

Contact

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Skills

Python, Pytorch, AI, Javascript, PHP, Java, Matlab, Pytorch, Scikitlearn Pandas, MySQL, MongoDB, Linux, Windows, Blender, D3, Seaborn.

These are some of the tools in my belt. I am continuosly learning more!

Human languages

Spanish

English



French





LESLIE SOLORZANO

Computer Scientist and Engineer, Visualization expert

Current work

Department of Immunology Genetics and Pathology Uppsala University Postdoctoral Researcher

Education

2021

Uppsala Universitet

PhD / Teknologie Doktor

Image Processing, Machine Learning and Visualization for Tissue Analysis

2016

Université Claude Bernard de Lyon 1 Master in Computer Graphics and Image Processing Visualization quantification and analysis of lung parenchyma in presence of ARDS

2015

Universidad de los Andes

Master in Computer Science and Engineering Visualization quantification and analysis of lung parenchyma in presence of ARDS

2013 🖒

Universidad Nacional de Colombia B.S in Computer Science and Engineering Human Computer Interaction Application for the

manipulation of software

Grants and awards

Karolinska Institutet

KID - Partial funding of Doctoral Education at Karolinska institutet 2022 Cosupervisor of PhD student 2024-2028

Selected publications







The ACROBAT 2022 Challenge: Automatic Registration Of Breast Cancer Tissue

Medical Image Analysis

P Weitz, M Valkonen, L Solorzano, ..., M Rantalainen



Improved breast cancer histological grading using deep learning Annals of oncology

Y Wang, B Acs, S Robertson, B Liu, L Solorzano, C Wählby, J Hartman, M Rantalainen





Artificial intelligence for diagnosis and grading of prostate cancer in biopsies: a population-based, diagnostic study The Lancet Oncology

P. Ström, K. Kartasalo, H. Olsson, L. Solorzano, B. Delahunt, et al.





TissUUmaps: Interactive visualization of large-scale spatial gene expression and tissue morphology data BioInformatics - OUP

L. Solorzano, G. Partel, C. Wählby

Analyze Imaging processing Segment Quantify Classify sources VisualizeInterpret

All publications

As seen in Google Scholar

Cytometry Part A

2024	The ACROBAT 2022 Challenge: Automatic Registration Of Breast Cancer Tissue P Weitz, M Valkonen, L Solorzano, C Carr, K Kartasalo, C Boissin,, M Rantalainen Medical Image Analysis
2024	Ensemble-based deep learning improves detection of invasive breast cancer in routine histopathology images L Solorzano, S Robertson, B Acs, J Hartman, M Rantalainen Heliyon
2023	A multi-stain breast cancer histological whole-slide-image data set from routine diagnostics P Weitz, M Valkonen, L Solorzano, C Carr, K Kartasalo, C Boissin,, M Rantalainen Nature Scientific Data
2022	<pre>Improved breast cancer histological grading using deep learning Y Wang, B Acs, S Robertson, B Liu, L Solorzano, C Wählby, J Hartman, M Rantalainen Annals of oncology</pre>
2022	Comparison of East-Asia and West-Europe cohorts explainsdisparities in survival outcomes and highlights predictivebiomarkers of early gastric cancer aggressiveness C Pereira, JH Park, S Campelos, I Gullo, C Lemos, L Solorzano et al. International Journal or Cancer
2021	Image Processing, Machine Learning and Visualization for Tissue Analysis (PhD Thesis) L Solorzano - May 12th 2021 - <u>Link to video</u> Acta Universitatis Upsaliensis
2021	Machine learning for cell classification andneighborhood analysis in glioma tissue L Solorzano, L Wik, T Olsson Bontell, Y Wang, AH Klemm, J Öfverstedt, Cytometry Part A
2020	Automated identification of the mouse brain's spatial compartments from in situ sequencing data G Partel, MM Hilscher, G Milli, L Solorzano, AH Klemm, M Nilsson, C Wählby BMC biology
2020	TissUUmaps: Interactive visualization of large-scale spatial gene expression and tissue morphology data L Solorzano, G Partel, C Wählby Bioinformatics
2020	Towards automatic protein co-expression quantification in immunohistochemical TMA slides L Solorzano, C Pereira, D Martins, R Almeida, F Carneiro, GM Almeida, IEEE journal of biomedical and health informatics
2020	Transcriptome-Supervised Classification of Tissue Morphology Using Deep Learning A Andersson, G Partel, L Solorzano, C Wählby 2020 IEEE 17th International Symposium on Biomedical Imaging (ISBI)
2020	Artificial intelligence for diagnosis and grading of prostate cancer in biopsies: a population-based, diagnostic study P Ström, K Kartasalo, H Olsson, L Solorzano, B Delahunt, DM Berney, The Lancet Oncology
2019	Deep learning in image cytometry: a review A Gupta, PJ Harrison, H Wieslander, N Pielawski, K Kartasalo, G Partel, L Solorzano

2019 Voxel-wise assessment of lung aeration changes on CT images using image registration: application to acute respiratory distress syndrome (ARDS) M Orkisz, AM Pinzón, JC Richard, C Guérin, L Solorzano, DF Sicaru, ... International journal of computer assisted radiology and surgery 2019 Quality Assurance and Local Regions for Whole Slide Image Registration. L Solorzano, C Wählby Journal of Pathology Informatics 2018 Whole slide image registration for the study of tumor heterogeneity L Solorzano, GM Almeida, B Mesquita, D Martins, C Oliveira, C Wählby Computational pathology and ophthalmic medical image analysis 2017 Decoding gene expression in 2D and 3D M Bombrun, P Ranefall, J Lindblad, A Allalou, G Partel, L Solorzano, ... Scandinavian Conference on Image Analysis Software Visualization to Simplify the Evolution of Software Systems 2012 D Montaño, L Solorzano, HR Umaña-Acosta Research topics in software evolution and maintenance

Academic experience

2009-2026

{Research groups}

- Vicković Innovation Lab (UU-NYGC) Immunology Genetics and Pathology (Python, JS, AI and visualization)
- Rantalainen Group (KI) Medical Epidemiology and Biostatistics (Python, JS, AI and visualization)
- Wählby Lab (UU) Visual Information and Interaction (Python, JS, AI and visualization)
- CREATIS (UCBL) Medical image processing lab (C++)
- IMAGINE (UNIANDES) Visualization, computer graphics and robotics lab (C++, shell)
- BioIngenium (UNAL) Bioengineering research group (Matlab, C++)
- ColSWE (UNAL) Software Engineering Colective (Java)
- EIDOS (UNAL) Linux and free software research group. (Unix)

2011-2022

{Teaching}

- ◆ Deployment of analytics solutions in Big Data (CO remote) 2022
- ◆ Scientific Data Visualization (SE) 2020
- ◆ Computer Graphics (SE) 2017-2018-2019-2020
- ◆ Programming Design and Data Structures (SE) 2017
- ◆ Image Processing (CO) 2015
- ◆ Software Engineering (CO) 2011
- ◆ Computer programming (C++) (CO) 2011

2016-2021

{Main project} {PhD program} {Sweden}

- ◆ TissUUmaps: Develop computational methods to combine spatially resolved information on tissue morphology with in situ RNA sequencing and protein detection and efficient visualization of results (Python, HTML, javascript)
- ◆ Reviewer ICASSP 2017, ISBI 2018-2019-2020
- ◆ Participated in the Deep learning bootcamp at Max Planck Institute 2018
- ◆ Participated in the QuPath Computational Pathology workshop at EPFL
- ◆ Information system for the Center for Image Analysis and annual report generation and edition

2016-2021

{Presentations} {PhD program} {Sweden}

- ◆ EMBO-EMBL (2019): Seeing is believing imaging the molecular processes of life, Symposium Presented: lightning talk, Poster: Exploratory analysis and visualization of in-situ
- sequencing data
 - SOFOSKO (2019): Talk at research summer school: Image analysis for microscopy, into the
- private life of tissue
- ◆ Soapbox Uppsala (2019): Women in STEM, popular science public presentation STorM (2019): The Swedish Tumor Microenvironment Meeting: Image analysis for the study of
- ◆ tumor micro environments
 - MICCAI (2018): International Workshop on Ophthalmic Medical Image Analysis: Whole Slide
- ◆ Image Registration for the Study of Tumor Heterogeneity
 - SOFOSKO (2018): Talk at research summer school: Image analysis for digital histopathology for personalized cancer treatment

2013-2015

{Project involvement} {Master Degree} {Colombia}

- Encuentro BigData 2015. Project for citizen public data acquisition, treatment and display for the Technology Ministry. (PHP, MongoDB, Python)
- WEBSIS project, website for the systems engineering department at Andes University and CMS module development (joomla, worpress).
- ◆ Support and maintenance of Linux servers, IMAGINE laboratory, Uniandes. (Fedora)

2007-2012

{National University of Colombia} {Undergraduate program}

- ◆ Computer Science and Engineering Programming, software engineering, mathematics,
- ◆ information theory
- Application for the manipulation of software graphical environments -Undergraduate thesis (Java)
- ◆ New Trends in Software Maintenance and Evolution Extracurricular course (Java)

 Trends, design and display in new generation information systems Extracurricular course.

 (C++)

Work experience

Sweden {2024-2026}

Sweden {2022-2024}

Sweden {2021-2022}

Sweden {2016-2021}

Colombia {2015-2016}

France {2013-2014}

United States {2012}

Colombia {2007-2009}

Vicković Innovation Lab - Department of Immunology Genetics and Pathology Uppsala University - Postdoctoral researcher

Rantalainen Group - Department of Medical Epidimiology and Biostatistics Karolinska Institutet - Postdoctoral researcher

Centrum för Bildanalys and Wählby Lab Research Engineer - Forskningsingenjör

Department of Information Technology - Wählby Lab Uppsala University - PhD Researcher

IMAGINE computer vision and robotics group Research engineer and linux system administrator

CREATIS Biomedical Imaging Research Lab Master student - Medical image registration

University of Delaware Summer research program at the Computer Science department Software testing methodologies (JVM)

National University of Colombia International affairs office Webmaster and support engineer (JSP)

Hobbies and interests

ThreeJS, OpenGL, WebGL, Blender, Inkscape Angular, Javascript General purpose programming in Python, Javascript Volunteer teaching in local school

