

# Package ‘lipidmaps’

March 24, 2017

**Type** Package

**Title** R Interface to the LIPID MAPS RESTful API

**Version** 0.1.0

**Author** Tom Wilson <tpw2@aber.ac.uk>

**Maintainer** Tom Wilson <tpw2@aber.ac.uk>

**Description** Simple searching of the LIPID MAPS database using Molecular Formula or an InChiKey as an input.

**Imports** RCurl, jsonlite

**Suggests** testthat

**License** GPL (>= 3)

**LazyData** TRUE

**RoxygenNote** 5.0.0

**NeedsCompilation** no

## R topics documented:

LMsearch . . . . .	<a href="#">1</a>
<b>Index</b>	<a href="#">3</a>

---

LMsearch	<i>Seach LIPID MAPS</i>
----------	-------------------------

---

## Description

Retrieve data from the LIPID MAPS database using either an InChiKey or Molecular Formula as an input

## Usage

LMsearch(x)

**Arguments**

x a character of a valid InChiKey or Molecular Formula

**Value**

a data.frame containing the following fields for each returned entry

**name** Systematic name

**id** LIPID MAPS database id

**mz** Monoisotopic accurate mass

**inchi** InChi Code

**inchi\_key** InChiKey

**smiles** Smiles

**formula** Molecular formula

**core\_class** core

**main\_class** main

**sub\_cls** sub

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
  LMsearch("C18H32O2")  
  LMsearch("ZKRPGPZHULJLKJ-JHRQRACZSA-N")  
  
## End(Not run)
```

# Index

LMsearch, [1](#)