

User: cgrgicak
Analysis Run: 13-Dec-2017 00:16:33 EST

Case Number:

Sample ID: A02_RD14-0003-31_32-1;1-M2c-0.03GF-Q2.0_01.25sec.hid

Sample File Name: RD14-0003_GF_25sec_GM_SE33F_2-5P.csv

Comments:

Calibration Name: C:\Users\cgrgicak\Desktop\RD14-0003(GF 25s)_Validation\RD14-0003(GF 25s)_Validation\CP_RD14 GF_25s_GM_F2.zip

Calibration MD5 Hash: f8ab7724feb215a2c856b60e3e314cfc

Population Name: FF_GF CAUC n=343

Population File Name: C:/Users/cgrgicak/Desktop/RD14-0003(GF 25s)_Validation/RD14-0003(GF 25s)_Validation/FF_GF CAUC n=343.csv

Population File MD5 Hash: a89f00d629d5cb216e29b444039d7ac1

Number of People: 343

Kit File: GlobalFiler_v1

Filtered: true

Pull-up Filter Height Ratio: 6.0

Pull-up Filter Size Range: 0.6

Complex Pull-up Filter Height Ratio: 6.0

Complex Pull-up Filter Sister Height Ratio: 50.0

Complex Pull-up Filter Size Range: 0.3

Minus-A Filter Height Ratio: 0.0

Minus-A Filter Size Range: 0.0

Analytical Thresholds:

D3S1358: 1

vWA: 1

D16S539: 1

CSF1PO: 1

TPOX: 1

AMEL: 1

D8S1179: 1

D21S11: 1

D18S51: 1

D2S441: 1

D19S433: 1

TH01: 1

FGA: 1

D22S1045: 1

D5S818: 1

D13S317: 1

D7S820: 1

SE33: 1

D10S1248: 1

D1S1656: 1

D12S391: 1

D2S1338: 1

Number of contributors: 2

Time taken: 1191.488 s

Number of samples: 100000

POI genotype ID: 31

Known genotypes IDs: None

log₁₀(POI LR): 20.3

p-value: 2.88e-34

Number of samples with LR > 1: 14

Pr(LR > 1): 1.40e-05

