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Development of bioinformatic tools and application of novel statistical methods in genome-wide analysis

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Document Version

Publisher's PDF, also known as Version of record

Publication date:
2017

[Link to publication in University of Groningen/UMCG research database](#)

Citation for published version (APA):

van der Most, P. J. (2017). *Development of bioinformatic tools and application of novel statistical methods in genome-wide analysis*. [Groningen]: University of Groningen.

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**Development of bioinformatic tools and
application of novel statistical methods in
genome-wide analysis**

Peter Johannes van der Most

The research reported in this thesis has been performed at the Department of Epidemiology, University Medical Center Groningen, University of Groningen, Groningen, the Netherlands.

The printing of this thesis was financially supported by the University of Groningen, the University Medical Center Groningen and the research institute Science in Healthy Ageing and healthcaRE (SHARE).

ISBN: 978-94-034-0148-5 (printed version)

ISBN: 978-94-034-0147-8 (electronic version)

Cover and layout design: Iliana Boshoven-Gkini

<http://www.AgileColor.com>

Printed by: Ridderprint

<https://www.ridderprint.nl>

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university of
 groningen

**Development of bioinformatic tools and
 application of novel statistical methods in
 genome-wide analysis**

PhD thesis

to obtain the degree of PhD at the
 University of Groningen
 on the authority of the
 Rector Magnificus Prof. E. Sterken
 and in accordance with
 the decision by the College of Deans.

This thesis will be defended in public on

Wednesday 8 November 2017 at 12.45 hours

by

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