

```
2023-02-27 12:43:55 [INFO] - Matching-between-runs: ipsc-50C-p3-
wt-4...
2023-02-27 12:43:55 [INFO] - Calculating correlations between all
other runs...
2023-02-27 12:43:55 [INFO] - Reading index from F:
\Proteomics\ipSC\ipSC-p3-50C\ipsc-50C-p3-wt-4.quantindex
2023-02-27 12:43:55 [INFO] - Transferring ipsc-50C-p3-ko-1...
2023-02-27 12:43:55 [INFO] - Transferring ipsc-50C-p3-ko-2...
2023-02-27 12:43:56 [INFO] - Transferring ipsc-50C-p3-ko-3...
2023-02-27 12:43:56 [INFO] - Transferring ipsc-50C-p3-ko-4...
2023-02-27 12:43:57 [INFO] - Transferring ipsc-50C-p3-wt-1...
2023-02-27 12:43:57 [INFO] - Transferring ipsc-50C-p3-wt-2...
2023-02-27 12:43:58 [INFO] - Transferring ipsc-50C-p3-wt-3...
2023-02-27 12:43:59 [INFO] - Training LDA models for all matched
features.
2023-02-27 12:43:59 [INFO] - Standardized coefficients:
2023-02-27 12:43:59 [INFO] - log10(intensity): 0.25600866747263157
2023-02-27 12:43:59 [INFO] - log10(KL): -0.03745909813258587
2023-02-27 12:43:59 [INFO] - abs(ppm): -0.8050711419740741
2023-02-27 12:43:59 [INFO] - RT diff: -0.5338311395466666
2023-02-27 12:43:59 [INFO] - Fitting a mixture model...
2023-02-27 12:44:00 [INFO] - Estimating match-between-runs FDR...
2023-02-27 12:44:02 [INFO] - With ion FDR 0.010000, ion probability
threshold is 0.979000
2023-02-27 12:44:02 [INFO] - With peptide FDR 1.000000, peptide
probability threshold is -0.000100
2023-02-27 12:44:02 [INFO] - With protein FDR 1.000000, protein
probability threshold is -0.000100
2023-02-27 12:44:02 [INFO] - Updating Philosopher's tables...
2023-02-27 12:44:15 [INFO] - Combining experiments and estimating
protein intensity...
2023-02-27 12:44:26 [INFO] - Done!
Process 'IonQuant' finished, exit code: 0
```