

Functional analysis



Miscellaneous

Quality control

Relative
quantification

MS
stats

artmsQualityControlEvidenceBasic()

artmsQualityControlEvidenceExtended()

artmsQualityControlsSummaryExtended()

antms@uantificati0n()

artmsAnalysisQuantifications()

artmsEvidenceToSAINTq()

artmsEvidenceToSaintExpress()

artmsPhosphateOutput()

artmsPhotonOutput()

etc



Input Files

keys.txt



evidence.txt

artMS: BASIC QUALITY CONTROL (evidence.txt based)

>> MERGING FILES

-- Plot: correlation matrices

---- by Biological replicates

---- by Conditions

-- Plot: intensity stats

---- PH PROCESSED

-- Plot: PTM PH stats

<< Basic quality control analysis completed!

artMS

**Analytical R Tools
for Mass Spectrometry**

<http://artms.org>



artMS

Analytical R Tools
for Mass Spectrometry

<http://artms.org>



Input Files

 evidence.txt

keys.txt

Quality control

artmsQualityControlEvidenceBasic()

artmsQualityControlEvidenceExtended()

artmsQualityControlSummaryExtended()

 Relative
quantification

artmsQuantification()

Functional analysis

artmsAnalysisQuantifications()

Miscellaneous

artmsEvidenceToSAINTq()
artmsEvidenceToSaintExpress()
artmsPhosphateOutput()
artmsPhotonOutput()
etc

Console

Terminal x

```
> library(artMS)
> artmsQualityControlEvidenceBasic(evidence_file = "evidence.txt",
+                                   keys_file = "keys.txt")
```

artMS: BASIC QUALITY CONTROL (evidence.txt based)

>> MERGING FILES

-- Plot: correlation matrices

---- by Biological replicates

---- by Conditions

-- Plot: intensity stats


---- PH PROCESSED

-- Plot: PTM PH stats

<< Basic quality control analysis completed!

artMS

Analytical R Tools
for Mass Spectrometry

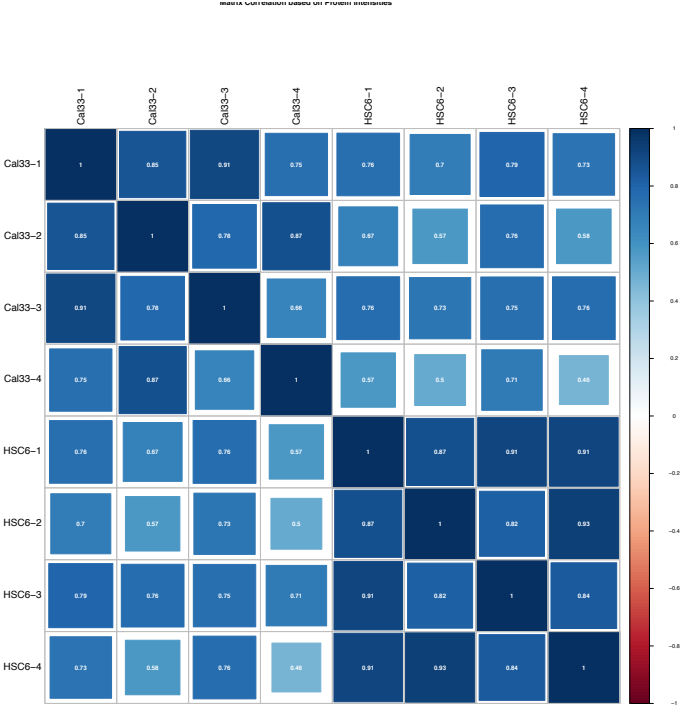
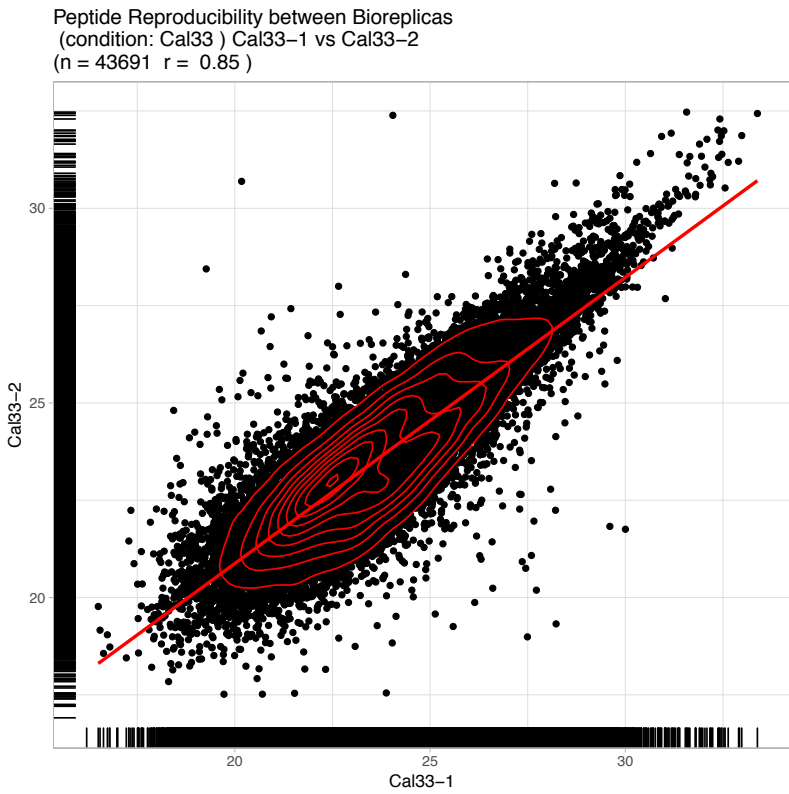
<http://artms.org>


Input Files

Mo

evidence.txt

keys.txt



Quality control

artmsQualityControlEvidenceBasic()

- Reproducibly
- MS Intensity
- Spectral Counts
- Contaminants
- PTMs

MS stats

Relative quantification

Functional analysis

Miscellaneous

