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nttps://svanuytven.github.io/

in svanuytven

**y** Svanuytven

Svanuytven Belgian

# Sebastiaan Vanuytven

Curriculum Vitae

#### Research Interests

- Single-Cell (Multi-)Omics
- Oncobiology
- Gene Therapy

- Bayesian Statistics
- Nuclear Architecture
- o HIV

## Experience

#### 2017-Present PhD Student Computational Biology,

Labratory of Reproductive Genomics, KU Leuven, Belgium.

Studying for the first time the complete phenotype of a single cancer cell in its natural environment by combining single-cell multi-omics techniques and spatial transcriptomics

### 2017-Present Visiting Scientist,

Cancer Genomics Laboratory, Francis Crick Institute, United Kingdom.

#### 2013–2014 Research Rotations.

KU Leuven, Belgium.

Internships in the research groups of Prof. Zeger Debyser, Prof. Stein Aerts and Prof. Johan Neyts

#### Education

2015–2017 Master of Science Bioinformatics, KU Leuven, Magna cum laude.

2013–2015 Master of Science Biomedical Sciences, KU Leuven, Magna cum laude.

2010–2013 Bachelor of Science Biomedical Sciences, KU Leuven, Cum laude.

#### Master thesis

#### Master of Science Bioinformatics

Title Single-cell sequencing to understand the biology of cellular heterogeneity in cancer

Supervisors Prof. Thierry Voet, Prof. Stein Aerts, Dr. Alejandro Sifrim, Dr. Daniel Brown

Description Development of novel analysis methods and robust pipelines for single-cell multiomics experiments and analysis of large-scale single-cell sequencing datasets of

primary breast cancers

#### Master of Science Biomedical Sciences

Title Development of safer MLV-based gene therapy vectors: studying BET proteinchromatin occupation and generation of p12-fusions

Supervisors Prof. Zeger Debyser, Prof. Rik Gijsbers, Dr. Jonas Demeulemeester

Description Analysis of BET protein ChIP-seq data and generation of p12-fusions to understand the genotoxicity of MLV-based gene therapy vectors

### Grants

2020 EACR travel fellowship

2018-2022 FWO Strategic basic research grant

2015 BBC2015 Student Fellowship

Travel grant to visit Crick Institute Personal Funding for PhD research

Travel grant to attend BBC2015 conference

#### **Publications**

#### Journal Articles

[1] Anna-Marei Böhm\*, Naomi Dirckx\*, Robert J. Tower\*, Nicolas Peredo, **Vanuytven**, **Sebastiaan**, Koen Theunis, Elena Nefyodova, Ruben Cardoen, Volkhard Lindner, Thierry Voet, Matthias Van Hul, and Christa Maes. Activation of skeletal stem and progenitor cells for bone regeneration is driven by pdgfrb signaling. *Developmental Cell*, 2019.

#### Conference Posters

- [1] Vanuytven Sebastiaan, Koen Theunis, Sarah Geurs, Peter Van Loo, Giuseppe Floris, and Thierry Voet. Single-Cell Multi-Omics Sequencing to Understand the Nature, Extent and Biology of Cellular Heterogeneity in Six Special Breast Cancer Cases. In *EACR Cancer Genomics 2019*, Cambridge, UK, June 2019.
- [2] Vanuytven Sebastiaan, Alejandro Sifirm, Mabel T Teng, Laura Mora Bitria, Daniel Brown, Elia Fernandez Gallardo, Koen Theunis, Sarah Geurs, Samira Majjaj, Hughes Duvilliers, Michail Ignatiadis, Francoisea Rothe, Christos Sotiriou, Peter Van Loo, Christine Desmedt, and Thierry Voet. Single-cell multi-omics sequencing to understand the nature, extent and biology of cellular heterogeneity in breast cancer. In Single Cell Biology Confernece, Wellcome Genome Campus, Hinxton, UK, Mar. 2018.
- [3] Vanuytven Sebastiaan, Alejandro Sifirm, Mabel T Teng, Laura Mora Bitria, Daniel Brown, Elia Fernandez Gallardo, Koen Theunis, Sarah Geurs, Samira Majjaj, Hughes Duvilliers, Michail Ignatiadis, Francoisea Rothe, Christos Sotiriou, Peter Van Loo, Christine Desmedt, and Thierry Voet. Single-cell multi-omics sequencing to understand the nature, extent and biology of cellular heterogeneity in breast cancer. In *Benelux Bioinformatics Conference*, Leuven, Belgium, Dec. 2017.
- [4] Vanuytven Sebastiaan, Jonas Demeulemeester, Zeger Debyser, and Gijsbers Rik. Studying BET protein-chromatin occupation to understand genotoxicity of MLV-based gene therapy vectors. In *Benelux Bioinformatics Conference*, Antwerp, Belgium, Dec. 2015.

#### **Talks**

- 05/2019 **EMBO Single-cell Biology**, *Tokyo*, Japan, Selected Speaker.

  Single-Cell Multi-Omics Sequencing to Understand the Nature, Extent and Biology of Cellular Heterogeneity in Six Special Breast Cancer Cases
- 11/2018 **Genomics for diagnostics of Rare Diseases TRAINMALTA meeting**, *Leuven*, Belgium, Invited Speaker.

  Single-cell multi-omics to detect genetic variants in cancer and development
- 06/2018 **European Human Genetics Conference 2018**, *Milan*, Italy, Selected speaker. Single-cell multi-omics sequencing to understand the nature, extent and biology of cellular heterogeneity in breast cancer
- 05/2018 **Bioinformatics Student Symposium 2018**, *Antwerp*, Belgium, Selected speaker. Single-cell multi-omics sequencing to understand the nature, extent and biology of cellular heterogeneity in breast cancer
- 02/2018 **Belgian Human Genetics Conference 2018**, *Ghent*, Belgium, Selected speaker. Single-cell multi-omics sequencing to understand the nature, extent and biology of cellular heterogeneity in breast cancer
- 12/2015 BBC2015 Student Conference, Antwerp, Belgium, Selected speaker. Studying BET protein-chromatin occupation to understand genotoxicity of MLV-based gene therapy vectors

## Reviewing activities

Journals Cell (junior reviewer)
Conferences ESHG 2018/2019

# Scientific leadership experiences

Organiser RGS Belgium student symposium 2019

## Languages

Dutch Native language

English Extensive Knowledge

French Good knowledge

German Basic knowledge

Swedish Basic knowledge

## Computer skills

General Microsoft Word, Microsoft Excel, Microsoft PowerPoint, LaTeX

Programming Python, Bash, Java, SQL, Git, Snakemake

Statistics R, Bioconductor, Stan, Matlab

Social Media Twitter, LinkedIn

## Bioinformatics

Single-Cell Seurat, Scater, SingleCellExperiment, Scanpy, SC3, SCENIC, MOFA+, Slingshot

Alignment BWA, Kallisto, STAR, Bowtie2, Samtools, bedtools, RSEM, HTseq, Featurecounts,

Cutadapt, GATK

Cancer ASCAT, MuTect2, Delly, Varscan, Sequenza, SomaticSignature, FreeBayes

ATAC MACS2, cisTopic, snapATAC

General IGV, Picard, Multigc, BEAST, PLINK

## Interests

2018-Present Volunteer and KU Leuven representative.

ISCB Regional Student Group Belgium

2018-Present **Scientific Blogger**, https://svanuytven.github.io/.

2010-Present Passionate mountaineer and runner.

2017-Present Instructor.

Teaching (single-cell) RNA-seq analysis to starting PhD students

2015-2017 Student representative and member of the education committee Bioinformatics KU Leuven.

Organiser buddy system first year students

#### References

#### Thierry Voet, Prof.

Laboratory of Reproductive Genomics, KU Leuven Thierry.Voet@kuleuven.be, +32 16 33 08 41

#### Jonas Demeulemeester, PhD.

Cancer Genomics Laboratory, The Francis Crick Institute Jonas.Demeulemeester@crick.ac.uk, +44 20 7269 3087

#### Alejandro Sifrim, PhD.

Laboratory of Reproductive Genomics, KU Leuven Alejandro.Sifrim@kuleuven.be

#### Rik Gijsbers, Prof.

Laboratory for Molecular Virology & Gene Therapy KU; Leuven Viral Vector Core, KU Leuven

Rik.Gijsbers@med.kuleuven.be, +32 16 332 170