

Scopira: An Open Source C++ Framework for Biomedical Data Analysis Applications – A Research Project Report

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Scopira

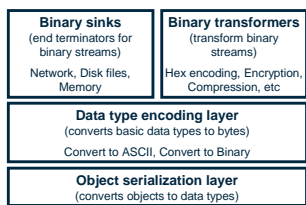
Scopira is an open source, object-oriented and generic C++ framework for scientific computing applications with emphasis on biomedical data analysis.

- Spectra and images (magnetic resonance, infrared, gene microarrays, Raman, mass spec).
- Visualization (2D and 3D via OpenGL).
- Computationally efficient.
- Parallel algorithm development (MPI and a dedicated agent facility).

Foundation

The tool subsystem provides a host of generic facilities and utilities for all types of Scopira based applications.

- Reference counting with "smart pointers".
- Threads and concurrent programming.
- Random number generators ("real" and pseudo) and distributions.
- Input/output & object serialization.



core Subsystem

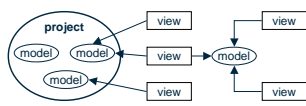
The core subsystem builds on the tool subsystem and provides non-graphical features useful for building applications.

- **basic_loop** main loop for parsing configuration options.
- Powerful plug-in loading systems.
- Flexible object registration system for registering objects for serialization and virtual construction.
- Dynamic model/view system.

Models and Views

Scopira provides a dynamic system (via run-time registration) for applications and plug-ins to declare models and views.

- Models are objects that are monitored by zero or more views. Views themselves can be graphical or non-graphical.
- A project is a model that organizes a collection of models in a tree like fashion. Users may use projects to save their data sets as related workspaces.

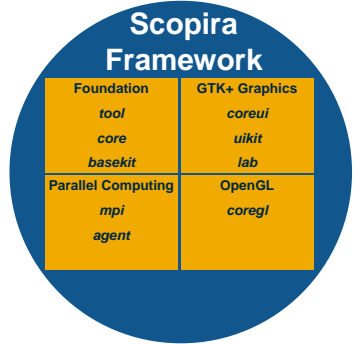
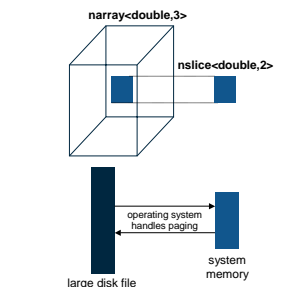


basekit Subsystem

The basekit subsystem contains various numerical routines and the core numerical data structure, **narray**.

- Generic, via C++ templates structure. Can be used with ANY data types (ints, floats, complex, etc) and ANY dimension (vector, matrix, cube, etc).
- Complementary **nslice** virtual sub-view, any size == **narray** dimensions.
- As-good-as-C performance via templates and inline methods.
- Range checked access via **assert()**, (debug mode only).
- STL-style iterators and thereby usable with STL algorithms.
- **DirectIO** back end, to directly access files as if they were in memory (via the operating system's **mmap** function)

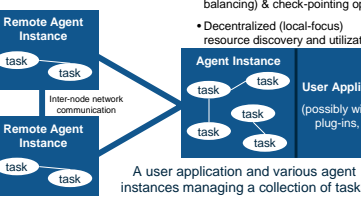
DirectIO: Memory Mapping



Parallel Computing

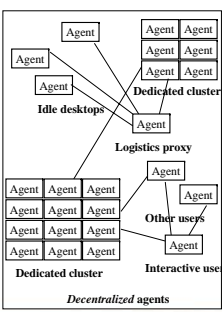
This subsystem provides a light (inline) helper layer for Scopira apps that use the MPI (Message Passing Interface) API.

- Aware of **narrays** and uses type information (via C++ traits classes) to deduce many typical parameters needed for many MPI functions.
- Drastically reduce the amount of information needed from the programmer.
- Tasks (agent-managed) that send messages to other tasks & migrate between agents over the network.



agent types

- **Local**: non-network aware, single machine (but multi-threaded).
- **Cluster** (under development): Used with dedicated, fully connected and persistent Beowulf-like clusters.
- Load balancing done at global level as resource allocation decisions.
- **Decentralized** (under development): For larger, complex agent networks.
- Allocate resources based on local information only (permits network scalability); may be used over unstable network links to possibly unreliable remote agents.
- Most dynamic agent, requiring many peer-to-peer like approaches to resource allocation/deployment.



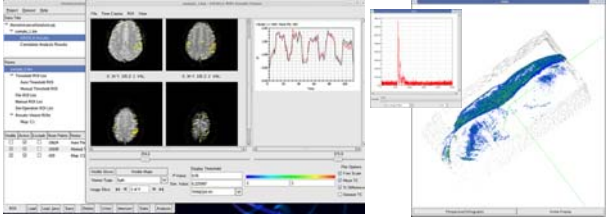
Applications

Scopira used to build a suite of applications and plug-ins, each with varying scopes of complexity.

- Algorithms prototyping, possibly with visualization (**BigVol**)
- Parallelized algorithms, via MPI, for the analysis and classification of biomedical data (**SFS**).
- Plug-ins for pattern recognition algorithms, visualization, and data projection (**RDP**).
- Full, stand-alone applications (**ScopiraPA**, **Evident**, **Opus**)

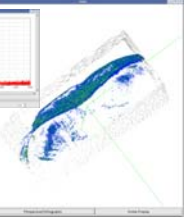
Evident

Project based application for the detection of "novel" activations (time courses) in functional magnetic resonance neuroimages.



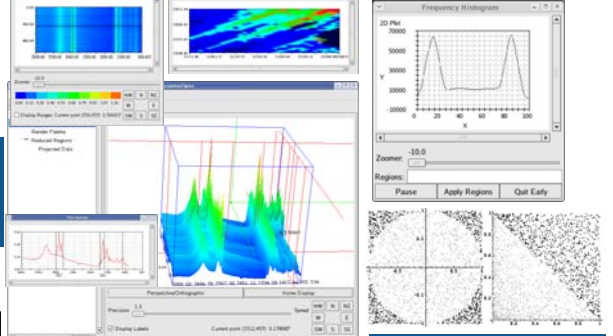
BigVol

A 64-bit and OpenGL-based 3D visualizer for large (>10 gigabytes) datasets.



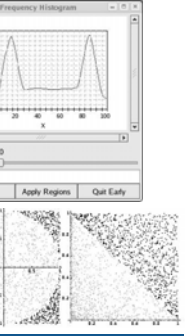
Opus

Project based application for processing/visualizing spectra.



SFS

Parallelized classification methodology for stochastic feature selection.



Graphics Subsystems

- **coreui**: basic **GTK+**-based widgets, window classes, and layout managers.
- **uikit**: builds on coreui to provide more complex visualization widgets and views (eg, plotters, image viewers, scalable matrix editor).
- **lab**: provides API for rapid development of algorithms that need graphical output. Allows main thread to do the computations while a background thread handles GUI event loop.
- **coregl**: builds on coreui to provide foundation for building 3D visualization widgets. Uses **GTKGLExt**, a small library that allows **GTK+** apps to use the industry standard **OpenGL** API.

