## Abstract

This provides a brief tour of how to use the XML parsing package. It starts by showing how to read an XML document into R and access the elements of the tree that represents the structured contents of the document. Next, it discusses how to govern the creation of the tree as it is being read from the file. And finally, it discusses the use of event-driven (or SAX) parsing.

Suppose we have the following text in a file named **as.Sxml**. A quick look at the contents shows that it contains some S-language functions and some documentation for each of them. This is similar to the self-documenting facility in S4.

Note also that each function definition consists of 3 elements: a name, some comments and the function definition itself. In this particular form, we have been able to specify the structure of the elements and not use the R or S syntax to assign the function definition to the name, i.e. as ¡- function(..) .... Why is this useful? Because we have expressed what mean – the relationships between the elements in a system-neutral format. I could add a fourth element to some or all of these functions which provides a Matlab implementation of the same function. The structure and syntax of the document would not have to change. Instead, the software that reads the document would know which bits to read.

Note also that the S syntax causes complications for XML because certain characters S uses (e.g. ;) have special significance in XML. To avoid processing these S language elements as XML, one must escape them. This is why the function definition (i.e. the right hand side) is enclosed within a ¡![CDATA[ ... ]]; construction. This indicates the XML parser that the text within the [ ] pair is to be read verbatim.

Note also that this is not needed in place. For example, the R processing instruction (¡?R ..¿ near the end of the document) does not require escaping the R command(s). This is because XML parsers recognize this tag type as being special and escape its contents automatically.

We are definitely giving up some readability in this format. For this to work, we must provide good tools that are easy to use to generate and work with these types of documents.

??

```
⟨ ??⟩≡
 <?xml version="1.0"?>
 <!DOCTYPE functions>
 <functions>
 <functionDef access="protected">
 <name>as</name>
 <comments>
  return a the version of this object coerced to be the given Class
  If the corresponding 'is' relation is true, it will be used.
                                                                   In particular,
  if the relation has a coerce method, the method will be invoked on 'object'.
  If the 'is' relation is FALSE, and the coerceFlag is TRUE,
  the coerce function will be called (which will throw an error if there is
  no valid way to coerce the two objects). Otherwise, NULL is returned.
 </comments>
 <def>
 <![CDATA[
   function(object, Class, coerceFlag = T) {
      thisClass <- Class(object)
      if(thisClass == Class)
       return(object)
      if(is(object, Class)) {
        ## look for coerce method or indirection
        thisClass <- Class(object)</pre>
       coe <- extendsCoerce(thisClass, Class)</pre>
        coe(object)
      else if(coerceFlag)
        coerce(object, new(Class, force=T))
```

```
else
      NULL
]]>
</def>
</functionDef>
<functionDef>
<name>extendsCoerce</name>
<comments>
  the function to perform coercion based on the is relation
 between two classes. May be explicitly stored in the metadata or
  inferred. If the latter, the inferred result is stored in the session
  metadata for fromClass, to save recomputation later.
</comments>
<def>
<![CDATA[
  function(fromClass, Class) {
    ext <- findExtends(fromClass, Class)</pre>
    f <- NULL
    if(is.list(ext)) {
      coe <- ext$coerce
      if(is.function(coe))
        return(coe)
      by <- list$by
      if(length(by) > 0)
        f <- substitute(function(object)</pre>
                         as(as(object, BY), CLASS), list(BY = by, CLASS=Class))
      ## else, drop through
    if(is.null(f)) {
      ## Because 'is' was TRUE, must be a direct extension.
      ## Copy slots if the slots are a subset. Else, just set the
      ## class. For VIRTUAL targets, never change the object.
      virtual <- isVirtualClass(Class)</pre>
      if(virtual)
        f <- function(object)object</pre>
      else {
        fromSlots <- slotNames(fromClass)</pre>
        toSlots <- slotNames(Class)
        sameSlots <- (length(toSlots) == 0</pre>
            | (length(fromSlots) == length(toSlots) &&
                !any(is.na(match(fromSlots, toSlots))))
      if(sameSlots)
          f <- substitute(function(object){Class(object) <- CLASS; object},</pre>
                           list(CLASS = Class))
      else
        f <- substitute(function(object) {</pre>
          value <- new(CLASS)</pre>
          for(what in TOSLOTS)
            slot(value, what) <- slot(object, what)</pre>
          value }, list(CLASS=Class, TOSLOTS = toSlots))
      ## we dropped through because there was no coerce function in the
```

```
## extends object. Make one and save it back in the session metadata
      ## so no further calls will require constructing the function
      if(!is.list(ext))
        ext <- list()
      ext$coerce <- f
      ClassDef <- getClass(fromClass)</pre>
      allExt <- getExtends(ClassDef)</pre>
      allExt$Class <- ext</pre>
      setExtends(ClassDef, allExt)
    }
    f
  }
]]>
</def>
</functionDef>
      x <- 1:10?>
</functions>
```

We parse the document and create the tree that contains the different elements within the XML document. Note that we are not interested in the DTD at this point, so we instruct the parsing function to omit translating it to S.

```
?? \langle ??\rangle+\equiv doc <- xmlTreeParse("/tmp/as.S", getDTD = F)
```

So now we want to get at the contents of the data. We get the top node of the document using the *xmlRoot()* function.

```
?? \langle ?? \rangle + \equiv
r \leftarrow xmlRoot(doc)
```

This is the node which is referenced in the DOCTYPE and whose name is functions. We can find its name using the *xmlName()* function.

```
?? \langle ??\rangle+\equiv xmlName(r)
```

We can determine how many sub-nodes this root node has by calling the function xmlSize().

```
?? \langle ?? \rangle + \equiv xmlSize(r)
```

In this case, the result is 3. That means that it has 3 children which are themselves *XMLNode* objects. We can access the different child nodes using the [ [ operator. For example, we can get the first child node with the command.

```
?? \langle ??\rangle +=
    r[[1]]

This is also an object of class XMLNode.
    We can ask it for its name and number of children, i.e. size.
?? \langle ??\rangle +=
    xmlName(r[[1]])
    xmlSize(r[[1]])
```

```
Most all XML nodes have attributes corresponding to the
       name="value" pairs within the tag start element. For
       example, the first function definition in our example (i.e. the first
       sub-child of the root node) has a access attribute with a
       value protected. We can retrieve a name character vector of
       a node's attributes via the function xmlAttrs().
     ⟨ ??⟩+≡
??
      xmlAttrs(r)
       Lets get the first function definition object.
       This is the first child of the top-level document object.
     ⟨ ??⟩+≡
??
      r[[1]]
     This is itself an object of class XMLNode and so has a tag
       name, attributes and children.
       It has $3$ children whose tag names are
       given by
22
     ⟨ ??⟩+≡
      > sapply(xmlChildren(r[[1]]), xmlName)
                                    def
                  comments
            name
           "name" "comments"
                                   "def"
     Applying an operation to children of a node is so common that we
       provide functions xmlApply() and xmlSApply() which
       are simple wrappers whose primary role is to fetch the list of
       children of the specified node. (The apply functions are not generic.)
??
     ⟨ ??⟩+≡
       xmlSApply(r[[1]], xmlName)
       xmlApply(r[[1]], xmlAttrs)
       xmlSApply(r[[1]], xmlSize)
     Let's look further at this functionDef element in the tree.
       As we see, it has three sub-nodes named name,
       comments and def.
       Let's grab the name
       element first.
     ⟨ ??⟩+≡
??
      r[[1]][[1]]
       Again, this is of class XMLNode.
     ⟨ ??⟩+≡
??
      > class(r[[1]][[1]])
      [1] "XMLNode"
```

and it has a single child whose class is XMLTextNode. This

```
basically means we have a leaf node. Objects of class
       XMLTextNode have no children (but they are XMLNode
       so they have a slot for children!)
     ⟨ ??⟩+≡
??
      > class(r[[1]][[1]][[1]])
      [1] "XMLTextNode" "XMLNode"
     This leaf node is not the text itself, but contains that text.
       We get it using the xmlValue()
       function.
22
     ⟨ ??⟩+≡
       xmlValue(r[[1]][[1]][[1]])
     We should not that the lengthy subscripting to access nodes within
       nodes $...$ is ugly and tedious.
       Of course, one can assign these intermediate nodes to variables
       and work on these
     ⟨ ??⟩+≡
??
       x < -r[[1]]
       x < -x[[1]]
       xmlValue(x[[1]])
     I personally find this sometimes more difficult to follow. But
       there are times that it is more readable. The key point to remember
       here is that these intermediate variables are copies of the
       element in the tree. This is obvious to users of the S language, but
       is slightly unexpected for those coming with backgrounds in C/C++,
       Java^{TM}, Perl, etc. These languages (can) use references to operate on
       XML trees and so changes to sub-nodes are reflected in the bigger tree
       in which that sub-node resides. S is not ideally suited to operating
       on highly recursive, deep objects. But it is more than sufficient.
     We have seen how we can extract individual sub-nodes from an object
       of class XMLNode using the [ [ operator and giving
       its index. It will come as no surprise that we can use the [
       ( a single bracket) operator to extract a list of nodes.
       For example,
     ⟨ ??⟩+≡
??
        r[[1]][1:2]
       returns the first two elements of the root node,
       i.e. the name and comments nodes.
       Similarly to using indices, one can identify
       nodes by name.
     ⟨ ??⟩+≡
??
       r[[1]]["comments"]
```

```
??
     ⟨ ??⟩+≡
        h <- xmlTreeParse(system.file("treeParseHelp.xml", pkg="XML"))
        xmlRoot(h)[c("name", "author")]
     ⟨ ??⟩+≡
??
       > names(xmlRoot(doc))
       [1] "name"
                          "title"
                                         "description" "usage"
                                                                      "arguments"
       [6] "details"
                          "value"
                                         "references" "author"
                                                                      "notes"
       [11] "seealso"
                          "examples"
                                         "keywords"
       > names(xmlRoot(doc)[["examples"]])
       [1] "example" "example" "example" "example"
     We find out which examples have explicit descriptions stanzas by
       looking at the number of sub-nodes they have.
     ⟨ ??⟩+≡
??
       > xmlSApply(xmlRoot(doc)[["examples"]], xmlSize)
       example example example example
             1
                  2
                           1
                                     1
       So we can get the last one and look at its description.
??
     ⟨ ??⟩+≡
      xmlRoot(doc)[["examples"]][[5]][["description"]]
      We can take the code from the example
       and execute it.
     ⟨ ??⟩+≡
??
       eg <- xmlRoot(doc)[["examples"]][[5]]</pre>
       canRun <- length(xmlAttrs(eg)) == 0 | is.na(match("dontRun", names(xmlAttrs(eg))))</pre>
       if(!canRun)
         canRun <- !as.logical(xmlAttrs(eg)["dontRun"])</pre>
       if(canRun)
          eval(parse(text=eg[[2]]))
??
     ⟨ ??⟩+≡
      xmlTagByName <-
       function(node, name)
       which <- (1:xmlSize(node))[sapply(node$children, xmlName) == name]</pre>
       node$children[which]
??
     ⟨ ??⟩+≡
       lapply(xmlTagByName(d$children[[1]], "functionDef"), function(x) x$children[[2]][[1]])
```

```
Reading the entire XML document into a tree and then processing this
              tree works well in most situations. There are cases however, where it
              is more convenient to process the nodes in the tree as they are being
              created and inserted into the tree. This allows us to modify the node
              or even to discard it from the tree altogether.
            ⟨ ??⟩+≡
      ??
               fileName <- system.file("data/mtcars.xml")</pre>
              doc <- xmlTreeParse(fileName, handlers = (function() {</pre>
                                               vars <- character(0);</pre>
                                              list(variable=function(x, attrs) {
                                                               vars <<- c(vars, xmlValue(x[[1]]));</pre>
                                                               NULL },
                                                    startElement=function(x,attr){
                                                                  NULL
                                                                 },
                                                   names = function() {
                                                            }
                                              })()
            Now, suppose we want to make post-processing the tree easier. We
              can start by providing additional class information. For example,
              when reading a R source code in XML format (see xml2tex.Sxml)
              document, we might process fragment chunks by giving them an
              additional class name XMLFragmentNode. This would then allow
              us to process the resulting objects in a simpler manner, dispatching
              to different functions.
      ??
            ⟨ ??⟩+≡
               h <- list(fragment=function(x, attr){ class(x) <- c("XMLFragmentNode", class(x))
                                                     })
               doc <- xmlTreeParse(file, handlers=h, asTree=T)</pre>
            1 Creating XML Nodes
            There are two styles that can be used for creating
              nodes:
a) top-down create a top-level object
            and assign children to it,
) bottom-up create child nodes and group them
            together into container/parent nodes, and recursively
```

work ones way up the tree.

```
??
     ⟨ ??⟩+≡
       a <- xmlNode("arg", attrs = c(default="T"), xmlNode("name", "foo"), xmlNode("defaultValue
       a$children[[3]] <- xmlNode("duncan")
     The resulting tree is
??
     ⟨ ??⟩+≡
       <arg default="T">
       <name>
       foo
       </name>
       <defaultValue>
       1:10
       </defaultValue>
       <duncan>
       </duncan>
       </arg>
        The worse form of generating a is
??
     ⟨ ??⟩+≡
       a <- xmlNode("arg", attrs = c(default="T"),</pre>
                       xmlNode("name", xmlTextNode("foo")), xmlNode("defaultValue",xmlTextNode("1
```

%

## 2 Writing XML Output

If one has a tree of XMLNode objects in S, then the basic print methods for these classes will generate XML that can be put in a file or generally used outside of S. But what about translating data in S into XML so that it can be used elsewhere, e.g sent to Matlab, put on the web, communicated directly to another application via SOAP, etc. How do we go about generating the XML text to represent an object?

Well, take a look at StatDataML