**\( \)** gencove

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MY FASTOS

& ACCOUNT

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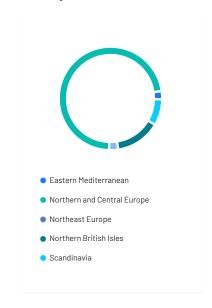
255fea15-fd24-4f50-92b4-dafd51547f57

Has data: 2020-03-07T06:32:13.042954Z
Running: 2020-03-07T06:32:13.563108Z
Succeeded: 2020-03-07T08:04:15.872670Z

## QC Overview

VALID FASTQ FORMAT	
Status	<ul> <li>Succeed</li> </ul>
BASES SEQUENCED, MINIMUM	1.41
Observed value	1,378,242,2
Threshold	330,000,0
Status	<ul> <li>Succeed</li> </ul>
BASES SEQUENCED, MAXIMUM	1.41
Observed value	1,378,242,2
Threshold	19,800,000,0
Status	<ul> <li>Succeed</li> </ul>
BASES DEDUPLICATED AND ALIGNED, MINIMUM	1.01
Observed value	1,042,357,8
Threshold	330,000,0
Status	<ul> <li>Succeed</li> </ul>
CONTAMINATION, MAXIMUM	0.4%
Observed value	0.4
	6.0
Threshold	
Threshold Status	<ul> <li>Succeed</li> </ul>
Status  KNOWN SITES COVERED BY A	• Succeed 21.01
Status  KNOWN SITES	

## Ancestry



## **Traits**



RAW DATA AVAILABILITY

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>