

Resource Summary Report

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NS-Forest version 2

DOI:10.17504/protocols.io.un7evhn

Type: Protocol

Proper Citation

Brian Aeversmann, Richard Scheuermann 2018. NS-Forest version 2. protocols.io
[dx.doi.org/10.17504/protocols.io.un7evhn](https://doi.org/10.17504/protocols.io.un7evhn)

Protocol Information

URL: <https://dx.doi.org/10.17504/protocols.io.un7evhn>

Authors: Brian Aeversmann, Richard Scheuermann

Group: Human Cell Atlas Method Development Community

Summary: NS-Forest is an algorithm that determines the minimum set of genes that are necessary and sufficient to define a cell type cluster derived from single cell RNAseq expression data. Development and stable releases can be found at :
<https://github.com/JCVenterInstitute/NSForest>

Affiliations: J. Craig Venter Institute, J. Craig Venter Institute

Version: 1

Publication Date: 2018

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