|  |  |
| --- | --- |
| TCGA-BRCA | 1,098 |
| Female | 1,085 |
| w/ 450k methylation array | 774 |
| CPE > 0.6 | 636 |
| Remove 2 tumors from patients that already had another tumor (see methods) | 634 |
| **SITES SELECTED** |  |
| Remove that had less than 5+% missing values of the Clock sites | 624 |
| Only ductal carcinomas | 400 |
| RNA-seq data | 400 |
| CNV data | 400 |