

Analysis Pipelines

This page represents samples with analyses from different pipelines.

Analysis Pipelines

Filter subsets e.g. by pre

Hierarchy Depth:

3 levels



No Selection

- ☐ ADM2: ADM2 (2092 samples, 8114 CNV profiles)
- ☐ Birdseye: Birdseye (270 samples, 2870 CNV profiles)
- ☐ Custom HMM algorithm: Custom HMM algorithm (1552 samples)
- ☐ DRAGEN-CNV: DRAGEN-CNV (3201 samples, 3201 CNV profiles)
- ☐ ELAND (PEM): ELAND (PEM) (1 sample, 16 CNV profiles)
- ☐ HMMSeg: HMMSeg (9 samples, 118 CNV profiles)
- ☐ PEM: PEM (1 sample)
- ☐ VAMP: VAMP (1 sample)
- ☐ era-EnsembleCNV: era-EnsembleCNV (2001 samples, 2001 CNV profiles)
- ☐ era-PennCNV: era-PennCNV (1993 samples, 1993 CNV profiles)
- ☐ era-QuantiSNP: era-QuantiSNP (2001 samples, 2001 CNV profiles)
- ☐ era-iPattern: era-iPattern (1866 samples, 1866 CNV profiles)
- ☐ labelSeg-based calibration: labelSeg-based calibration (16671 samples, 16671 CNV profiles)
- ☐ mrFAST (Read Depth): mrFAST (Read Depth) (3 samples, 16 CNV profiles)

era-EnsembleCNV (era-EnsembleCNV)

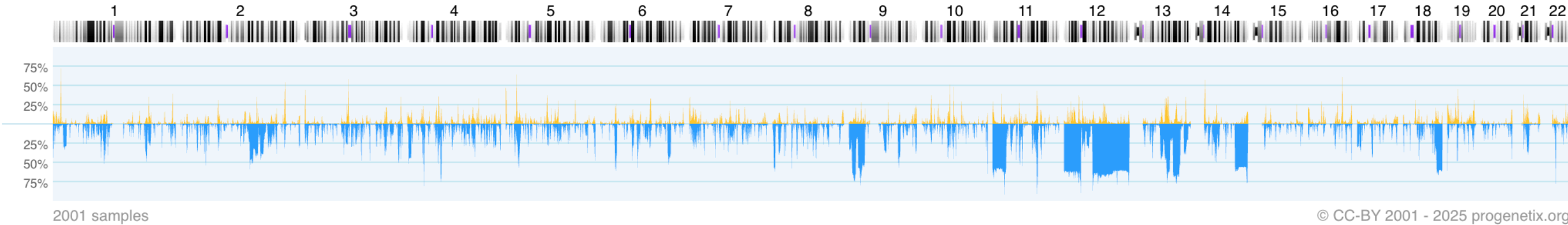
Subset Type

- alphanumeric [era-EnsembleCNV](#)

Sample Counts

- 2001 samples
- 2001 direct era-EnsembleCNV code matches
- 2001 CNV analyses

CNV Histogram



[Download SVG](#) | [Go to era-EnsembleCNV](#) | [Download CNV Frequencies](#)

[Download CNV frequencies](#)

Download CNV frequency data for genomic 1Mb bins.

[Search for era-EnsembleCNV Samples](#)

Select samples through the search form, e.g. by querying for specific genomic variants or phenotypes.