Lecture08 - "Fun" with indices

Kevin Bonham, PhD

Final Project Details

_ab Grading

Indices vs values

Writing code in notebooks

Lab08 == Lab07

What's left in the course?

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2021-07-02

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Outline

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Final project: Code + analysis

Final Contents

- Code repository with resuable functions (largely built from labs and assignments)
 - including documentation
 - including test suite
- Analysis repository with details, descriptions of code, and plots
- Default project: sequence analysis of Sars-CoV genomes

Earlier components

- If proposing alternate project: Analysis proposal
- Analysis plan
- First draft
 - Code, tests, docs should be complete
 - Analysis should be complete

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Dates

School Deadlines

- Summer Term Final Projects Due: July 23
- Grades Due: July 30

Project Deadlines

- Proposals for alternate projects Due: July 9
- Analysis plans Due: July 12
- First drafts code / notebooks Due: July 16

Missing these deadlines will have grading consequences

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Getting caught up

I tried to sort out labs, but it's a bit confusing

- Please update github pull request titles with:
 - Your name
 - Lab Partner's name (if applicable)
 - Lab number
- Should have separate PRs for
 - Lab03
 - Lab04 / 5
 - Lab06
 - Lab07 / 8
- You are not (yet) getting credit unless you have a PR with your name in the title and a "complete" badge.

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Confusion between "location" and "thing"

```
julia> myvec = [1.2, 2.3, 3.4]
3-element Vector{Float64}:
    1.2
    2.3
    3.4
julia> x = myvec[2]; # 2 is the index
julia> x # this is the value stored at index 2
2.3
```

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```
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Indices must be integers, values can be anything
                                                                   with indices
                                                                  Kevin Bonham,
                                                                     PhD
 julia> myvec[1]
 1.2
                                                                 Indices vs values
 julia> myvec[1.2]
 ERROR: ArgumentError: invalid index: 1.2 of type Float64 What's left in the
 #...
 julia> othervec = ["something", 'A', 2.2];
 julia> map(typeof, othervec)
 3-element Vector{DataType}:
  String
  Char
  Float64
                                    ◆□▶ ◆□▶ ◆□▶ ◆□▶ □ のQ@
```

Specialized functions can find things inside vectors

```
julia> newvec = rand(5)
5-element Vector{Float64}:
 0.516023786589465
 0.4465775523061499
 0.21788789287837185
 0.08900106348786951
 0.7016481961587768
julia> findfirst(<(0.5), newvec)</pre>
2
julia> findall(<(0.5), newvec)</pre>
3-element Vector{Int64}:
```

2 3 4

julia> newvec[findall(<(0.5), newvec)] # index based on result 3-element Vector{Float64}: 0.4465775523061499

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Mixing code, results, and descriptions

- can use comments, but they have limited expressiveness
- using "notebook" environments allows including results "inline"
- Many options for notebooks,
 - markdown (R Markdown, Weave.jl)
 - Jupyter notebooks
 - Pluto.jl

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What's left in the course?

For scientific coding, code is usually ad-hoc

- Can't write unit tests for a specific plot
- But you can for the plot function itself
- Functions, packages, etc, are like "protocols"
- Use code notebooks for "experiments"

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Alignment tracing for NW and SW

start from M_(i,j) where i and j are

- the last indices in 1st and 2nd dimension for NW
- the indices for the matrix with the maximum score in SW

Check the score from

- 1. $M_{(i,j-1)}$ (cell to the left), a gap score
- 2. $M_{(i-1,j)}$ (cell above), a gap score
- 3. $M_{(i-1,j-1)}$ (cell from diagonal), a match or mismatch
- If any match your current cell, push correct characters to alignments
 - 1. push gap to seq1, character at j to seq2
 - 2. push character at i to seq1, gap to seq2
 - 3. push character at i to seq1, character at j to seq2

update indices

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Special considerations

Be mindful of what happens when you hit the first row or first column

- i-1 or j-1 may throw bounds error
- When should your loop stop?

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Coming up next

Lectures

- Lecture 9: using packages, plotting functions
- Lecture 10: custom structs / advanced programming
- Lecture 11: Working with code from other languages
- Lecture 12: Wrap-up

Labs

- Lab 9: Gap open and gap-extension scores
- Lab 10: Multiple sequence alignment algorithms
- Lab 11-12: Work on Final Projects

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