# Sparse Matrix Methods and Fields

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### URL

### Outline

What are sparse matrices? Why should we use sparse matrices? What are sparse matrix formats?



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What is spam? Sparse matrices in statistics Solving linear systems Determinants and Cholesky factorization

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What is spam? Sparse matrices in statistics Solving linear systems Determinants and Cholesky factorization

fields Upon spam Finally, Teal examples Beyond large sparse matrices

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What is "sparse" or a sparse matrix?

According to Wiktionary/Wikipedia:

Sparse: (Adjective)1. Having widely spaced intervals2. Not dense; meager

Sparse matrix: a matrix populated primarily with zeros.

- R> n <- 15
- $R > A <- array(runif(n^2), c(n,n)) + diag(n)$
- R > A[A < 0.75] <- 0



column

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R> AtA <- t(A) %\*% A



Why should we use sparse matrices?



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1. Savings in storage

2. Savings in computing time



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To exploit the savings need to exploit the sparsity.

We need a clever storage format and fast algorithms.

Let  $\mathbf{A} = (a_{ij}) \in \mathbb{R}^{n \times m}$  and z the number of its nonzero elements.

1. Naive/ "traditional" /classic format: one vector of length  $n \times m$  and a dimension attribute.



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4. and about 10 more . . .









Naive/traditional/classic: 1, .4, 0, .7, .9, .1, 2, 0, .8, 0, 0, 0, 3, 0, .0, .2, .5, 0, 4, 0, .3, 0, .6, 0, 5







#### **Compressed Sparse Row Format**

- 1. the nonzero values row by row
- 2. the (ordered) column indices of nonzero values
- 3. the position in the previous two vectors corresponding to new rows, given as pointers
- 4. the column dimension of the matrix.



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Savings in storage and computation for sparse matrices Loss in storage and computation for full matrices Intuitive

3. Compressed sparse row (CSR) format: Apart from intuitive, same as triplet Faster element access Available algorithms
Arbitrary choice for "dimension"

### Implications

With a new storage format new "algorithms" are required ...

Is it worthwhile???



Setup:

- R> timing <- function(expr)</pre>
- + as.vector( system.time( for (i in 1:N) expr)[1])
- R > N < -1000 # how many operations
- R> n <- 999 # matrix dimension
- R> cutoff <- 0.9 # what will be set to 0

 $R > A <- array(runif(n^2), c(n,n))$ 

- R > A[A < cutoff] <- 0
- R> S <- somecalltomagicfunctiontogetsparseformat( A)

Compare timing for different operations on A and S.

```
R> timing(A + sqrt(A))
[1] 0.058
R> timing(S + sqrt(S))
[1] 0.061
```

```
R> timing(AtA <- t(A) %*% A)
[1] 0.467
R> timing(StS <- t(S) %*% S)
[1] 4.222</pre>
```

```
R> timing(A[1,2] <- .5)
[1] 0.007
R> timing(A[n,n-1] <- .5)
[1] 0.001</pre>
```

```
R> timing(S[1,2] <- .5)
[1] 0.018
R> timing(S[n,n-1] <- .5)
[1] 0.012</pre>
```

R> timing(solve(AtA, rep(1,n)))
[1] 1.116
R> timing(solve(StS, rep(1,n)))
[1] 1.51

R> timing(chol(AtA))
[1] 0.488
R> timing(chol(StS))
[1] 1.504

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Is it really worthwhile? What is going on?



0



100

-72

With cutoff 0.99:

R> timing(AtA <- t(A) %\*% A)
[1] 0.106
R> timing(StS <- t(S) %\*% S)
[1] 0.089</pre>

R> timing(chol(AtA))
[1] 0.494
R> timing(chol(StS))
[1] 0.551



With cutoff 0.99:

R> timing(AtA <- t(A) %\*% A)
[1] 0.059
R> timing(StS <- t(S) %\*% S)
[1] 0.002</pre>

R> timing(chol(AtA))
[1] 0.466
R> timing(chol(StS))
[1] 0.007



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Especially since

**spam**: R package for sparse matrix algebra.

#### Some slides about spam?

... see inlet one.

### **Sparse Matrices in Statistics**

Where do large matrices occur?

- Location matrices
- Design matrices



### **Sparse Matrices in Statistics**

Where do large matrices occur?

- Location matrices
- Design matrices
- Covariance matrices
- Precision matrices

### **Sparse Matrices in Statistics**

- Covariance matrices: Compactly supported covariance functions Tapering
- Precision matrices: (Gaussian) Markov random fields (Tapering???)

We have symmetric positive definite (spd) matrices.

#### Some slides about tapering?

... see inlet two.

### **Positive Definite Matrices**

A (large dimensional) covariance (often) appears in:

- drawing from a multivariate normal distribution
- calculating/maximizing the (log-)likelihood
- linear/quadratic Discrimination analysis
- PCA, EOF, ...

But all boils down to solving a linear system and possibly calculating the determinant ...

Sparse PCA is sparse in a different sense . . .

### **Solving Linear Systems**

To solve the system Ax = b, we

- perform a Cholesky factorisation  $\mathbf{A} = \mathbf{U}^{\mathsf{T}}\mathbf{U}$
- solve two triangular systems  $\mathbf{U}^{\mathsf{T}}\mathbf{z} = \mathbf{b}$  and  $\mathbf{U}\mathbf{x} = \mathbf{z}$

But we need to "ensure" that **U** is as sparse as possible!

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But we need to "ensure" that **U** is as sparse as possible! Permute the rows and columns of  $\mathbf{A}$ :  $\mathbf{P}^{\mathsf{T}}\mathbf{A}\mathbf{P} = \mathbf{U}^{\mathsf{T}}\mathbf{U}$ .

spam performs Cholesky factorization very efficiently!

### Determinant

0

$$det(\mathbf{C}) = det(\mathbf{U}^{\mathsf{T}}) det(\mathbf{U}) = \prod_{i=1}^{n} \mathbf{U}_{ii}^{2}$$

#### Sparse Matrices and fields

- fields is not bound to a specific sparse matrix format
- All heavy lifting is done in mKrig Or Krig.engine.fixed
- For a specific sparse format, requires the methods: chol, backsolve, forwardsolve and diag as well as elementary matrix operations need to exist
- If available uses operators to handle diagonal matrices quickly
- $\rightsquigarrow$  The covariance matrix has to stem from particular class.

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fields uses spam as default package!

#### Example

With appropriate covariance function:

R> x <- USprecip[ precipsubset,1:2] # locations
R> Y <- USprecip[ precipsubset,4] # anomaly</pre>

```
R> lon <- seq(-125, to=-68, by=blat)
R> lat <- seq( 24.5, to=49, by=blat)</pre>
```

R> pred.x <- expand.grid(lon,lat)</pre>

R> out <- mKrig(x,Y, m=1, cov.function="wendland.cov")
R> pred.out <- predict(out, xnew=pred.x)</pre>

## Example



### How Big is Big?

Upper limit to create a large matrix is the minimum of:

- (1) available memory (machine and OS/shell dependent)Error: 'cannot allocate vector of size'
- (2) addressing capacity  $(2^{31} 1)$ Error: 'cannot allocate vector of length'

However, R is based on passing by value, calls create local copies (often 3–4 times the space of the object is used).

R> help("Memory-limits")

### And Beyond?

Parallelization: nws, snow, Rmpi, ...

Memory "Outsourcing": Matrices are not (entirely) kept in memory: ff, filehash, biglm, ...

(S+ has the library BufferedMatrix)

# And Now?



# Mixer!!



