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Unraveling proteoform complexity by native liquid chromatography-mass spectrometry

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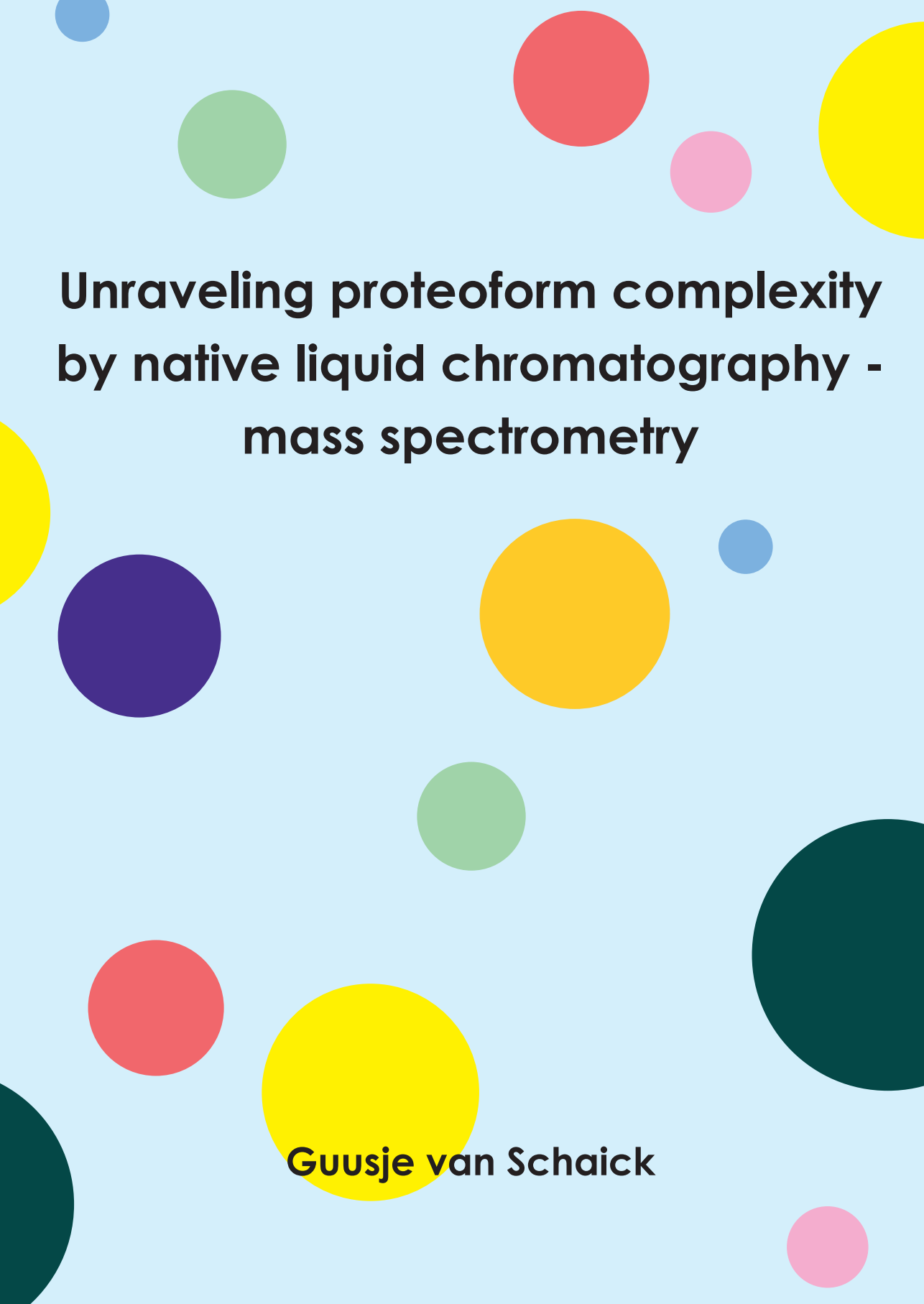
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