SPASE Resource Tools

Version 2.0.0

Table of Contents

[Overview 1](#_Toc272906088)

[Installation 1](#_Toc272906089)

[System Requirements 1](#_Toc272906090)

[Installation 1](#_Toc272906091)

[Tools 2](#_Toc272906092)

[Collator 3](#_Toc272906093)

[Downloader 4](#_Toc272906094)

[Profiler 5](#_Toc272906095)

[Validator 7](#_Toc272906096)

[RefCheck 8](#_Toc272906097)

[XMLGrep 9](#_Toc272906098)

[Appendix A: Release Notes 10](#_Toc272906099)

[Releases 10](#_Toc272906100)

# Overview

The SPASE (Space Physics Archive Search and Extract) Toolkit contains a set of command-line applications which can be used to generate, validate, referentially check, use and organize resource descriptions written in SPASE XML. The toolkit is written in Java.

# Installation

The SPASE toolkit is written entirely in Java and should run on any system with a Java Runtime Environment (JRE). Any required extensions are included in the SPASE toolkit installation package. The toolkit is packaged as a self-contained executable "jar" file.

## System Requirements

* Java JRE 1.5 or higher

## Installation

* Download the SPASE toolkit jar file from the SPASE web site.  
  The most recent release can be found at:

http://www.spase-group.org/tools/toolkit/spase-tools-2.0.0.jar

* Place the jar file in any directory that you want.

# Tools

All tools are written in Java and are part of the "org.spase.tools" package. Each tool has a common name which can be used with the executable jar package. A tool can run with a command like:

[node] java –jar spase-tools-2.0.0.jar [toolname] [tooloptions]

Each tool also has a Java class which can be called by other programs. Tool class names are in proper case.

## Collator

Separate each SPASE resource description in a file into a separate file stored in a folder tree according to the Resource ID.

Optionally recursively scan a directory for all files with a given extension

and process each file.

Usage:

collator [options] file

-or-

java org.spase.tools.Collator [options] file

Options:

|  |  |
| --- | --- |
| -b,--base <arg> | Base path for collated output (default: .). |
| -h,--help | Dispay this text |
| -k,--check | Check files, but do not write collated output. |
| -r,--recurse | Recursively process all files starting at path. |
| -v,--verbose | Verbose. Show status at each step |
| -x,--ext <arg> | File name extension for filtering files when processing folders (default: .xml) |

Acknowledgements

Development funded by NASA's VMO project at UCLA.

Example

Suppose a file (example.xml) contains three SPASE descriptions like:

<Spase>

<Version>2.0.0</Version>

<NumericalData>

<ResourceID>spase://VMO/NumericalData/GeoTail/LEP/PT60S</ResourceID>

... Details omitted ...

</NumericalData>

<NumericalData>

<ResourceID>spase://VMO/NumericalData/GeoTail/MGF/PT60S</ResourceID>

... Details omitted ...

</NumericalData>

<NumericalData>

<ResourceID>spase://VMO/NumericalData/GeoTail/CPI/PT60S</ResourceID>

... Details omitted ...

</NumericalData>

</Spase>

Then a command like:

collator example.zip

will generate three files with the path names of:

./VMO/NumericalData/GeoTail/LEP/PT60S.xml

./VMO/NumericalData/GeoTail/MGF/PT60S.xml

./VMO/NumericalData/GeoTail/CPI/PT60S.xml

With each file containing the description for the corresponding resource.

## Downloader

Obtains a list of URLs associated with a resource by querying a registry server, then downloads and packages all the source files. The collection of files is packaged into a zip file and written to the output file.

Usage:

org.spase.tools.Downloader [options] id

Options:

|  |  |
| --- | --- |
| -b, --startdate | The start date of the desired time span. |
| -e, --stopdate | The stop date of the desired time span. |
| -h,--help | Display this text |
| -c, --check <arg> | Check a download package. |
| -o,--output <arg> | Output filename (default: resource.zip). |
| -s,--service <arg> | The URL to the registry service which generates download packages (default: http://www.spase-group.org/registry/downloader). |
| -v,--verbose | Verbose. Show status at each step |

Acknowledgements

Development funded by NASA's VMO project at UCLA.

Example

To download all Granules associated with the resource with the SPASE ResourceID of:

spase://VMO/NumericalData/AMPTE\_UKS/Plasma/SWI\_PT5S

use the command:

downloader -o example.zip spase://VMO/NumericalData/AMPTE\_UKS/Plasma/SWI\_PT5S

which will collect all data files (granules) associated with the resource, package them in a zip file and write the file to "example.zip".

To download only granules that span a time range use a command like:

downloader -o example.zip spase://VMO/NumericalData/AMPTE\_UKS/Plasma/SWI\_PT5S \

-b 1990-12-01T00:00:00 -e 2000-01-01T00:00:00

which will collect only those data files (granules) which contain data between   
1990-12-01T00:00:00 and 2000-01-01T00:00:00.

To check a package to make sure it is complete and intact:

downloader -c example.zip

## Profiler

Profile generator. Create resource profiles for SPASE resource descriptions.

Profiles all have a common schema which can be used in a solr search engine.

Usage:

java org.spase.tools.Profiler [options] [file...]

Options:

|  |  |
| --- | --- |
| -f,--file <arg> | File. File containing a list of file names to scan. |
| -h,--help | Display this text |
| -i,--id <arg> | ID. The registry ID to set for each resource |
| -l,--lookup <arg> | Lookup. THe URL to the resource lookup service to resolve resource IDs. Default: http://www.spase-group.org/registry/ |
| -o,--output <arg> | Output. Output generated profiles to {file}. Default: System.out. |
| -r,--recurse | Recurse. Process all items in the current folder. Recurse into sub-folders. |
| -v,--verbose | Verbose. Show status at each step. |
| -x,--extension <arg> | Extension. The file name extension for files to process (default: .xml) |

Acknowledgements:

Development funded by NASA's VMO project at UCLA.

Example

To create profiles for all resources in the current directory and below use the command:

profiler -o /temp/vmo.xml -r \*

The profiles will be written to the file "/temp/vmo.xml". The profiles can then be posted the appropriate solr search engine.

The expected schema for the solr index is:

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Field Name | Type | Indexed | Stored | Required | Multi-valued |
| resourceid | string | true | true | true |  |
| registryid | string | true | true | true |  |
| resourcetype | string | true | true |  |  |
| resourcename | text | true | true |  |  |
| measurementtype | text | true | true |  |  |
| phenomenontype | text | true | true |  |  |
| observedregion | text | true | true |  |  |
| observatoryid | string | true | true |  |  |
| observatoryname | text | true | true |  |  |
| observatorytype | text | true | true |  |  |
| observatorygroup | text | true | true | true |  |
| instrumentid | string | true | true |  |  |
| instrumentname | text | true | true |  |  |
| instrumenttype | text | true | true |  |  |
| releasedate | date | true | true |  |  |
| startdate | date | true | true |  |  |
| stopdate | date | true | true" |  |  |
| cadence | string | true | true" |  |  |
| latitude | string | true | true |  |  |
| longitude | string | true | true" |  |  |
| description | text | true | true |  |  |
| authority | text | true | true |  |  |
| association | text | true | true |  | true |
| word | text | true | false |  | true |

## Validator

SPASE Resource Description grammar checker.

Checks a resource description for compliance to a specified version of the SPASE data model.

Usage:

validator [options] file

-or-

java org.spase.tools.Validator [options] file

Options:

|  |  |
| --- | --- |
| -h,--help | Dispay this text |
| -n,--version <arg> | Version of standard schema to use available from www.spase-group.org (default: 1.2.1) |
| -r,--recurse | Recursively process all files starting at path. |
| -s,--schema | Path to the XML schema document (XSD) to use for checking files. |
| -v,--verbose | Verbose. Show status at each step |
| -x,--ext <arg> | File name extension for filtering files when processing folders (default: .xml) |

Acknowledgements

Development funded by NASA's VMO project at UCLA.

Example

To validate the SPASE description in the file "example.xml" to version 2.0.0 of the SPASE schema use the command:

validator -n 2.0.0 example.xml

The schema will be loaded from the web site [www.spase-group.org](http://www.spase-group.org). If you are not one a network you can use a local XML Schema document with the command:

validator -s spase-2\_0\_0.xsd example.xml

## RefCheck

SPASE Resource Description reference checker.

Can check both resource identifiers and URLs for referential integrity.

Can also produce lists of identifiers and URLs.

Usage:

refcheck [options] file

-or-

java org.spase.tools.RefCheck [options] file

Options:

|  |  |
| --- | --- |
| -h,--help | Dispay this text |
| -i,--identifier | Check each identifier in the resource description. |
| -l,--list | List all identifiers and URLs. Do not perform referential checks. |
| -r,--recurse | Recursively process all files starting at path. |
| -s,--service <arg> | The URL to the registry service to look-up resource identifiers (default: http://www.spase-group.org/registry/lookup). |
| -u,--urlcheck | Check each all URLs in the resource description. |
| -v,--verbose | Verbose. Show status at each step |
| -x,--ext <arg> | File name extension for filtering files when processing folders (default: .xml) |

Acknowledgements

Development funded by NASA's VMO project at UCLA.

Example

To check that all Resource IDs in all files can be resolved:

refcheck -i -r \*

To check all Resource IDs and URLs use the command:

refcheck -i -r \*

To see detailed information while checking add the verbose flag (-v):

refcheck -v -i -r \*

## XMLGrep

XML Parser, XPath generator and search tool.

Parses an XML file and flattens the document content.

Values can be retrieved using an XPath that can contain regular expressions.

All values in the XML file can be listed with a corresponding XPath.

Usage:

java org.spase.tools.XMLGrep [options] file

Options:

|  |  |
| --- | --- |
| -e,--extract <arg> | Extract. Extract all nodes with a given XPath |
| -f,--find <arg> | Find. Locate the value associated with an XPath |
| -h,--help | Display this text |
| -n,--nodes <arg> | Nodes. List all nodes at the given XPath |
| -v,--verbose | Verbose. Show status at each step |

Acknowledgements:

Development funded by NASA's VMO project at UCLA.

Example

To find all items in an XML file that have an XPath ending in "ResourceID" use the command:

xmlgrep -f ".\*/ResourceID" example.xml

will output something like:

/Spase/NumericalData/ResourceID: spase://VMO/NumericalData/IMP8/MAG/PT15.36S

To list the XPath tagged list of all content in an XML file use the command:

xmlgrep example.xml

will generate a list like:

/Spase/Version: 2.0.0

/Spase/NumericalData/ResourceID: spase://VMO/NumericalData/IMP8/MAG/PT15.36S

/Spase/NumericalData/ResourceHeader/ResourceName: IMP 8 Magnetic Field

/Spase/NumericalData/ResourceHeader/ReleaseDate: 2008-06-02T18:53:44Z

... and so on ...

# Appendix A: Release Notes

All Tools written by Todd King

# Releases

Release 1.0.0: Initial release.

Release 2.0.0: Aligned with current SPASE registry services.