

# Validation of CellScanner 2.0

Data sets described in: <https://www.nature.com/articles/s43705-022-00123-6>

Monocultures mixed in known ratios in vitro in stationary phase

Monocultures paired with corresponding cocultures, e.g. RI\_5.fcs with FP\_95.fcs for co-culture RI-FP\_5\_95.fcs

Blank file: control\_PBS.fcs

Data sets available at: <https://flowrepository.org/id/FR-FCM-Z3TM>

## Settings

CellScanner 2.0 run with default values:

1000 events

Scaling constant: 150

number of neighbors: 50

UMAP min dist: 0

number of required blank neighbors: 20

number of required non-blank neighbors: 25

folds: 0

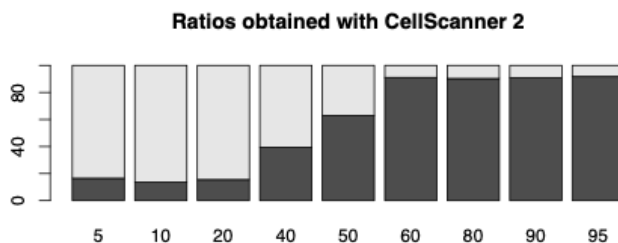
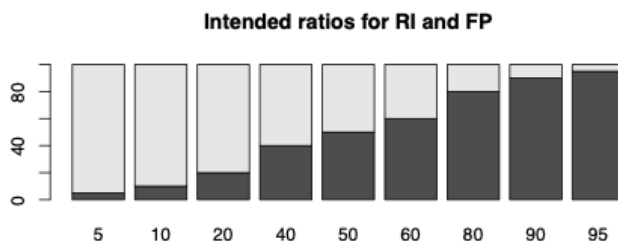
epochs: 50

early stopping patience: 10

batch size: 32

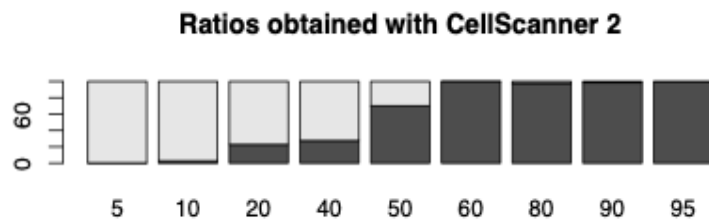
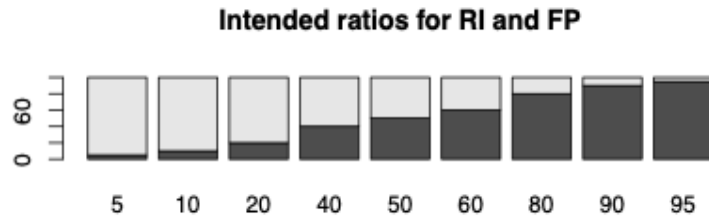
## Result

***Roseburia intestinalis* (RI) and *Faecalibacterium prausnitzii* (FP)**



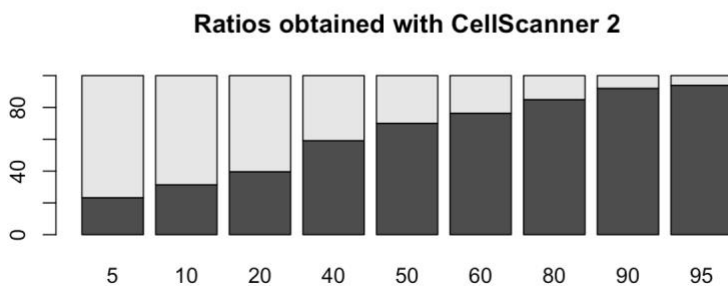
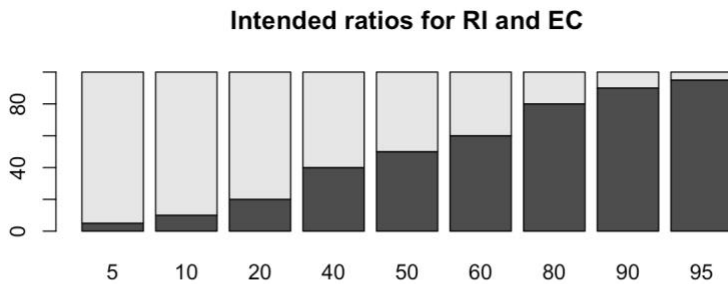
### ***Roseburia intestinalis* (RI) and *Faecalibacterium prausnitzii* (FP) with uncertainty filtering**

Uncertainty filtering: events with entropy above a threshold set automatically are removed

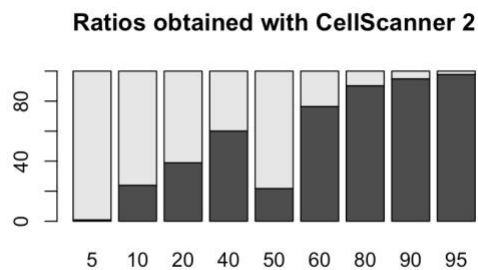
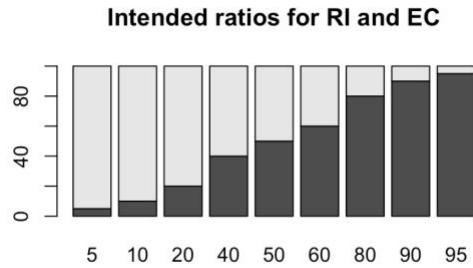


### ***Roseburia intestinalis* (RI) and *Escherichia coli* (EC)**

This is a positive control since *E. coli* is labelled with mCherry.



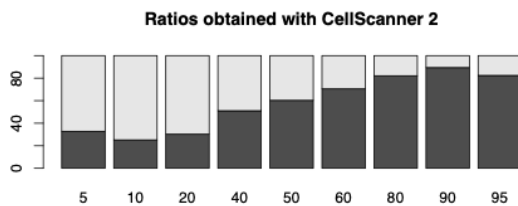
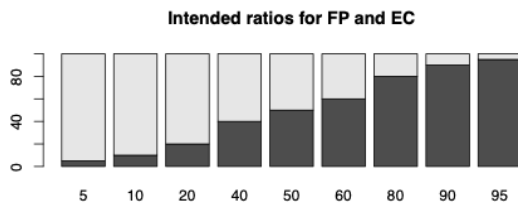
### ***Roseburia intestinalis* (RI) and *Escherichia coli* (EC) with uncertainty filtering**



The outlier at 50:50 in the results with uncertainty filtering goes to show that this filtering step is only effective if uncertain events are evenly distributed between species. Here, *E. coli*, being labelled with mCherry, had only few uncertain events, and thus *R. intestinalis* was over-represented in the removed events, which biased the predicted proportion.

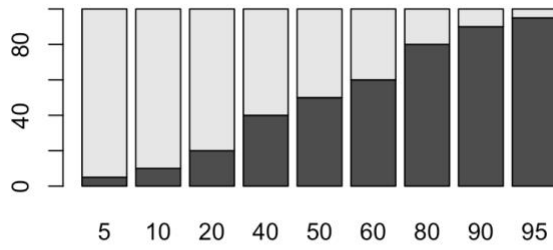
### ***Faecalibacterium prausnitzii* (FP) and *Escherichia coli* (EC)**

This is another positive control since *E. coli* is labelled with mCherry.



### ***Faecalibacterium prausnitzii* (FP) and *Escherichia coli* (EC) with uncertainty filtering**

**Intended ratios for FP and EC**



**Ratios obtained with CellScanner 2**

