

Pipeline

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## Probe filtering

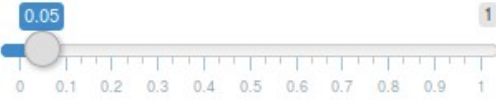
Here you can filter the probes with regard to p-value and limit.

This is done by lumi function `detectionCall` with Th parameter as p-value and filtering this data by limit value.

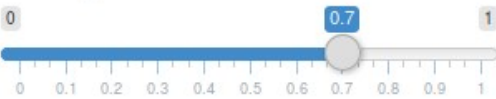
☒ Enabled

Default values are 0.05 for p-value and 0.7 for filtering limit.

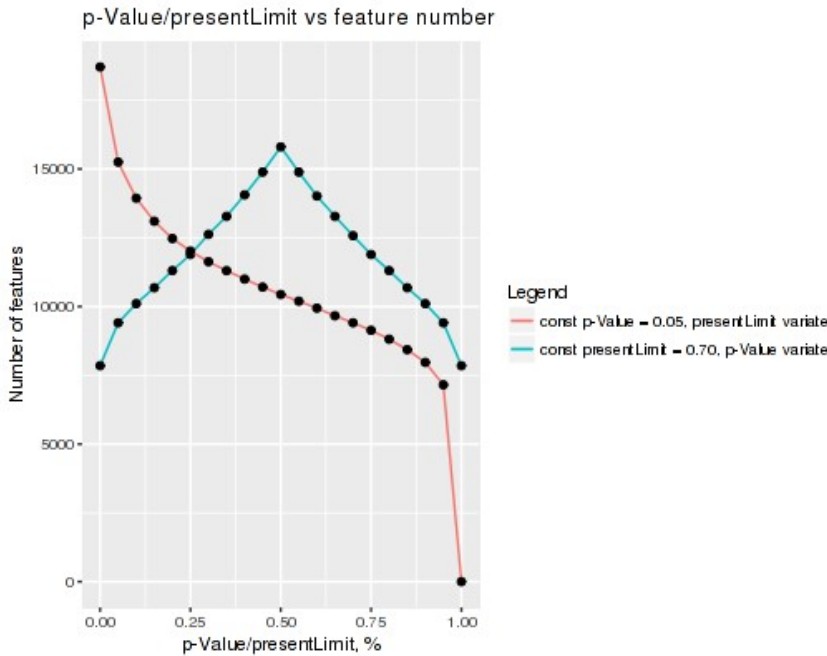
### P-value



### Filtering limit



☒ Show plot



Continue

Previous step

To final step

## Project information:

### Primary dataset information

Dataset: Uterus HiScan  
prospective  
168 samples with 29377 features

### Settings

Outlier removal: Enabled  
Exclude control-case transitions:  
Not enabled  
Background correction: Enabled  
P value: 0.05  
Filtering limit: 0.7  
Normalization method: Not  
enabled  
Include questionnaire variables:  
Not enabled

### Dataset properties after processing

Samples: 160  
Features: 9408