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The opm R package facilitates management, visualisation and statistical analysis of PM data and similar data such as growth curves. Raw measurements can easily be input into R, combined with relevant meta-information and accordingly analysed. The kinetics can be aggregated by estimating curve parameters using several methods. Some of them

~~opm: An R package for analysing phenotype microarray and ...~~

The opm package for R is a comprehensive software for analysing phenotype microarray and growth-curve data. For more information,

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see the opm R-Forge site or the main tutorial for opm. Installation. The package can be used on Windows, Mac and Linux/UNIX systems. As a prerequisite, one needs to obtain the statistical computing environment R.

~~OPM and its Helper Packages — gocker.org~~
Summary: opm is an R package designed to analyse multidimensional OmniLog® phenotype microarray (PM) data. opm provides management, visualization and statistical analysis of PM data, including curve-parameter estimation and discretization, dedicated and customizable plots, metadata

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management, automated generation of textual and tabular reports, mapping of substrates to databases, batch conversion of files and export to phylogenetic software in the YAML markup language.

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~~(PDF) opm: An R Package for Analysing
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Package for analysing OmniLog(R) phenotype
microarray (PM) data, as well as similar
kinetic data such as growth curves.

opm.package: The 'opm' package in opm:
Analysing Phenotype Microarray and Growth
Curve Data

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SUMMARY: opm is an R package designed to
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In opm: Analysing Phenotype Microarray and Growth Curve Data. Description Details See

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Also Examples. Description. Classes whose members can be generated and manipulated by an opm user: OPM, OPMA, OPMD, OPMS and MOPMX. Details. OPM is an acronym for 'OmniLog(R)

~~OPM: Real classes of the 'opm' package in
opm: Analysing ...~~

* OPM's regulations permit an agency to set an employee's rate of basic pay based on a single factor after a superior qualifications or special need determination has been made. However, if more than one factor supports the pay-setting decision, the agency should document all factors that apply. A

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determination based on more than one factor may ...

~~Superior Qualifications and Special Needs Pay
... — OPM.gov~~

Welcome to opm.gov. The U.S. Office of Personnel Management (OPM) announced today that for plan year 2021, the average total premiums for current non-Postal employees and retirees enrolled in plans under the Federal Employees Health Benefits (FEHB) Program will increase 3.6 percent.

~~OPM.gov~~

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The generic accessor functions `coefficients`, `fitted.values`, `residuals`, `logLik`, and `df.residual` can be used to extract various useful features of the value returned by `opm`. Note Dummy time variables exist as an additional column for each wave of data, excluding the first and second wave (i.e., at `\(t=0\)` and `\(t=1\)` using the terminology from ...

~~opm function | R Documentation~~

Tools for analysing OmniLog (R) Phenotype Microarray (PM) data as produced by the device distributed by BIOLOG Inc. and similar

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kinds of data such as growth curves are available in the opm package. The package opmextra contains functions for special purposes and thus have not been placed in opm.

~~R-Forge: opm and opmdata: R Development Page~~
General-purpose optimization. General-purpose optimization wrapper function that calls other R tools for optimization, including the existing optim () function. Also tries to unify the calling sequence to allow a number of tools to use the same front-end. Note that optim () itself allows Nelder--Mead, quasi-

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Newton and conjugate-gradient algorithms as well as box-constrained optimization via L-BFGS-B.

~~opm function | R Documentation~~

Summary: `opm` is an R package designed to analyse multidimensional OmniLog® phenotype microarray (PM) data. `opm` provides management, visualization and statistical analysis of PM data, including curve-parameter estimation and discretization, dedicated and customizable plots, metadata management, automated generation of textual and tabular reports, mapping of substrates to

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databases, batch conversion of files and export to phylogenetic

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SUMMARY: opm is an R package designed to analyse multidimensional OmniLog® phenotype microarray (PM) data. opm provides management, visualization and statistical analysis of PM data, including...

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The opm R package facilitates management,

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visualization and statistical analysis of Phenotype Microarray data. Raw measurements can be easily input into R, combined with relevant meta-information and accordingly analysed. The kinetics can be aggregated by estimating curve parameters using several methods. Some of them have been specifically

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~~U.S. Office of Personnel Management — OPM.gov~~

The opm R package facilitates management, visualization and statistical analysis of Phenotype Microarray data. Raw measurements can be easily input into R, combined with relevant meta-information and accordingly analysed. The kinetics can be aggregated by estimating curve parameters using two distinct methods.

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SUMMARY opm is an R package designed to
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