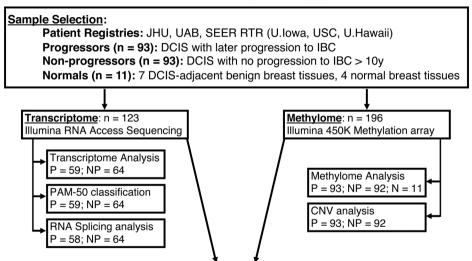
DCIS Study Design



Multi-Omic analyses: P = 59; NP = 63.

A total of 122 samples for which there is both transcriptome and methylome data.