## Package 'CCl4'

April 13, 2022	
itle Carbon Tetrachloride (CCl4) treated hepatocytes	
Version 1.32.0	
uthor Audrey Kauffmann, Wolfgang Huber	
<b>Description</b> NChannelSet for rat hepatocytes treated with Carbon Tetrachloride (CCl4) data from LGC company.	
faintainer Audrey Kauffmann <audrey@ebi.ac.uk></audrey@ebi.ac.uk>	
icense Artistic-2.0	
<b>Depends</b> R (>= 2.6.0), Biobase (>= 2.5.5), limma	
iocViews ExperimentData, Genome, Rattus_norvegicus_Data, MicroarrayData	
it_url https://git.bioconductor.org/packages/CCl4	
it_branch RELEASE_3_14	
it_last_commit 60e6be4	
it_last_commit_date 2021-10-26	
Pate/Publication 2022-04-13	
R topics documented:	
CC14	1
ndex	3
CC14 Carbon Tetrachloride (CCl4) treated hepatocytes	_
Description	
These datasets contain raw intensities from Agilent <i>Rat Whole Genome</i> microarrays. Total RN from rat hepatocytes treated by Carbon Tetrachloride or DMSO was extracted. Purified RNA fro	

9.7, degraded RNA RIN values of 5.0 and 2.5 were achieved.

Carbon Tetrachloride treated cells was degraded by ribonuclease A in order to study the effect of RNA quality on the expression data. From an initial Agilent RIN (RNA Integrity Number) value of

2 CC14

## Usage

```
data("CC14")
data("CC14_RGList")
```

### **Format**

 ${\tt CC14}$  is a  ${\tt NChannelSet}$  with 44290 probes and 18 arrays.  ${\tt CC14\_RGList}$  is an  ${\tt RGList}$  with the same data.

#### Source

Carole Foy, Holger Laux, Timothy Wilkes from LGC company in London, UK.

## **Examples**

```
data("CC14")
show(CC14)

data("CC14_RGList")
show(CC14_RGList)
```

# **Index**

```
* datasets
CC14, 1
CC14, 1
CC14_RGList (CC14), 1
NChannelSet, 2
RGList, 2
```