STATS 769 Data Formats

Paul Murrell

The University of Auckland

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Overview

- In this lecture we will discuss some modern data formats.
 - JSON.
 - XML (and XPath).
- The aim is to become familiar with these formats and to be able to work with them in R.

JSON

- JavaScript Object Notation.
- Text format.
- JSON Data Types: null, true, false, number, string.
- JSON Data Containers: ordered (and unnamed) arrays [], named (and unordered) arrays {}.
- Can mix types within arrays.

JSON in R

- The **jsonlite** package has from JSON() (and to JSON()).
- The result is the simplest structure possible, in the order: vector, matrix, data frame, list.
- Nested JSON objects can produce nested data frames; the flatten() function turns these into normal 2-D data frames.
- Use prettify() to eyeball raw JSON and str() to explore nested list structures.

XML

```
<?xml version="1.0" encoding="UTF-8"?>
<response>
  <row>
    <row _uuid="00000000-0000-0000-AF9A-401551B08E58">
      <month>Jan</month>
      <pets_adopted>129</pets_adopted>
    </row>
    <row _uuid="00000000-0000-0000-F7B9-E37345BC66E7">
      <month>Mar</month>
      <pets_adopted>126</pets_adopted>
    </row>
  </row>
</response>
```

XML in R

- The xml2 package has read_xml()
- The result is NOT a data frame.
- Extract elements of interest using xml_find_all() and XPath expressions
- Use xml_text() to extract text content from elements.
- Use xml_attr() to extract attribute values from elements.
- NOTE that all content and attribute values are character values.

Language for expressing subsets of an XML document.

 An XPath expression consists of some combination of location paths of the form ...

```
axisname::nodetest[predicate]
... but usually just ...
  nodetest
... or ...
  nodetest[predicate]
```

An XPath expression is formed by combining several location paths, separated by a forward slash, /.

• If the expression **begins** with a forward slash, matching starts from the document root node.

```
/a/b/c
```

 A double forward slash, //, is short for /descendant-or-self::node()/. /a//c

The nodetest is commonly just the name of an element or @name to match an attribute.

 It can also be a wildcard, *, which matches any element, or @*, which matches any attribute.

```
/a/b
/a/@id
/a/*/c
```

The predicate is like a subsetting expression.

• It can be a simple integer.

```
/a/b[1]
```

• It can be a comparison.

```
/a/b[@year > 2000]
/a[@lang = "en"]
```

• It can be a special function.

```
/a/b[last() - 1]
/a/b[contains(@id, "paul")]
```

The axis is relative to the current node in the XML document. The default axis is child, which means to search children of the current node (this is what happens if we do not specify an axis in a location path).

 The following-sibling axis can force the search to look at siblings of the current node.

```
/a/b/following-sibling::*
```

 The parent or ancestor axes can force the search to look back up the hierarchy of XML nodes.

```
/a/b/c/ancestor::a
```

• These can be useful within a predicate.

```
//c[ancestor::b@lang = "en"]
```

Resources

- 'jsonlite' mapping between R objects and JSON http://cran.rproject.org/web/packages/jsonlite/vignettes/jsonmapping.pdf
- w3schools XPath tutorial http://www.w3schools.com/xpath/