

opm: An R Package for Analysing OmniLog® Phenotype MicroArray Data

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Abstract

here some abstrac-text. This is the tutorial of **opm** in the version of September 2, 2013.

Keywords: Bootstrap, Cell Lines, **grofit**, Growth Curves, **lattice**, Metadata, Microbiology, Respiration Kinetics, Splines, YAML, JSON, CSV.

1. Introduction

1.1. Preamble for “eager to start” readers

2. Acknowledgements

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