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

van der Most, Peter Johannes

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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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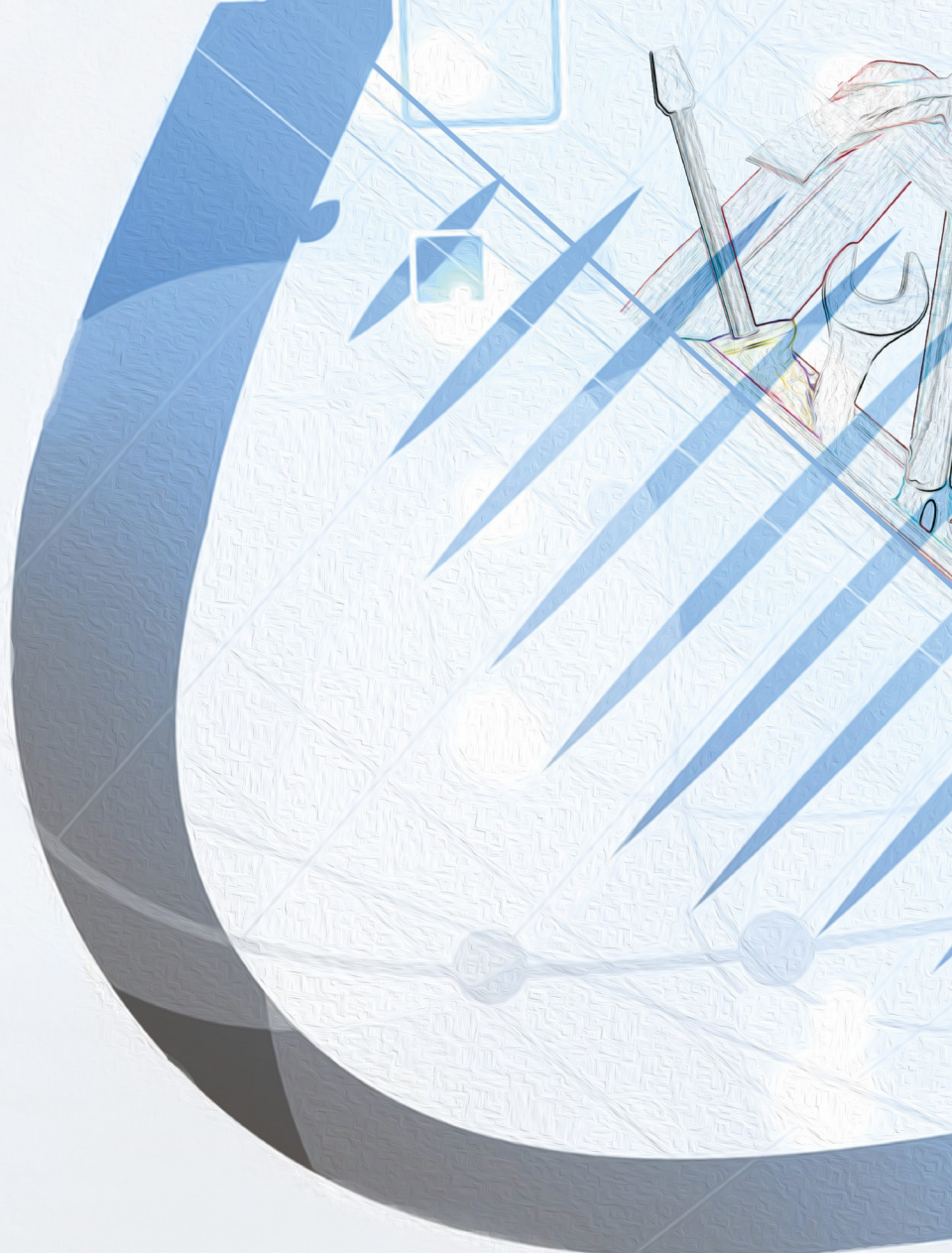
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