Mapping of EGFR treatment effects and uncovering DNA repair mechanisms using quantitative proteomics

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The studies described in this thesis were performed at the Department of Medical Oncology of the University Medical Center Groningen in Groningen, The Netherlands.

ISBN: 978-94-034-0298-7
Cover: JoWeir.com
lay-out design: H.R. de Boer & N. van den Tempel
Printing: Ipskamp Printing

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Mapping of EGFR treatment effects and uncovering DNA repair mechanisms using quantitative proteomics

Proefschrift

ter verkrijging van de graad van doctor aan de
Rijksuniversiteit Groningen
op gezag van de
rector magnificus prof. dr. E. Sterken
en volgens besluit van het College voor Promoties.

De openbare verdediging zal plaatsvinden op

woensdag 13 december 2017 om 12:45 uur

door

Harmen Rudolf de Boer

geboren op 16 mei 1986
te Oosterhout

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