# **GUSTAVO SCHETTINI**

175 West Campus Drive, 24061, Blacksburg, VA, United States

gschettini@vt.edu \( \rightarrow \frac{gschettini.github.io}{} \)

# **EDUCATION**

Virginia Polytechnic Institute and State University, Virginia Tech

Ph.D. in progress in Animal Sciences

2022 - Present Blacksburg-VA, USA

Sao Paulo State University, UNESP

M.Sc. in Genetics and Animal Breeding

2019 - 2021 Jaboticabal-SP, Brazil

Federal University of Bahia, UFBA

B.S. in Veterinary Medicine

2013 - 2018 Salvador-BA, Brazil

#### **COMPLEMENTARY EDUCATION**

**Graduate Teaching Assistant Workshop**, Virginia Polytechnic Institute and State University.

Credit Hours: -

2022.

Systems Biology and Gene Networks Inference: Application to Livestock Breeding and Genetics, Sao Paulo University (ESALQ-USP).

Credit Hours: 40h

2019.

Special Topics: Data Analysis in R Software, Sao Paulo State University

Credit Hours: 24h

(UNESP). 2019.

Special Topics in Applied Genomic Selection in Livestock: Methods and Tools, Sao Paulo State University (UNESP).

Credit Hours: 24h

2019.

Introduction to Using Software R, Federal University of Bahia (UFBA).

Credit Hours: 20h

2015.

### TEACHING EXPERIENCE

**Teaching Assistant**. Course: Genomics (APSC 4054). Virginia Polytechnic Institute

Fall 2024

and State University. Instructor: Fernando Biase

**Teaching Assistant**. Course: Genomics (APSC 4054). Virginia Polytechnic Institute

and State University. Instructor: Fernando Biase

Fall 2023

Spring 2023

Teaching Assistant. Course: Animal Breeding and Genetics (ALS 3104). Virginia

Polytechnic Institute and State University. Instructor: Gota Morota

**Teaching Assistant**. Course: Special Topics in Biotechnology in Animal Production (MEVB10). Federal University of Bahia (UFBA). Instructor: Barbara M. P. da Silva Souza

Fall 2020

- Biotechnologies applied to animal breeding I: Principles of DNA sequencing and DNA polymorphisms' identification.
- Biotechnologies applied to animal breeding II: Principles of RNA sequencing and Gene (Co-) expression analyses overview.

#### RESEARCH EXPERIENCE

Graduate Research Assistant, Virginia Polytechnic Institute and State University	2022- Present
Graduate Research Assistant, Sao Paulo State University (UNESP)	2019-2021
Undergraduate Research Internship, Sao Paulo State University (UNESP)	2018-2018
Undergraduate Research Internship, Federal University of Bahia (UFBA)	2018-2018
Rural Extension Projects Internship, Federal University of Bahia (UFBA)	2018-2018
<b>Technological Development and Innovation Internship</b> , Federal University of	2017-2018
Bahia (UFBA)	
Scientific Research Initiation, Federal University of Bahia (UFBA)	2015-2017

#### RESEARCH SUPPORT

1. Graduate Research Development Program (GRDP).

\$1000

Transcriptome profile of in vitro-produced bovine blastocyst associated with hatching rate.

PI: Schettini, G.P.

09/30/2022-10/01/2023

**2.** AG2PI support to develop genomic workflows.

\$8000

Integrating Illumina and ONT sequence workflow.

PI: Schettini, G.P.

09/30/2023-05/31/2024

3. Graduate Research Development Program (GRDP).

\$1000

Integrating Illumina and ONT sequence workflow.

PI: Schettini, G.P.

09/13/2024-01/31/2025

#### PEER-REVIEWED JOURNAL PAPERS

- 1. Schettini G. P.; Morozyuk, M.; Biase, F.H. Identification of novel cattle (*Bos taurus*) genes and biological insights of their function in pre-implantation embryo development. BMC Genomics, v. 25, 775, 2024. DOI
- 2. Kalbfleisch, T.S., McKay, S.D., Murdoch, B.M. *et al.* The Ruminant Telomere-to-Telomere (RT2T) Consortium. Nature Genetics, v. 56, 1566–1573, 2024. DOI

- 3. Biase F. H.; Schettini G. P. Protocol for the electroporation of CRISPR-Cas for DNA and RNA targeting in Bos taurus zygotes. STAR Protocols, v. 5, 2024.
- 4. Nix, J. L.; Schettini, G. P.; Speckhart S. L.; Ealy, A. D.; Biase, F. H. Ablation of OCT4 function in cattle embryos by double electroporation of CRISPR-Cas for DNA and RNA targeting (CRISPR-DART). PNAS Nexus, pgad343, 2023. DOI
- 5. Nix, J. L.; Schettini, G. P.; Biase, F. H. Sexing of cattle embryos using RNA-sequencing data or polymerase chain reaction based on a complete sequence of cattle chromosome Y. Frontiers in Genetics, v. 14, 2023. DOI
- 6. Toro-Ospina, A.; Herrera Rios, A. C.; dos Santos, W. B.; Schettini, G. P.; Vallejo Aristizabal, V. H.; Tovar Claros, G.; Ortiz Morea, E. G. Genetic Architecture and Signatures of Selection in the Caqueteño Creole (Colombian Native Cattle). Diversity, v. 14, 828, 2022. DOI
- 7. Toro-Ospina, A.; Herrera Rios, A. C.; **Schettini, G. P.**; Vallejo Aristizabal, V. H.; dos Santos, W. B.; Zapata, C. A.; Ortiz Morea, E. G. **Identification of Runs of Homozygosity Islands and Genomic Estimated Inbreeding Values in Caqueteño Creole Cattle (Colombia)**. Genes, v. 7, 1232, 2022. DOI
- 8. Schettini, G. P.; Peripolli, E.; Alexandre, P. A.; Dos Santos, W. B.; Pereira, A. S. C.; De Albuquerque, L. G.; Baldi, F.; Curi, R. A. Transcriptome Profile Reveals Genetic and Metabolic Mechanisms Related to Essential Fatty Acid Content of Intramuscular Longissimus tho-racis in Nellore Cattle. Metabolites, v. 12, 471, 2022. DOI
- Schettini, G. P.; Peripolli, E.; Alexandre, P. A.; Dos Santos, W. B.; da Silva Neto, J. B.; Pereira, A. S. C.; De Albuquerque, L. G.; Curi, R. A.; Baldi, F. Transcriptomic profile of longissimus thoracis associated with fatty acid content in Nellore beef cattle. Animal Genetics, v. 53, p. 264-280, 2022. DOI
- 10. Dos Santos, W. B.; Schettini, G. P.; Maiorano, A. M.; Bussiman, F. O.; Balieiro, J. C. C.; Ferraz, G. C.; Pereira, G. L.; Baldassini, W. A.; Neto, O. R. M.; Neto, H. N. O.; Curi, R. A. Genome-wide scans for signatures of selection in Mangalarga Marchador horses using high-throughput SNP genotyping. BMC genomics, v. 22, p. 737-754, 2021. DOI
- 11. **Schettini, G. P.**, Lambert, S. M.; Da Silva Souza, B. M. P.; Costa, R. B.; De Camargo, G. M. F. **Genetic potential of Sindhi cattle for A2 milk production**. Animal Production Science, v. 60, p. 893-895, 2020. DOI
- 12. Dos Santos, W. B.; Schettini, G. P.; Fonseca, M. G.; Pereira, G. L.; Chardulo, L. A. L.; Neto, O.; Baldassini, W. A.; Oliveira, H. N.; Curi, R. A. Fine-scale estimation of inbreeding rates, runs of homozygosity and genome-wide heterozygosity levels in the Mangalarga Marchador horse breed. Journal of Animal Breeding and Genetics, v. 00, p. jbg.12508, 2020. DOI
- 13. Da Silva Neto, J. B.; Peripolli, E.; Da Costa e Silva, E. V.; Espigolan, R.; Neira, J. D. R.; Schettini, G. P.; Da Costa Filho, L. C. C.; Barbosa, F. B.; Macedo, G. G.; Costa-Brunes, L.; Lobo, R. B.; Pereira, A. S. C.; Baldi, F. Genetic correlation estimates between age at puberty and growth, reproductive, and carcass traits in young Nelore bulls. Livestock Science, v. 241, p.104266, 2020. DOI

# **SKILLS**

- 1. Laboratory-related techniques
  - DNA and RNA extraction
  - Library construction for Next Generation Sequencing
  - Oxford Nanopore Technologies Sequencing
  - *In vitro* embryo production (bovine)
- 2. Programming
  - Unix/Linux
  - Bash
  - awk
  - R
  - Markdown
  - Docker/Conda
  - Git

# **LANGUAGES**

- 1. English Full professional proficiency
- 2. Spanish Professional working proficiency
- **3.** Portuguese Native