# **Accuracy Check, CrIrCntRepMnTots.Rdata**

## See hand-written sheet for family tree

Notes on checks, starting with the base document ✓✓✓✔︎✔︎✔︎✔︎

R 100x\_RawCount\_R.csv and 400x\_RawCount\_R.csv: Raw data, checked

R propCntd\_100\_final.xlsx and .csv

R propCntd\_400\_final.xlsx and .csv

R raw100\_final, raw400\_final.Rdata

vol100.Rdata: Check YBP2 diatom centric 40x40µm, counts:40, pres factor: 0.914, volume settled, ml : 20, proportion of sample counted: 0.5, volume per cell, µm3 : 50265 total volume of all 40 diatoms µm3 : 201619