

# Unravelling vascular tumors : combining molecular and computational biology

IJzendoorn, D.G.P. van

### Citation

IJzendoorn, D. G. P. van. (2020, January 16). Unravelling vascular tumors : combining molecular and computational biology. Retrieved from https://hdl.handle.net/1887/82754

Version:	Publisher's Version
License:	<u>Licence agreement concerning inclusion of doctoral thesis in the</u> <u>Institutional Repository of the University of Leiden</u>
Downloaded from:	https://hdl.handle.net/1887/82754

Note: To cite this publication please use the final published version (if applicable).

Cover Page



## Universiteit Leiden



The handle <u>http://hdl.handle.net/1887/82754</u> holds various files of this Leiden University dissertation.

Author: IJzendoorn, D.G.P. van

**Title:** Unravelling vascular tumors : combining molecular and computational biology **Issue Date:** 2020-01-16

### Stellingen

Behorend bij het proefschrift

#### Unravelling Vascular Tumors: Combining Molecular and Computational Biology

- 1. Truncation of the FOS-protein, resulting from different translocations, drives the tumorigenesis of epithelioid hemangioma. *This thesis.*
- 2. Epithelioid hemangioma occurring at multiple locations is a result of locoregional metastatic spread. *This thesis.*
- 3. CRISPR/Cas9 is an excellent genome editing tool to generate models for tumors driven by balanced translocations. *This thesis.*
- 4. Pseudomyogenic hemangioendothelioma displays predominant endothelial differentiation, therefore endothelial cells are the best basis for in vitro models of this tumor. *This thesis.*
- 5. Open-access and well annotated next-generation sequencing datasets enable the use of analysis techniques such as machine learning to draw clinically relevant conclusions. *This thesis.*
- For many types of cancer, gene specific fusions remain to be discovered that could give insight into the tumorigenesis. Mertens, F., Johansson, B., Fioretos, T., and Mitelman, F. (2015). The emerging complexity of gene fusions in cancer. Nat. Rev. Cancer 15, 371–381.
- 7. Endothelial cells derived from Human Induced Pluripotent Stem Cells display the morphology and phenotype of human endothelial cells, and can therefore be used as a replacement for Human Umbilical Vein Endothelial Cells. Orlova, V. V., Van Den Hil, F.E., Petrus-Reurer, S., Drabsch, Y., Ten Dijke, P., and Mummery, C.L. (2014). Generation, expansion and functional analysis of endothelial cells and pericytes derived from human pluripotent stem cells. Nat. Protoc. 9, 1514–1531.
- 8. Computational biology -a core part of biomedical research- is essential for understanding biology. Markowetz, F. (2017). All biology is computational biology. PLoS Biol. 15.
- 9. Artificial intelligence in digital pathology will become a valuable tool to aid the pathologist in diagnostics. Bera, K., Schalper, K.A., Rimm, D.L., Velcheti, V., and Madabhushi, A. (2019). Artificial intelligence in digital pathology — new tools for diagnosis and precision oncology. Nat. Rev. Clin. Oncol.