# My Projects / Personal IcWGS / 951023c1725b4b52b150c46469121abd

CW



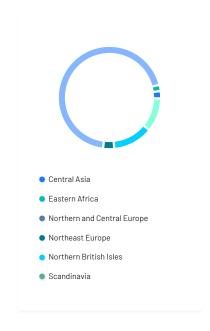
gencove

- MY PROJECTS
- MY FASTOS
- ACCOUNT

### QC Overview

#### VALID FASTQ FORMAT Status Succeeded BASES SEQUENCED, 689.1M MINIMUM Observed value 689, 121, 116 Threshold 330,000,000 Status Succeeded BASES SEQUENCED, 689.1M MAXIMUM 689,121,116 Observed value 19,800,000,000 Threshold Status Succeeded BASES DEDUPLICATED 570.6M AND ALIGNED, MINIMUM 570,580,664 Observed value 330,000,000 Threshold Status Succeeded CONTAMINATION, 0.3% MUMIXAM 0.30% Observed value Threshold 6.00% Status Succeeded KNOWN SITES COVERED BY A 12.5M READ, MINIMUM 12,472,708

## **Ancestry**



#### **Traits**



(?) HELP

LOG OUT

Observed value

Threshold

Status

5,000,000

Succeeded

Alignment, BAI file	<u>_</u>
Alignment, BAM file	<u></u>
Ancestry, JSON file	<u>↓</u>
CNV analysis, CNR file	<u>↓</u>
CNV analysis, CNS file	<u></u>
CNV analysis, PDF file	<u>↓</u>
Input, FASTQ file (R1)	<u></u>
Input, FASTQ file (R2)	<u>↓</u>
Imputation, CSI file	<u></u>
Imputation, TBI file	<u>↓</u>
Imputation, VCF file	<u></u>
Polygenic traits, JSON file	<u></u>