# **R** documentation

of 'trainChemPC.Rd'

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trainChemPC trainChemPC

## Description

This function performs learning step of GP and EI.

# Usage

trainChemPC( trainData, targetVector)

# Arguments

trainData specifies a data frame including an array of a training data with

dimension of n\*m. This data is used in to find a hyperlog vector.

targetVector targetVector is a one dimensional array with n rows which is equal to number

of rows in trainData.

#### **Details**

This function performs training step of GP or EI by finding a loghyper using gpr package. A loghyper can be used to predict potent compounds.

#### Value

It returns a vector that holds a calculated loghyper.

#### Author(s)

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#### References

1. Predicting Potent Compounds via Model-Based Global Optimization, Journal of Chemical Information and Modeling, 2013, 53 (3), pp 553-559, M Ahmadi, M Vogt, P Iyer, J Bajorath, H Froehlich. 2. Software MOE is used to calculate the numerical descriptors in data sets. Ref: http://www.chemcomp.com/MOE-Molecular\_Operating\_Environment.htm 3. ChEMBL was the source of the compound data and potency annotations in data sets. Ref: https://www.ebi.ac.uk/chembl/

## **Examples**

```
library(gpr)
library(SimuChemPC)
a = as.data.frame(array(1:10, dim=c(2,5)))
b = array(1:2)
myloghyper = trainChemPC( a, b)
```

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