



## Teemu Daniel Laajala

FICAN Cancer Researcher



b. 1987



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## Scientific focus

- Applied math & machine learning
- Prognostic & predictive modeling
- Preclinical & clinical oncology
- Multi-'omics & translatability
- Open science, data, and source code
- R-programming & CRAN-packages

## Other skills

### IT

- R, C#, Python, Java, C, Fortran
- LaTeX, SQL, XML, JSON, Git
- MS Office, Inkscape, GIMP, Adobe suite, HTML/CSS

### Languages

- Professional: English & Finnish
- Conversational: Swedish

### Linked profiles

- [ResearchGate](#)
- [Google Scholar](#)
- [Stack Overflow](#)
- [GitHub](#)
- [LinkedIn](#)
- [Twitter](#)

## Research experience

- 2018 – FICAN West (Läntinen Syöpäkeskus, UTU) Post-doc
- 2018 – University of Colorado – Costello Lab (UCDenver) Post-doc
- 2013 – 2018 Department of Mathematics & Statistics (UTU) PhD student  
Multiple research topics mainly focused on oncology with a PhD thesis focus on advanced prostate cancer.
- 2013 – 2018 Turku Centre for Biotechnology (UTU, ÅA) PhD student  
PSA-molecule research and models for predicting progression of prostate and kidney cancer.
- 2013 – 2018 Institute for Molecular Medicine Finland (FIMM, HY) Project researcher  
Head bioinformatician in multiple collaborative projects.
- 2010 – 2012 Department of Mathematics & Statistics (UTU) Project researcher  
ChIP-seq method development and analysis of cancer studies.
- 2008, 2009 Turku Centre for Biotechnology (UTU, ÅA) Summer intern  
ChIP-seq sequence and diabetes microarray analyses.

## Education

- 2013 – 2018 PhD ('with honours' special mention) University of Turku  
Topic of thesis: *Modeling and Prediction of Advanced Prostate Cancer*
- 2006 – 2012 MSc (Tech) (Bioinformatics, 'exceptional' honors) Aalto University

## Selected publications (First or shared first author)

IF '21-'22

- 6.937 Laajala TD\*, Murtojärvi M\*, et al. *ePCR: an R-package for survival and time-to-event prediction in advanced prostate cancer, applied to real-world patient cohorts*. *Bioinformatics*. 2018 Nov 15;34(22):3957-3959.
- 5.482 Huvila J\*, Laajala TD\*, et al. *Combined ASRGL1 and p53 immunohistochemistry as an independent predictor of survival in endometrioid endometrial carcinoma*. *Gynecol Oncol*. 2018 Apr;149(1):173-180.
- 41.316 Guinney J\*, Wang T\*, Laajala TD\*, et al. *Prediction of overall survival for patients with metastatic castration-resistant prostate cancer: development of a prognostic model through a crowdsourced challenge with open clinical trial data*. *Lancet Oncol*. 2017 Jan;18(1):132-142.
- 4.379 Laajala TD, et al. *Optimized design and analysis of preclinical intervention studies in vivo*. *Sci Rep*. 2016 Aug 2;6:30723.
- 12.531 Laajala TD, et al. *Improved statistical modeling of tumor growth and treatment effect in preclinical animal studies with highly heterogeneous responses in vivo*. *Clin Cancer Res*. 2012 Aug 15;18(16):4385-96.

\* = Equal contribution

## Researcher

### Awards & Funding

Participated in two DREAM challenges resulting in DREAM 9.5 mCRPC Challenge Top Performer (2015) & DREAM Anti-PD1 Response Prediction Challenge Top Performer (2021, ongoing). Rewarded Elias Tillandz -prize 2017 (best scientific publication in Turku BioCity) as first author; again in 2019 as non-first author.

FICAN Cancer Researcher (Finnish Cancer Institute, 2020-2022); NIH grant "Curated prostate cancer data for novel and reproducible prognostic modeling" (2020-2022); Finnish Cultural Foundation (central fund 2014, VS-regional 2018, 2019); DRDP doctoral programme (2014); NIH/NCI DREAM/mCRPC-sponsorship (2016).

### Overview

Brief statistics 04/2022: Citation count 1168 & h-index 17 (Google Scholar); ResearchGate score 34.03 (best >10% quantile) & h-index 15; 30+ peer-reviewed articles in PubMed, of which 5+ in >10 IF journals; Stack Overflow best >25% quantile, >3% quantile in R.