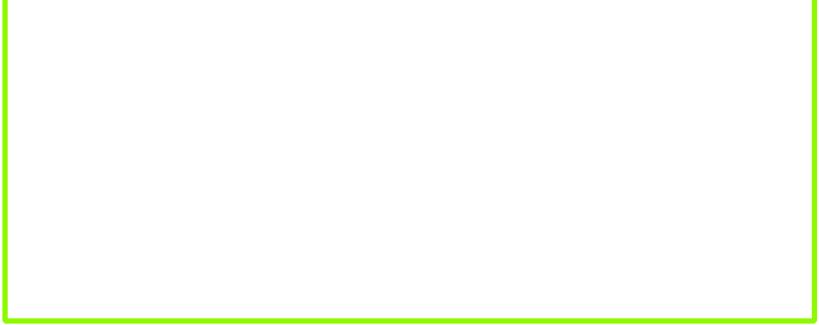
```
config-phglobal.yaml
                           ×
      files:
♦
        evidence: evidence.txt
        keys: keys.txt
 3
 4
        contrasts: contrast.txt
 5
        summary: summary.txt
 6
        output: phglobal_results/phglobal-results.txt
 7
      qc:
        basic: 1
 8
 9
        extended: 1
        extendedSummary: 1
10
      data:
11
        enabled: 1
12
        fractions:
13
           enabled: 0
14
15
        silac:
           enabled: 0
16
        filters:
17
           enabled: 1
18
           contaminants: 1
19
20
           protein_groups: remove
        modifications: PH sample_plots: 1
21
22
      msstats:
23
24
        enabled: 1
25
        msstats_input:
        profilePlots: none
normalization_method: equalizeMedians
26
27
        normalization_reference:
28
        summaryMethod: TMP
29
30
        censoredInt: NA
        cutoffCensored: minFeature
MBimpute: 1
31
32
         feature_subset: all
33
34
      output_extras:
        enabled: 1
35
        annotate:
  enabled: 1
36
37
           species: HUMAN
38
        plots:
39
           volcano: 1
40
           heatmap: 1
41
           LFC: -0.58 0.58
42
           FDR: 0.05
43
44
          heatmap_cluster_cols: 0
           heatmap_display: log2FC
45
```

∢▶

config.yaml

Configuration





Relative Quantification

Outputs / extras



Quality control



Relative quantification

Functional analysis



artMS

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