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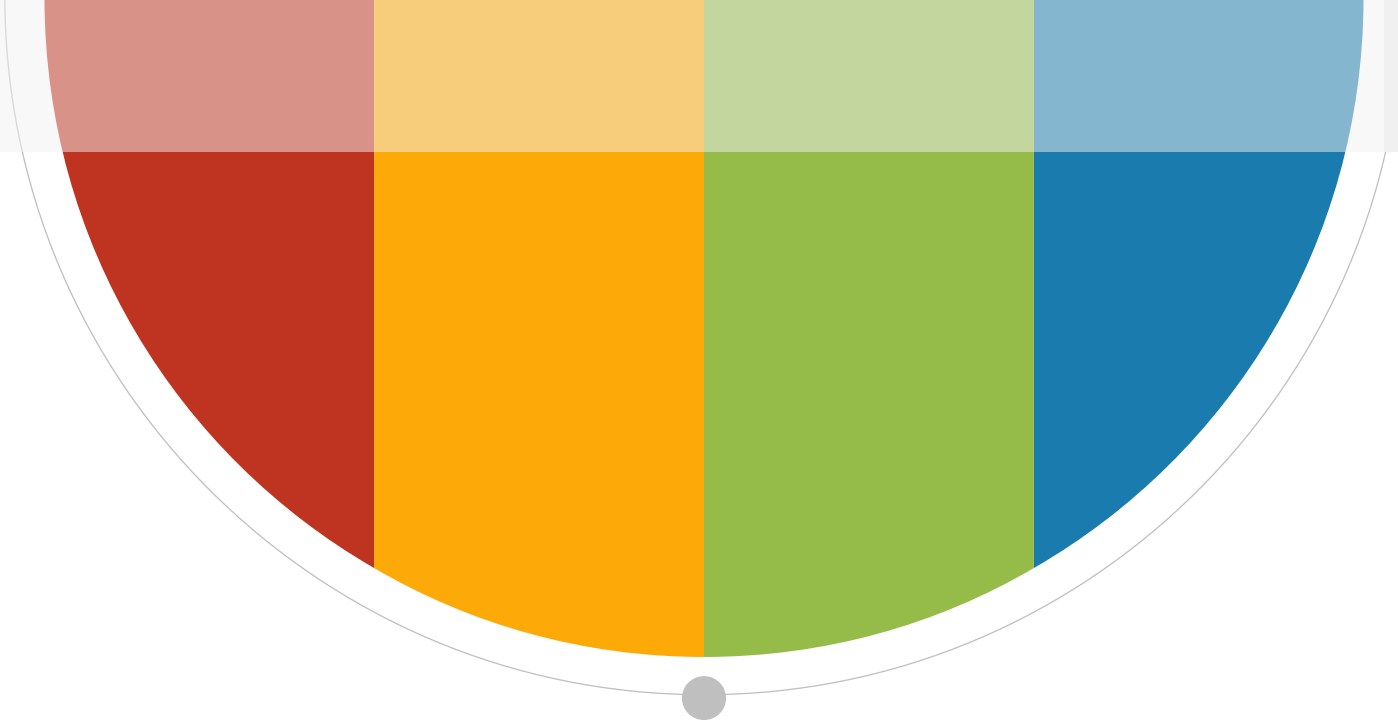

CAFE: a Collaborative Analysis Framework for distributed Environmental data

Hao Xu
On behalf of the CAFE Team

Dec.7th 2016

Outline

01 CAFE Introduction

02 System Design

- ✓ Logical design
- ✓ Modularity
- ✓ Extensibility

03 Prototype and Use Case

04 Conclusions

Introduction-about CAFE

CAFE is a dedicated software package for **collaborative analysis** of large volumes of **distributed environmental data.**

Key features:

1) Computing near the data;
2) data is logically grouped, while physically distributed;
3) Analytic tasks are divided as subtasks, and then fulfilled on corresponding nodes;
4) Easy way to enrich the built-in analytic functions;
5) Open source projects on github

Introduction-why CAFE?

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Typical workflow

- *Download data from multiple nodes*
- *Data subsetting*
- *Data regridding*
- *Data averaging*
- *Writing analytic scripts*
- *Executing codes locally*
- *Result visulization*
- *Result analysis*
- *.....*

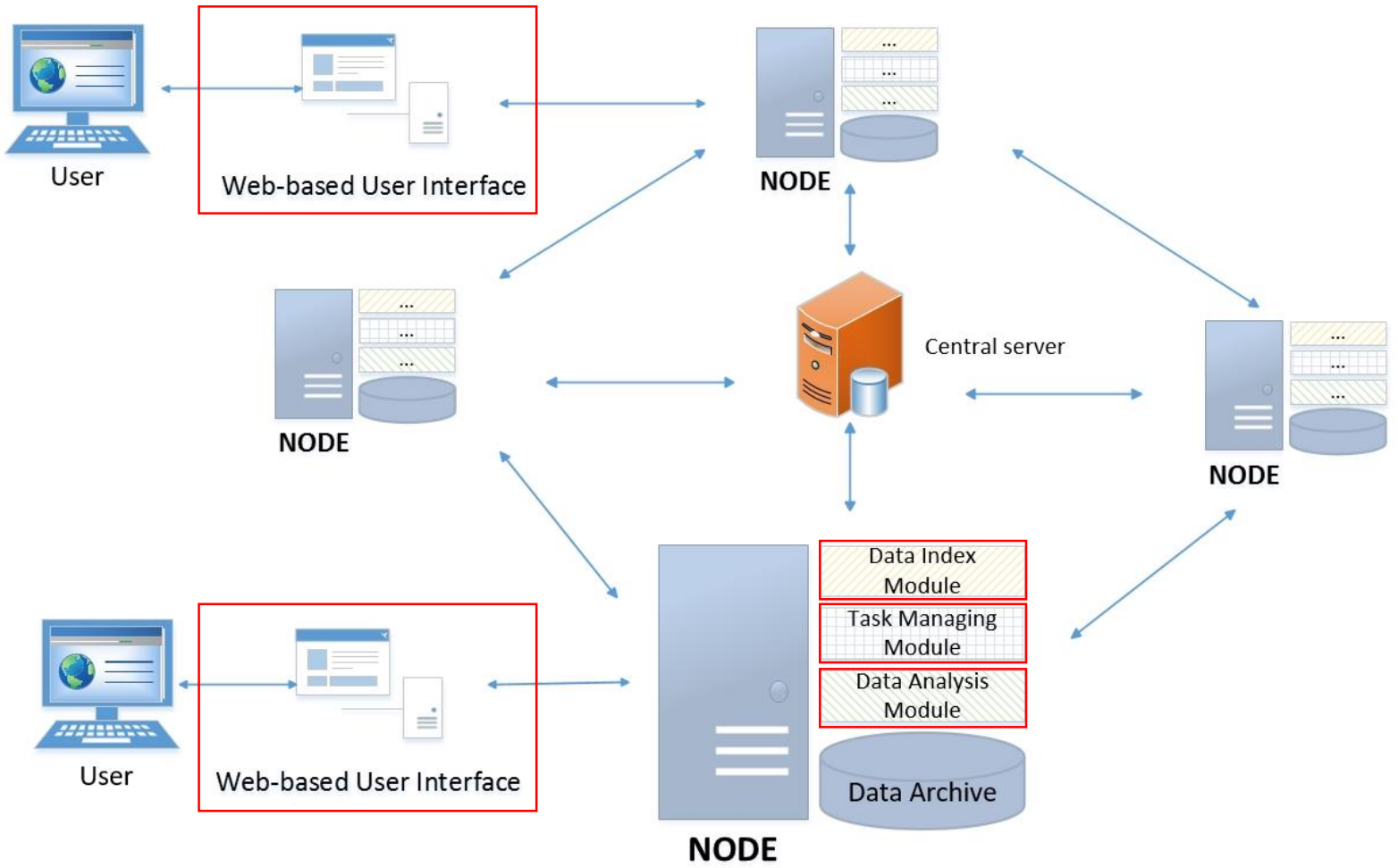
CAFE Features

- *Web-based UI provided*
- *REST APIs availbale*
- *One-stop service for data discovery, visualization, and analysis*
- *User-transparent*
- *Task management*
- *Multi-node collaboration*
- *Support for data intercomparison*
- *Multiple built-in analytic functions*
- *Easy way for extensions*
- *Batch analysis*

.....

System design-logical design

A p2p architecture



System design-modularity

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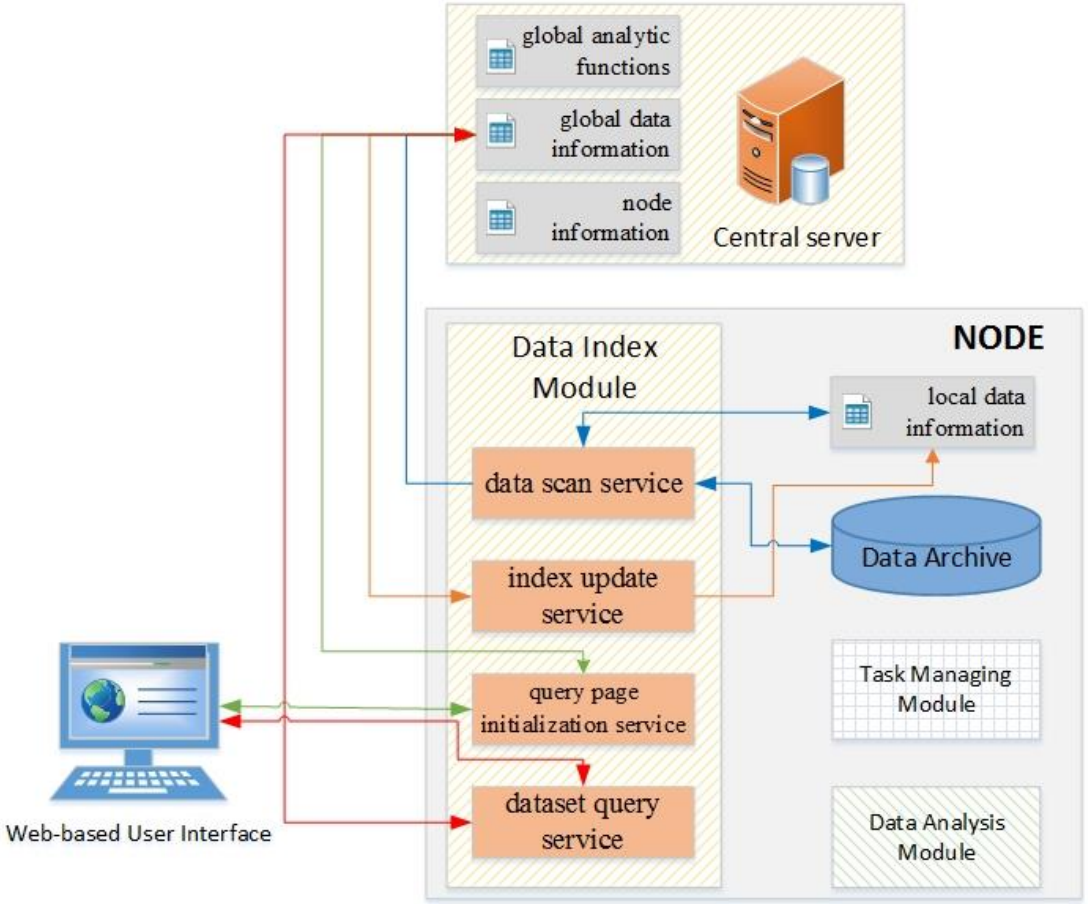
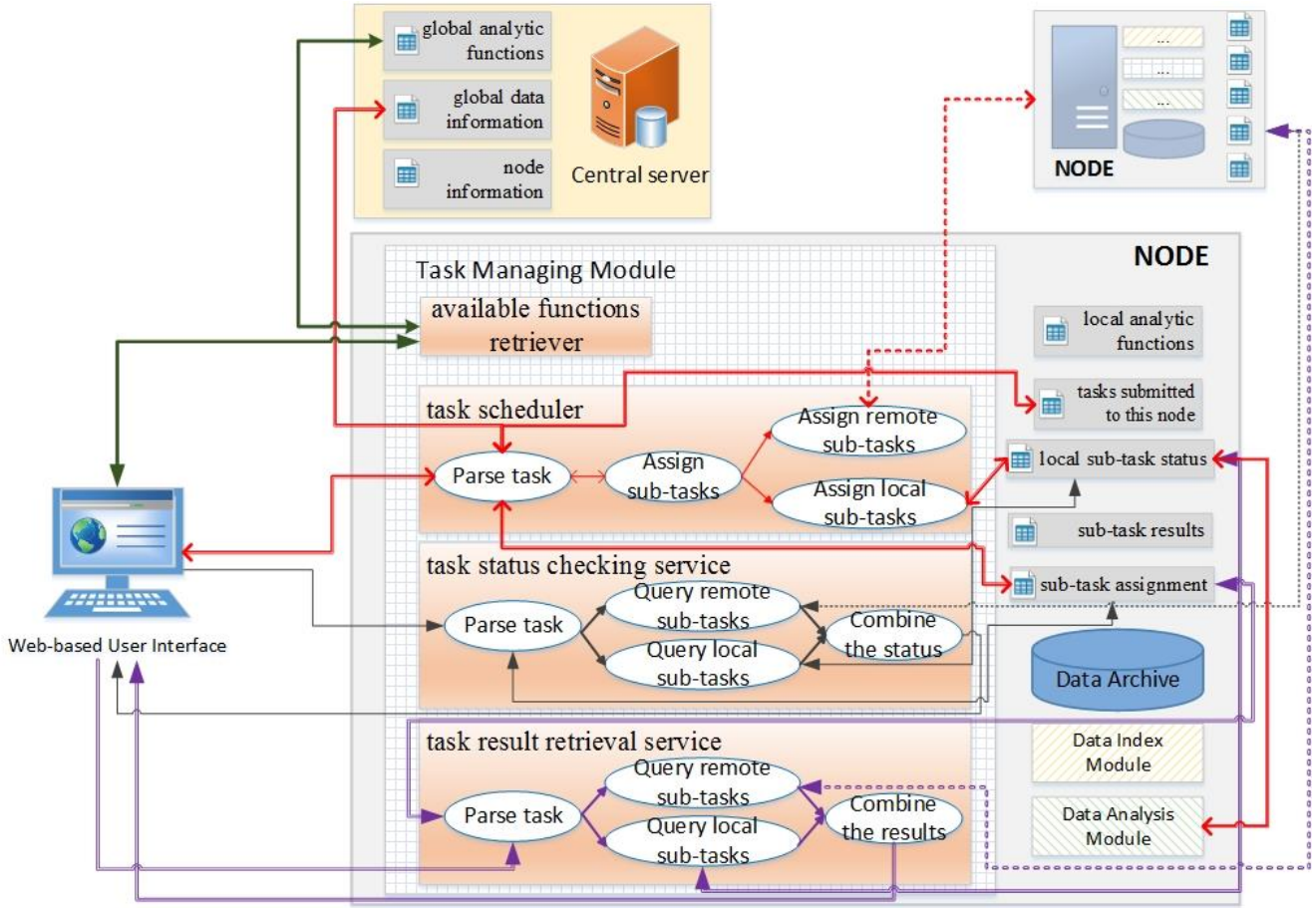
Data Index Module



Task Managing Module



System design-modularity

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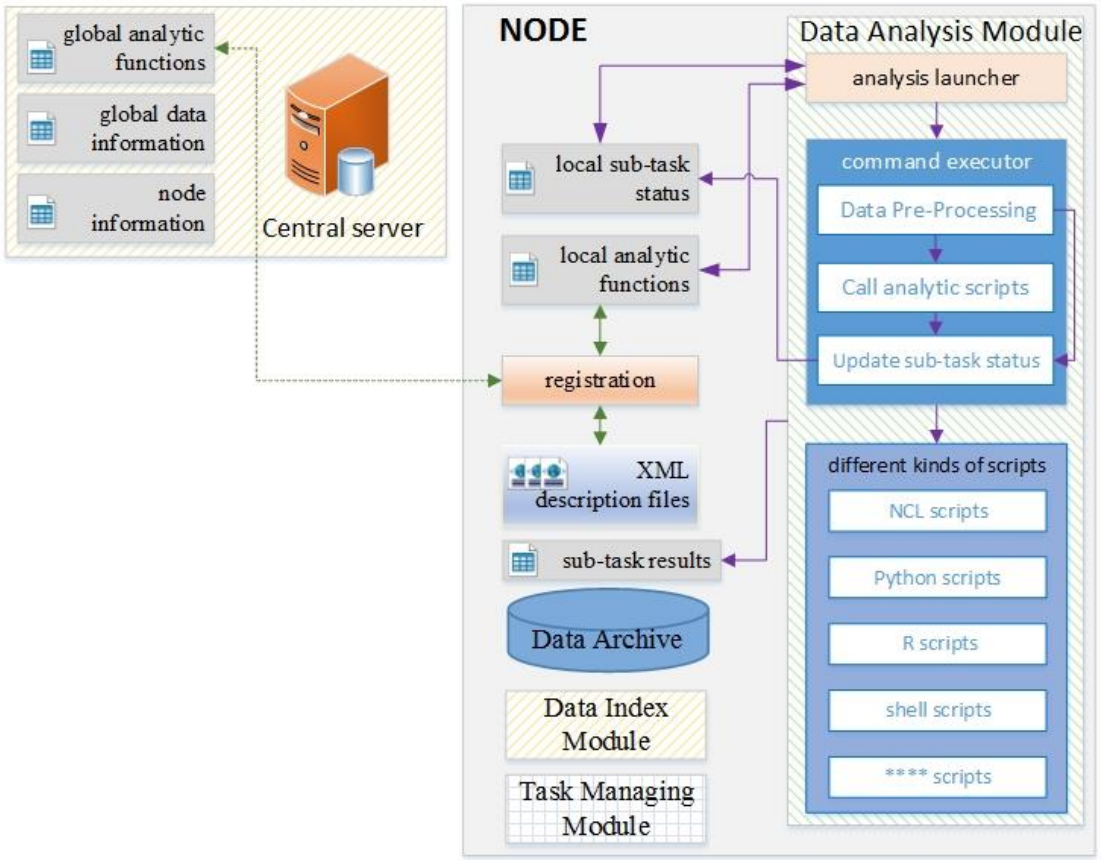
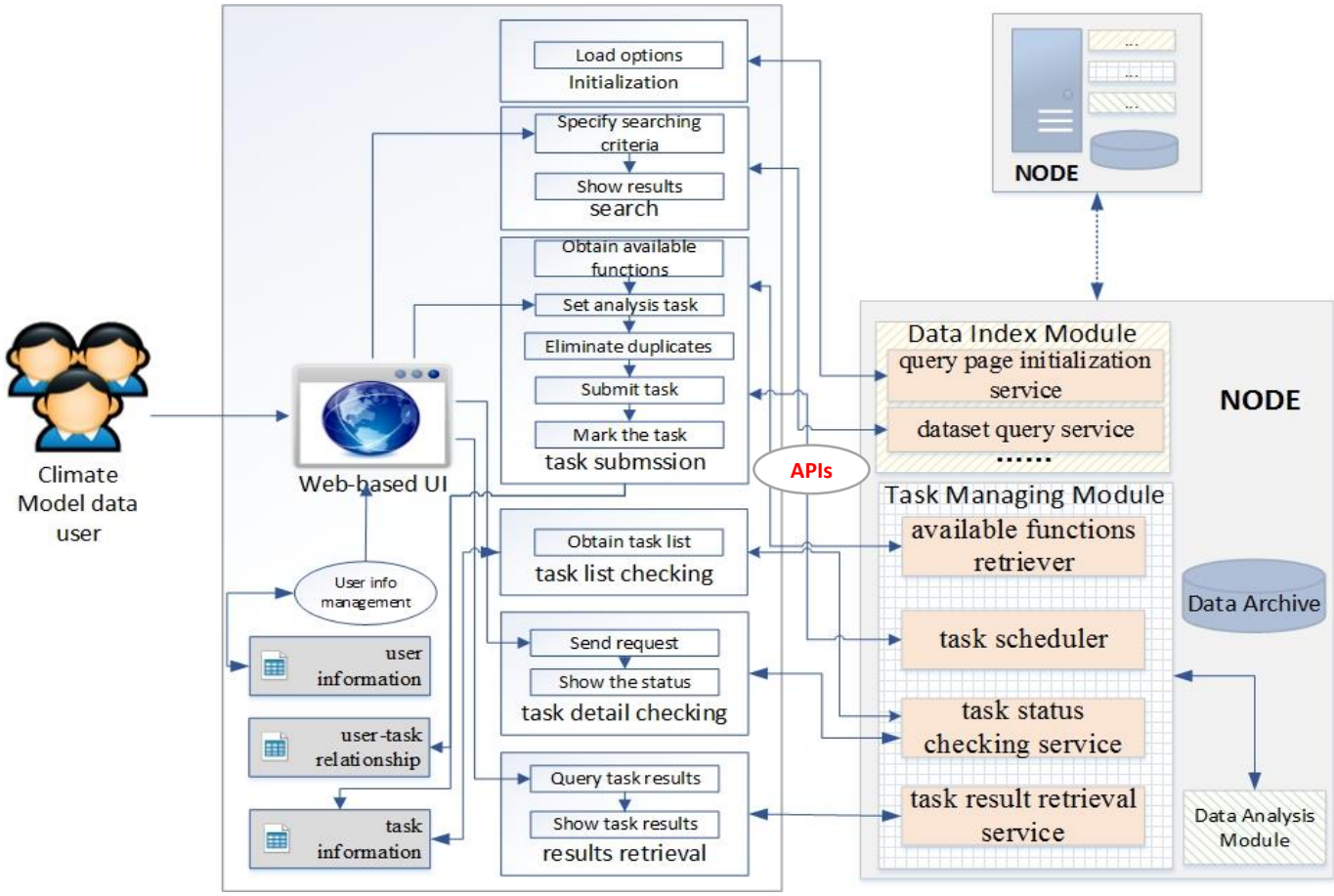
Data Analysis Module



Web-based User Interface



System design-extensibility


```
<?xml version="1.0"?>
<function lang="NCL">  <!--defining the language of the analytic function-->
    <name>PolarNHEOF</name>  <!--the name of the analytic function-->
    <script>PolarNHEOF.ncl</script> <!--the file name the script-->
    <description>xxx</description> <!--description of the function-->
    <isGlobalFunction>true</isGlobalFunction>
    <!--defining if the function can be distributed to all the nodes-->
    <MultiInputFiles>false</MultiInputFiles>
    <!--defining if the function can be distributed to all the nodes-->
    <InputFileFormat>netCDF</InputFileFormat> <!--setting format of the input file-->
    <properties>
        <!--defining which datasets can use this function, type can be "include" or "exclude"-->
        <!--the contributor can define model,frequency,modelingRealm and variableName values for filtering-->
        <Model type="include"><!--refer to PCMDI documents to get acceptable values-->
            <value>xxx</value>
            ...<!--multiple values-->
        </Model>
        ... <!--frequency,modelingRealm,variableName-->
    </properties>
    <Controls>  <!--setting the parameters that input from the webpage-->
        <parameter>
            <!--defining name,description,tag,type and value range of the parameter-->
        </parameter>
        ... <!--multiple parameters-->
    </Controls>
    <InputFileParameters>  <!--setting the parameters about the input file(s)-->
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    </InputFileParameters>
    <OutputFileParameters>  <!--setting the parameters about the output file(s)-->
        <result>
            <filetype>xxx</filetype> <!--png/nc/txt...-->
            <filecount>1</filecount>
            <parameter>
                <!--defining name,description,tag,type and value range of the parameter-->
            </parameter>
            ...
        </result>
        ... <!--multiple kinds of results-->
    </OutputFileParameters>
    <pre-processing> <!--defining pre-processing type before invoking the script-->
        <pre-processing-type>Latlon</pre-processing-type> <!--Origin/Latlon/YearAvg/SeasAvg/LTM/Subset-->
        <AddToResults>false</AddToResults>
        <!--defining if the pre-processing result need to be added to the result files-->
    </pre-processing>
</function>
```

properties

controls

Input/Output related

Pre-processing

Easy way for extensions. Only the analytic script and its XML description are needed.

The script should have an I/O interface for command line and can be invoked by Java.

```
#Example: for NCL
if(.not.isvar("date")) then
   date=19000101
end if
if(.not.isvar("filename")) then
    filename="test.nc"
end if   print(date)
print(filename)
#command invoking: ncl test.ncl
date=19000120'filename="test1.nc"'
```

System design-extensibility


```xml
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```

properties

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Input/Output related

Pre-processing

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Shell:

```
#test.sh
    #!/bin/sh
    date=$1
    filename=$2
    echo "date:${date} filename:${filename}"
#command invoking: sh test.sh 19000101 test.nc
```

System design-extensibility


```
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```

properties

controls

Input/Output related

Pre-processing

Easy way for extensions. Only the analytic script and its XML description are needed.

The script should have an I/O interface for command line and can be invoked by Java.

Python:

```
#test.py
    import sys
    date=sys.argv[1]
    filename=sys.argv[2]
    print date,filename
#command invoking: python test.py 19000101 test.nc
```

System design-extensibility


```xml
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<function lang="NCL">  <!--defining the language of the analytic function-->
    <name>PolarNHEOF</name>  <!--the name of the analytic function-->
    <script>PolarNHEOF.ncl</script> <!--the file name the script-->
    <description>xxx</description> <!--description of the function-->
    <isGlobalFunction>true</isGlobalFunction>
    <!--defining if the function can be distributed to all the nodes-->
    <MultiInputFiles>false</MultiInputFiles>
    <!--defining if the function can be distributed to all the nodes-->
    <InputFileFormat>netCDF</InputFileFormat> <!--setting format of the input file-->
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            <filecount>1</filecount>
            <parameter>
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            </parameter>
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        </result>
        ... <!--multiple kinds of results-->
    </OutputFileParameters>
    <pre-processing> <!--defining pre-processing type before invoking the script-->
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        <AddToResults>false</AddToResults>
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    </pre-processing>
</function>
```

properties

controls

Input/Output related

Pre-processing

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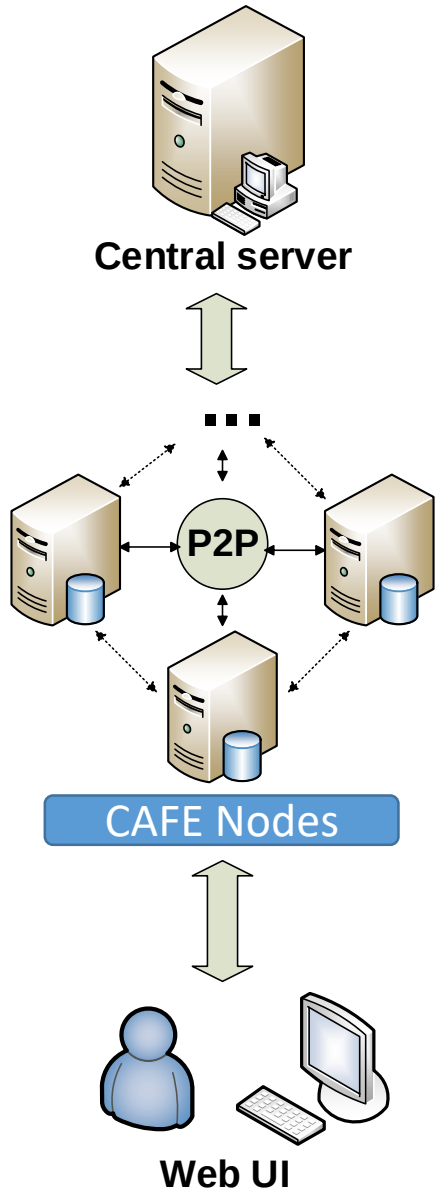
The script should have an I/O interface for command line and can be invoked by Java.

R:

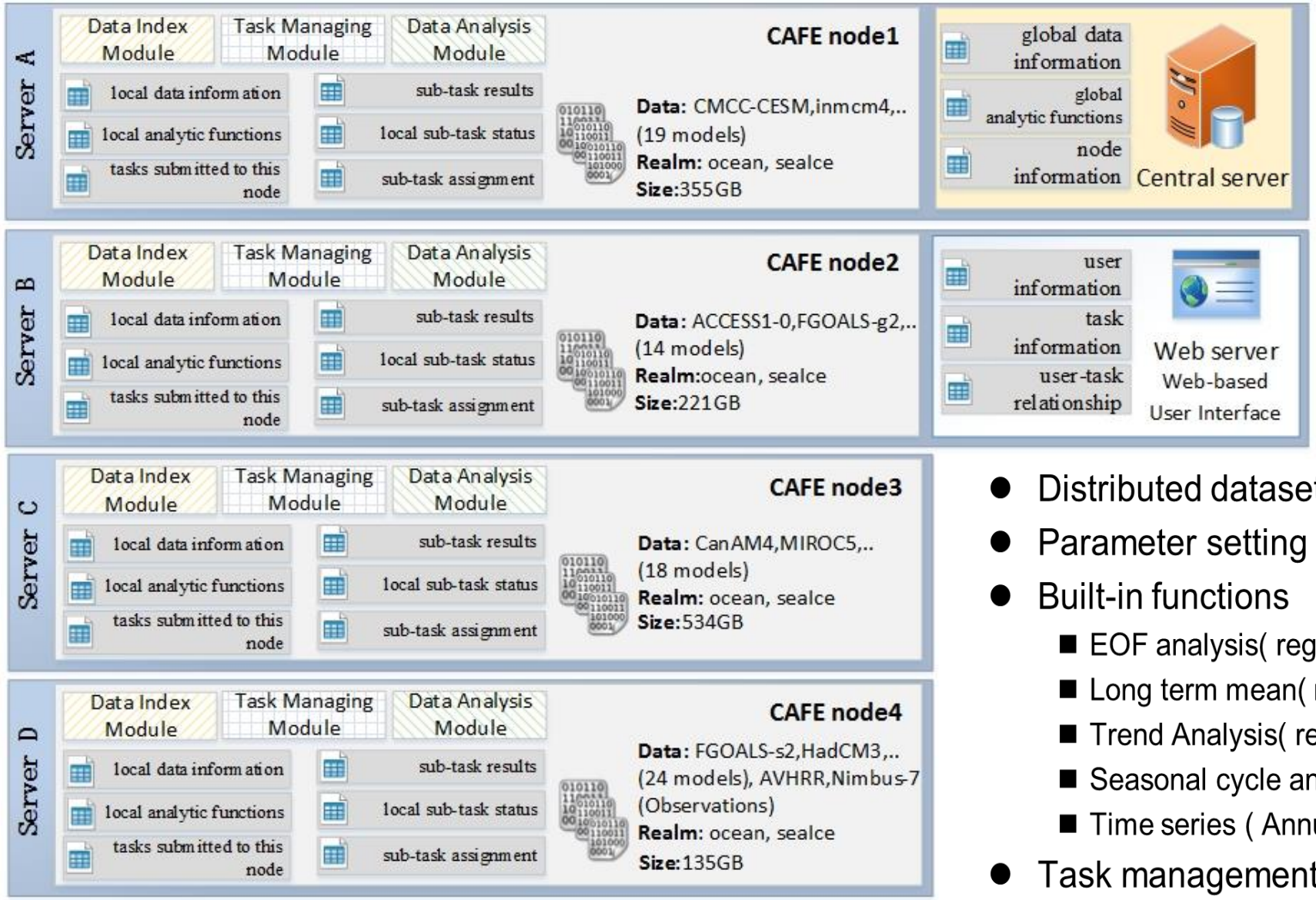
```r
#test.R
    args <- commandArgs(trailingOnly = TRUE)
    date<- as.numeric(args[1])
    filename<- as.character(args[2])
    print(date)
    print(filename)
#command invoking: Rscript test.R 19000101 test.nc
```

Prototype and Use Case-prototype system

- **CAFE nodes:**
  - Data layer:mybatis3.2.3+mysql
  - Service layer: Spring4.0
  - Interaction: Spring MVC4.0+REST API
- **Analytic scripts:**
  - NCL+NCO+CDO
- **Web-based UI:**
  - Service: pHp+yii
  - Database: mysql
  - Web Server: apache2



Prototype and Use Case-prototype system



- Distributed dataset access and searching
- Parameter setting and task submitting
- Built-in functions
  - EOF analysis(region, NH, SH)
  - Long term mean(region, NH, SH)
  - Trend Analysis(region, NH, SH)
  - Seasonal cycle analysis(NH, SH)
  - Time series (Annual, Seasonal)
- Task management, result display and downloading

Prototype and Use Case-use case

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- Compare and analyze sea surface temperature data(tos) among different model output and the observation data
- Model: GFDL-CM3 and IPSL-CM5B-LR
- Observation data: AVHRR
- Time range: 198501~200512
- Spatial range: 0° N~90° N , 110° E~280° E
- Analysis method: EOF Analysis (Specified regions)

Prototype and Use Case-use case

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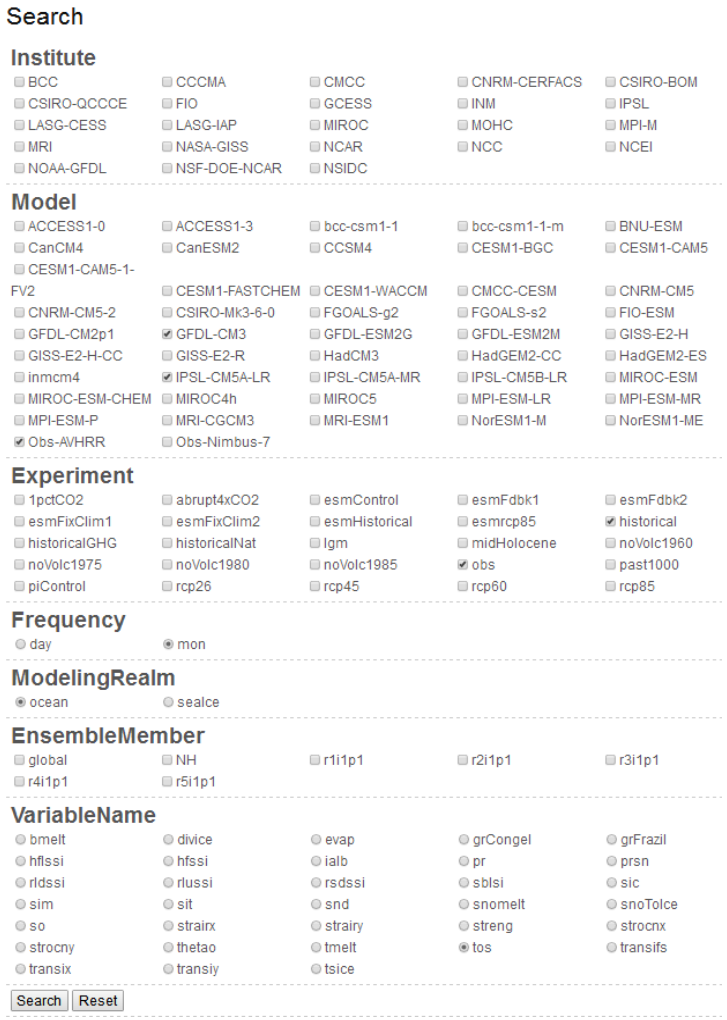
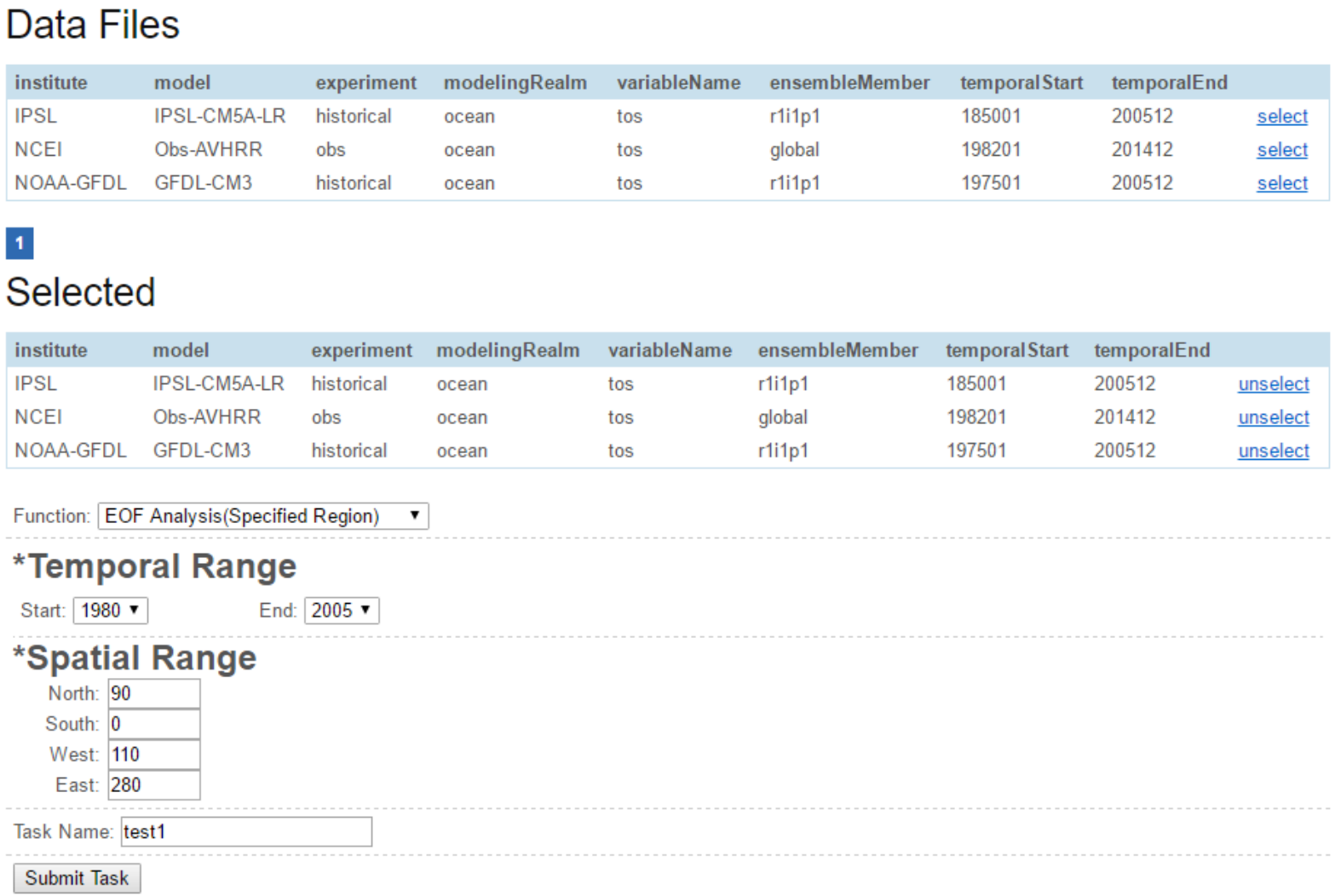
- At first, user can search for data by specifying querable attributes. Selected three datasets may be archived on several nodes.



Prototype and Use Case-use case

- At first, user can search for data by specifying querable attributes. Selected three datasets may be archived on several nodes.



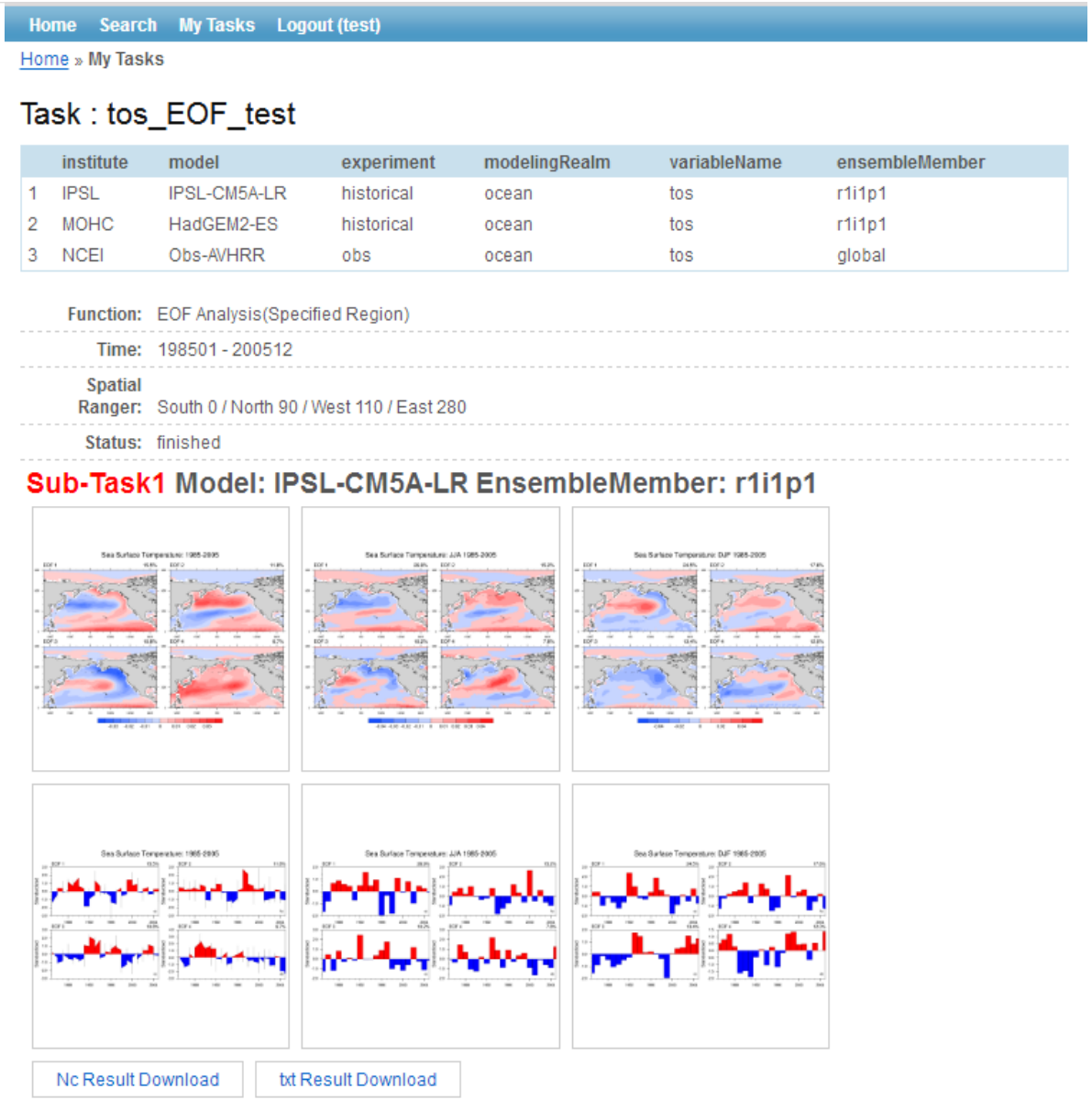
Prototype and Use Case-use case

- After submitting the task, user can check status of each task.
- User can obtain the results when the status is *finished*.

Prototype and Use Case-use case

- Maps and time series graph are provided.

- Results as NetCDF files and Text files are provided for downloading.



Conclusion

- CAFE can support web-based online batch analysis of distributed environmental data, as well as multi-node collaboration for data intercomparison.
- CAFE can dramatically decrease the amount of data need to be transferred from data centers to data users.
- CAFE can be easily extended to support more data analytics functions and more data formats.
- CAFE is very promising in facilitating overall research efficiency when dealing with large volume of environmental data that are distributedly maintained.

Future Work

- Look forward to a close collaboration with the development of ESGF
- Support for OPeNDAP, WMS and WPS
- Support for user work space and management of tasks
- Autogeneration of XML descriptions of the analytic functions from user-input web page forms.
- Tracing data provenance and provide intermediate results
- Adding a third-party authentication mechanism and encryption for the APIs between the Web-based UI and CAFE nodes

THANKS FOR YOUR ATTENTION

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