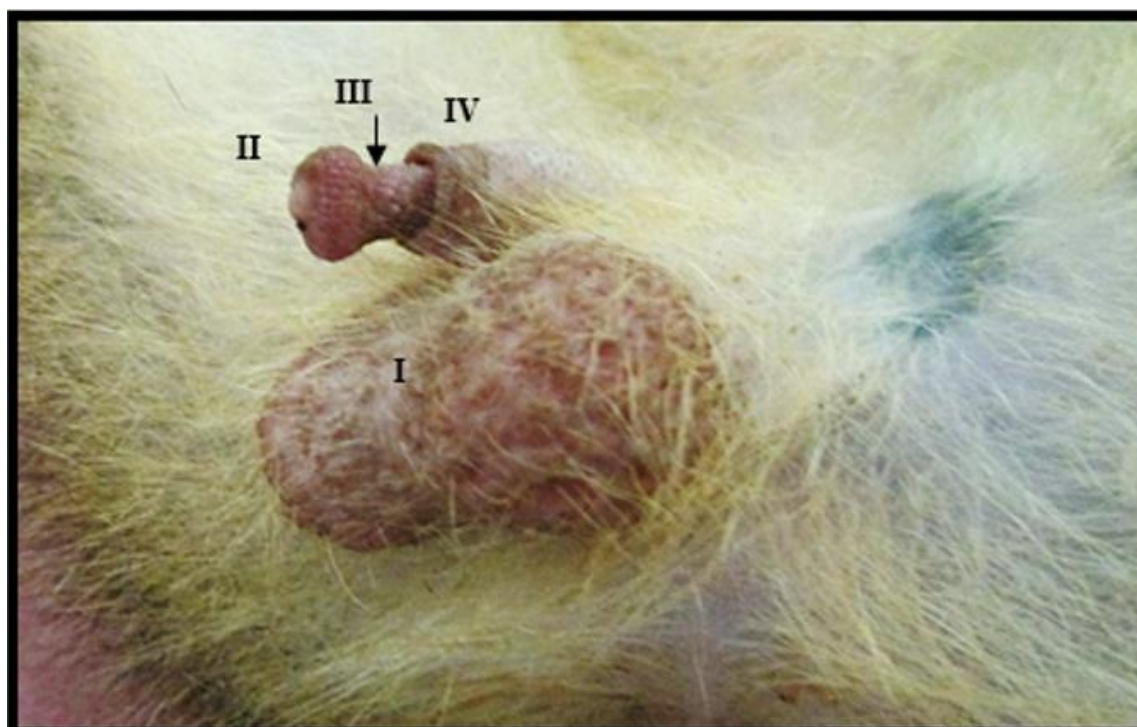
Entretanto, Chen et al. (1981) avaliaram a influência da sazonalidade na qualidade seminal em *S. sciureus* mantidos em cativeiro, e verificaram que não há diferença na concentração e motilidade espermática desses animais durante 13 meses. Contudo, durante a estação reprodutiva, há um aumento no volume do ejaculado, no peso corporal e nos níveis de testosterona circulante (CHEN et al., 1981).

Esses estudos em *Saimiri* sp. demonstraram que a sazonalidade reprodutiva pode ou não influenciar na produção espermática e qualidade seminal em animais de cativeiro, mas, pouco se sabe em relação aos componentes moleculares, a exemplo das proteínas que também podem ser influenciadas por esse processo. Vale ressaltar que em *S. collinsi*, assim como em outras espécies do gênero *Saimiri*, ainda não há estudos que correlacionem os diferentes parâmetros espermáticos e o proteoma, com a sazonalidade reprodutiva.

#### **1.4 O sistema reprodutor do macho de *Saimiri* sp.**

O pênis de *Saimiri* tem comprimento médio de 21 mm (Paulino, G. Dados não publicados), encoberto por prepúcio retrátil, osso peniano e algumas espículas queratinizadas vestigiais laterais ao seu corpo (STEINBERG et al., 2005). O escroto é semi pendulosa e assimétrica, proporcionalmente grande considerando-se o porte desses animais (STEINBERG et al., 2005) (Figura 4). O testículo e o epidídimo juntos formam uma massa relativamente pequena e globular. As glândulas seminais são tubulares e não ramificadas, e a próstata localiza-se na parte proximal da uretra, imediatamente distal à bexiga (HILL, 1960). Quanto a biometria testicular, em *S. collinsi*, foi descrita pela primeira vez por Oliveira et al. (2015), conforme dados da tabela 1.

Figura 4 - Genitália externa do macho de *S. collinsi*: I- escroto e testículo; II- glande; III- corpo do pênis; IV- Prepúcio. Seta: espículas laterais do corpo do pênis.



Fonte: A autora.

Tabela 1 - Biometria testicular de *S. collinsi* segundo Oliveira et al., 2015.

Testículo	Comprimento (cm)	Largura (cm)	Altura (cm)	Volume (cm <sup>3</sup> )
Direito	1,86 ± 0,21	1,23 ± 0,11	1,16 ± 0,05	1,40 ± 0,26
Esquerdo	1,83 ± 0,24	1,34 ± 0,14	1,18 ± 0,09	1,55 ± 0,42

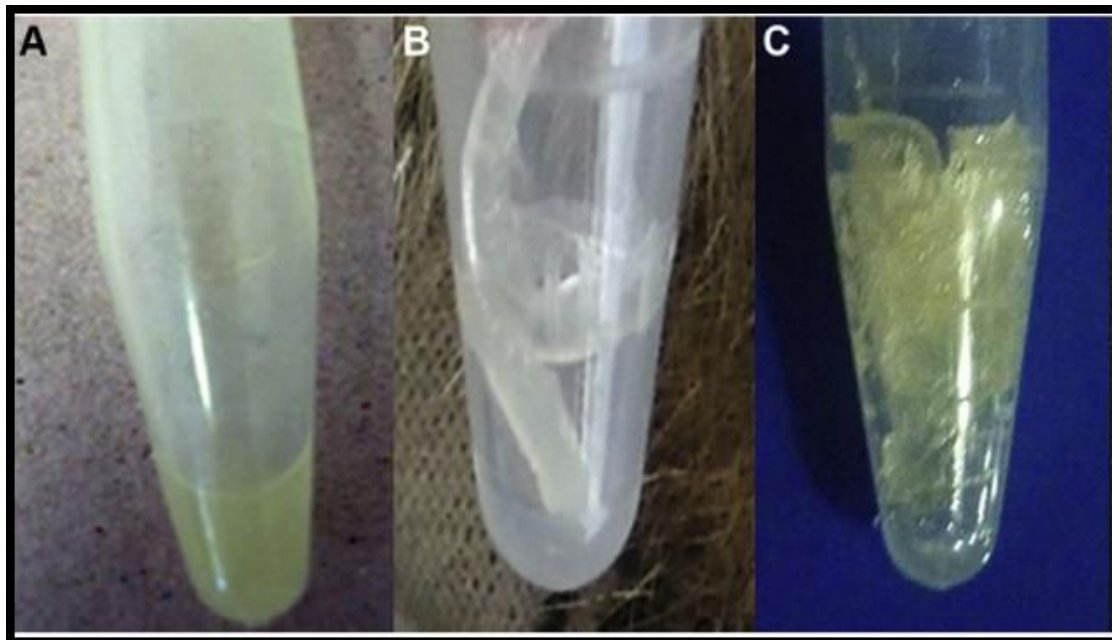
### 1.5 O sêmen de *Saimiri collinsi*

O sêmen de *Saimiri collinsi* é caracterizado por possuir duas frações, uma líquida e outra coagulada (Figura 5). Nesta espécie, a ejaculação quase sempre é iniciada pela fração líquida que é total ou parcialmente coagulada após 10 segundos no ambiente (OLIVEIRA et al., 2015).

A fração coagulada possui aparência filamentar ou amorfa, sendo que ambas as frações podem apresentar-se incolores, esbranquiçadas ou amareladas, transparentes ou opaca. O volume seminal obtido por eletroejaculação é de  $51,8 \pm 49,5 \mu\text{L}$  (5–200  $\mu\text{L}$ ) e

$304 \pm 283,6 \mu\text{L}$  (10–1100  $\mu\text{L}$ ), para a fração líquida e coagulada respectivamente. A concentração espermática do sêmen é cerca de  $6,48 \times 10^6$  espermatozoides/mL (OLIVEIRA et al., 2015).

Figura 5 - Sêmen de *S. collinsi* coletado por eletroejaculação: (A) fração líquida opaca, (B) coágulo filamentar e (C) coágulo amorfo.



Fonte: Oliveira et al. (2015).

## 1.6 O espermatozoide de *Saimiri* sp.

A espermatogênese dos macacos-de-cheiro dura cerca de 39 dias, sendo 30,5 dias da condição de espermatogônia até o estágio de espermátide, restando 8,5 dias para a diferenciação celular, culminando na formação dos espermatozoides (BARR, 1973).

A morfometria dos espermatozoides de macaco-de-cheiro foi primeiramente descrita por Bennett (1967), onde o comprimento da cabeça mediu 6  $\mu\text{m}$ , o comprimento da peça intermediária 12  $\mu\text{m}$  e o comprimento da cauda 60  $\mu\text{m}$ , posteriormente outros pesquisadores também descreveram a morfometria dos espermatozoides, algumas delas estão demonstradas na tabela 2. Os espermatozoides de *Saimiri* sp., são caracterizados por apresentar cabeça achatada lateralmente com intumescimento apical, área superficial pequena e margem posterior do acrossomo com

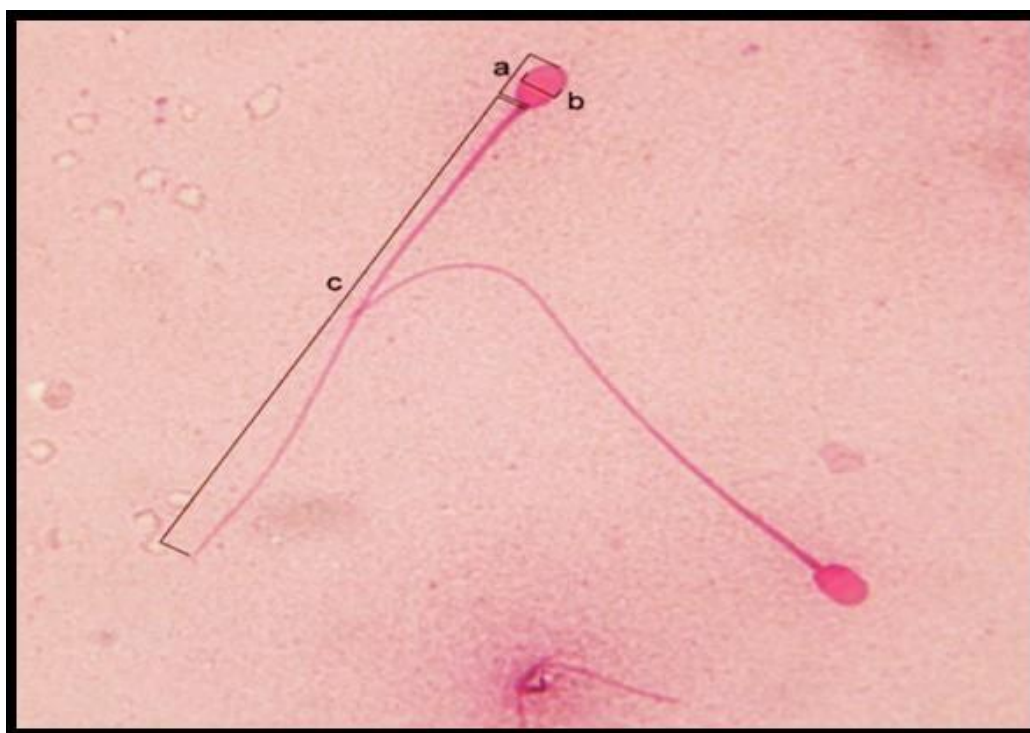
aparência serrilhada ou de microvilosidades, além da peça intermediária inserir-se de forma excêntrica na porção posterior da cabeça (Figura 6).

Tabela 2- Morfometria dos espermatozoides de espécies do gênero *Saimiri* sp.

Espécie	Dimensões lineares do espermatozoide ( $\mu\text{m}$ ) - Média $\pm$ DP				
	Largura da cabeça	Comprimento da cabeça	Comprimento da PI*	Comprimento total da cauda	Comprimento total
	$3,51 \pm 0,01^1$	$5,11 \pm 0,02^1$	$9,03 \pm 0,12^1$	$55,34 \pm 0,13^1$	$69,24 \pm 0,15^1$
<i>S. sciureus</i>	$3,6 \pm 0,3^2$	$5,5 \pm 0,03^2$	$10 \pm 0,5^2$	$55,1 \pm 2,7^2$	$70,6 \pm 2,8^2$
	-	$5,1^3$	$9,0^3$	$55,3^3$	$69,2^3$
<i>S. boliviensis</i>	$3,76 \pm 0,45^4$	$5,71 \pm 0,45^4$	$12,2 \pm 0,45^4$	$65,68 \pm 0,45^4$	$71,39 \pm 0,45^4$
<i>S. collinsi</i>	$4,3 \pm 0,01^5$	$6,2 \pm 0,01^5$	-	$70,5 \pm 0,19^5$	$76,7 \pm 0,19^5$

\*PI: Peça intermediária. <sup>1</sup>DUKELOW, 1983; <sup>2</sup>LAVERDE-CORREA; ROBLES-MEDINA, 2001; <sup>3</sup>GAGE; FRECKLETON, 2003; <sup>4</sup>STEINBERG ET AL., 2005; <sup>5</sup>SAMPAIO et al., 2017.

Figura 6 - Morfometria do espermatozoide de *S. collinsi* (a: comprimento da cabeça, b: largura da cabeça e c: comprimento da cauda) corados com eosina-nigrosina.



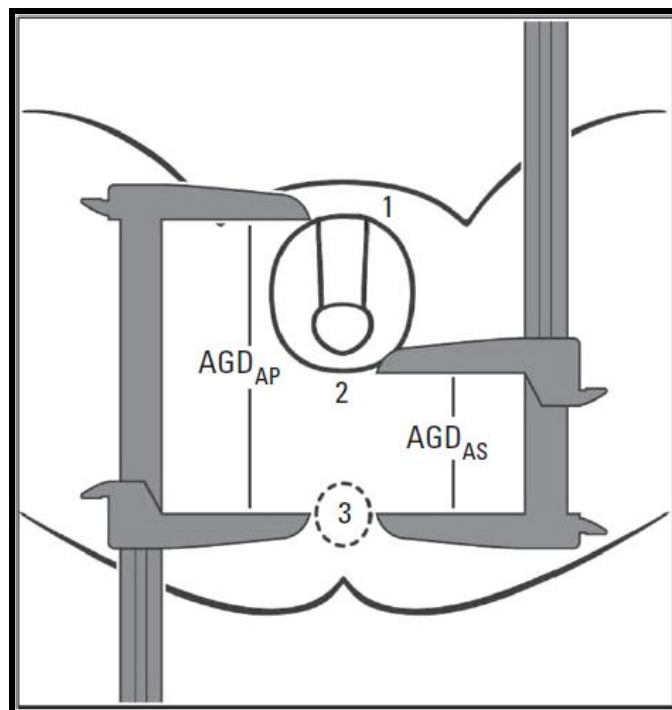
Fonte: SAMPAIO, W.V. (2014).



### 1.7 A distância anogenital

A distância do ânus aos órgãos genitais (distância anogenital; AGD) é considerada uma característica sexualmente dimórfica em mamíferos, uma vez que os machos têm uma AGD quase duas vezes maior que as fêmeas (MACLEOD et al., 2010; SWAN, 2008; THANKAMONY et al., 2009;). As variantes do AGD medidas com paquímetro universal, são:  $AGD_{AP}$  – que é a distância da inserção cefálica do pênis ao centro do ânus, e  $AGD_{AS}$  que é a distância da base posterior (primeira dobra) do escroto ao centro do ânus (Figura 7; MENDIOLA et al., 2011)

Figura 7 - Desenho esquemático das duas medidas de AGD:  $AGD_{AP}$ , da inserção cefálica do pênis até o centro do ânus (ponto 1 ao ponto 3); e  $AGD_{AS}$ , da base posterior (primeira dobra) do escroto ao centro do ânus (ponto 2 ao ponto 3).



Fonte: MENDIOLA et al. (2011).

Em ratos, as medidas de AGD foram associadas a um tamanho menor de pênis, testículos, próstata e vesículas seminais (MACLEOD et al., 2010; SCOTT et al., 2008; VAN DEN DRIESCHE et al., 2011). O tamanho da AGD é determinado pela quantidade de andrógeno à qual um feto masculino é exposto no desenvolvimento fetal durante a janela de programação da masculinização (WELSH et al., 2008). Da mesma forma, em humanos, foi relatado que a AGD é determinada no útero como em ratos, e

isso influencia o desenvolvimento genital e a função testicular do adulto, assim como a baixa qualidade seminal (MENDIOLA et al., 2011).

Nesse contexto, o AGD tem sido relatado como um biomarcador que determina a síndrome de disgenesia testicular (DEAN; SHARPE, 2013), como criptorquidia e hipospádia (GILBOA et al., 2017; JAIN; SINGAL, 2013), bem como um determinante de uma concentração alta ou baixa de espermatozóides (LÓPEZ-ESPÍN et al., 2017; MENDIOLA et al., 2011, 2015).

Embora existam estudos que avaliam a correlação AGD com uma qualidade seminal em humanos, essa medida pode auxiliar na determinação da etiologia dos homens com azoospermia (EISENBERG et al., 2012). Não se sabe se o tamanho da AGD também está associado à qualidade seminal e à biometria testicular em outras espécies de primatas como o *Saimiri collinsi*, pois, de acordo com a literatura, essa associação só foi estudada em homens (DEAN; SHARPE, 2013; EISENBERG et al., 2011, 2012; MENDIOLA et al., 2011; PARRA et al., 2016; ZHOU et al., 2015). Se confirmada, a medida de AGD pode substituir exames andrológicos invasivos durante a avaliação e seleção de machos *Saimiri collinsi* em programas de biotecnologia de reprodução animal para conservação de espécies ameaçadas.

## **1.8 O estudo da micromorfologia e ultraestrutura dos espermatozoides de primatas**

Na reprodução animal, a análise seminal é uma importante ferramenta para avaliar o potencial de fertilidade do macho (PARIZ et al., 2014). Contudo, a análise espermática sob a microscopia óptica possui limitação no tocante à resolução, visto que não permite a identificação e análise de defeitos morfológicos que podem acometer as organelas dos espermatozoides, como defeitos da teca perinuclear, estruturas mitocondriais e dos axonemas, os quais são frequentemente associadas com patologias espermáticas como a teratozoospermia, a astenozoospermia e infertilidade idiopática, que influenciam negativamente na capacidade fecundante da célula espermática (MORETTI; COLODELL, 2012; PARIZ et al., 2014).

Nesse contexto, a microscopia eletrônica de varredura possibilita a avaliação tridimensional dos espermatozoides, e a de transmissão possui um alto poder de resolução em relação ao microscópio óptico, visando a análise de estruturas extra e

intracelulares, respectivamente, proporcionando uma visão de alto alcance, em dimensões nanométricas, que auxiliam à descrição celular e à detecção de possíveis anormalidades espermáticas (MORETTI; COLODELL, 2012). Essas técnicas já foram utilizadas em diversos mamíferos como touros (DUCHA et al., 2012; OLIVEIRA et al., 2015), cavalos (BLOTTNER et al., 2001; PESCH et al., 2006), carneiros (LÓPEZ-ARMENGOL et al., 2012), cães (SILVA et al., 2009), tatus (SOUSA et al., 2013), ursos (BRITO et al., 2010), capivara (BATALHA; OBA, 2006) e primatas não-humanos (OKADA et al., 2001; RUTLLANT; POMMER; MEYERS, 2003; SHABDULYA et al., 1982; STEINBERG et al., 2009).

Por meio da análise por microscopia eletrônica de varredura, Zaneveld et al. (1974) demonstraram diferenças entre os aspectos morfológicos superficiais do coágulo seminal de humanos e de macacos *Rherus* (*Macaca mulatta*), determinando a relação entre esses aspectos e a taxa de liquefação do coágulo. Nesse mesmo estudo, as eletromicrografias mostraram que os coágulos de *Macaca* sp. consistem de fibras desorganizadas de espessura variável, formando uma estrutura sólida, e que o coágulo humano se apresentou constituído por uma fina rede extensivamente organizada de longos e finos filamentos fibrosos que confinam os espermatozoides. Em ambas as espécies, essas fibras tornam-se desorganizadas e frouxas à medida que o coágulo se liquefaz, havendo a liberação gradativa dos espermatozoides. Um fato interessante em sêmen de *Macaca* sp. é que a maioria dos espermatozoides foi visualizada na superfície do coágulo, sendo possível a existência de alguma propriedade de adesão. Também, em macacos *Rhesus*, Sivashanmugam et al. (1997) verificaram, por meio da microscopia eletrônica de transmissão, alterações na condensação da cromatina e nos componentes da membrana plasmática de espermatozoides durante a maturação epididimária, e relataram uma reorganização dos lipídios da membrana plasmática durante o trânsito pelo epidídimo, no qual a restrição de domínios lipídicos sobre a região acrossomal pode ser importante para a capacitação e subsequente reação acrossômica.

Já em primatas neotropicais, Grabner (2016) avaliou por microscopia eletrônica de transmissão, a morfologia espermática de três espécies (*Leontopithecus chrysomelas*, *Alouatta caraya* e *A. guariba clamitans*), e verificou que os espermatozoides desses primatas não-humanos apresentaram os componentes básicos já descritos para o homem, entretanto existe variação na forma e no tamanho dos mesmos, sendo que na extensão do acrossomo, bem como a largura, comprimento e organização da peça

intermediária demonstram as diferenças mais marcantes da superfamília Hominoidea para o gênero *Homo*. Em os macacos-de-cheiro (*S. sciureus*), os estudos utilizando técnicas de microscopia eletrônica de varredura e de transmissão foram utilizados há décadas também com o objetivo de analisar a morfologia espermática (GOULD, 1980; MARTIN, GOULD, 1975). Steinberg et al. (2009) obtiveram, com auxílio de imagens captadas por microscopia eletrônica de varredura, dados morfométricos da cabeça dos espermatozoides (comprimento da cabeça:  $4,93 \pm 0,41 \mu\text{m}$  e largura da cabeça:  $3,83 \pm 0,26 \mu\text{m}$ ) como ferramenta para identificação taxonômica mais precisa da espécie *S. boliviensis*.

Nesse contexto, para um maior conhecimento sobre os aspectos mais intrínsecos, e ainda não relatados em sêmen de *S. collinsi*, faz-se necessária a descrição em detalhes nanométricos de componentes da matriz extracelular, de organelas celulares e das membranas dos espermatozoides, possíveis somente por análise em microscopia eletrônica de varredura e de transmissão.

## 1.9 O estudo das proteínas na reprodução

A proteômica busca fornecer informações a partir da análise sistemática de macromoléculas em um tecido ou célula (COX; MANN, 2007). Técnicas de proteômica têm sido utilizadas no intuito de identificar constituintes expressos no sistema reprodutor masculino em diferentes espécies, tais como bovinos (REGO et al., 2014; THEPPARAT et al., 2012), equinos (SWEGEN et al., 2015), caprinos (HE et al., 2015; VAN TILBURG et al., 2013), ovinos (RODRIGUES et al., 2013), suínos (GONZÁLEZ-CADAVID et al., 2014), catetos (SANTOS et al., 2014), PHN (LUNDWALL; OLSSON, 2001; SKERGET et al., 2013; VALTONEN-ANDRÉ et al., 2005) e no homem (FRAPSAUCE et al., 2014). Tais pesquisas trazem resultados importantes das interações entre proteínas do plasma seminal e espermatozoides. Atualmente, marcadores proteicos têm sido descritos no intuito de detectar propriedades biológicas relacionadas com a qualidade seminal (FRAZER et al., 2006; PARK et al., 2012; THEPPARAT et al., 2012), e assim, desenvolver novos critérios para o prognóstico e aumento na taxa de fertilidade dos machos (AGARWAL et al., 2016; LÉGARÉ et al., 2014; NARESH; ATREJA, 2015).

Além disso, em PNH, algumas pesquisas têm oferecido informações básicas acerca da composição proteica do plasma seminal relacionada ao processo de coagulação seminal como demonstrado em sagui-do-tufo-branco (*Callithrix jacchus*) (VALTONEN-ANDRÉ et al., 2005) e sagui-da-cabeça-de-algodão (*Saguinus oedipus*) (LUNDWALL; OLSSON, 2001), assim como para a caracterização das proteínas espermáticas do gênero *Macaca* (KAWASE; CAO; XUAN, 2015; SKERGET et al. 2013; ZHOU et al., 2015).

### 1.10 Proteínas espermáticas

Durante a espermatogênese, os espermatozoides perdem a capacidade de sintetizar e secretar proteínas (SHABANOWITZ; KILLIAN, 1987), e ao final deste processo estão funcionalmente imaturos e imóveis (YOSHIDA et al., 2008). Mas, durante o trânsito epididimário (ou espermição), os espermatozoides passam pelo processo de maturação em que ocorre uma série de mudanças, incluindo aquisição de motilidade, remodelação da membrana plasmática, alterações no pH intra-espermático, e fosforilação de tirosina (DACHEUX et al., 2009). Durante a ejaculação, o gameta masculino entra em contato com as secreções das glândulas sexuais acessórias resultando na aderência de outras proteínas à membrana plasmática (VARRICCHIO et al., 1996), dando início a mudanças bioquímicas e estruturais relacionadas a hiperomotilidade, capacitação e reação acrossômica, que tornam os espermatozoides aptos à fecundação (MOREAU et al., 1999).

Essa série de alterações funcionais dependem integralmente de mudanças na composição proteica dos espermatozoides, que inclui a remoção e adição de proteínas, além de uma complexa matriz de modificações pós-translacionais, as quais são responsáveis por conduzir alterações estruturais e funcionais dos espermatozoides em função da maturação e capacitação espermática (BAKER et al., 2012; DACHEUX; DACHEUX, 2014). Além disso, as proteínas espermáticas possuem papel importante na integridade, morfologia e funcionalidade da célula espermática ao qual incluem a motilidade, capacitação, fecundação, ativação oocitária e desenvolvimento embrionário (PARISI et al., 2014).

No tocante a motilidade espermática, sabe-se que as proteínas associadas a esse parâmetro são classificadas em (i) proteínas enzimáticas relacionadas com a energia por

vias mitocondriais e glicolíticas; (ii) proteínas estruturais, tais como as fibras densas externas e proteínas que ancoram quinase A (AKAPs) nos flagelos; e (iii) proteínas de ativação como as proteínas quinase A transdutoras de sinal (PKA) e fosfatases de serina-treonina-tirosina-quinase (MURATORI et al., 2009). Em ratos, a proteína das fibras densas externas 2 (ODF2) foi identificada como um componente estrutural da cauda espermática, estando envolvida na regulação da motilidade dos espermatozoides (TARNASKY et al., 2010).

Em espermatozoides epididimários humanos foram identificadas 35 proteínas com massa molecular entre 15 e 160 kDa. Parte dessas proteínas está localizada na região subcelular desses gametas, e foram associadas a diferentes estruturas como o citoesqueleto, citoplasma, núcleo e mitocôndrias, sendo correlacionadas com diferentes funções biológicas (sinalização, transporte e ciclo celular, apoptose e resposta ao estresse, síntese proteica e metabolismo) (PILATZ et al., 2014). Algumas proteínas foram previamente reportadas apresentando relação com a capacitação espermática (SECCIANI et al., 2009) e também com a infertilidade (HOSSEINIFAR et al., 2013).

Na membrana plasmática espermática humana foi identificado, na região que abrange o acrossoma, um componente proteico denominado de 5'- nucleotidase (5 – NT; 125 kDa). Esta enzima, a priori, quando purificada e adicionada em um sistema de incubação *in vitro* para espermatozoides não teve nenhum efeito sobre a motilidade. No entanto, quando inibidores da 5'-nucleotidase foram incubados nesse mesmo sistema, foi observada uma inibição clara da motilidade dos espermatozoides, de um modo dependente da dose. Este resultado é interpretado como uma indicação de que a 5'-nucleotidase possui um papel significativo na regulação da motilidade espermática (AUMULLER et al., 1997).

Em espermatozoides epididimários de macaco-japonês (*Macaca fuscata*), foram identificadas 154 proteínas por meio da eletroforese bidimensional seguida da espectrometria de massa. Estas foram agrupadas em proteínas de membrana plasmática e mitocondrial, citoplasmáticas, de matriz mitocondrial e núcleo, e estão envolvidas na sinalização celular, motilidade, transporte espermático através do trato reprodutor da fêmea, hiperativação e reação acrossômica, eventos estes de suma importância para que ocorra a fecundação (KAWASE; CAO; XUAN, 2015). Zhou et al. (2015) avaliaram a composição proteica dos espermatozoides de primatas do gênero *Macaca*, por meio da cromatografia líquida acoplada à espectrometria de massas, e identificaram 2.044

proteínas espermáticas das quais 1.948 são ortólogas ao homem, indicando assim que esses PNH possuem a composição proteica similar comparada ao espermatozoide humano.

#### 1.10.1 Métodos para estudo das proteínas espermáticas

O recente progresso metodológico no estudo do proteoma tem aberto novos caminhos para obtenção de informações descritivas e quantitativas acerca dos processos biológicos que ocorrem no organismo animal e humano (BARBOSA et al., 2012). Sabe-se que as metodologias empregadas na proteômica podem ser classificadas em *bottom-up*, *middle-down* e *top-down*. A proteômica tipo *bottom-up* é caracterizada por um processo de separação por cromatografia líquida dos peptídeos obtidos após digestão enzimática, seguida da análise por espectrometria de massas (MS do inglês *Mass Spectrometry*). Essa técnica também é conhecida por proteômica *Shotgun*, contudo nesse processo as misturas proteicas são fracionadas por cromatografia líquida livre de gel. Já na proteômica tipo *middle-down* as proteínas são submetidas à digestão enzimática, entretanto, geram fragmentos de peptídeos maiores quando comparado a técnica *bottom-up*, que passará em seguida pela análise por MS, e por fim proteômica tipo *top-down*, no qual as proteínas (e não os peptídeos) são submetidas à análise por espectrometria de massa, essa estratégia de análise foca na caracterização completa da proteína intacta e de suas modificações pós-traducionais (ARMIROTTI; DAMONTE, 2010; AHRENS et al., 2010; WU et al., 2012; ZHANG et al., 2013). A combinação dessas abordagens com outros processos, como fracionamento subcelular ou precipitação de proteínas, tem sido utilizado como uma alternativa efetiva no enriquecimento de amostras compostas de baixas concentrações proteicas (KOSAKO et al., 2011).

Contudo, para que se tenha sucesso nesse processo, inicialmente é de suma importância a extração adequada das proteínas, o qual deve se levar em consideração o tipo e origem das amostras biológicas. Logo, o procedimento de extração necessita de otimização individual, mas em geral, as proteínas precisam ser solubilizadas, desagregadas, desnaturadas e submetidas a tratamento com agentes redutores de pontes dissulfeto (DE MARQUI et al., 2006).

Técnicas de eletroforese uni (1D) e bidimensional (2D) têm sido comumente utilizadas com o objetivo de separação proteica. Na eletroforese 1D ocorre a separação das proteínas de acordo com o peso molecular, em que há a formação de bandas, enquanto que na eletroforese 2D elas são separadas em duas etapas, em que primeiramente há a separação de proteínas utilizando focalização isoelétrica (etapa I), seguida por uma eletroforese em gel de poliacrilamida na presença de SDS (SDS-PAGE do inglês *Sodium dodecyl sulfate polyacrylamide gel electrophoresis*) visando a separação numa segunda dimensão, ou seja, de acordo com o peso molecular (etapa II), onde há a formação dos *spots* (HACHEY; CHAURAND, 2004). Posteriormente, para tornar visíveis as bandas ou *spots* proteicos (eletroforese 1D e 2D, respectivamente), os géis são comumente corados com Azul de *Coomassie*, nitrato de prata ou outros corantes comerciais (DOWSEY et al., 2003; GHARAH DAGHI et al., 1999; SHEVCHENKO et al., 1996;). Em seguida há a excisão das bandas ou *spots* de proteínas individuais do gel e digestão com uma protease (ex. tripsina), seguindo-se de uma identificação por espectrometria de massas (DELAHUNTY; YATES, 2005).

A eletroforese 1D e 2D já foi utilizada na análise de proteínas em espermatozoides de mamíferos tais como: homem (JOHNSTON et al., 2005; PILCH; MANN, 2006), touros (PARK et al., 2012), búfalos (ASADPOUR et al. 2007), carneiros (YUE et al., 2009), suínos (CORCINI et al., 2012), catetos (SANTOS et al., 2014), PNH (KAWASE; CAO; XUAN, 2015) e cães (AQUINO-CORTEZ et al., 2017).

Neste cenário, a MS têm sido comumente empregada nos últimos anos devido a possibilidade de identificação direta dos constituintes individuais de complexos proteicos envolvidos em uma ampla gama de funções fisiológicas, sendo assim, tornou-se uma plataforma padrão na proteômica (PLATT et al., 2008; SCHILLER et al., 2000; SWEGEN et al., 2015; ZERBINATI et al., 2017). MS é uma ferramenta analítica que identifica proteínas ou peptídeos medindo as massas de moléculas convertidas em íons por meio da sua relação massa sobre carga ( $m/z$ ). Entretanto, essa técnica depende ou não da digestão proteolítica de proteínas em peptídeos antes da introdução no espectrômetro de massa (PLESSIS et al., 2011).

No estudo das proteínas espermáticas, duas das técnicas MS tem sido mais comumente empregadas para caracterizar *spots* proteicos de um gel de eletroforese previamente executado, que são: a ionização e dessorção a laser assistida por matriz – analisador do tipo tempo de voo, conhecida como MALDI-TOF (do inglês *Matrix*

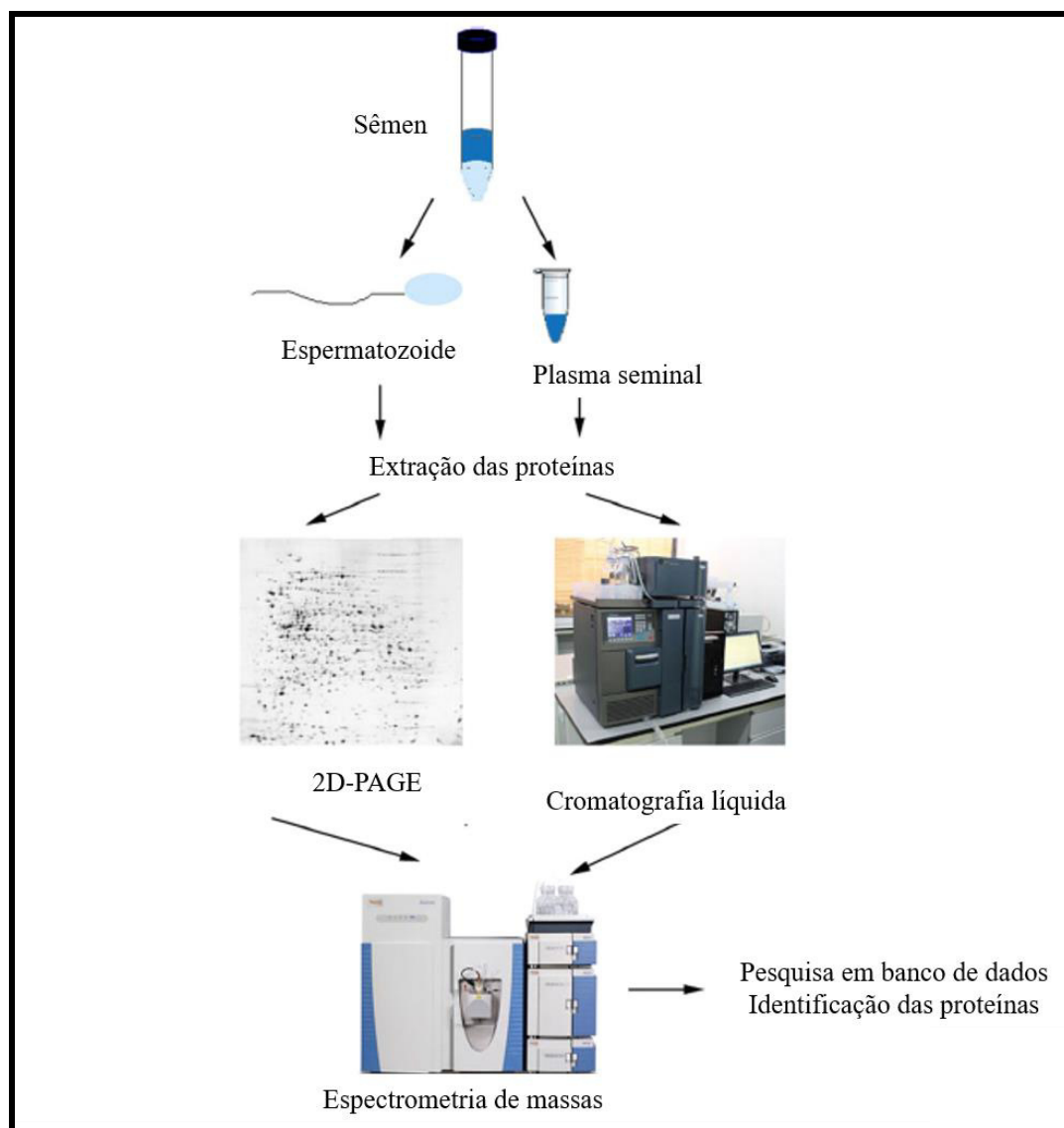


*Associated Laser Desorption-Ionization - Time of Flight*), e a cromatografia líquida com um espectrômetro de massa em tandem (LC-MS/MS do inglês *Liquid Chromatography–Mass Spectrometry*). Os dados espectrais obtidos a partir da MS proporcionam a relação  $m/z$  da proteína e sua intensidade associada, o que ajuda a determinar a expressão diferencial proteica e suas modificações (MITULOVIC; MECHTLER, 2006; PLESSIS et al., 2011).

Na análise MALDI-TOF, a determinar da relação  $m/z$  de uma proteína é realizada após a digestão das bandas de proteínas do gel com proteases (PLESSIS et al., 2011). A digestão de proteínas utilizando enzimas resultam em fragmentos peptídicos que são facilmente ionizados no espectrômetro de massa (MCLUCKEY, 1992). Depois que as massas peptídicas foram registradas, elas são combinadas com conjuntos de massas de referência de proteína teoricamente digeridas a partir de um banco de dados para identificação proteica (mapeamento de massa de peptídeo).

Por outro lado, o LC-MS/MS, é uma técnica analítica altamente sensível e específica que combina a separação física de peptídeos gerados pela digestão enzimática de um *spot* por cromatografia líquida de alta eficiência (do inglês *High performance liquid chromatography* - HPLC) seguida da determinação da massa e subsequente sequenciamento em MS/MS (MITULOVIC; MECHTLER, 2006) (Figura 8).

Figura 8- Diagrama esquemático mostrando a análise proteômica do plasma seminal e da célula espermática.



Fonte: Adaptado de Cao et al., (2018).

Contudo, nos últimos anos, a proteômica *Shotgun* tornou-se o método de escolha para identificar e quantificar proteínas em larga escala (ALVES et al., 2007; WASHBURN; WOLTERS; YATES, 2001), já sendo utilizada na identificação proteica de espermatozoides de camundongos (BAKER et al., 2008), humano (NIXON et al., 2011), bovinos (CANIO et al., 2014), primatas não-humanos (ZHOU et al., 2015). Esta estratégia baseia-se na digestão de proteínas em peptídeos (sem termos previamente a etapa de separação proteica em gel de eletroforese), seguido do sequenciamento usando espectrometria de massa em tandem (LC-MS/MS) e pesquisa automatizada de banco de dados (WASHBURN; WOLTERS; YATES, 2001). Em comparação com os métodos

identificação baseada em MS, como os géis 1D e 2D, na proteômica *Shotgun* há uma melhor sensibilidade à detecção de proteínas e com isso uma maior taxa de transferência de dados (GORG; WEISS; DUNN, 2004).

### 1.11 Hipóteses

- A distância anogenital pode ser utilizada como um biomarcado no exame andrológico para a seleção de machos de *S. collinsi*.
- A micromorfologia e ultraestrutura dos espermatozoides de *S. collinsi* demonstrada por eletromicroscopia de varredura e transmissão, é semelhante do que foi descrito em outros primatas do gênero Saimiri.
- As proteínas associadas com a qualidade seminal em macacos-de-cheiro (*S. collinsi*) são comuns para outros primatas.

### 1.12 Objetivos

#### 1.12.1 Geral

- Estudar a morfofisiologia reprodutiva de macaco-de-cheiro (*S. collinsi*).

#### 1.12.2 Específicos

- Associar a distância anogenital com os parâmetros espermáticos e hormônios reprodutivos em macacos-de-cheiro (*S. collinsi*).
- Descrever as características micromorfológicas e ultraestruturais dos espermatozoides de macacos-de-cheiro (*S. collinsi*) por meio da microscopia eletrônica de varredura e transmissão.
- Caracterizar o perfil proteico dos espermatozoides de macaco-de-cheiro (*S. collinsi*).
- Associar as expressões de proteínas espermáticas e as características espermáticas de macaco-de-cheiro (*S. collinsi*).

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**The use of anogenital distance as a noninvasive predictor of seminal quality in captive squirrel monkey (*Saimiri collinsi* Osgood, 1961)**

**Anogenital distance in squirrel monkey**

Danuza Leite Leão<sup>1\*</sup>, Wlaysia Vasconcelos Sampaio<sup>1,2,3</sup>, Patrícia da Cunha Sousa<sup>1</sup>, Irma Caroline Oskam<sup>4</sup>, Regiane Rodrigues dos Santos<sup>1</sup>, Sheyla Farhayldes Souza Domingues<sup>1,2</sup>

<sup>1</sup> Laboratory of Wild Animal Biotechnology and Medicine, Federal University of Pará, Belém, Pará, Brazil

<sup>2</sup> Postgraduate Program in Animal Health and Production in the Amazon, Federal Rural University of the Amazon, Belém, Pará, Brazil

<sup>3</sup> Federal Rural University of the Amazon, Parauapebas, Pará, Brazil

<sup>4</sup> The Livestock Production Research Centre, Norwegian University of Life Sciences, Oslo, Norway

\*Corresponding author: Danuza Leite Leão. Laboratory of Wild Animal Biotechnology and Medicine, Federal University of Pará, BR 316 Km 61, CEP 68740–970, Castanhal, Pará, Brazil. Tel.: +55 91 33114707.

E-mail: danleao.88@gmail.com

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**Text figures:** 1

**Tables:** 3

## ABSTRACT

**Background:** Anogenital distance is considered a noninvasive measure to assess the development and functionality of sexual organs in different animal species. Hence, this measurement could potentially be used during the selection of non-human primates for reproductive biotechnology programs. The aim of this study was to assess the correlation between anogenital distance and reproductive parameters in captive *Saimiri collinsi*.

**Methods:** Eight mature *S. collinsi* males were evaluated. Body weight, reproductive hormone levels, testicular volume and seminal parameters were determined and their relationship with anogenital distance measurements was assessed.

**Results:** Anogenital distance was correlated with seminal volume, sperm motility, vigour, and plasma membrane integrity, but not with body weight, reproductive hormones, and testicular volume.

**Conclusion:** The determination of anogenital distance is a noninvasive method to predict seminal quality. This procedure has the advantage of providing andrologic information without a negative impact on animal welfare.

**Keywords:** anogenital distance, spermatozoa, neotropical primates, testes.

## 1 INTRODUCTION

Anogenital distance (AGD) is an measure and has been used as a noninvasive method to evaluate the development and functionality of male sex organs in rats<sup>1</sup> and humans<sup>1,2,3,4,5</sup>. Deficient androgen exposure during a critical period of testis development defined as “masculinisation programming window - MPW” (8–14 weeks of gestation in humans) was associated with outcomes such as reduced penile length, testicular dysgenesis syndrome, and the reduction in AGD<sup>6,7,8,9,10,11</sup>.

In adult human studies, AGD is measured in two different ways: in man, from the posterior base of the scrotum to the center of the anus (AGD<sub>AS</sub>) and from the cephalad insertion of the penis to the center of the anus (AGD<sub>AP</sub>)<sup>9,3,12</sup>. In women, the measurement is made from the anterior clitoral surface to the center of the anus (AGD<sub>AC</sub>) and from the posterior fourchette to the center of the anus (AGD<sub>AF</sub>)<sup>13</sup>. Since increased androgen concentrations lead to increased AGD in utero, males have AGDs almost twice as long those in females, and this is considered a sexually dimorphic characteristic in humans<sup>14</sup>, rodents<sup>10,15</sup> and rhesus monkeys<sup>16</sup>. In adult rats, it has been reported that AGD may be mediated by modulation of local androgen/estrogen action.<sup>17</sup> However, only in men, studies have shown that a short AGD predicts small testes size and poor seminal quality.<sup>9,8,18,3</sup> In an Old World Primate, the Rwenzori Angolan colobus monkeys (*Colobus angolensis ruwenzorii*), the AGD<sub>AS</sub> measurement was considered a noninvasive method to assess male competitive ability in the wild because it correlated with male dominance rank, but no information was given on seminal quality<sup>19</sup>.

*Saimiri collinsi*, popularly known as squirrel monkey, is a Neotropical primate endemic to the Amazon in Brazil<sup>20</sup>, and was proposed as an experimental model for reproductive biotechnology for the conservation of free-living species in the genus *Saimiri*.<sup>21,22</sup> Although, in squirrel monkeys there is documented a positive correlation between

testicular volume and the liquid fraction of the ejaculate<sup>23</sup>, nothing is known about relationship between testosterone- and estradiol levels, testicular volume and sperm quality and the AGD. Given the biological importance of this potential biomarker, our overall hypothesis is that *S. collinsi* testicular function is related to anogenital length, and could predict seminal quality in nonhuman primates. In this way, the AGD measurement could be used as an auxiliary noninvasive parameter in andrologic examination for the selection of non-human primates in biotechnology programs for the conservation of endangered species, as well as studies on reproductive physiology. Thus, the objective of this study was to evaluate the correlation between AGD values and hormone levels (testosterone and estradiol) with body weight, testicular biometry, seminal volume and sperm parameters in *S. collinsi* kept in captivity.

## **2 MATERIALS AND METHODS**

### **2.1 Ethical Committee and place of execution**

This study was approved by the Ethical Committee in Animal Research (no. 02/2015/CEPAN/IEC/SVS/MS) and by the System of Authorization and Information in Biodiversity (SISBIO/ICMBio/MMA no.47051-2). All procedures were performed under the supervision of a veterinarian. Semen collections were carried out during 12 months at the National Primate Center (CENP, Ananindeua, Pará, Brazil).

### **2.2 Animals**

*Saimiri collinsi* males maintained in captivity (indoors) at the CENP (1°22'58"S and 48°22'51"W), were used for collecting semen. We selected (N=8) sexually mature and healthy males. The animals were housed in mixed groups (males and females varied numbers) in collective cages of 4.74 m x 1.45 m x 2.26 m (length, width and height, respectively), under a natural photoperiod (i.e. 12 h of light and 12 h of dark). The climate is humid and tropical,

with an average annual temperature of 28 °C. The diet consisted of fresh fruits, vegetables, commercial pellet chow specific for neotropical non-human primates (P18, 18% protein, 6.5% fiber, Megazoo®, Minas Gerais, Brazil) and cricket larvae (*Zophobas morio*). Vitamins, minerals and eggs were supplied once a week and water was available *ad libitum*.

For biometric measurements, blood and semen collection, the animals were first physically restrained by a trained animal caretaker wearing leather gloves. After this, the animals were anesthetized with ketamine hydrochloride (20 mg/kg; IM; Vetanarcol, König S.A., Avellaneda, Argentina) and xylazine hydrochloride (1 mg/kg; IM; Kensol, König S.A.), and monitored by a veterinarian. All data (biometric measurements, blood and semen collection) were collected once a month for 12 months.

### **2.3 Genital measurements**

After total anesthetic effect, the animals were weighed using a digital weighing balance (Toledo Prix 3 Plus 30 kg Digital Balance, Toledo do Brasil, São Bernardo do Campo, SP, Brazil) and the testicular biometry (length, width, height and circumference) was measured using a universal caliper. The testicular volume was calculated by the ellipsoid formula (length x width x height x 0.524) according to Oliveira et al.<sup>21</sup>

Two variants of AGD were assessed as described for humans<sup>9,4</sup> using a universal caliper: the first was measured from the cephalad insertion of the penis to the center of the anus (AGD<sub>AP</sub>), and the second was measured from the posterior base (first fold) of the scrotum to the center of the anus (AGD<sub>AS</sub>) (Figure 1).

### **2.4 Hormonal assays**

For the blood collection (1 mL), the femoral vein was punctured with hypodermic needles (20 mm × 0.55 mm; 24 G × 3/4") coupled to a 3 mL syringe, and blood was transferred to tubes containing the anticoagulating EDTA. Samples were centrifuged (3000 rpm for 5 minutes) (Modelo Combate, Celm, Brazil) and the obtained serum was frozen (-80 °C) for

hormonal dosage. The plasma levels of Testosterone (ng/mL) and 7 $\beta$ -estradiol (pg/mL) were measured by chemiluminescence assay using an immunoassay system (VITROS® ECIQ, Ortho Clinical Diagnostics, New Jersey, United States of America). All the protocols were applied according to the manufacturer.

## **2.5 Semen collection and analyses**

The genital region was sanitized with a mild soap and distilled water (1:10) and gauze. The prepuce was retracted with the thumb and index finger for a more efficient cleaning of the penis with saline solution. Animals were stimulated with rectal electro-ejaculation (EEJ) procedure described in Oliveira et al.<sup>21,22</sup> In brief, an EEJ (Autojac-Neovet, Uber-aba, Brazil) rectal probe was smeared with a sterile lubricant jelly (KY™ Jelly, Johnson & Johnson Co., Arlington, TX, USA), introduced in the rectum (~2.5 cm deep) and electrical stimuli were delivered. The stimulation session consisted of three series (7–8 min), composed of 35 electrical stimuli (12.5 – 100 mA) within an interval of 30s between series. For each animal, the intervals between semen collections were 30 days. If a male did not ejaculate after the session, no further attempts were made to collect semen.

Ejaculates were collected into microtubes (1.5 mL) and placed in a water bath at 37 °C immediately after ejaculation. Seminal volume (liquid fraction) was evaluated in a graduated tube. Appearances were assessed subjectively, i.e., color (colorless, yellowish, or whitish) and opacity (opaque or transparent). Seminal pH was measured with a pH strip (Merk Pharmaceuticals, Darmstadt, Germany). Sperm concentration was determined in a Neubauer chamber, 1  $\mu$ L semen was diluted in 99  $\mu$ L formalin solution 10%. All evaluations were performed under a light microscope (Nikon E400, Japan) at a magnification of 100 $\times$ . Sperm morphology was evaluated by a smear prepared by adding 5  $\mu$ L of eosin-nigrosin stain (Vetec, Rio de Janeiro, Brazil) to 5  $\mu$ L of semen on a pre-warmed (37 °C) glass slide.



Morphologic defects detected in sperm were classified as major or minor. Major defects are those that affect fertility, and minor defects are of less importance.<sup>22</sup>

## 2.6 Statistical analyses

All data are expressed as the mean  $\pm$  standard error (SE) and analyzed by the StatView 5.0 program (SAS Institute Inc., Cary, NC, USA). Data were checked for normality using the Kolmogorov-Smirnov test. Correlation coefficients (Spearman correlation) were used to determine the relationship in *S. collinsi* between AGD measurements (AGD<sub>AP</sub> and AGD<sub>AS</sub>) and the hormone values (testosterone and estradiol) along with total testicular volume, seminal volume, sperm concentration, motility, vigor, normal sperm, major and minor sperm defects.  $P \leq 0.05$  was considered statistically significant.

## 3 RESULTS

The mean ( $\pm$  SE) values in adult male *S. collinsi* of body weight was  $860 \pm 882$  (694-1,152 g min-max), AGD<sub>AP</sub> was  $3.3 \pm 0.06$  (2.5-4.1 cm min-max), AGD<sub>AS</sub> was  $2.3 \pm 0.04$  (1.7-2.9 cm min-max), testosterone level was  $19.2 \pm 2.6$  (467-50,900 ng/dl; min-max) and estradiol level was  $40.6 \pm 6.0$  (80.7-98.9 pg/ml; min-max) (Table 1).

Seminal liquid fractions were colorless, whitish or yellowish, transparent or opaque and the pH (Mean  $\pm$  SE) of the ejaculates was  $7.5 \pm 0.06$ . The mean value of seminal parameters are summarized in Table 1.

Correlation analyses for body weight, reproductive hormones (testosterone and estradiol), testicular biometry and semen parameters with AGD measurements are depicted in Table 2. There was no significant correlation between body weight, testosterone and estradiol levels with AGD<sub>AP</sub> and AS. On the other hand, the AGD<sub>AP</sub> was positively and significantly correlated with the sperm parameters of motility ( $\rho = 0.56$ ;  $Z = 2.57$ ;  $P = 0.01$ ) and vigour ( $\rho = 0.59$ ;  $Z = 2.72$ ;  $P = 0.006$ ). In addition, it was possible to verify the positive and significant correlation

between the AGD<sub>AS</sub> with seminal volume ( $\rho = 0.35$ ;  $Z = 2.06$ ;  $P = 0.03$ ) and sperm plasma membrane integrity ( $\rho = 0.46$ ;  $Z = 2.09$ ;  $P = 0.03$ ) (Table 2).

The hormonal analysis showed a negative and significant correlation between testosterone levels and the weight of the animals ( $\rho = -0.35$   $Z = -2.32$   $P = 0.02$ ), however there was no significant correlation between the estradiol levels and the weight, testicular biometry and seminal parameters analyzed (Table 3).

#### 4 DISCUSSION

In the present study, we examined two variants of AGD (AGD<sub>AS</sub> and AGD<sub>AP</sub>) and it was possible to show that AGD measurements might be an important noninvasive parameter to select *Saimiri* monkeys, with good fertility to participate in assisted reproduction programs.

Human male AGD can be measured in a few different ways<sup>23</sup>, because the measurements depend on different landmarks on the soft tissue. It was reported that in men, the AGD<sub>AS</sub> is effortless to measure due to the easy identification of soft tissue boundaries.<sup>24</sup> Additionally in adult men, AGD<sub>AS</sub> correlates most strongly with fertility parameters.<sup>9</sup> In Rwenzori Angolan colobus, only AGD<sub>AS</sub> was measured and found to be a predictor of male competitive ability.<sup>19</sup> However, there is no gold standard method for measuring AGD in adult squirrel monkeys, so we cannot establish whether one measure is better than the other, thus we verified the relationship of both measures with reproductive parameters. In *S. collinsi*, both measurements of AGD had a positive and significant correlation with seminal parameters. Thus, it is worth analyzing if the same occurs in other species of non-human primates.

Our results showed that AGD measurements were not correlated with body weight, reproductive hormones (testosterone and estradiol), nor testicular volume in *S. collinsi*. In men, AGD measurements are also not correlated with body weight<sup>25</sup>, but significant positive

correlations between body mass index and AGD<sub>AP</sub> and AGD<sub>AS</sub> was previously shown<sup>5</sup>. Body mass is influenced by the accumulation of fat in the anterior region of the pubic symphysis, an area that is included in the AGD<sub>AP</sub> measurement, but not at AGD<sub>AS</sub>.<sup>9</sup> AGD<sub>AS</sub> is considered to be the most reliable and repeatable measurement, in men, because this variant is unaffected by obesity and age.<sup>26,27,28</sup> Male *Saimiri* sp. gain weight seasonally about three months before, as well as during the copulation season. This increase is a result of the deposition of fat and water especially in the thorax, arms, and shoulders<sup>29,30</sup>, thus not influencing the measures AGD<sub>AP</sub> and AGD<sub>AS</sub>.

During sexual development, the immature genital precursors migrate ventrally via an androgen mediated pathway, and endocrine disruptors during the MPW (to 8–14 weeks of gestation in humans) and can permanently alter genital development, growth and function.<sup>31</sup> On the other hand, it was shown in rats that, when the endocrine action occurs in the postnatal period there is less damage in the genital development.<sup>14,32,33</sup> Although studies with rats and humans have shown the relationship between small AGD size with low serum testosterone level and testicular size<sup>9,1,34</sup>, we did not find correlation between AGD and testicular volume or reproductive hormones (testosterone and estradiol serum level) in *S. collinsi*. Jain et al.<sup>35</sup> observed, in humans, a correlation of AGD with testosterone levels, however, with increasing gestational age (22 to 44 weeks), in utero, this correlation was lost. Therefore, the authors suggest that AGD is influenced by testosterone early in gestation during the MPW, but with advancing gestational age, AGD becomes independent of testosterone levels. Thus, the increase in AGD over time is related to the growth of the body and not to testosterone levels. Liu et al.<sup>36</sup> measured cord blood testosterone and estradiol using radioimmunoassay in human term neonates and found no correlation with AGD. However, in adults, associations of AGD with testosterone levels have been inconclusive, since studies have shown an association<sup>7</sup>, while others have not<sup>4</sup>.

Previous studies have shown that AGD is related to fertility and sperm production in men.<sup>9,37</sup> The current report establishes that AGD measurements might also help determine genital development and function in squirrel monkeys. We showed a positive and significant correlation between AGD<sub>AP</sub> with motility and vigour, and AGD<sub>AS</sub> with seminal volume and sperm plasma membrane integrity in *S. collinsi*, thus showing that the increase in AGD measurements are indicative of better seminal quality. In humans, Mendiola et al.<sup>9</sup> evaluated men's sperm parameters in relation to AGD<sub>AP</sub> and AGD<sub>AS</sub>, and found a significant association only for AGD<sub>AS</sub> with sperm parameters (sperm concentration, motility, morphology, total sperm count and total motile count). Rodent studies indicate that seminal parameters may be associated with AGD measurements, since seminal volume may reflect seminal vesicle and prostate function, and the motility may reflect germ cell number and testicular function.<sup>14,32</sup>

Parra et al.<sup>4</sup> and Zhou et al.<sup>5</sup> reported in men that both AGD<sub>AS</sub> and AGD<sub>AP</sub> were not associated with any semen parameters. The authors reported the possible reason for these conflicting results may be due to the fact that most of these studies in men were conducted on patients attending in andrology practice for infertility assessment.<sup>8,37,3</sup> Thus, infertile men may have infertility reasons that can cause testis dysfunction and changes in sperm parameters, so therefore, the AGD may also have changed as a consequence of infertility.<sup>5</sup> In the present study, all seminal parameters analyzed are in accordance with the values described for healthy *S. collinsi*<sup>21</sup>, as expected since animals were selected without a history of reproductive problems.

Despite the variable results in humans, it is worth mentioning that the men included these in studies were all submitted to a period of sexual abstinence prior to semen collection in order to obtain optimal sperm quality.<sup>7,3,37</sup> Nevertheless, in our experiment, for the sake of best animal welfare in captivity, all monkeys were housed in mixed groups of males and females. Therefore, it was not possible to isolate the animals and prevent copulation or male

masturbatory activity, which is part of the natural behavioral repertoire in nonhuman primate species.<sup>38,39</sup> These factors (copulation and masturbation) directly influence seminal quality and may explain our negative but not significant correlation results found between AGD<sub>AS</sub> and sperm concentration, motility, vigour and major sperm defects.

Sperm quality has a direct effect on the fertilization and developmental competence of embryos in mammals<sup>40</sup>, due to the paternal factors contributing to embryo quality. In humans, lower cleavage rates and blastocyst formation rates have been noted when morphologically abnormal sperm were used to fertilize oocytes in in vitro embryo production.<sup>41</sup>

Squirrel monkeys (*Samiri* sp.) are Neotropical primates that show strict reproductive seasonality.<sup>42</sup> In addition to weight gain as previously mentioned during the breeding season (fattening phenomenon), an increase of testosterone hormone levels during the breeding season in *Saimiri* sp. has been described<sup>29,43</sup> Chen et al.<sup>44</sup> evaluated seven adult Bolivian squirrel monkeys (*Saimiri sciureus*) kept in captivity for 13 months, and showed there were seasonal changes in body weight, plasma testosterone and seminal volume in these animals. However, the authors report that seasonal changes in seminal volume appear independent of seasonal testosterone changes. In captive *S. collinsi*, we showed a decrease in testosterone levels with the increase in the animals weight, and no correlations between testosterone and estradiol levels and testicular biometry and seminal parameters were found. Nevertheless, it is still necessary to determine the reproductive and fattening period in captivity of these animals, as well as the influence of a social hierarchy in this period and the relationship with the levels of reproductive hormones, testicular biometry and seminal quality.

In conclusion, our results indicate that AGD size predicts seminal quality in *S. collinsi*, and the measurement of AGD may provide a new metric for andrologic examination in assisted reproduction programs in nonhuman primates. Furthermore, this noninvasive and

rapid approach can be considered as a refinement to support the field control of free-living monkeys that are sometimes exposed to environmental contaminants or other stress sources that may affect their reproductive function.

## CONFLICT OF INTEREST STATEMENT

None of the authors have any conflict of interest to declare.

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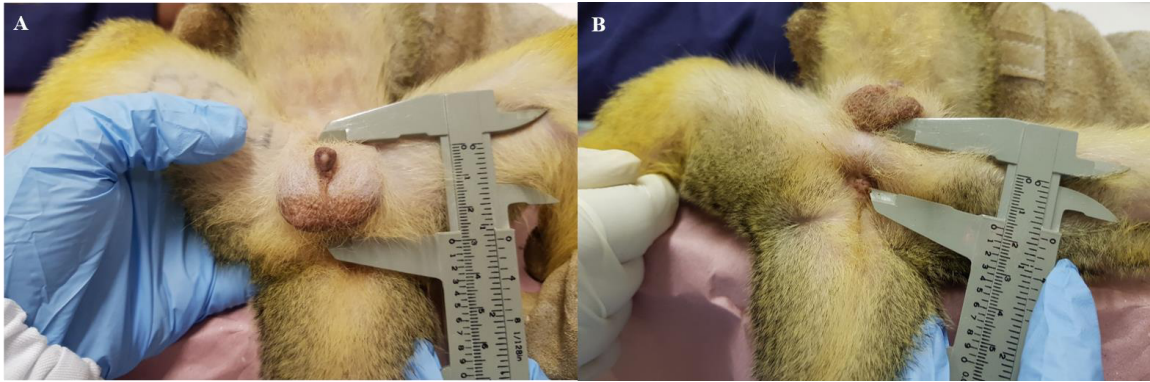


FIGURE 1 Representative image showing the measurement of the anogenital distances: (A)  $AGD_{AP}$  - from the cephalad insertion of the penis to the center of the anus and (B)  $AGD_{AS}$  - from the posterior base of the scrotum to the center of the anus.

TABLE 1 Mean  $\pm$  standard error (SE) of body weight, hormone levels (testosterone and estradiol), anogenital distance (AGD<sub>AP</sub> and AGD<sub>AS</sub>) and seminal parameters in adult squirrel monkeys (*Saimiri collinsi*; N=8).

Variable	Mean $\pm$ SE	Minimum-Maximum
Body weight (g)	860 $\pm$ 882	694-1,152
<i>Reproductive Hormones</i>		
Testosterone (ng/dl)	19,2 $\pm$ 2,6	467-50,900
Estradiol (pg/ml)	40,6 $\pm$ 6,0	80,7-98,9
<i>Anogenital distance and testicular biometry</i>		
AGD <sub>AP</sub> (cm)	3.3 $\pm$ 0.06	2.5-4.1
AGD <sub>AS</sub> (cm)	2.3 $\pm$ 0.04	1.7-2.9
Testicular volume (cm <sup>3</sup> )	2.3 $\pm$ 0.1	0.69-5.2
<i>Semen parameters</i>		
Seminal Volume ( $\mu$ L)	179 $\pm$ 18.9	10-600
Sperm concentration ( $\times 10^6$ /mL)	127 $\pm$ 164	25-45
Motility (%)	60.6 $\pm$ 4.0	0-100
Vigour (1-5)	3.0 $\pm$ 0.2	0-5
Plasm membrane integrity (%)	60.4 $\pm$ 3.5	0-100
Normal sperm (%)	72.1 $\pm$ 2.3	37-96
Major sperm defects (%)	11.5 $\pm$ 1.8	0-46
Minor sperm defects (%)	5.2 $\pm$ 1.0	0-40

TABLE 2 Correlation of anogenital distance (AGD<sub>AP</sub> and AGD<sub>AS</sub>) with hormone levels, body weight and seminal parameters in adult squirrel monkeys (*Saimiri collinsi*; N=8).

Parameters	AGD <sub>AP</sub>	AGD <sub>AS</sub>
Testosterone	$\rho = -0.07$ Z= -0.15 P = 0.88	$\rho = -0.07$ Z= -0.15 P = 0.88
Estradiol	$\rho = -0.57$ Z= -1.15 P = 0.25	$\rho = -0,57$ Z= -1.15 P = 0.25
Body weight	$\rho = 0.17$ Z= 1.12 P = 0.26	$\rho = -0.27$ Z= -1.78 P = 0.07
Testicular volume	$\rho = 0.21$ Z= 1.38 P = 0.16	$\rho = 0.10$ Z= 0.64 P = 0.51
Seminal Volume	$\rho = 0.11$ Z= 0.67 P = 0.50	$\rho = 0.35$ Z= 2.06 P = 0.03
Sperm concentration	$\rho = 0.12$ Z= 0.57 P = 0.56	$\rho = -0.29$ Z= -1.3 P = 0.18
Motility	$\rho = 0.56$ Z= 2.57 P = 0.01	$\rho = -0.07$ Z= -0.33 P = 0.73
Vigour	$\rho = 0.6$ Z= 2.72 P = 0.006	$\rho = -0.06$ Z= -0.27 P = 0.78
Plasm membrane integrity	$\rho = 0.22$ Z= 1.01 P = 0.30	$\rho = 0.46$ Z= 2.09 P = 0.03
Normal sperm	$\rho = 0.32$ Z= 1.51 P = 0.13	$\rho = 0.4$ Z= 1.87 P = 0.06
Major sperm defects	$\rho = -0.01$ Z= -0.06 P = 0.95	$\rho = -0.2$ Z= -1.07 P = 0.28
Minor sperm defects	$\rho = 0.07$ Z= 0.41 P = 0.67	$\rho = 0.06$ Z= 0.34 P = 0.73

TABLE 3 Correlation of testosterone and estradiol levels with body weight and seminal parameters in adult squirrel monkeys (*Saimiri collinsi*; N=8).

Parameters	Testosterone	Estradiol
Body weight	$\rho = -0,35$ Z= -2,32 P = 0,02	$\rho = -0,05$ Z= -0,36 P = 0,71
Testicular volume	$\rho = -0,05$ Z= -0,37 P = 0,70	$\rho = -0,20$ Z= -1,27 P = 0,20
Seminal Volume	$\rho = -0,07$ Z= -0,49 P = 0,62	$\rho = -0,05$ Z= -0,33 P = 0,73
Sperm concentration	$\rho = -0,15$ Z= -0,96 P = 0,33	$\rho = 0,07$ Z= 0,46 P = 0,63
Motility	$\rho = -0,02$ Z= -0,15 P = 0,87	$\rho = 0,16$ Z= 0,877 P = 0,38
Vigour	$\rho = -0,02$ Z= -0,12 P = 0,89	$\rho = 0,14$ Z= 0,80 P = 0,42
Plasm membrane integrity	$\rho = -0,05$ Z= -0,27 P = 0,78	$\rho = -0,142$ Z= -0,77 P = 0,43
Normal sperm	$\rho = 0,11$ Z= 0,71 P = 0,47	$\rho = 0,04$ Z= 0,26 P = 0,79
Major sperm defects	$\rho = 0,05$ Z= 0,32 P = 0,74	$\rho = 0,04$ Z= 0,24 P = 0,80
Minor sperm defects	$\rho = 0,12$ Z= 0,6 P = 0,50	$\rho = -0,13$ Z= -0,75 P = 0,44

1 **Micromorphological and ultrastructural description of squirrel monkeys (*Saimiri***  
2 ***collinsi* Osgood, 1916) sperm**

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4 Danuza L. Leão<sup>a,b\*</sup>, Wlaysia V. Sampaio<sup>a,b,c</sup>, Patrícia C. Sousa<sup>a</sup>, Arlindo A. A. Moura<sup>d</sup>, Irma C.  
5 Oskam<sup>e</sup>, Regiane R. Santos<sup>a,f</sup>, Sheyla F. S. Domingues<sup>a,b,g</sup>

6  
7 <sup>a</sup> Laboratory of Wild Animal Biotechnology and Medicine, Federal University of Pará, Belém,  
8 Pará, Brazil

9 <sup>b</sup> Postgraduate program in Animal Health and Production in the Amazon, Federal Rural  
10 University of the Amazon, Belém, Pará, Brazil

11 <sup>c</sup> Federal Rural University of the Amazon, *Campus* Parauapebas, Pará, Brazil

12 <sup>d</sup> Laboratory of Animal Physiology, Federal University of Cearpa, Ceará, Brazil

13 <sup>e</sup> The Animal Production Experimental Center, Norwegian University of Life Sciences, Ås,  
14 Norway

15 <sup>f</sup> Schothorst Feed Research, Lelystad, Netherlands

16 <sup>g</sup> Faculty of Veterinary Medicine, Federal University of Pará, Castanhal, Pará, Brazil

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20  
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22  
23 \*Corresponding author: Danuza Leite Leão. Laboratory of Wild Animal Biotechnology and  
24 Medicine, Federal University of Pará, BR 316 Km 61, CEP 68740–970, Castanhal, Pará,  
25 Brazil. Tel.: +55 91 33114707.

26 E-mail: danleao.88@gmail.com

27 **Abstract**

28 *Saimiri collinsi* is an animal model in biotechnology research for conservation of species from  
29 the genus *Saimiri*. However, the development of biotechnologies depends on a proper  
30 knowledge of the sperm morphology to understand the basic aspects of the sperm physiology,  
31 since the potential male fertility depends of different cellular sperm's structures. Whit this  
32 purpose, this study characterized the micro morphological and ultrastructural characteristics  
33 of squirrel monkeys (*Saimiri collinsi*) sperm using scanning- and transmission eléctron  
34 microscopy (SEM and TEM, respectively). The SEM electromyography revealed that a  
35 normal *Saimiri collinsi* sperm measures  $71.7 \pm 0.7 \mu\text{m}$  with lateral tail insertion, a paddle-  
36 shaped flattened head and an acrosome occupied most of the head. The TEM also showed that  
37 the middle piece is characterized by a central 9 + 2 microtubule axoneme surrounded by nine  
38 dense fibres, and the mitochondria were juxtaposed forming the mitochondrial sheath. In  
39 conclusion, we provide the first micro morphological and ultrastructure description of *S.*  
40 *collinsi* sperm.

41 **Keywords:** Sperm, electron microscopy, scanning, transmission, neotropical primates

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## 53 Introduction

54 The squirrel monkey *Saimiri collinsi*, a neotropical primate endemic from Amazon-  
55 Brazil, is an animal model for conservational and biotechnology studies for species from the  
56 genus *Saimiri* (Oliveira et al., 2015; 2016ab). Among them, *S. oerstedii* and *S. vanzollini* are  
57 listed as vulnerable, while the *S. ustus* is almost threatened to extinction (IUCN, 2019).

58 Despite the relevant studies conducted to investigate the male reproductive physiology  
59 in this non-human primates using classical semen analysis (Oliveira et al., 2015; 2016ab;  
60 Sampaio et al., 2017; Almeida et al., 2018), knowledge on the morphology of sperm from *S.*  
61 *collinsi* is still limited. The seminal analysis or spermogram is an important tool to evaluate  
62 and determine the sperm's morphological aspects that are directly related to the potential male  
63 fertility (Visco et al., 2010) and is crucial for the processes of artificial insemination, in vitro  
64 fertilization, and embryo development (Ozkavukcu et al., 2008).

65 At a routine basis, sperm morphology from neotropical primates is examined in semen  
66 smears with the main criteria for normality relying on morphological parameters of the sperm  
67 head, middle-piece and flagellum (Arakaki et al., 2017, 2018; Sampaio et al., 2017; Swanson  
68 et al., 2016; Oliveira et al., 2015, 2016ab; Leão et al., 2015). In *S. collinsi*, the sperm  
69 morphology was only described using eosin–nigrosine stain by optical microscopy (Oliveira  
70 et al., 2015; 2016ab; Sampaio et al., 2017). However, sperm analysis using optical  
71 microscopy has low resolution, and does not allow the identification and analysis with more  
72 details than the micro-anatomical aspect that can affect the sperm and decrease the sperm  
73 capacity of fertilizing the oocyte (Visco et al., 2010).

74 In this context, the scanning- and transmission electron microscopy are technologies  
75 providing a high-range view, in nanometric dimensions, that can aid the morphological  
76 description and the detection of possible abnormalities in the sperm (Nussdorfer et al., 2018).  
77 These techniques have already been used in domestic animals as horses (Pesch et al., 2006),

78 sheep (López-Armengol et al., 2012), and wild animals as six-banded armadillo (Sousa et al.,  
79 2013), collared peccaries (Bezerra et al., 2018), jaguar (Silva et al., 2019) and some non-  
80 human primates (Bedford & Nicander et al., 1971; Martin et al., 1975; Gould, 1980; Steinberg  
81 et al., 2009; Nakazato et al., 2015). Regarding these microscopy technologies in the genus  
82 *Saimiri*, only the scanning electron microscopy was used to characterize the sperm  
83 morphology in squirrel monkeys (*S. sciureus*) (Martin et al., 1975; Gould, 1980), or obtained  
84 morphometric sperm data as a tool for more accurate taxonomic identification of the *S.*  
85 *boliviensis* (Steinberg et al., 2009). In *S. collinsi* nothing is known about the sperm  
86 components in higher resolution. Thus, the aim of the present study is to describe the micro  
87 morphological and ultrastructural characteristics of squirrel monkeys (*S. collinsi*) sperm using  
88 scanning- and transmission electron microscopy.

89

## 90 **Material and Methods**

### 91 **Ethical aspects and place of execution**

92 This study was approved by the Ethical Committee in Animal Research (no.  
93 02/2015/CEPAN/IEC/SVS/MS) and by the System of Authorization and Information in  
94 Biodiversity (SISBIO/ICMBio/MMA no.47051-2). All procedures were performed under the  
95 supervision of a veterinarian. Semen collections were carried out at the National Primate  
96 Center (CENP, Ananindeua, Pará, Brazil), the scanning electron microscopy was conducted at  
97 the Analytical Center of the Universidade Federal do Ceará (UFC, Fortaleza, Ceará, Brazil),  
98 and transmission electron microscopy was conducted at the Laboratory of Electron  
99 Microscopy of the Evandro Chagas Institute (IEC, Belém, Pará, Brazil).

### 100 **Animals and semen collection**

101 *Saimiri collinsi* males originated from the Marajó Archipelago (0°58'S and 49°34'W)  
102 and maintained in captivity at the CENP (1°22'58"S and 48°22'51"W) were used for collecting

103 semen. The experimental group (n=5 males, ~15 years old) was selected by their physical  
104 characteristics, and clinical parameters such as complete hemogram, hepatic and renal  
105 function.

106 Animals were collectively housed in mixed groups (males and females in a varied  
107 number of members) in cages of 4.74 m x 1.45 m x 2.26 m (length, width and height,  
108 respectively), under natural photoperiod (i.e. 12 h of light and 12 h of dark). The climate is  
109 humid and tropical, with an average annual temperature of 28°C. The diet consisted of fresh  
110 fruits, vegetables, commercial pellet chow specific for neotropical non-human primates  
111 (Megazoo®P18, 18% protein, 6.5% fibre, Brazil) and cricket larvae (*Zophobas morio*).  
112 Vitamins, minerals and eggs were supplied once a week, and water was available ad libitum.

### 113 **Semen collection**

114 Semen was collected at the same period of the day (in the morning before feeding).  
115 Physical restraint was performed by a trained animal caretaker wearing leather gloves. All  
116 animals were anesthetized with ketamine hydrochloride (20 mg/kg; IM; Vetanarcol, König  
117 S.A., Avellaneda, Argentina) and xylazine hydrochloride 1 mg/kg; IM; Kensol, König S.A.),  
118 and monitored by a veterinarian.

119 Achieved total anaesthetic effect, the animals were placed in lateral recumbency,  
120 genital region was then sanitized with a mild soap and distilled water (1:10) and gauze. The  
121 prepuce was retracted with the thumb and index fingers for a more efficient cleaning of the  
122 penis with saline solution. Animals were stimulated with rectal electroejaculation (EEJ)  
123 procedure described by Oliveira et al. (2015). Ejaculates (liquid and coagulated fractions)  
124 were collected into microtubes (1.5 mL), then, were placed in a water bath at 37°C. Scanning  
125 and transmission electron microscopy were only performed in the liquid fraction of the  
126 ejaculate.

127

**128 Sperm preparation**

129           The sample (liquid fraction) were centrifuged at 500g/5 min, and the supernatant was  
130 discarded. The pellet formed was washed three times in sodium cacodylate and fixed in  
131 Karnovsky (4% paraformaldehyde, 2.5% glutaraldehyde in 0.1M sodium cacodylate buffer,  
132 pH 7.2).

**133 Semen evaluation**

134           Seminal volume (liquid fractions) was evaluated in a graduated tube, with the aid of a  
135 pipette. Appearance and consistency were assessed subjectively, i.e., colour (colourless,  
136 yellowish, or whitish) and opacity (opaque or transparent) (Oliveira et al., 2015). Sperm  
137 motility, vigour, and morphology were evaluated according to Oliveira et al. (2015; 2016ab).  
138 Sperm morphology and plasma membrane integrity were evaluated by a smear prepared  
139 adding 5  $\mu$ L eosin 1% (Vetec) and 5  $\mu$ L nigrosine 1% (Vetec) to 5  $\mu$ L of semen on a pre-  
140 warmed (37°C) glass slide. Sperm concentration was determined in a Neubauer chamber after  
141 dilution of 1  $\mu$ L semen in 99  $\mu$ L formalin solution 10%. Morphologic defects detected in  
142 sperm were classified as primary or secondary according Bloom (1973). Plasma membrane  
143 functionality was assessed by hypoosmotic swelling test (HOST) after dilution of 5  $\mu$ L of  
144 semen in 45  $\mu$ L of hypoosmotic solution (0.73 g sodium citrate, 1.35 g fructose and 100 mL  
145 ultrapure water; pH 7.2 and 108 mOsm/L). After 45 min incubation in water bath (37 °C),  
146 number of sperm tail coiled was assessed by placing 10 $\mu$ L of this solution in a pre-warmed  
147 (37 °C) glass slide with cover slip, and at least 200 spermatozoa were counted. Spermatozoa  
148 with functional plasma membrane were those presenting coiled tail. All evaluations were  
149 performed under a light microscope (Nikon E400, Japan) at a magnification of 100 $\times$ .

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153 **Scanning electron microscopy**

154 Sperm samples were placed in a coated coverslip with poly-D-lysine, air-dried it onto  
155 a filter and dehydrated in a series of acetone solution (50, 70, 90 and 100% x 3, 5 min each).  
156 Afterwards, the samples were dried by the critical point method (EMS 850, Quorum  
157 Technologies, Lewes, United Kingdom) and then attached to supports using aluminium stubs  
158 with carbon tape. The samples were covered with a thin (20 nm) layer of gold (Q150T ES,  
159 Quorum Technologies, Lewes, United Kingdom) and observed under scanning electron  
160 microscopy (Quanta 450-FEG, Thermo Fisher Scientific, Massachusetts, EUA). The  
161 description of *S. collinsi* sperm by SEM was conducted according to that previously reported  
162 for non-humans primates by Martin et al. (1975).

163 **Transmission electron microscopy**

164 For transmission electron microscopy, the sperm sample were washed in 0.05 M  
165 cacodylate buffer (40 min/3x), post-fixed with osmium tetroxide (OsO<sub>4</sub>) for 60 min and  
166 washed in 0.05 M cacodylate buffer with 0.08% potassium ferrocyanide (40 min/5x).  
167 Subsequently, the contracting block was performed with 2.5% uranyl acetate in 50% acetone  
168 for 1h. Then, the sample were dehydrated in series of acetone solution (70, 80, 90 and 100%)  
169 for 10 min each, the solution of 100% acetone was carried out 3 times. After dehydration, the  
170 samples were infiltrated with epoxy resin (EponPolibed), using increasing series of resin in  
171 acetone (dilution 1:2; 1:1; 2:1), until 100% Epon + DMP-30 (2,4,6-Trisdimethylaminomethyl)  
172 phenol). Polymerization of the resin was performed at 60 °C/ 48h. Semi-thin cuts of  
173 approximately 70 nm thickness were obtained in ultramicrotome with glass cutters (Leica EM  
174 UC7, Leica Microsystems, Wetzlar, Germany) and contrasted with 5% uranyl acetate for 20  
175 min in 60 °C. The observations of the sperm ultrastructure's were performed in transmission  
176 electron microscopy (EM 900, Carl Zeiss do Brasil Ltda, São Paulo, Brazil).

177

## 178 **Results**

### 179 **Semen evaluation**

180 Mean ( $\pm$ SEM) of seminal volume was  $339 \pm 61 \mu\text{L}$  (15–500  $\mu\text{L}$ ; min–max). liquid  
181 fractions were whitish or yellowish, and opaque. The sperm concentration was  $1.718 \pm 610 \times$   
182  $10^6$  sperm/ml. In addition, the samples presented a mean of,  $67 \pm 11\%$  motility,  $3 \pm 0.3$   
183 vigour,  $38 \pm 13\%$  and  $43 \pm 8\%$  functional and intact sperm membranes, respectively. The  
184 evaluation of sperm morphology through light microscopy was  $56 \pm 5\%$  normal sperm,  $13 \pm 4$   
185 bent tail,  $27 \pm 4$  coiled tail and  $3.6 \pm 1.2$  strongly coiled tail.

### 186 **Scanning electron microscopy**

187 The scanning electron microscopy (SEM) revealed that the head of *S. collinsi* sperm is  
188 paddle-shaped flattened with a uniform thickness throughout; measuring  $5.14 \pm 0.08 \mu\text{m}$  in  
189 length and  $3.36 \pm 0.03 \mu\text{m}$  in width, with an intact and uninterrupted surface, and a acrosome  
190 occupied most of the head with length of  $3.56 \pm 0.06 \mu\text{m}$  (Fig. 1A). The insertion of the tail is  
191 lateral (abaxial; Figure 1A), with a length of middle piece of  $9.49 \pm 0.38 \mu\text{m}$ . The squirrel  
192 monkeys' sperm presents a total length of  $71.7 \pm 0.7 \mu\text{m}$  (Fig. 1B).

### 193 **Transmission electron microscopy**

194 The transmission electron microscopy (TEM) images showed the *S. collinsi* sperm  
195 presents a flattened head in a longitudinal cut, containing a very electron-dense nucleus,  
196 which is larger at the base, narrowing as it goes to the tip of the head. There is a diffuse small  
197 electronic spot. The structures present in the sperm head are the nucleus, the acrosome, and  
198 membranous envelopes (Fig. 3A-B). The large nucleus covered by the acrosome (ac), which  
199 occupies one-half to two-thirds of the anterior portion of the head. The acrosome is thickest  
200 around the anterior margin of the sperm head, and thinnest from the equator to the posterior  
201 third of the nucleus where it terminates (Fig. 3B). The neck area (na), which is the region  
202 located between the head and the middle piece, is characterized by the presence of a centriole,

203 the proximal centriole (oriented transversally to the cell; pc), and segmented columns (Figure  
204 3B). The middle piece is characterized by a central 9 + 2 microtubule axoneme surrounded by  
205 nine dense fibres. This set is surrounded by a plasmalemma and a helical mitochondrial sheath  
206 (Fig. 3C-D), where was identified the presence of an intact axoneme as well as outer dense  
207 fibres (ODFs). The mitochondria were juxtaposed, thus forming the mitochondrial sheath,  
208 which was surrounded by the plasmalemma with some undulations (Figure 3E). It is possible  
209 to visualize about ~72 mitochondria.

210

## 211 **Discussion**

212 Our study represents a valuable basis for the comprehension of sperm physiology  
213 through the description of micro morphological and ultrastructural aspects in *S. collinsi*. Over  
214 the years, the evaluation of sperm morphology has become much more precise, and the  
215 rigorous analysis of these parameters is of pivotal importance in semen analysis, since it  
216 permits the detailed study of different sperm abnormalities that predict the fertile capacity of  
217 the male gamete (WHO, 2010).

218 In the genus *Saimiri*, the SEM was performed only in *S. sciureus* (Martin et al., 1975)  
219 and *S. boliviensis* (Stembeirg et al., 2009). In *S. collinsi*, the abaxial tail insertion was also  
220 observed by SEM, as previously described as a normal morphology characteristic in the  
221 morphometric analysis of sperm stained with eosin-nigrosine for *S. collinsi* and *S. vanzollini*  
222 (Sampaio et al., 2017). The average total length of the *S. collinsi* sperm was similar described  
223 for other squirrel monkeys species in the stained samples (Nakazato et al., 2015; Sampaio et  
224 al., 2017). The morphology of sperm head showed a thick, paddle-shaped flattened, which  
225 looks similar to other primates, including humans (Martin et al., 1975; Gould & Martin,  
226 1978). The measurements of head morphology using SEM agree with values obtained  
227 previously via optical microscopy for *S. collinsi* (Sampaio et al., 2017).

228 For the first time, the ultrastructure description of *S. collinsi* sperm by TEM was  
229 reported. Thus, the results presented here will serve as a parameter for future morphological  
230 studies involving the preparation and evaluation of the semen in nonhuman primates, as well  
231 as for studies using morphological data for the diagnosis of sperm alterations, including  
232 acquired damages post-thawing, which is a relevant factor for the development of  
233 reproduction (reproductive )biotechnology (Sousa et al., 2013). The TEM might be an  
234 additional diagnostic tool in the presence of asthenozoospermia or the absence of motility,  
235 which is important since the pattern of axoneme structure can be visualized in the longitudinal  
236 and cross sections with this technique. In humans, TEM have been used in the diagnosis of  
237 sperm alterations with a possible genetic origin, known as systematic defects (Baccetti et al.,  
238 2001). The observation of sperm head ultrastructure revealed a large and obvious nucleus.  
239 The acrosome is the only cytoplasmic element in the sperm head. Similar as described for  
240 human sperm, in *S. collinsi*, the acrosome is a relatively inconspicuous, which covers the  
241 anterior two-thirds to half of the head, caplike structure containing enzymes that are essential  
242 for fertilization (Bartoov et al., 1980). In Rhesus monkeys sperm, the acrosome is thickest  
243 around the anterior margin of the sperm head and thinnest from the equator to the posterior  
244 third of the nucleus where it terminates, and the thin portion of the acrosome is usually  
245 referred to as the equatorial segment. As was shown in human and Rhesus monkey (Zamboni  
246 et al., 1971), the plasma membrane surrounding the anterior portion of the head, is continuous  
247 and often undulated in *S. collinsi*.

248 The sperm head and tail are connected via the sperm neck, a region formed by the  
249 centriole and connecting piece (Bornens, 2012). In Rhesus macaque, middle piece sheath  
250 consists of 84 to 86 mitochondria (Zamboni et al., 1971). In our study, the exact number of  
251 mitochondria in *S. collinsi* could not be measured due to the absence of longitudinal sections  
252 showing the middle piece to its full extent. These organelles function to provide energy for



253 the movement of the flagella, since the midpiece accommodates more mitochondria to  
254 produce more energy in order to fuel motility (Pesch et al., 2006), and generates greater  
255 propulsion forces in the principal piece (Gomendio & Roldan, 1991). However, it was  
256 demonstrated that the swimming speed is not dependent on the size of the tail, but on the head  
257 size, which in return may retard the propulsion given by the sperm tail (Humphries et al.,  
258 2008).

259       Regarding morphological defects in light microscopy, according to Bloom (1973)  
260 classification, some secondary defects were identified in the *S. collinsi* sperm, including bent  
261 tail, coiled tail and strongly coiled tail. Such defects were also verified in the ejaculates of  
262 neotropical primates (Arakaki et al., 2017; 2018; Leão et al., 2015; Oliveira et al., 2015;  
263 Swanson et al., 2016). In our study compared to others already performed for the *S. collinsi*  
264 species (Oliveira et al., 2015; 2016ab; Sampaio et al., 2017), we observed a high percentage  
265 of sperm pathologies, which for Sampaio et al. (2017) is characterized as a group of animals  
266 with low seminal quality. On the other hand, it is noteworthy that we only worked with the  
267 liquid fraction for sperm characterization, since the dilution process could imply changes in  
268 sperm morphology. It is known that the liquid fraction is the fraction of lower seminal volume  
269 and sperm concentration (Oliveira et al., 2015), which may also contain sperm with higher  
270 percentage of sperm pathologies as compared with coagulated fraction.

271       *Saimiri collinsi* ejaculates present a normal range for sperm morphology similar to that  
272 accepted for the genus *Saimiri*. Additionally, they present a general sperm structure, but with  
273 peculiarities such as abaxial middle piece insertion showed by SEM.

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278 **Conflict of interest statement**

279 The authors declare that there is no conflict of interest that can be perceived as  
280 prejudicing the impartiality of the research reported.

281

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289

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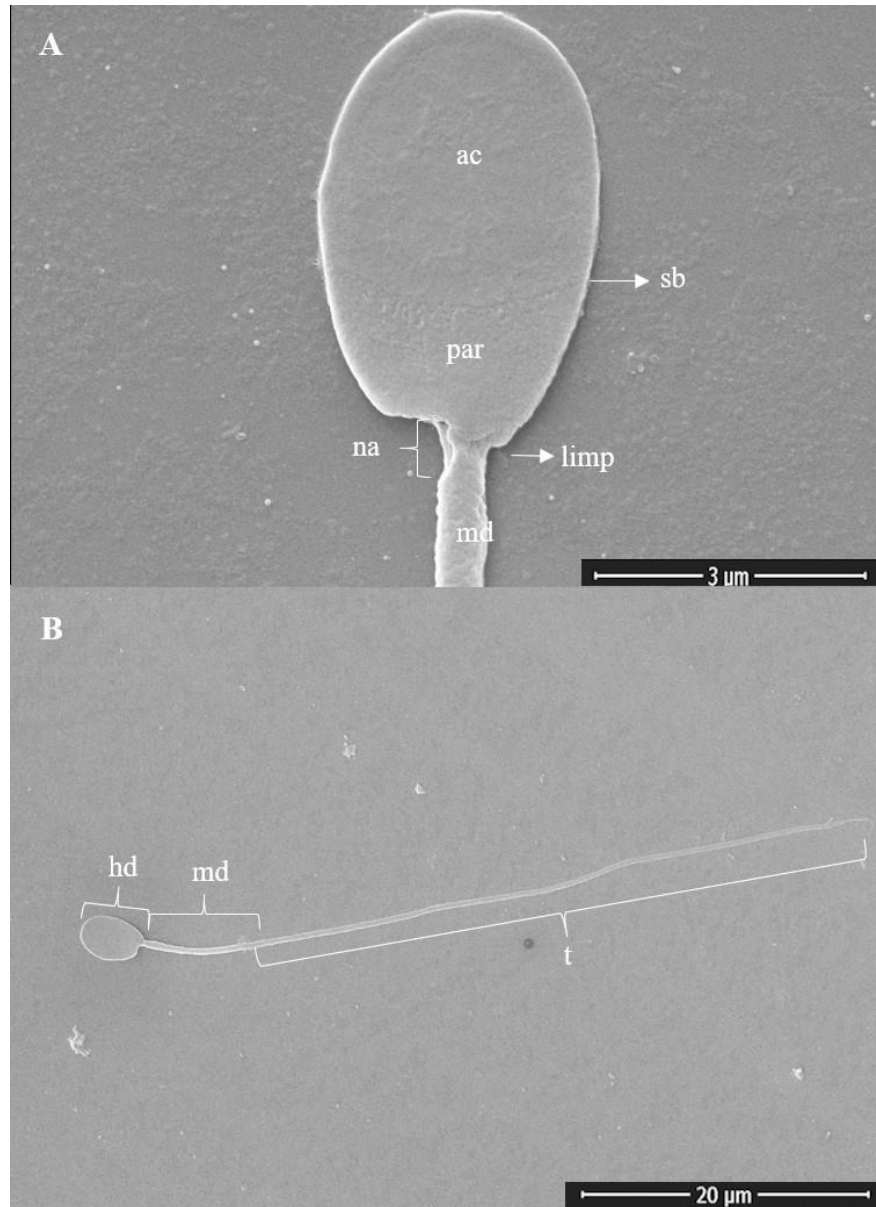
400 **Figure captions**

401

402 **Fig. 1:** Scanning electron micrograph of *Saimiri collinsi* sperm: A - Sperm head with plasma  
403 membrane integrity, acrossomal region (ac), serrated band (sb); post-acrosomal region (par),  
404 neck area (na), lateral insertion of the middle piece (limp) and middle piece (md); B -Normal  
405 sperm: head (hd), middle piece (md) and tail (t).

406

407 **Fig. 2:** Transmission electron micrograph of the squirrel monkey (*Saimiri collinsi*) sperm: (A)  
408 hd: head; md: middle piece; an: annulus; t: tail. (B) oam: outer acrosome membrane; ac:  
409 acrosome; pl: plasmalemma; nu: nucleus; nm: nuclear membrane; pas: postacrosomal sheath;  
410 pc: principal centriole; na: neck area; md: middle piece. (C) pl: plasmalemma; nm: nuclear  
411 membrane; hd: head; na: neck area; md: middle piece; ax: axonema; ofd: outer dense fibers;  
412 ms: mitochondrial spiral. (D) pl: plasmalemma; ax: axonema; ofd: outer dense fibers; ms:  
413 mitochondrial spiral; \*: longitudinal cut. (E) ofd: outer dense fibers; ax: axonema; pl:  
414 plasmalemma, ms: mitochondrial spiral.



415

416 Fig. 1: Scanning electron micrograph of *Saimiri collinsi* sperm: A - Sperm head with plasma  
 417 membrane integrity, acrossomal region (ac), serrated band (sb); post-acrosomal region (par),  
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 419 sperm: head (hd), middle piece (md) and tail (t).

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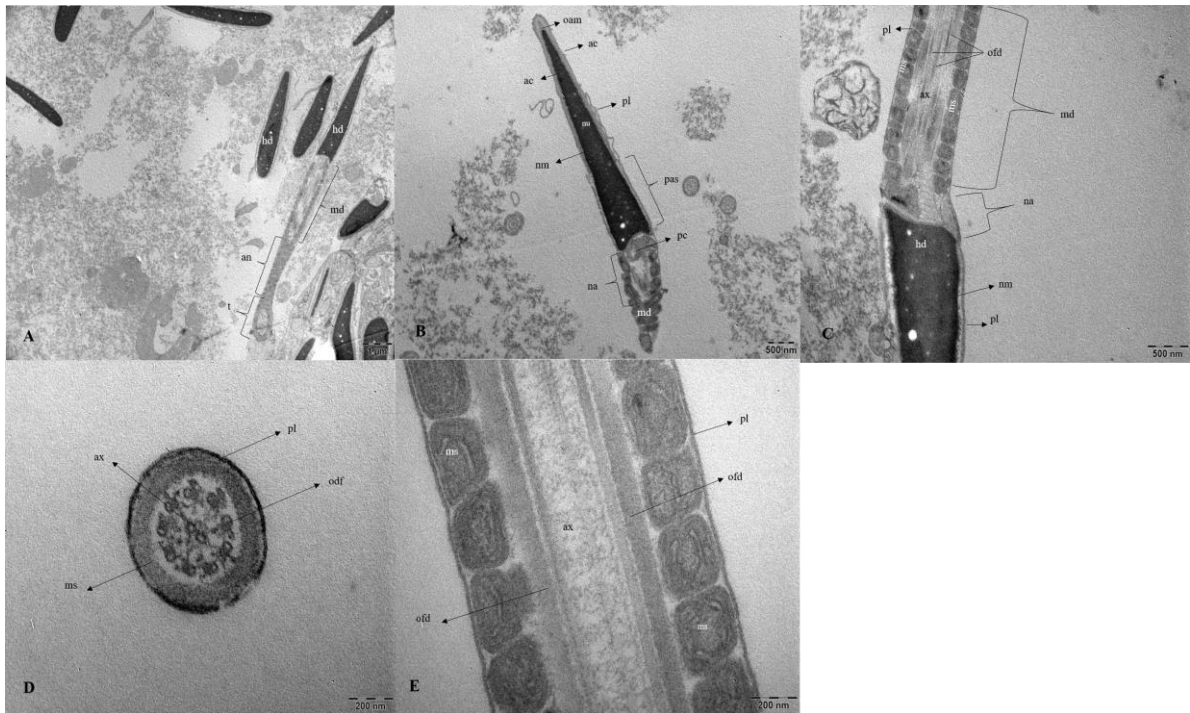
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63x87mm (300 x 300 DPI)

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425 Fig. 2: Transmission electron micrograph of the squirrel monkey (*Saimiri collinsi*) sperm: (A)  
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 427 acrosome; pl: plasmalemma; nu: nucleus; nm: nuclear membrane; pas: postacrosomal sheath;  
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 429 membrane; hd: head; na: neck area; md: middle piece; ax: axonema; odf: outer dense fibers;  
 430 ms: mitochondrial spiral. (D) pl: plasmalemma; ax: axonema; odf: outer dense fibers; ms:  
 431 mitochondrial spiral; \*: longitudinal cut. (E) odf: outer dense fibers; ax: axonema; pl:  
 432 plasmalemma, ms: mitochondrial spiral.

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174x93mm (300 x 300 DPI)

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1           **GLOBAL PROTEOMIC ANALYSIS OF SPERMATOZOA FROM CAPTIVE**  
 2           **AMAZON SQUIRREL MONKEYS (*Saimiri collinsi* Osgood, 1916)**

3  
 4           Danuza Leite Leão<sup>ab\*1</sup>, Sheyla Farhayldes Souza Domingues<sup>a</sup>, Patrícia da Cunha  
 5           Sousa<sup>a</sup>, Wlaises Vasconcelos Sampaio<sup>ab</sup>, Fábio Roger Vasconcelos<sup>c</sup>, Arlindo Alencar Moura<sup>c</sup>,  
 6           Regiane Rodrigues dos Santos<sup>a</sup>, Morten Skaugen<sup>d</sup>, Irma Caroline Oskam<sup>e</sup>

7  
 8                           <sup>a</sup>*Laboratory of Wild Animal Biology and Medicine, Faculty of Veterinary*  
 9                           *Medicine, Federal University of Pará, Castanhal, Pará, Brazil, CEP 68746-360.*

10                           *wlaises.sampaio@gmail.com, pattbio13@hotmail.com, danleao.88@gmail.com,*  
 11   *shfarha@gmail.com*

12                           <sup>b</sup>*Postgraduate Program in Animal Health and Production in the Amazon, Federal*  
 13                           *Rural University of Amazonia, Belém, Pará, Brazil, CEP 66077-530.*

14   *wlaises.sampaio@gmail.com, danleao.88@gmail.com.*

15                           <sup>c</sup>*Laboratory of Animal Physiology, Department of Animal Science, Federal*  
 16                           *University of Ceará, Fortaleza, Ceará, Brazil, CEP 60021-970. arlindo.moura@gmail.com,*

17   *fr.vasconcelos@yahoo.com.br.*

18                           <sup>d</sup>*Faculty of Chemistry, Biotechnology and Food Science, Norwegian University of*  
 19                           *Life Sciences, Ås, Norway, P.O. Box 5003. morten.skaugen@nmbu.no.*

20                           <sup>e</sup>*The Animal Production Experimental Center, Norwegian University of Life*  
 21                           *Sciences, Ås, Norway, P.O. Box 5003. irma.caroline.oskam@nmbu.no.*

22  
 23           \*Corresponding author: Danuza Leite Leão

24           <sup>1</sup>Present Address: Laboratory of Wild Animal Biotechnology and Medicine, Federal  
 25           University of Pará, BR 316 Km 61, CEP 68740–970, Castanhal, Pará, Brazil.

26           Email: danleao.88@gmail.com

27 **Abstract**

28           The aims of this study were to describe the global sperm proteomics in *Saimiri*  
29 *collinsi*, a Neotropical primate endemic to the Amazon in Brazil, which is used as a biological  
30 model for reproductive research on the genus *Saimiri*. The study approach based on bottom-up  
31 proteomics allowed the identification of 2,343 proteins present in the sperm samples. Of the 39  
32 proteins that were related to spermatogenesis, sperm motility, capacitation, fecundation, and  
33 defense systems against oxidative stress were upregulated in the dry season. Knowledge on the  
34 sperm proteins provides crucial information for elucidating the underlying mechanisms  
35 associated with sperm functionality. Thus, our results help to advance our understanding of the  
36 reproductive physiology of *S. collinsi*, providing valuable information for the improvement of  
37 protocols used in assisted reproduction techniques for the conservation of endangered *Saimiri*  
38 species.

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## 56 1. Introduction

57 Mammalian male fertility depends on physiological events that begin with  
58 spermatogenesis and culminate with successful adhesion/signaling between the sperm  
59 membrane and the extracellular coat of the oocyte, followed by adhesion/fusion between the  
60 oocyte and sperm membranes during fertilization in the female reproductive tract [1, 2].  
61 Proteins expressed by spermatozoa and those from the seminal plasma that bind to the sperm  
62 plasma membrane render the spermatozoa capable of fertilizing a mature oocyte [3, 4].

63 Studies in animals and humans have described sperm proteins that have significant  
64 associations with sperm motility (i.e., L-lactate dehydrogenase and dynein heavy chain 1  
65 (DNAH1)) [5, 6], sperm capacitation (i.e., clusterin, spermadhesin, and mitochondrial  
66 peroxiredoxin-5) [7, 8], and fertility (i.e., enolase 1, ropporin- 1-like protein (ROPN1), and  
67 Izumo sperm–egg fusion 1 (IZUMO1)) [9, 10]. In non-human primates, sperm proteomics has  
68 been carried out only in Old World primates, in the genus *Macaca*, for characterization of the  
69 sperm protein profile [3, 11-15]. Knowledge about the absence, presence, underexpression, or  
70 overexpression of these sperm proteins could help to further our understanding of the  
71 mechanisms behind the reduction in the fertilization ability of sperm [4, 16].

72 The squirrel monkey (*Saimiri collinsi*), a Neotropical primate endemic to the Amazon  
73 in Brazil [17], was used as an experimental model for reproductive research on the genus  
74 *Saimiri* [18-20]. According to the International Union for Conservation of Nature's Red List of  
75 Threatened Species, two *Saimiri* species are ranked as vulnerable (*Saimiri oerstedii* and  
76 *Saimiri vanzolini*) and one species as almost threatened (*Saimiri ustus*) to extinction [21].  
77 Defining the sperm protein profiles of *Saimiri collinsi* may provide us with a better  
78 understanding about the reproductive physiology of these animals, as well as whether the  
79 sperm cells could be used in assisted reproduction techniques throughout. Therefore, the aims  
80 of this study were to describe the global sperm proteomics in *S. collinsi* and evaluate the  
81 potential correlation between the expression of the sperm proteins and the seminal quality in *S.*  
82 *collinsi*.

83

84

## 85 **2. Material and methods**

### 86 *2.1 Study design*

87 We conducted a global proteomic analysis of spermatozoa collected from adult squirrel  
88 monkeys (*S. collinsi*), in the Brazilian Amazon. The seminal coagulum was collected by  
89 electroejaculation and liquefied in a powdered coconut water extender (ACP-118; ACP  
90 Biotecnologia, Fortaleza, Ceará, Brazil). After 1 h in the ACP-118 extender, the viable sperm  
91 cells were separated on Percoll density gradient media and washed. Then, the sperm proteins  
92 were extracted and subjected to tryptic digestion, followed by liquid chromatography-tandem  
93 mass spectrometry. Computational biology was used for the identification of the proteins,  
94 categorization of the proteins, and *in silico* analysis of the protein network.

95

### 96 *2.2 Animal ethics statement and study location*

97 The animal study was approved by the Ethical Committee in Animal Research  
98 (Approval No. 02/2015/CEPAN/IEC/SVS/MS) and by the System of Authorization and  
99 Information in Biodiversity (SISBIO/ICMBio/MMA No. 47051-2), and carried the license of  
100 the Convention on International Trade in Endangered Species of Wild Fauna and Flora  
101 (CITES/IBAMA/Permit No. 17BR025045-DF). All procedures were performed under the  
102 supervision of a veterinarian.

103 Semen collections were performed at the National Primate Center (CENP, Ananindeua,  
104 Pará, Brazil) and the characterization of the protein profile was performed in the Faculty of  
105 Chemistry, Biotechnology and Food Science of Norwegian University of Life  
106 Sciences (NMBU, Ås, Norway).

107

### 108 *2.3 Animals and housing conditions*

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110 *S. collinsi* males (n = 4) maintained in captivity at the Centro Nacional de Primatas,  
111 Brazil (1°22'58"S and 48°22'51"W) were used for the semen collection. The animals were  
112 housed collectively in cages (4.74 m × 1.45 m × 2.26 m), with 12 h of natural light each day.  
113 The mixed animal groups typically consisted of three males and three females and their

114 juvenile offspring. The animals were fed fresh fruits, vegetables, commercial pellet chow  
115 specific for Neotropical non-human primates (18% protein, 6.5% fiber; Megazoo, Minas  
116 Gerais, Brazil), and cricket larvae (*Zophobas morio*). Vitamins, minerals, and eggs were  
117 supplied once a week, and water was available *ad libitum*.

118

#### 119 *2.4 Body weight, testicular biometry, and semen collection*

120 Semen was collected every morning before feeding, making up a total of 48 semen  
121 collections. For the semen collection, physical restraint of each animal was performed by a  
122 trained animal caretaker wearing leather gloves, and all animals were anesthetized with  
123 ketamine hydrochloride (20 mg/kg; intramuscularly (IM); Vetanarcol, König S.A.,  
124 Avellaneda, Argentina) and xylazine hydrochloride (1 mg/kg; IM; Kensol, König S.A.) and  
125 monitored by a veterinarian. After anesthesia, the animals were weighed using a weight  
126 balance, and the testicular length, width, height, and circumference were measured using a  
127 universal caliper. The testicular volume was calculated according to the method described by  
128 Oliveira et al. [20]. After the animal had been placed in dorsal recumbency, the genital region  
129 was sanitized with a mild soap and distilled water (1:10) and the prepuce was retracted for a  
130 more efficient cleaning of the penis with saline solution. The animal was then stimulated  
131 according to the rectal electroejaculation procedure described by Oliveira et al. [18-20]. In  
132 brief, an electroejaculator (Autojac-Neovet, Uberaba, Brazil) rectal probe was smeared with a  
133 sterile lubricant gel (KY Jelly, Johnson & Johnson Co., Arlington, TX, USA) and introduced  
134 into the rectum (~2.5 cm deep) and electrical stimuli were then delivered. The stimulation  
135 session consisted of three series (7 and 8 min), composed of 35 electrical stimuli (12.5 and 100  
136 mA), with an interval of 30 s between the series. The ejaculates (liquid and coagulated  
137 fractions) were collected into microtubes (1.5 mL).

138

#### 139 *2.5 Semen evaluation*

140 The 1.5-mL conical microtubes containing the semen were placed in a water bath at  
141 37°C immediately after collection for evaluation of the seminal volume, color, and viscosity.  
142 The semen appearance was assessed subjectively for color (colorless, yellowish, or whitish)

143 and opacity (opaque or transparent) [18-20]. The sperm motility, vigor, and morphology were  
144 evaluated according to the methods described by Oliveira et al. [18-20]. For evaluation of the  
145 normal sperm morphology and plasma membrane integrity, a smear sample was prepared by  
146 adding 5  $\mu$ L of 1% eosin (Vetec, Rio de Janeiro, Brazil) and 5  $\mu$ L of 1% nigrosine (Vetec, Rio  
147 de Janeiro, Brazil) to 5  $\mu$ L of semen on a prewarmed (37°C) glass slide. The plasma  
148 membrane functionality was assessed with the hypoosmotic swelling test after the dilution of 5  
149  $\mu$ L of semen in 45  $\mu$ L of hypoosmotic solution (0.73 g of sodium citrate, 1.35 g of fructose,  
150 and 100 mL of ultrapure water; pH 7.2 and 108 mOsm/L). After a 45-min incubation in a  
151 water bath (37°C), 10  $\mu$ L of this solution was placed on a prewarmed (37°C) glass slide and  
152 covered with a coverslip, and at least 200 spermatozoa were counted to determine the number  
153 with coiled tails (indicative of spermatozoa with a functional plasma membrane). All  
154 evaluations were performed under a light microscope (E400; Nikon, Tokyo, Japan) at a  
155 magnification of 100 $\times$ . The semen was assessed after dilution in ACP-118.

156

#### 157 *2.6 Sperm separation and freeze-drying*

158 Owing to the occurrence of seminal coagulation in *S. collinsi*, the semen sample was  
159 diluted 1:1 in ACP-118 (300 mOsm/kg and pH 6.42), incubated in a water bath (Biomatic,  
160 Porto Alegre, Rio Grande do Sul, Brazil) at 37°C for 1 h, and then separated on 45%/90%  
161 Percoll gradient media (centrifugation at 10,000 *g*, 15 min, 12°C). Thereafter, the samples  
162 were washed in Tris-NaCl medium (centrifugation at 8000 *g*, 5 min, 12°C), and the separated  
163 sperm fraction (pellet) was stored in microtubes, together with Tris-NaCl and a protease  
164 inhibitor (1:1000; P8340 catalog, Sigma-Aldrich, St. Louis, MO, USA), in liquid nitrogen or a  
165 -80°C freezer. For lyophilization, the frozen sperm samples were placed in a freeze dryer  
166 (FreeZone 2.5 Liter Benchtop Freeze Dry System; Labconco, Kansas City, MO, USA) for 10 h  
167 at a temperature of -55°C and vacuum pressure of 0.025 mbar.

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172 *2.7 Liquid chromatography-mass spectrometry*

173 Each individual dried sperm sample was resuspended in 50  $\mu$ L of lysis buffer (0.1 M  
174 Tris-Cl (pH 8.0), 4% sodium dodecyl sulfate, and 10 mM dithiothreitol) and centrifuged at  
175 5000 *g* for 1 h at 4°C. The supernatant was reserved for the preparation of suspension samples  
176 for bottom-up proteomic analysis with tryptic digestion, using the method established by  
177 Zougman et al. [22]. The extracted peptides were analyzed on an UltiMate 3000 RSLCnano/Q-  
178 Exactive system (Thermo Fisher Scientific, Bremen, Germany) that was set up with a  
179 Nanospray Flex ion source. The tryptic peptides (~1  $\mu$ g loaded) were separated on a 50 cm  $\times$   
180 75  $\mu$ m (i.d.) column (Thermo Fisher Scientific) using a 120 min gradient of 12–45%  
181 acetonitrile. The mass spectrometry (MS) and tandem mass spectrometry (MS/MS) data were  
182 recorded using a standard data-dependent acquisition method, with the following conditions:  
183 *m/z* range of 300–1600; Automatic Gain Control targets of  $3 \times 10^6$  (MS) and  $5 \times 10^4$  (MS/MS);  
184 resolutions of 70 K (MS) and 35 K (MS/MS); dynamic exclusion set to 20 s, and normalized  
185 collision energy set to 28. Xcalibur software (v. 3.1; Thermo Fisher Scientific) was used to  
186 evaluate the raw data, which were converted to *mgf* format (for Mascot database searching)  
187 using the MS convert module of ProteoWizard (v. 3.0.9016). The Mascot (v. 2.6) searches  
188 were performed on an in-house server against an online *Saimiri boliviensis boliviensis*  
189 (Bolivian squirrel monkey) database (National Center for Biotechnology Information,  
190 Bethesda, MD, USA). MaxQuant software (v.1.6.1.0) [23] was used for the label-free  
191 quantification.

192

193 *2.8 Protein categorization*

194 The protein information obtained by Mascot was analyzed using the STRuctural  
195 Analysis Programs (STRAP) for searching annotations of proteins. STRAP automatically  
196 obtains Gene Ontology (GO) terms associated with proteins in an identification list of results  
197 based on homology search analysis using various freely accessible databases [24].

198

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200



### 201 2.9 *In silico* protein network analysis

202 Protein–protein networks were retrieved from the STRING database (v. 10.0), which  
203 consists of known and predicted protein interactions collected from direct (physical) and  
204 indirect (functional) associations. The database quantitatively integrates interaction data from  
205 four sources: a genomic context, high-throughput experiments, co-expression data, and  
206 previous knowledge from research publications [25]. The STRING program was set to show  
207 no more than 10 interactions and medium confidence. Pathways not described for *S.*  
208 *boliviensis boliviensis* were analyzed using those for other non-human primate species and  
209 *Homo sapiens*.

210

211

### 212 3. Results

213 Mean ( $\pm$  SE) body weight was  $853.7 \pm 8.6$  g (762 – 1,022 g; min–max), and testicular  
214 width, length, height, and volume were  $1.12 \pm 0.1$  (0.8 – 1.8 cm; min-max),  $1.7 \pm 0.2$  (0.8 – 2.2  
215 cm, min-max),  $1.2 \pm 0.4$  cm (0.7 – 2.2 cm, min-max) and  $2.2 \pm 0.1$  ( $1.08 - 4.1$  cm<sup>3</sup>, min-max),  
216 respectively, without significant differences between the left and right testis. Semen collection  
217 was successful in 42 of the 48 attempts (88%); of these, 39 samples were used for the  
218 experiments, because three ejaculates did not contain sperm. The highest percentage of  
219 ejaculates in both the liquid and coagulated fractions was 59%. With regard to the semen color  
220 and opacity, 10% of the samples were colorless, 33% were whitish, 57% were yellowish, 46%  
221 were transparent, and 54% were opaque. The seminal parameters of *Saimiri collinisi* were  
222 represented in Table 1.

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230 Table 1

231

	Mean ( $\pm$ SEM)
<b>Motility</b>	45 $\pm$ 5.5 <sup>b</sup>
<b>Vigor</b>	2 $\pm$ 0.2 <sup>b</sup>
<b>PMF</b>	65.4 $\pm$ 4.1
<b>PMI</b>	71.1 $\pm$ 3.4
<b>Normal sperms</b>	73.1 $\pm$ 2.6

232 Mean ( $\pm$ SEM) values of sperm motility (%),, vigor, sperm plasma membrane functionality  
 233 (PMF; %), sperm plasma membrane integrity (PMI; %), and normal sperm morphology (%) of  
 234 diluted semen in *Saimiri collinsi*.

235

236 The approach based on bottom-up proteomics allowed the identification of 2,343  
 237 proteins in sperm samples after seminal coagulation dilution in ACP-118<sup>®</sup> extender  
 238 (Supplemental Table 1). Of the total proteins identified, 223 were determined to participate in  
 239 important reproductive events, such as spermatogenesis (67 proteins), sperm motility (42  
 240 proteins), capacitation/acrosome reaction (20 proteins), and fertilization (32 proteins)  
 241 (Supplemental Table 2).

242 On the basis of the GO analysis, the proteins were grouped according to biological  
 243 process, molecular function, and cellular component (i.e., localization) classes (Figure 1). In  
 244 the cellular component class, most of the proteins identified were associated with the  
 245 cytoplasm (12.3%), cytoskeleton (9.4%), and nucleus (8.9%) (Figure 1A). The most common  
 246 biological processes associated with the proteins were cellular processes (41.6%), regulation  
 247 (17.6%), and metabolic processes (11.4%) (Figure 1B). Binding (42.8%) and catalytic activity  
 248 (42.9%) corresponded to the most frequent molecular functions for the proteins (Figure 1C).

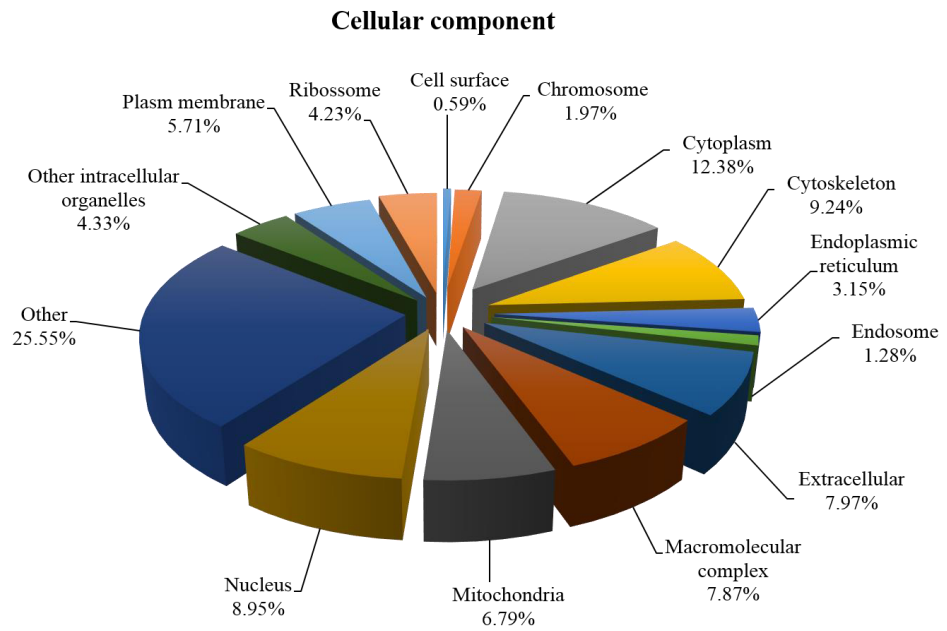
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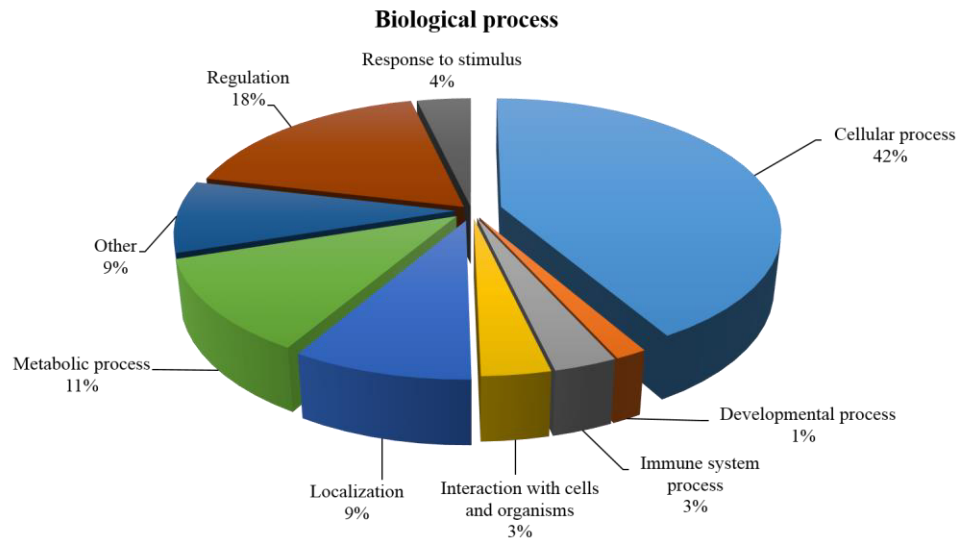
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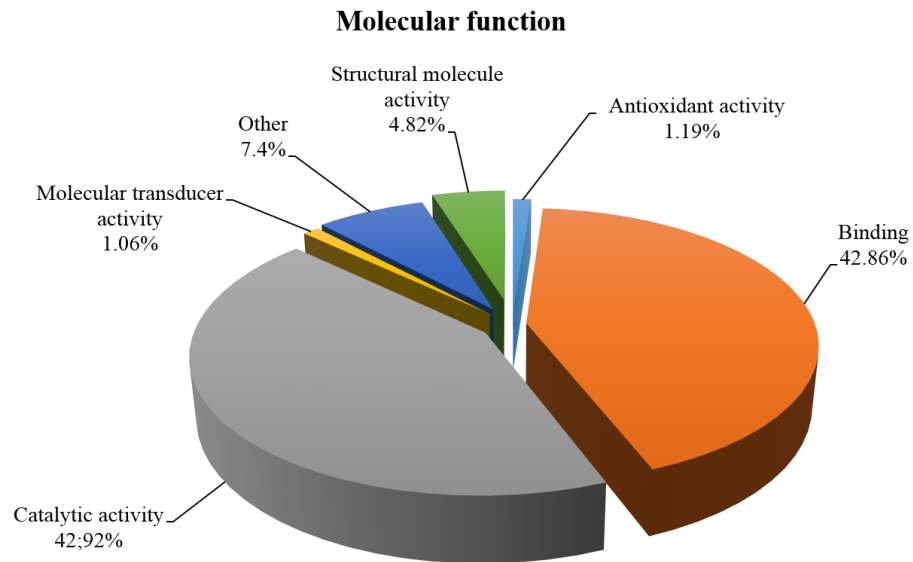
253 Figure 1



254



255



256

257 Cellular component, biological process and molecular function of identified proteins by  
 258 nanoLC-QExactive spectrometry analyzed by STRAP. Gene ontology terms were obtained  
 259 from the UniProtKB database.

260

261 In silico analysis indicated that semenogelin I, heat shock proteins members (60kDa,  
 262 70kDa and 90-alpha), sperm acrosome membrane-associated family member (3 and 4), and  
 263 izumo sperm-egg fusion family members (1 and 4) interacted with other 10 proteins. Among  
 264 these interactions, the heat shock proteins 60kDa and 70kDa proteins interacted with each  
 265 other and heat shock proteins 90-alpha, as well as izumo sperm-egg fusion 1 proteins also  
 266 interact with each and sperm acrosome membrane-associated protein 4. The sperm acrosome  
 267 membrane-associated protein 4 interacts with both sperm acrosome membrane-associated  
 268 protein 1 and 3 (Figure 02).

269

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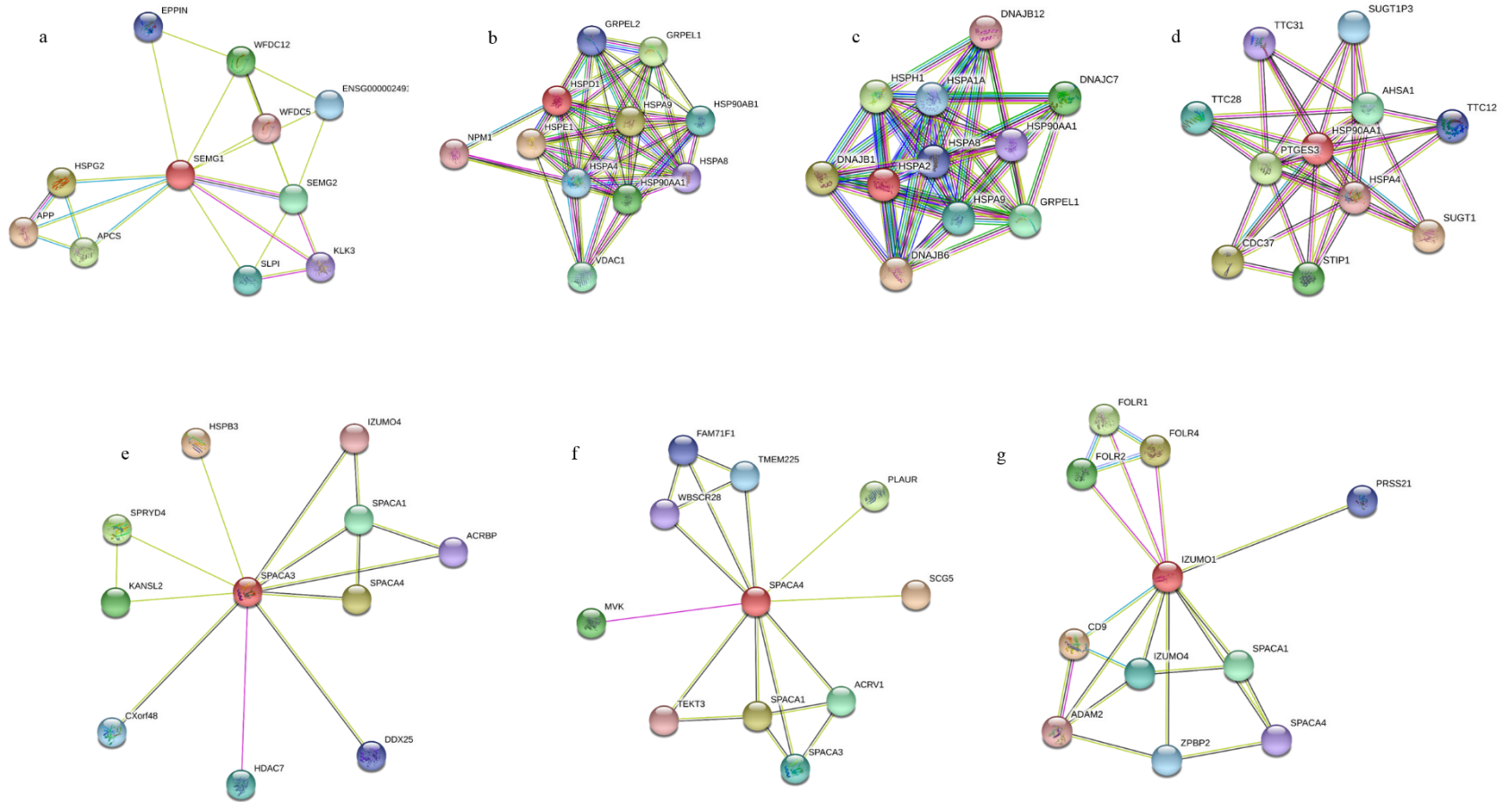
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Figure 2



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277 Protein interaction analysis. Proteins were analyzed with the web-based STRING software.  
 278 Analyzed proteins were: a- semenogelin I; b- 60kDa heat shock proteins; c- Heat shock proteins  
 279 70kDa protein 2; d- Heat shock proteins 90-alpha; e- Sperm acrosome membrane-associated 3; f-  
 280 Sperm acrosome membrane-associated 4; g- Izumo sperm-egg fusion protein 1. Different line  
 281 colors represent the types of evidence for the association. Yellow textming; black coexpression;  
 282 blue databases; and pink experiments. APP Amyloid beta (A4) precursor protein; HSPG2  
 283 Heparan sulfate proteoglycan 2; APCS Amyloid P component, serum; WFDC12 WAP four-  
 284 disulfide core domain 12; SEMG2 Semenogelin II; SLPI Secretory leukocyte peptidase inhibitor;  
 285 ENSG00000249139 EPPIN-WFDC6 readthrough; EPPIN Epididymal peptidase inhibitor; KLK3  
 286 Kallikrein-related peptidase 3; WFDC5 WAP four-disulfide core domain 5; HSPE1 Heat shock  
 287 10kDa protein 1 (chaperonin 10); HSPA9 Heat shock 70kDa protein 9 (mortalin); GRPEL1  
 288 GrpE-like 1, mitochondrial (*E. coli*); HSP90AA1 Heat shock protein 90kDa alpha (cytosolic);  
 289 VDAC1 Voltage-dependent anion channel 1; HSP90AB1 Heat shock protein 90kDa alpha  
 290 (cytosolic), class B member 1; HSPA4 Heat shock 70kDa protein 4; GRPEL2 GrpE-like 2,  
 291 mitochondrial (*E. coli*); HSPA8 Heat shock 70kDa protein 8; NPM1 Nucleophosmin (nucleolar  
 292 phosphoprotein B23, numatrin); DNAJB6 DnaJ (Hsp40) homolog, subfamily B, member 6;  
 293 DNAJB1 DnaJ (Hsp40) homolog, subfamily B, member 1; HSPH1 Heat shock 105kDa/110kDa  
 294 protein 1; DNAJC7 DnaJ (Hsp40) homolog, subfamily C, member 7; HSPA1A Heat shock  
 295 70kDa protein 1A; DNAJB12 DnaJ (Hsp40) homolog, subfamily B, member 12; SUGT1 SGT1,  
 296 suppressor of G2 allele of SKP1 (*S. cerevisiae*); CDC37 Cell division cycle 37 homolog (*S.*  
 297 *cerevisiae*); PTGES3 Prostaglandin E synthase 3 (cytosolic); STIP1 Stress-induced-  
 298 phosphoprotein 1; AHSA1 AHA1, activator of heat shock 90kDa protein ATPase homolog 1  
 299 (yeast); TTC28 Tetratricopeptide repeat domain 28; SUGT1P3 Suppressor of G2 allele of SKP1  
 300 (*S. cerevisiae*) pseudogene 3; TTC12 Tetratricopeptide repeat domain 12; TTC31  
 301 Tetratricopeptide repeat domain 31; HSPB3 Heat shock 27kDa protein 3; Inhibitor of actin  
 302 polymerization; SPACA4 Sperm acrosome associated 4; SPRYD4 SPRY domain containing 4;  
 303 KANSL2 KAT8 regulatory NSL complex subunit 2; SPACA1 Sperm acrosome associated 1;  
 304 HDAC7 Histone deacetylase 7; CXorf48 Chromosome X open reading frame 48; DDX25 DEAD  
 305 (Asp-Glu-Ala-Asp) box helicase 25; ACRBP Acrosin binding protein; IZUMO4 IZUMO family  
 306 member 4; SCG5 Secretogranin V (7B2 protein); PLAUR Plasminogen activator, urokinase  
 307 receptor; MVK Mevalonate kinase; ACRV1 Acrosomal vesicle protein 1; SPACA3 Sperm  
 308 acrosome associated 3; TMEM225 Transmembrane protein 225; FAM71F1 Family with  
 309 sequence similarity 71, member; WBSCR28 Williams-Beuren syndrome chromosome region  
 310 28;TEKT3 Tektin 3; Structural component of ciliary and flagellar microtubules; CD9 CD9  
 311 molecule; FOLR4 Folate receptor 4 (delta) homolog (mouse); FOLR1 Folate receptor 1 (adult);  
 312 FOLR2 Folate receptor 2 (fetal); ZBP2 Zona pellucida binding protein 2; PRSS21 Protease,  
 313 serine, 21 (testisin); ADAM2 ADAM metallopeptidase domain 2.

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#### 321 4. Discussion

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323 In the present study, we used a liquid chromatography mass spectrometry approach  
324 (Shotgun proteomics) to characterize the protein profile of spermatozoa from the Amazon  
325 squirrel monkey (*Saimiri collinsi*) kept in captive. The seminal coagulum and the parameters,  
326 such as appearance, semen volume, sperm concentration, motility, vigor and sperm morphology  
327 in Amazon squirrel monkey were similar to those previously reported for sperm from coagulated  
328 fraction after dilution in ACP-118<sup>®</sup> extender [18-20].

329 This proteomic strategy allowed the identification of 2,343 proteins and represents a  
330 major contribution to the understanding of spermatozoa composition in *Saimiri collinsi* species.  
331 In *S. collinsi*, it was possible to verify important proteins that participated in spermatogenesis and  
332 sperm motility, such as ROPN1L, HSPA2, cat eye syndrome critical region protein 5, and  
333 phospholipid hydroperoxide glutathione. In mice, the loss of ROPN1L impairs sperm motility,  
334 cAMP-dependent protein kinase phosphorylation, and fibrous sheath integrity [26]. ROPN1L is a  
335 sperm flagellar protein that binds A-kinase anchoring protein (AKAP) 3 and 4, which are primary  
336 components of the sperm fibrous sheath. The fibrous sheath is a flagellar cytoskeletal structure  
337 unique to sperm that surrounds the outer dense fibers and axoneme [26, 27]. The degradation of  
338 AKAP3 and subsequent dephosphorylation of tyrosine result in sperm capacitation [28].

339 Heat shock proteins (HSPs) are chaperone proteins that are expressed in response to cell  
340 stress [29, 30]. Several HSP family members are expressed in the sperm, such as HSP 70 kDa  
341 (HSP70), which appears in the acrosome membranes. HSP 60 kDa (HSP60) is located primarily  
342 in the sperm midpiece, in association with the mitochondria, whereas HSP 90-alpha  
343 (HSP90AA1) is located in the sperm flagellum [31]. HSP60, HSP70, and HSP90AA1 are known  
344 components of sperm in different species, such as humans [32], rams [33], bulls, stallions, cats,  
345 and dogs [34]. The acrosomal HSP70 has a role in gamete interaction and fertilization [35],  
346 whereas HSP90AA1 expression has been correlated with the resistance of sperm to freezing [36,  
347 37] since this protein is characterized as a ubiquitous molecular chaperone that provides  
348 protection and protein folding during thermal stress and resistance against cell oxidative stress  
349 [38].

350 HSPA2, which is a molecular chaperone that assists in the folding, transport, and  
351 assembly of proteins in the cytoplasm, mitochondria, and endoplasmic reticulum and is a testis-

352 specific member of the 70-kDa family [39], has been suggested to be crucially involved in  
353 spermatogenesis and meiosis [40]. In humans, the downregulation of HSPA2 mRNA was  
354 observed in testes with abnormal spermatogenesis, and the protein expression was high in normal  
355 spermatogenesis and low in spermatogenesis arrest [41]. Human HSPA2 was shown to regulate  
356 the expression of the sperm surface receptors involved in sperm-oocyte recognition [42], and its  
357 depression in the testes was also associated with spermatogenic impairment and the fertilization  
358 rate in men with azoospermia who were treated with intracytoplasmic sperm injections [43].

359 Sperm motility is driven mainly by the energy produced by the mitochondria present in  
360 the intermediate piece of the male gamete [44]. However, the axoneme is another important  
361 cellular component that is directly associated with sperm motility. The dynein heavy chains have  
362 been annotated as subunits of the axonemal dynein complexes, which are multisubunit axonemal  
363 ATPase complexes that generate the force for cilia motility and govern the beat frequency [45].  
364 DNAH1 is related to spermatogenesis and cell proliferation [46]. In humans, mutations in  
365 DNAH1 cause multiple morphologic abnormalities of the sperm flagella, leading to male  
366 infertility [6]. The radial spoke proteins play a key role in regulating dynein activity and flagellar  
367 motility [47, 48].

368 In this context, Imai et al. [49] showed that the failure to express phospholipid  
369 hydroperoxide glutathione peroxidase (GPX4) caused human male infertility, with 30% of men  
370 diagnosed with oligoasthenozoospermia showing a significant decrease in the level of the  
371 enzyme. Those authors also found a significantly lower number of spermatozoa in the semen and  
372 significantly lower motility of the spermatozoa than those seen in fertile men. GPX4 is an  
373 intracellular antioxidant that directly reduces peroxidized phospholipids and is strongly expressed  
374 in the mitochondria of the testis and spermatozoa. In bulls, GPX4 is considered a unique marker  
375 for seminal quality analysis owing to the direct correlation between the selenoperoxidase and the  
376 progressive motility of the sperm [50].

377 The acrosome, which is a membrane-bound exocytotic vesicle that is located over the  
378 anterior portion of the nucleus, contains the hydrolytic enzymes that are required for the  
379 acrosome reaction, binding of the zona pellucida (ZP), penetration through the ZP, and sperm-  
380 egg membrane fusion, all of which are indispensable events during the fertilization process [51].  
381 In the acrosome membrane (internal and external membranes), the sperm acrosome membrane-  
382 associated family (i.e., SPACA3, SPACA1, and SPACA4) [52, 53] are sperm surface membrane



383 proteins that may be involved in the adhesion and fusion of the sperm to the egg prior to  
384 fertilization [54]. SPACA1 and SPACA3 are localized in the acrosomal matrix, including the  
385 principal segment and equatorial segment, and are proteins characterized as membrane antigens  
386 [55, 56, 57]. SPACA1 may be involved in sperm fusion with the oölemma, since treatment of  
387 human sperm with the anti-SPACA1 antibody prevented sperm penetration into zona-free  
388 hamster eggs [54]. Fujihara et al. [56] demonstrated that the SPACA1 protein was indispensable  
389 for the normal shaping of the sperm heads during spermiogenesis in mice. In humans, this protein  
390 was identified as a sperm membrane antigen, with a molecular mass ranging from 32 to 34 kDa  
391 [54].

392 Membrane fusion is a key event in the fertilization process that culminates in the merger  
393 of the male–female gamete membranes and cytoplasm and fusion of the genomes, thereby  
394 initiating embryonic development [57]. In humans, a change in the expression of the sperm  
395 proteins may be a major cause of subfertility in men with normozoospermia [58]. In this context,  
396 research has been focusing on the identification of the key molecular players and their functions,  
397 and several proteins in the egg or the spermatozoa have been found to be essential for  
398 fertilization.

399 Until now, IZUMO1 has been found to be the essential protein on the sperm side for the  
400 fusion process. As a testis-specific protein, IZUMO was discovered on the equatorial segment of  
401 the acrosome-reacted mouse spermatozoa through proteomic analysis of the antigen recognized  
402 by the monoclonal anti-mouse sperm antibody [59]. IZUMO is present in both mouse (~56 kDa  
403 protein) and human (~38 kDa protein) sperm [60]. In mice, immunization with the IZUMO  
404 protein caused a contraceptive effect in females, which was due to the significantly inhibited  
405 fusion of sperm to the zona-free mouse eggs with the anti-PrimeB antibody. However, no effect  
406 on sperm motility was observed [61]. IZUMO2, IZUMO3, and IZUMO4 have significant  
407 homology with the N-terminal domain of IZUMO1 [62]. Inoue et al. [9] showed the interaction  
408 between angiotensin-converting enzyme-3 located on the sperm acrosomal cap and IZUMO1 in  
409 the fertilization process. However, it was reported that angiotensin-converting enzyme-3  
410 disappears from the membrane after the acrosome reaction. Nevertheless, the *in silico* protein  
411 interaction analysis of IZUMO1 revealed its association with the CD9 molecule, folate receptor 4  
412 (delta) homolog (mouse), folate receptor 1 (adult), folate receptor 2 (fetal), SPACA1, SPACA4,  
413 IZUMO family member 4, zona pellucida binding protein 2, and metallopeptidase domain 2.

414           After the acrosome reaction, the C-terminal calmodulin domain (20 kDa) of SPA17  
415 (located on the external side of the sperm plasma membrane) is proteolytically cleaved to 17 kDa  
416 and then binds to the extracellular matrix of the oocyte. This C-terminus of SPA17 plays a role in  
417 cell–cell adhesion [63, 64].

418

## 419 **5. Conclusions**

420

421           The present study is a comprehensive overview of the sperm proteome in the Amazon  
422 squirrel monkey, and is the broadest inventory (investigation) of the sperm proteome in the genus  
423 *Saimiri* as well as in Neotropical primates thus far. The knowledge acquired about the sperm  
424 proteins is a significant step forward in helping toward our understanding of the reproductive  
425 biology of the genus *Saimiri*, as it provides crucial information for the elucidation of the  
426 underlying mechanisms associated with sperm function. In this way, our study amplifies the  
427 advances in biotechnological research on animal reproduction for the conservation of endangered  
428 species, and provides a reference for similar studies on other Neotropical primates.

429

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694 **Supporting information**

695 **S1 Table. Spectral count of *Saimiri collinsi* sperm protein throughout an entire year (.XLS).**

696 **S2 Table. Sperm proteins of *Saimri collinsi* that participate in important reproductive**  
697 **events (.XLS).**

## CONCLUSÕES GERAIS

De um total de 2.342 proteínas identificadas por meio da cromatografia líquida acoplada à espectrometria de massas em tandem dos espermatozoides de *Saimiri collinsi*. Essas proteínas foram relacionadas a importantes eventos reprodutivos como a espermatogênese, motilidade espermática, capacitação e fecundação.

No tocante as eletromicrografias obtidas por microscopia eletrônica de varredura e transmissão, essas imagens nos permitiram caracterizar e avaliar de forma mais ampla os espermatozoides de *S. collinsi*, fator este que nos auxilia na detecção de possíveis anormalidades espermáticas que podem afetar e diminuir a capacidade fecundante dos espermatozoides.

O conhecimento adquirido sobre as proteínas espermáticas juntamente com a análise micromorfológica e ultraestrutural da célula espermática, é um avanço significativo no sentido de ajudar a compreender a biologia reprodutiva do gênero *Saimiri*, pois fornece informações cruciais para a elucidação dos mecanismos subjacentes associados à função espermática, proporcionando ampliar os avanços da pesquisa biotecnológica em reprodução animal para a conservação de espécies ameaçadas de extinção e fornece uma referência para estudos semelhantes sobre outros primatas neotropicais.

Nesse sentido, complementarmente, a busca por parâmetro não invasivo, como a distância anogenital, que auxiliam no exame andrológico para a seleção de primatas não-humanos é de suma importância, visto que podem predizer a baixa qualidade seminal e inviabilizaria o uso destes animais em programas de biotecnologia para a conservação de espécies ameaçadas de extinção.