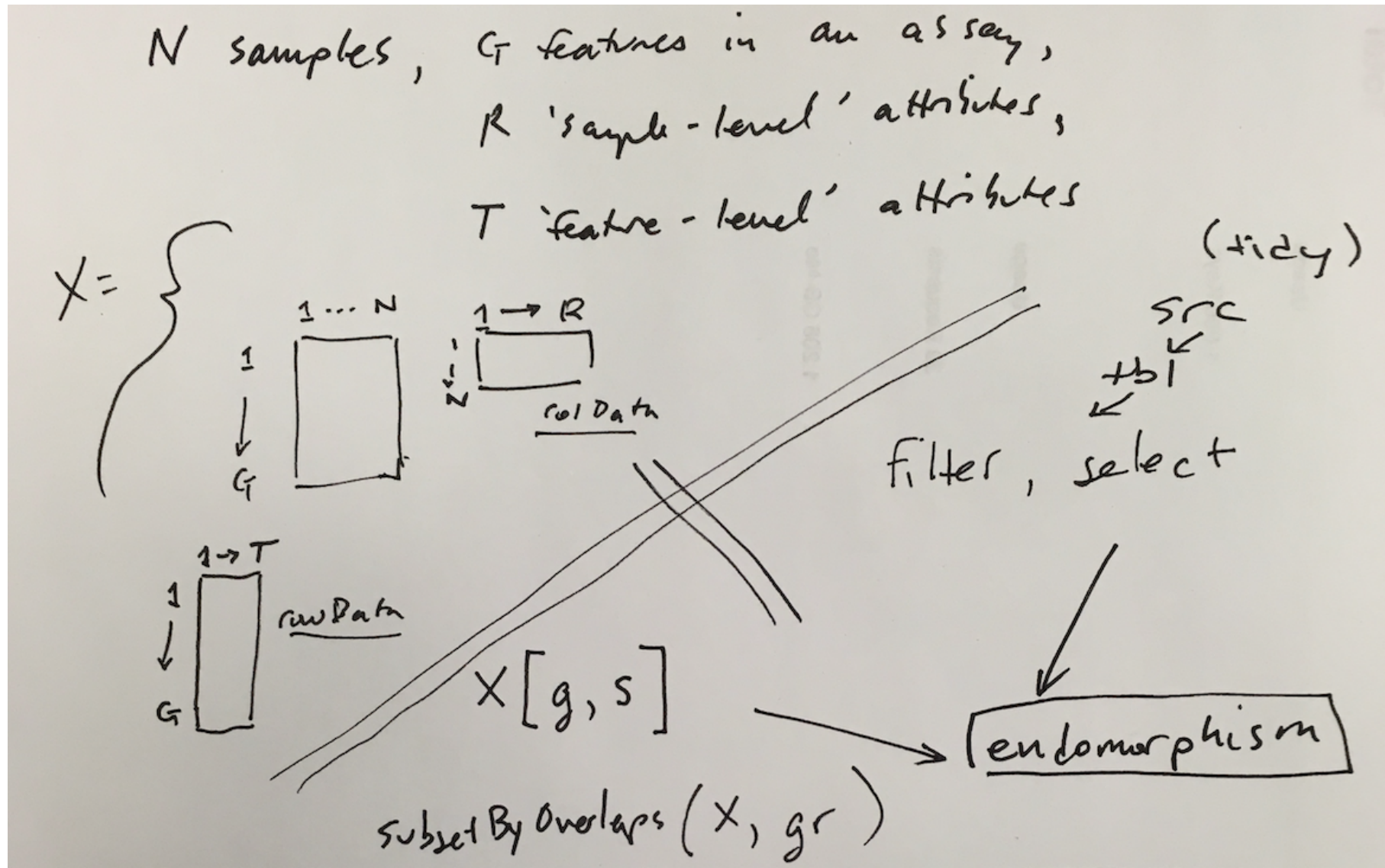


# Recent developments in MultiAssayExperiment

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Levi Waldron [lwaldron.research@gmail.com](mailto:lwaldron.research@gmail.com),  
Martin Morgan [martin.morgan@roswellpark.org](mailto:martin.morgan@roswellpark.org), and others

# Rich containers, "tidy" processes



Prototype: YRI Coriell cell lines with RNA-seq, 450K DNA Meth, 1.5M DHS scores, and all 1000 Genome VCF

```
library(MultiAssayExperiment)
API()
# GEUVADIS -- ExperimentHub?
library(geuvPack); data(geuFPKM)
# Gilad 450k
library(yriMulti); data(banovichSE)
# Degner DnaseI hyp. QTL
library(dsQTL); data(DHStop5_hg19)
# 1000 genomes in S3 VCF (in cloud)
library(ldblock); st = stack1kg()
el = ExperimentList(list(YRIexp=geuFPKM,
  YRImeth=banovichSE, YRIidhs = DHStop5_hg19,
  YRIsnp=st))
YRImult = MultiAssayExperiment(el,
  pData=colData(geuFPKM))
```

□

# Show method has hints

```
[> YRImult
```

```
A MultiAssayExperiment object of 4 listed  
experiments with user-defined names and respective classes.
```

```
Containing an ExperimentList class object of length 4:
```

```
[1] YRIexp: RangedSummarizedExperiment with 23722 rows and 462 columns
```

```
[2] YRImeth: RangedSummarizedExperiment with 329469 rows and 43 columns
```

```
[3] YRIidhs: RangedSummarizedExperiment with 1465442 rows and 50 columns
```

```
[4] YRIisnp: VcfStack with 22 rows and 445 columns
```

```
To access:
```

```
experiments() – to obtain the ExperimentList instance
```

```
pData() – for the primary/phenotype DataFrame
```

```
sampleMap() – for the sample availability DataFrame
```

```
metadata() – for the metadata object of ANY class
```

```
See also: subsetByAssay(), subsetByRow(), subsetByColumn()
```

```
□
```

```
library(gQTLstats); example(tqbrowser,ask=FALSE)
```

## cytoband chooser

**cytoband**

17q21.1 ▼

**SNP**

rs72832966 ▼

**celltype**

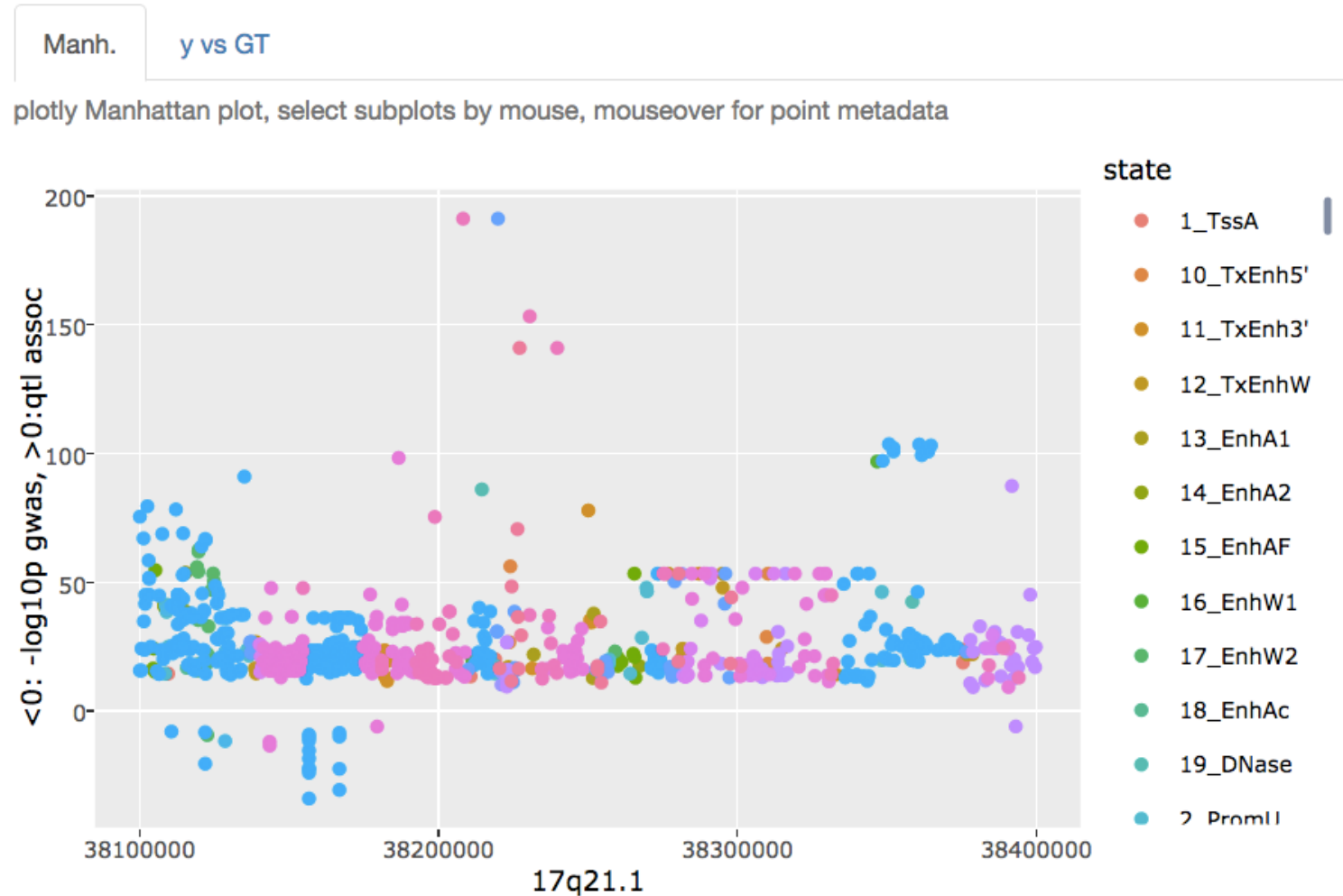
Primary B cells from peripheral blood ▼

**rank**

1

**# to label**

5





# AWS S3-resident representations of TCGA

MAEO URLs					
File Edit View Insert Format Data Tools Add-ons Help Last edit was made on September 12 by Levi Waldron					
Comments Share					
fx MAEO (MultiAssayExperiment Object) URLs					
	A	B	C	D	E
1	MAEO (MultiAssayExperiment Object) URLs				
2					
3	Requirement:	<a href="https://bioconductor.org/packages/MultiAssayExperiment/">https://bioconductor.org/packages/MultiAssayExperiment/</a>			
4	Bug reports:	<a href="https://github.com/vjcitn/MultiAssayExperiment/issues">https://github.com/vjcitn/MultiAssayExperiment/issues</a>			
5	Tiny URL	<a href="http://tinyurl.com/MAEOurls">http://tinyurl.com/MAEOurls</a>			
6	Google Link	<a href="https://docs.google.com/spreadsheets/d/1lh64DDS5mqDIYFzDyCY9HAUnxvl1b6hapKP_akFuNPY/edit?usp=sharing">https://docs.google.com/spreadsheets/d/1lh64DDS5mqDIYFzDyCY9HAUnxvl1b6hapKP_akFuNPY/edit?usp=sharing</a>			
7	Scripts	<a href="https://github.com/waldronlab/MultiAssayExperiment-TCGA">https://github.com/waldronlab/MultiAssayExperiment-TCGA</a>			
8		<a href="https://github.com/waldronlab/MultiAssayExperiment-CCLE">https://github.com/waldronlab/MultiAssayExperiment-CCLE</a>			
9	Cohort #	Cohort Name	Dataset Name	MAEO Name	MAEO URL
10	1	Adrenocortical carcinoma	ACC	accMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/accMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/accMAEO.rds</a>
11	2	Bladder Urothelial Carcinoma	BLCA	blcaMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/blcaMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/blcaMAEO.rds</a>
12	3	Breast invasive carcinoma	BRCA	brcaMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/brcaMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/brcaMAEO.rds</a>
13	4	Cervical squamous cell carcinoma and endocervical adenocarcinoma	CESC	cescMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/cescMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/cescMAEO.rds</a>
14	5	Cholangiocarcinoma	CHOL	cholMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/cholMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/cholMAEO.rds</a>
15	6	Colorectal adenocarcinoma	COADREAD	N/A	N/A
16	7	Colon adenocarcinoma	COAD	coadMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/coadMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/coadMAEO.rds</a>
17	8	Lymphoid Neoplasm Diffuse Large B-cell Lymphoma	DLBC	dlbcMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/dlbcMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/dlbcMAEO.rds</a>
18	9	Esophageal carcinoma	ESCA	escaMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/escaMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/escaMAEO.rds</a>
19	10	FFPE Pilot Phase II	FPPP	N/A	N/A
20	11	Glioma	GBMLGG	N/A	N/A
21	12	Glioblastoma multiforme	GBM	gbmMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/gbmMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/gbmMAEO.rds</a>
22	13	Head and Neck squamous cell carcinoma	HNSC	hnscaMAEO.rds	<a href="http://s3.amazonaws.com/multiassayexperiments/hnscaMAEO.rds">http://s3.amazonaws.com/multiassayexperiments/hnscaMAEO.rds</a>

# Conclusions

- MAEs fairly easy to understand, construct
- Hybrid in-memory/out-of-memory representations are coordinated
- Deployed against TCGA and CCLE
- Supplement with annotation components defining feature and sample relationships
- Can get large; HDF5 or other back end schemes under study
- Multi-omic SIG should communicate on strategies and solutions