

MABT – MultiAssay big table experiments

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```
suppressPackageStartupMessages({
library(MABT)
})
bq = getBQ2()
bq
## src: bigquery [cgc-05-0009:yriMulti]
## tbls: banovichSE_expressionData, banovichSE_rowRanges, geuFPKM_colData,
##      geuFPKM_expressionData, geuFPKM_rowRanges
bano =
  RangedBT("cgc-05-0009", bq,
    "banovichSE_expressionData",
    "banovichSE_rowRanges")
bano
## rangedBT instance.
## assay colnames include:
##      cg_Methyly450 NA18498 ... NA18489 NA18909
## rangeData colnames include:
##      seqnames start ... probeEnd probeTarget
nrow(bano)
## [1] 329469
```