## Package 'healthyFlowData'

### February 4, 2025

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Type Package		
Title Healthy datase	et used by the flowMatch package	
Version 1.44.0		
<b>Date</b> 2013-10-29		
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<b>Description</b> A heal	thy dataset with 20 flow cytometry samples used by the flowMatch package.	
License Artistic-2.0		
biocViews FlowCyt	tometryData	
<b>Depends</b> R (>= 2.1	5.0), flowCore	
Imports methods		
git_url https://git.bi	ioconductor.org/packages/healthyFlowData	
git_branch RELEA	ASE_3_20	
git_last_commit 4d	lb75f2	
git_last_commit_da	nte 2024-10-29	
Repository Biocon	ductor 3.20	
Date/Publication 2	025-02-04	
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hd	Healthy donor (HD) dataset	_
Description		
_	ntains 20 samples from four healthy individuals.	
Usage		
data(hd)		

2 hd

#### **Format**

A flowSet named hd.flowSet containing 20 FC samples. Each sample is represeted by a flowFrame.

#### **Details**

Peripheral blood mononuclear cells (PBMC) were collected from four healthy individuals. Each sample was divided five repplicates and each replicate was stained using labeled antibodies against CD3, CD4, CD8, and CD19 protein markers. Therefore we have total 20 samples from four healthy subjects. Each sample was compensated and transformed in order to stabilize per-channel variance. Each sample is then gated on the forward and side scatter to identify lymphocites. Hence the samples contain only lymphocites cells.

This is a part of a larger dataset consisting of 65 samples. Please ask the author if you would like to obtain the complete dataset.

#### References

Azad, Ariful and Khan, Arif and Rajwa, Bartek and Pyne, Saumyadipta and Pothen, Alex (2013); "Classifying Immunophenotypes With Templates From Flow Cytometry", proceedings of ACM BCB, Washington DC.

#### **Examples**

data(hd)
summary(hd.flowSet)

# Index

```
* datasets
hd, 1
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```