Sample size

Data source

Data type

Histopathological data
Tumor grade, molecular subtype, HER2 and hormone receptor status, miRNA profiles

Clinical data
Treatment modality, cancer stage, cancer-related death, distant metastases, relapse
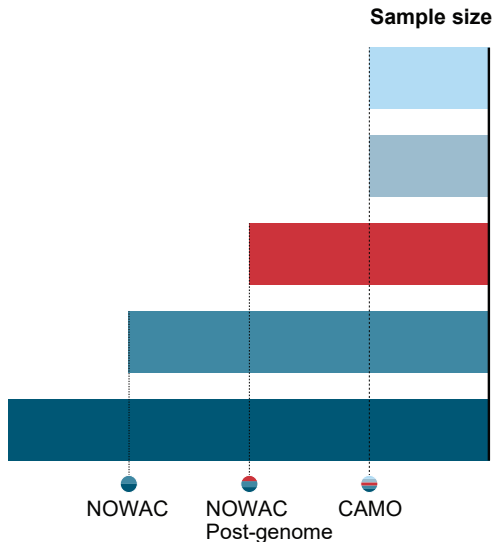
Omics data and biomarkers
Whole-blood mRNA gene expression profiles, plasma miRNA profiles, plasma metabolomics

Epidemiological data
Reproductive factors, lifestyle factors, family history of breast cancer

Registry data
Diagnosis, stage, receptor status, dates and causes of death

Tumor tissue

Medical records

Blood samples

Questionnaires

National registries

NOWAC
n=172 000

NOWAC
Post-genome
n=50 000

CAMO
n=388

Sample size

Data source

Tumor tissue

Medical records

Blood samples

Questionnaires

National registries

Data type

Histopathological data
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NOWAC

NOWAC
Post-genome

CAMO