

# History:

V0.42 – 2013-11-12

added duplicated peptide sequence statistic (dynamic exclusion window check)

fixed some axis labels

V0.41 – 2013-10-24

first official release

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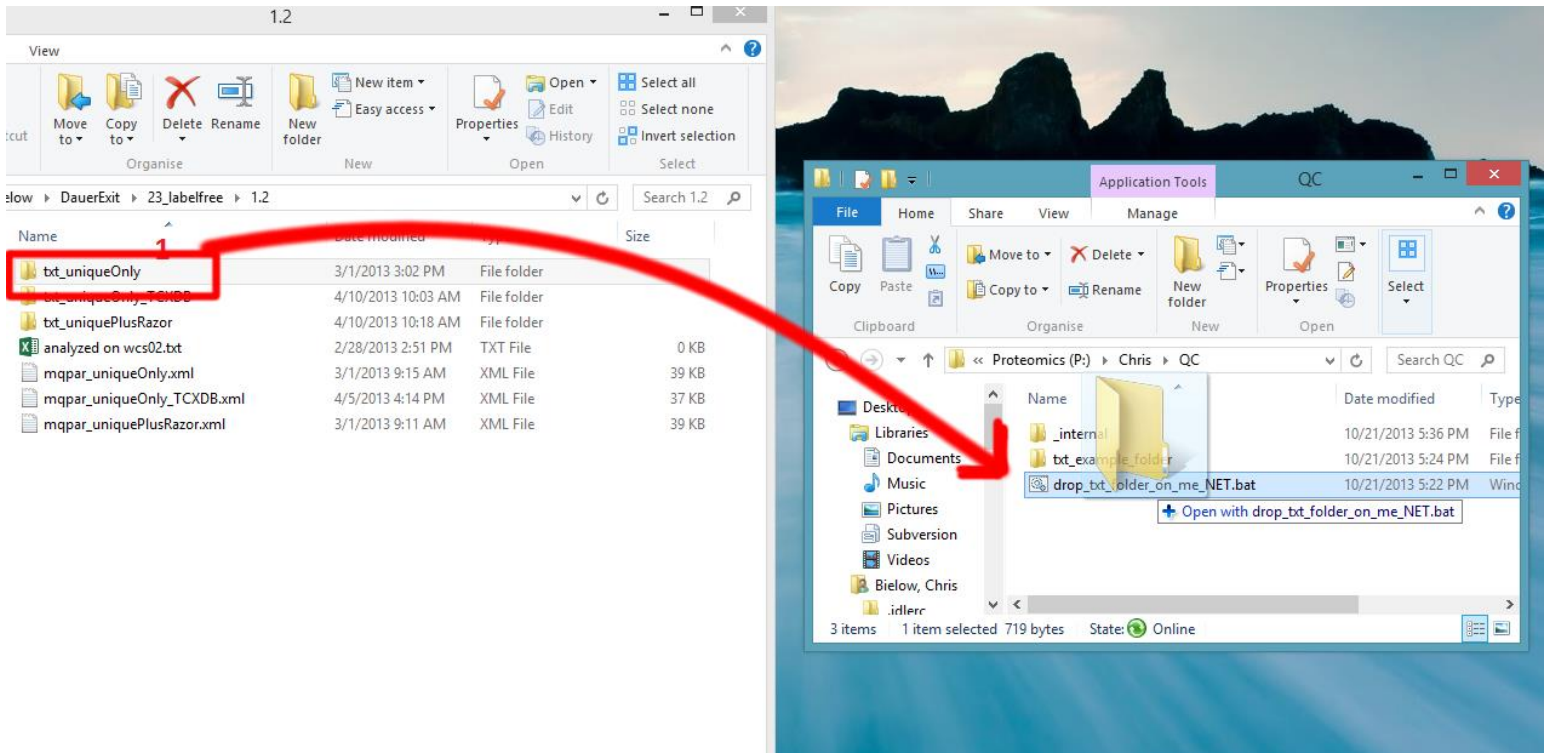
# Create a Quality control report for your MaxQuant output

## Requirements:

Any Windows System (no R required!)

## Steps

1. open a Windows Explorer and find your txt folder
2. open **another** Windows Explorer window with P:\Chris\Exchange\QC (where P==proteomics network share)
3. drag'n'drop the txt folder onto drop\_txt\_folder\_on\_me\_NET.bat



4. A console will pop up and show stuff. Wait... if all went well, there is a pdf report within your txt folder now. It has a version attached to it (here v0.41), which will increase every time there is an update (big change in version number means big change in functionality)

```
C:\Windows\system32\cmd.exe

Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Reading file msms.txt ...
Keeping 2 of 53 columns!
Read 385641 entries from msms.txt .
Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Scale for 'x' is already present. Adding another scale for 'x', which will replace the existing scale.
Reading file msmsScans.txt ...
Keeping 5 of 37 columns!
Read 849824 entries from msmsScans.txt .
Loading required package: plyr
Report file created at

S:\chielow\DauerExit\23_labelfree\1.2\txt_uniqueOnly\report_v0.41.pdf

There were 20 warnings (use warnings()) to see them
Press any key to continue . . .
```

If you have suggestions on how to improve the report (other metrics/plots, axis labels etc), let me know.