Statistical Scientific programming: challenges in converting R to C++

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R to C++?

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Conclusion



Conclusion

CluePoints

CluePoints is the premier provider of Risk-Based Monitoring and Data Quality Oversight Software. Our products utilize unique statistical algorithms to determine the quality, accuracy, and integrity of clinical trial data both during and after study conduct.

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Conclusion

Clinical Trials

Why?

- Is it working?
- Is it safe?
- ► Scope, dosage, ...
- ► Approuval from FDA, EMA, ...



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Clinical Trials

How?

- Clinical protocol
- Study conducted at sites
- Patients are enrolled
- Data is collected: demographics, medical history, vital signs, adverse events, labs, patient journals, ...
- Data is verified and analyzed



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Clinical Trials

\$\$\$?

- 1.5-2.5 billion on 10-plus years
- ▶ 30% for sending investigators on sites

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Statistical tests



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Conclusion

Initially developed in the R language by the R&D team



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Conclusion

- Initially developed in the R language by the R&D team
- Very good for research purposes

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Conclusion

- Initially developed in the R language by the R&D team
- Very good for research purposes
- Not so much for production



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Conclusion

- Initially developed in the R language by the R&D team
- Very good for research purposes
- Not so much for production
- Need for something reliable, robust and fast

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The R language



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The R language

The R language

- R is a programing language for statisticians created by statisticians
- R is weakly/dynamically typed
- R operates on named data structures: vector, matrix, array, data frame, factors, lists, objects, functions
- It is very concise
- Lot of statistical libraries



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Some examples (I)

1 w = !is.na(d[[field]]); 2 ctr = factor(d\$center[w]); 3 npat = unclass(table(ctr)); 4 v = d[w, field]; 5 v = rowsum(v, ctr);

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Some examples (I)

```
1 w = !is.na(d[[field]]);
2 ctr = factor(d$center[w]);
3 npat = unclass(table(ctr));
4 v = d[w, field];
5 v = rowsum(v, ctr);
```

- 1. Select the rows where the values of the column "field" are not missing
- 2. Get the values as factor of the column "center" for the selected rows, i.e. the list of centers
- 3. Count the number of rows associated to the different centers, i.e. the number of patients per center
- 4. Get the values for the column "field" for the selected rows
- 5. Sum by center the values from (4)

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Some examples (I)

1 w = !is.na(d[[field]]); 2 ctr = factor(d\$center[w]); 3 npat = unclass(table(ctr)); 4 v = d[w, field]; 5 y = rowsum(v, ctr);

center	xyz
ctr01	
ctr02	1
ctr01	2
ctr03	3
ctr05	4
ctr02	
ctr02	5
ctr02	



ctr	npat	У
ctr01	1	2
ctr02	2	7
ctr03	5	10

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Some examples (II)

- 1 xc = x offset;
- 2 v = tapply(xc, ctr, mean, na.rm=T);
- 3 Sn = unclass(table(ctr)):
- 4 Sn2 = tapply(sid, ctr, function(i) sum(table(i)²)); 5 sigma = sqrt(Sn*vc[3]² + Sn2*vc[2]² + Sn²*vc[1]²)/Sn;
- 6 p = pnorm(v, sd=sigma)

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Some examples

Some examples (II)

```
1 xc = x - offset;
2 v = tapply(xc, ctr, mean, na.rm=T);
3 Sn = unclass(table(ctr));
4 Sn2 = tapply(sid, ctr, function(i) sum(table(i)^2));
5 sigma = sqrt(Sn*vc[3]^2 + Sn2*vc[2]^2 + Sn^2*vc[1]^2)/Sn;
6 p = pnorm(v, sd=sigma)
```

- 1. Apply an offet to x
- 2. Compute per center the mean of xc
- 3. Number of records per center
- Compute per center the sum of the squares of the number of values per patient
- 5. Compute sigma
- 6. Compute the p-values for each center based on a normal distribution

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- 6 p = pnorm(v, sd=sigma)

center	subjid	x
ctr01	s01001	
ctr02	s02001	1
ctr01	s01001	2
ctr03	s03001	3
ctr05	s05001	4
ctr02	s02002	
ctr02	s02001	5
ctr02	s02001	



ctr	Sn	Sn2	sigma	р
ctr01	1	1	0.37	0.21
ctr02	3	5	0.25	0.55
ctr03	5	5	0.19	0.06

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Some examples

Some examples (III)

- 1 dd = duplicated(d\$subjid);
- 2 v = d[[field]]
- 3 w = dd & c(FALSE, v[1:(length(v)-1)]==1);
- 4 x10 = rowsum(1-v[w], ctr[w]);
- 5 N10 = unclass(table(ctr[w]));
- 6 x10Max = rowsum(as.integer(!c(TRUE, w[1:(length(w)-1)])[w]), ctr[w]);

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- 1. Create a boolean vector indicating if a subjid is duplicated or not
- 2. Get the values of the column "field"
- 3. Do some wierd selection
- 4. Get the number of transitions $1 \rightarrow 0$ per center for each patient
- 5. Get the number of potential transitions $1 {\rightarrow} 0$ per center
- 6. Get the maximum number of valid transitions $1 \rightarrow 0$ per center

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- 6 x10Max = rowsum(as.integer(!c(TRUE, w[1:(length(w)-1)])[w]), ctr[w]);

center	subjid	visit	x
ctr01	s01001	v01	1
ctr01	s01001	v02	0
ctr01	s01001	v03	1
ctr01	s01001	v04	0
ctr01	s01002	v01	1
ctr01	s01002	v02	1
ctr02	s02001	v03	0
ctr02	s02002	v01	1
ctr02	s02002	v02	1
ctr02	s02002	v03	0
ctr03	s03001	v01	1
ctr03	s03001	v02	0
ctr03	s03001	v03	1



ctr	X10	N10	x10max
ctr01	2	3	3
ctr02	1	2	1
ctr03	1	1	1

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 Straightforward approach: Recode each R function in C++ PRO C++ and R codes are similar CON Too many combinations of parameters/structures



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onclusion

- Straightforward approach: Recode each R function in C++ PRO C++ and R codes are similar CON Too many combinations of parameters/structures
- ▶ Hard: understanding what the researcher wanted to do PRO Faster code CON C++ and R codes can be very different 1 line in R $\rightarrow \pm 30$ lines in C++

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- Straightforward approach: Recode each R function in C++ PRO C++ and R codes are similar CON Too many combinations of parameters/structures
- Hard: understanding what the researcher wanted to do PRO Faster code
 CON C++ and R codes can be very different
 1 line in R → ±30 lines in C++
- Hardest: changing the data structure
 PRO Less ressource/faster code
 CON C++ and R codes are even more different

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- ► Hard: understanding what the researcher wanted to do

- Hardest: changing the data structure
 PRO Less ressource/faster code
 CON C++ and R codes are even more different
- Recoding model fitting algorithms is a huge (tremendous) task. It's easier to call the R function from the C++ code
 PRO Updates of the fitting model code
 CON Added dependencies

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- Straightforward approach: Recode each R function in C++ PRO C++ and R codes are similar CON Too many combinations of parameters/structures
- \blacktriangleright Hard: understanding what the researcher wanted to do

 $\begin{array}{ll} \mbox{PRO} & \mbox{Faster code} \\ \mbox{CON} & \mbox{C}{++} \mbox{ and } \mbox{R codes can be very different} \\ & 1 \mbox{ line in } \mbox{R} \longrightarrow \pm 30 \mbox{ lines in } \mbox{C}{++} \end{array}$

- Hardest: changing the data structure
 PRO Less ressource/faster code
 CON C++ and R codes are even more different
- Recoding model fitting algorithms is a huge (tremendous) task. It's easier to call the R function from the C++ code
 PRO Updates of the fitting model code
 CON Added dependencies
- Beware of Numerical (in)accuracy
- Testing and testing and testing (no data, invalid data, NaN, Inf, ...)

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Scientific programming challenges



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Measure

Software architecture

Data structure

Smart pointers, Pimpl, Factories

Fail-fast/Fail-safe

Numerical errors

Accumulators

std::algorithms, boost, GSL BLAS, LAPACK, ...

Conclusion

Scientific programming challenges

- Requirements include low response time and memory usage, minimizing numerical errors and error propagation.
- Testing
- Software architecture
- Data structure
- Fail-fast/Fail-safe idioms
- Exceptions
- RAII
- Pimpl idiom and smart pointers
- Factory pattern
- Iterator pattern and accumulators
- ▶ std::algorithms, boost, GSL, BLAS, LAPACK, ...



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Testing



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Conclusion

Testing

- Framework
- Unit testing, Integration testing, ...
- Test Driven Development
- Behavior Driven Development to replicate the documentation specification
- Continuous Integration

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Testing

- Framework
- Unit testing, Integration testing, ...
- Test Driven Development
- Behavior Driven Development to replicate the documentation specification
- Continuous Integration
- Each bug must be tested

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Code for Testing

- If you cannot test your code, rewrite it
- If you cannot test your code easily, rewrite it
- If you cannot test your code independently, rewrite it

▶ ...

Tools like clang static analyzer and gcov/lcov code coverage are a great help

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Measure!!!



THE REASON I AM SO INEFFICIENT

Credit xkcd

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Conclusion
Measure!!!

- Select between different data structures
- Select between different algorithms
- Use generated data
- Use real data
- Use data of different sizes

▶ ...

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Measure!!! - an example

Context: Originally, an algorithm has to be applied on vectors: f(x, y)



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Measure!!! – an example

Context: Originally, an algorithm has to be applied on vectors: f(x, y)Then only on some filtered elements: f(x, y, w)

X	Y	w	
42.5	100	true	
		true	
		false	
		true	
		false	
		true	
		true	
		false	

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Measure!!! - an example

Context: Originally, an algorithm has to be applied on vectors: f(x, y)Then only on some filtered elements: f(x, y, w)

- Modify the algorithm to take into account only the filtered vectors'elements: *filter algo*
- Create pseudo vectors with the filtered elements: filter vector
- Create new vectors with the filtered elements: copy vector

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Measure!!! - an example

Context: Originally, an algorithm has to be applied on vectors: f(x, y)Then only on some filtered elements: f(x, y, w)

- Modify the algorithm to take into account only the filtered vectors'elements: *filter algo*
- Create pseudo vectors with the filtered elements: filter vector
- Create new vectors with the filtered elements: copy vector

Option	Timing (s)				
	$N = 10^{2}$	$N = 10^{4}$	$N = 10^{6}$	$N = 10^{8}$	
filter algo	0.0003	0.008	0.9	100	
filter vector	0.0003	0.006	0.8	98	
copy vector	0.0006	0.015	4.6	/	

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Software architecture & Data structure



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std::algorithms, boost, GSL BLAS, LAPACK, ...

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Software architecture & Data structure

Important points to consider

- Input/output data structure?
- Computational units?
- Simple but not too simple!
- Which doors are you closing?
- Expressiveness

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Software architecture & Data structure

Important points to consider

- Input/output data structure?
- Computational units?
- Simple but not too simple!
- Which doors are you closing?
- Expressiveness

For this project

- Data is organized in datasets, i.e. tables in which each column represents a particular variable or key variable, and each row corresponds to a given record. There may also be missing values.
- Statistical tests are the computational units.

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Conclusion

Levels of abstraction

- The most important good pratice
- Divide and conquer
- Top down design
- Bottom up design
- Separation of concerns
- $\blacktriangleright \textit{ Modularity: low coupling} \longleftrightarrow \textit{ high cohesion}$
- Design review

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Levels of abstraction – mathematical formula

WHAT PART OF $i\hbar\frac{\partial}{\partial t}\Psi(\vec{r},t) = \left(-\frac{\hbar^2}{2m}\nabla^2 + V(\vec{r},t)\right)\Psi(\vec{r},t)$ DON'T YOU UNDERSTAND?

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Conclusion

Levels of abstraction – mathematical formula

- Very tempting to code one mathematical formula into one function.
- Decompose the formula into meaningful steps, e.g. numerator, denominator, partial sums, ...
- Transform the function into a class
- Transform each step into a struct



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Conclusion

Abstraction levels – Example

Sample variance – Standard formula

$$s_N^2 = \frac{1}{N-1} \sum_{k=1}^N (x_k - \bar{x})^2$$

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Abstraction levels – Example

Sample variance - Standard formula



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Abstraction levels – Example

```
namespace MATH INTERNAL {
 2
     template<typename T=double>
     struct sample variance {
 3
       T s2{std::numeric limits<T>::quiet NaN()};
 5
 6
       template<typename Container>
       sample variance(const Container X) { if(X.size()>1) s2 = frac(X)*sum(X, mean(X)); }
 7
 8
9
       operator T() const { return s2; }
10
11
       template<typename Container>
       static T frac(const Container& X) { return ONE/(X.size()-ONE); }
12
13
14
       template<typename Container>
       static T mean(const Container& X) { return std::accumulate(X.begin(), X.end(), ZERO)/X.size(); }
15
16
17
       struct square of difference {
18
         const T xbar:
         square of difference(const T mean) : xbar(mean) {}
19
         T operator()(const T x) const { return (x-xbar)*(x-xbar); }
20
       }:
21
22
23
       template<typename Container>
       static T sum(const Container& X, const T xbar) {
24
25
         const square_of_difference d(xbar);
26
         return std::accumulate(X.begin(), X.end(), ZERO, [d](const T s, const T x) { return s + d(x); });
27
28
     };
29
   template<typename Container>
30
31
   inline typename Container::value type sample variance(const Container& X)
32
   Ł
33
     return MATH_INTERNAL::sample_variance<typename Container::value_type>(X);
34 }
```

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```
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```

```
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std:nagorithms, boost,
Bd:na_logorithms, bd:na_logorithms, bd:na_logorithms, bd:na_logorithms, bd:na
```

Data structure

- Performance requires well thought data structure
- Cache usage
- Prefetching
- Lazy evaluation
- Sparse representation

▶ ...

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Duplicate patients: comparing patient's fingerprints

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N numerical vectors of the same length M
 Typical cases: N = 1000 - 40000 and M = 20 - 20000



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Conclusion

- N numerical vectors of the same length M
 Typical cases: N = 1000 40000 and M = 20 20000
- $N \times (N-1)/2$ scalar products

$$X \cdot Y = \sum_{i}^{M} X_i \times Y_j$$

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Sparse vectors!!!

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- $N \times (N-1)/2$ scalar products

$$X \cdot Y = \sum_{i}^{M} X_{i} \times Y_{j}$$

- Sparse vectors!!!
- Performance results

	N =	841	N = 35613		
	M = 1060		M = 14304		
	Memory	Timing	Memory	Timing	
R		± 1 m		/	
C++ (normal vectors)	57MB	0.68s	9.8GB	29m	
C++ (sparse vectors)	34MB	0.45s	6.6GB	38s	

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Smart pointers, Pimpl. Factories

Pointer to implementation or Private implementation

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Conclusion

Pointer to implementation or Private implementation

PROS

- Separate interface from implementation
- Decrease recompilation cycles
- Binary compatibility of shared libraries



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Pointer to implementation or Private implementation

PROS

- Separate interface from implementation
- Decrease recompilation cycles
- Binary compatibility of shared libraries

CONS

- Increase in memory usage
- Increase in maintenance effort
- Performance loss
- Doesn't work well with templates

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- std::unique_ptr or std::shared_ptr?
- Mutable or non mutable objects?
- Access to the objects, how often?
- Multiple inheritance, virtual inheritance (diamond problem)?
- Template member functions, template classes?
- Objects in a coherent state!

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Smart pointers, Pimpl, Factory pattern : Inheritance



Questions, Remarks?

Statistical

Smart pointers, Pimpl, Factory pattern: Diamond problem



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Conclusion

Smart pointers, Pimpl, Factory pattern: Template members

api.hpp

```
1 class MyPublic : public Pimpl {
  public:
2
3
    class impl:
4
5
    MyPublic(...);
6
    template<typename Tp> Tp as() const;
7
8
 };
```

api.cpp

1 2

4 5 6

7

```
MyPublic::MyPublic(...) : Pimpl(impl::build(...)) {}
3
    template<>
    double MyPublic::as() const { return _valid_ptr() ? _get_ptr<impl>()->asNumber() : NaN(); }
    template<>
    std::string MyPublic::as() const { return valid ptr() ? get ptr<impl>()->asString() : std::string(); }
```

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The R language

Smart pointers, Pimpl. Factories

Pimpl: use

```
1 typedef std::vector<Test> TESTS;
 2
   // Create the tests
 3
   TESTS tests:
 4
 5 tests.push_back(BetaBinomial(...));
 6 tests.push_back(CountField(...));
 7 tests.push back(CountField(...));
 8 tests.push_back(CNR_ByCenter(...));
 9
10
11
   // Export the results
12
   json ostream os(...);
13
   print_results(os, tests);
14
15
   void print_results(json_ostream& os, const TESTS& tests)
16
17
   ſ
     for(const auto& test: tests) {
18
19
20
21
   }
```

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Pimpl: use



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Conclusion

Check constraints on input/output

1 double foo(const std::vector<size_t>& l, const std::vector<double>& x, const std::vector<bool>& w)
2 {
3 CP_ASSERT(l.size() == x.size());
4 CP_ASSERT(l.size() == w.size());
5 // Rest of the code
6 }

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Check constraints on input/output

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2 {
3 CP_ASSERT(l.size() == x.size());
4 CP_ASSERT(l.size() == w.size());
5 // Rest of the code
6 }

Fitting of statistical models might fails

```
try {
1
2
    fit = vglm("cbind(a,b)~1",
               Named("family", family),
3
               Named("data", dateframe).
4
5
               Named("control", control(Named("criterion", "coef"),
6
                                         Named("stepsize", 0.5))));
7
  } catch(std::exception& e) {
    // Retry with other parameters
8
9
  }
```

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Check constraints on input/output

1 double foo(const std::vector<size_t>& l, const std::vector<double>& x, const std::vector<bool>& w)
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Fitting of statistical models might fails

```
try {
1
2
    fit = vglm("cbind(a,b)~1",
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                Named("data", dateframe).
4
5
                Named("control", control(Named("criterion", "coef"),
6
                                         Named("stepsize", 0.5))));
7
  } catch(std::exception& e) {
    // Retry with other parameters
8
9
  3
```

Propagate the error message

- Rethrow the exception
- Store the exception as an error message inside the object

▶ ...

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Conclusion

Numerical instabilities



Credit xkcd

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Conclusion
Numerical instabilities

Test on standard deviations

▶ P values computed from the integration of two functions:

 $f_1(x) = pchisq(s/x^2; N, left.tail) \times dgamma(x; scale, shape)$

 $f_2(x) = pchisq(s/x^2; N, right.tail) \times dgamma(x; scale, shape)$



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Numerical instabilities

calcPsd: test on standard deviations

• $f_1(x)$ is unstable in case shape < 1



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Numerical instabilities

calcPsd: test on standard deviations

- $f_1(x)$ is unstable in case shape < 1
- $f_1(x)$ can be rewritten by using the integration by parts theorem $\int_0^a u dv = [uv]_0^a \int_0^a v du$

$$f_{1'}(x) = rac{2s}{x^3} imes dchisq(s/x^2; N) imes$$
 pgamma(x; scale, shape, left.tail)



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Conclusion

Sample variance – Standard formula

$$s_N^2 = \frac{1}{N-1} \sum_{k=1}^N (x_k - \bar{x})^2$$

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Sample variance – Standard formula

$$s_N^2 = \frac{1}{N-1} \sum_{k=1}^N (x_k - \bar{x})^2$$

Can be implemented as a 2 pass algorithm, first the mean x
, and the variance s² afterwards.



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Conclusion

Sample variance – Standard formula

$$s_N^2 = \frac{1}{N-1} \sum_{k=1}^N (x_k - \bar{x})^2$$

• Can be implemented as a 2 pass algorithm, first the mean \bar{x} , and the variance s^2 afterwards.

BUT the number of items *N* can be huge

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Sample variance – Standard formula

$$s_N^2 = \frac{1}{N-1} \sum_{k=1}^N (x_k - \bar{x})^2$$

• Can be implemented as a 2 pass algorithm, first the mean \bar{x} , and the variance s^2 afterwards.

BUT the number of items *N* can be huge

Have a one pass algorithm

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Sample variance – Standard formula

$$s_N^2 = \frac{1}{N-1} \sum_{k=1}^N (x_k - \bar{x})^2$$

► Can be implemented as a 2 pass algorithm, first the mean x̄, and the variance s² afterwards.

BUT the number of items *N* can be huge

- Have a one pass algorithm
- Compute the variance for increasing *N* to observe convergence.

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Conclusion

Sample variance – One pass algorithm: Sum of squares method

$$s_N^2 = \frac{1}{N(N-1)} \left(N \sum_{k=1}^N x_k^2 - \left(\sum_{k=1}^N x_k \right)^2 \right)$$

One pass algorithm but the formula is unstable:

- ► float precision: for {10000f, 10001f, 10002f}, the result is -1.0666667e+01 instead of 1.
- double precision: for {10000000, 10000001, 10000002}, the result is 0 instead of 1.

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BLAS, LAPACK, ...

onclusion

Sample variance - Iterative algorithm: Welford's recursion method

with $M_0 = 0$ and $S_0 = 0$, and then

$$s_N^2=\frac{S_N}{N-1},$$

This stable algorithm with can be easily turned into an accumulator

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Accumulators



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Conclusion

Accumulators

- Separation of the operations on the elements from the iteration leads to smaller testable code.
- Statisticals tests involve operations (agregation, sum, average, variance, ...) on one or more variables based on one or more several key variables.
 E.g.: Preprocess involves taking the mean by visits or the sum by patients, the count of non missing values per center, ...

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Conclusion

Accumulators as a OO pattern

Mean of the elements of a vector

- without accumulator
- 1 double sum{0};
- 2 for(const auto x: myvector) sum += x;
- 3 const double mean = sum / myvector.size();

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std::algorithms, boost, GSL BLAS, LAPACK, ...

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Accumulators as a OO pattern

Mean of the elements of a vector

without accumulator

double sum{0};

1

- 2 for(const auto x: myvector) sum += x;
- 3 const double mean = sum / myvector.size();

with accumulator

1 using namespace boost::accumulators; 2 accumulator_set<double, stats<tag::mean>> acc; 3 for(const auto x: myvector) acc(x);

4 mean(acc);

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Accumulator implementation

Sample variance – Welford's recursion method

```
template<typename T> class Variance {
      size t k: /**< Number of elements */
 2
      T m: /**< 0th order moment, i.e. average */
 3
 4
      T s; /**< 1st order moment */
 5 public:
      Variance() : k(0), m(0), s(0) {}
 6
      void operator()(const T x) {
 7
       if(std::isnan(x)) return;
 8
       ++k:
 9
       const T pm(m);
10
11
       m += (x-pm) * (ONE/k);
       s += (x-pm) * (x-m);
12
13
      3
      T average() const noexcept { return m; }
14
      T s2() const no except { return k > 1 ? s / (k-1) : ZERO; }
15
16 };
```

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std::algorithms, boost, GSL, BLAS, LAPACK, ...



Credit Big Bang Theory

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Conclusion

std::algorithms, boost, GSL, BLAS, LAPACK, ...

- <algorithm>
- ▶ boost
 - Statistics, . . .
 - Logging facilities
 - System (command line arguments, ...)
 - Thread, MPI, Serialization
 - ▶ ...
- GNU Scientific Library
 - Optimization, minimization, ...
- BLAS and LAPACK
 - Operations on matrices
- Numerical Recipes
 - Lots of algorithms



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onclusion

In probability theory and statistics, the beta-binomial distribution is a family of discrete probability distributions on a finite support of non-negative integers arising when the probability of success in each of a fixed or known number of Bernoulli trials is either unknown or random.



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$$f(k; n, \alpha, \beta) = \binom{n}{k} \frac{B(k + \alpha, n - k + \beta)}{B(\alpha, \beta)}$$

- k and n are positive integers with $k \leq n$
- $\blacktriangleright \ \alpha$ and β are strictly positive numbers
- Binomial coefficient

$$\binom{n}{k} = \frac{n!}{k!(n-k)!} = \frac{\Gamma(n+1)}{\Gamma(k+1)\Gamma(n-k+1)}$$

Beta function

$$B(x,y) = \frac{\Gamma(x) + \Gamma(y)}{\Gamma(x+y)}$$

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$$f(k; n, \alpha, \beta) = \binom{n}{k} \frac{B(k + \alpha, n - k + \beta)}{B(\alpha, \beta)}$$

- \blacktriangleright Numerically fine as long as α and β are small
- When α and β are not small, $B(\alpha, \beta)$ tends toward zero.

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Conclusion

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- \blacktriangleright Numerically fine as long as α and β are small
- When α and β are not small, $B(\alpha, \beta)$ tends toward zero.

Trick: Do the calculation in the log scale:

$$f(k; n, \alpha, \beta) = \exp\left(\log\binom{n}{k} + \log B(k + \alpha, n - k + \beta) - \log B(\alpha, \beta)\right)$$

$$\log \binom{n}{k} = l_binomial_coefficient(n, k)$$
$$= lgamma(n+1) - lgamma(k+1) - lgamma(-k+n+1)$$

$$log B(x, y) = lbeta(x, y)$$

= lgamma(x) + lgamma(y) - lgamma(x + y)

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Minimize with GSL

- ► GSL is a C library
- Use wrapper classes
- Pointer to the minimizer created/owned by GSL
- Pointer to the function definition struct

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Conclusion

enum class M_TYPE {

NO GRADIENT,

};

template<M_TYPE> struct M_API; /**< Template for the C API */
template<M_TYPE> class M_FCT; /**< Template for defining the function to minimize */
template<M_TYPE> class IMinimizer; /**< Template for the minimizer */</pre>

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Conclusion

```
/// Specialized template defining the function to minimize (no gradient)
 2 template<>
 3 class M FCT<M TYPE::NO GRADIENT> {
   public:
 4
     friend class IMinimizer<M TYPE::NO GRADIENT>;
 5
 6
 7
     /// Virtual base class for the implementation of the function to minimize
 8
     class Base : public boost::noncopyable {
 9
     public:
     virtual ~Base() {}
10
     virtual double evaluate(const double x) = 0;
11
12
     };
13
     typedef std::unique_ptr<Base> PTR; /**< Type of the pointer to the instance of the function to minimize */
14
15
     typedef gsl function DEF; /**< Type for the definition */
16
17
     M FCT(PTR fct, const NUMBER minimum, const NUMBER lower, const NUMBER upper); ....
18
19
     double evaluate(const double x) { return m fct->evaluate( x); }
20
     double get lowest bound() const { return m f lower < m f upper ? m lower : m upper: }
     double get lowest f bound() const { return std::min(m f lower, m f upper); }
21
22
23
   private:
24
25
   };
```

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Conclusion

```
template<M TYPE Type> class IMinimizer : public boost::noncopyable {
 2 public:
     typedef M_FCT<_Type> FCT;
 3
 4
 5
     explicit IMinimizer(const std::string& type, FCT& fct, const double epsabs, const double epsrel) { ... }
 6
     ~IMinimizer() { ... }
 7
 8
     bool iterate(const size t maxiter) {
 9
       if(m can minimize) {
         for(size_t iter = 0; iter<_maxiter && next(); ++iter) {</pre>
10
           if(converged()) return true;
11
12
         }
13
       3
       return false:
14
15
     }
16
17
     std::string name() const
18
     { return m ptr ? M API< Type>::name(m ptr) : std::string(); }
19
     bool next()
20
     { return m ptr && m can minimize ? M API< Type>::iterate(m ptr) == 0 : false; }
21
     bool converged() const
     { return m_ptr && m_can_minimize ? M_API<_Type>::converged(m_ptr, m_epsabs, m_epsrel) : false; }
22
23
     double x() const
24
     { return m_ptr ? (m_can_minimize ? M_API<_Type>::x_minimum(m_ptr) : m_fct.get_lowest_bound()) : 0; }
25
     double y() const
     { return m_ptr ? (m_can_minimize ? M_API<_Type>::f_minimum(m_ptr) : m_fct.get_lowest_f_bound()) : 0; }
26
27
   private:
28
29
   }:
   typedef IMinimizer
30
```

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Conclusion

```
template<M TYPE> struct M API; /// Generic template holding the API
 2
 3 template<>
   struct M API<M TYPE::NO GRADIENT> {
     typedef gsl min fminimizer* PTR; /**< Type of the pointer to the minimizer */
 5
 6
     typedef gsl function* DEF: /**< Type of the pointer to the definition */
 7
 8
     static PTR alloc(const STRING& type) { return gsl min fminimizer alloc(type( type)); }
 9
     static void free(PTR p) { gsl min fminimizer free( p); }
10
11
     static const gsl min fminimizer type* type(const std::string& type);
12
     static std::string name(PTR p) { return gsl min fminimizer name( p); }
13
14
     static bool set(PTR _p, DEF _fct, const double _minimum, const double _lower, const double _upper)
     { return gsl min fminimizer set( p, fct, minimum, lower, upper) != GSL EINVAL; }
15
     static bool set(PTR p. DEF fct, const double minimum, const double fminimum, const double lower, const
16
            double _flower, const double _upper, const double _fupper)
17
     { return gsl min fminimizer set with values( p, fct, minimum, fminimum, lower, flower, upper, fupper)
            != GSL EINVAL; }
18
19
     static int iterate(PTR p) { return gsl min fminimizer iterate( p); }
     static bool converged(PTR p, const double epsabs, const double epsrel)
20
21
     { return gsl_min_test_interval(x_lower(_p), x_upper(_p), _epsabs, _epsrel) == GSL_SUCCESS; }
22
23
     static double x minimum(PTR p) { return gsl min fminimizer x minimum( p); }
24
     static double x_upper(PTR _p) { return gsl_min_fminimizer_x_upper(_p); }
25
     static double x_lower(PTR _p) { return gsl_min_fminimizer_x_lower(_p); }
     static double f minimum(PTR p) { return gsl min fminimizer f minimum( p); }
26
     static double f_upper(PTR _p) { return gsl_min_fminimizer_f_upper(_p); }
27
     static double f lower(PTR p) { return gsl min fminimizer f lower( p); }
28
29 }:
```

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```
class MyFctNoGradient : public gsl::MinimizerNoGradient::FCT::Base {
2 public:
     MyFctNoGradient(...) { ... }
 3
 4
 5
     double evaluate(const double x) override { ... }
   };
 6
 7
8
   double minimize_my_fct(...)
9
   ſ
     gsl::MinimizerNoGradient::FCT f(new MyFctNoGradient(...), .5*(low+hi), low, hi);
10
     gsl::MinimizerNoGradient minimizer("Brent", f, 0.1, 0.1);
11
12
     minimizer.iterate(10);
13
     return minimizer.x();
14 }
```

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Conclusion

- Fast production code
- No task is impossible
- Seek expertise
- Testing: Don't trust your code
- Have fun and keep learning



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Questions, Remarks?



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Questions, Remarks?

Thank you for your attention!