

# Amira 5.4

## Specification Sheet

### Amira – Base Package

The base package is required for all applications of Amira. It provides advanced modules for general 3D/4D analysis and visualization. When multidimensional data needs to be processed and explored, this package offers a rich set of powerful tools with intuitive graphical user interfaces to get detailed insight into complex data sets. Many modules are parallelized and thus achieve a significant speedup with modern multicore processors; calculations in the background reduce waiting times. Besides high visualization requirements, this package satisfies the needs for advanced analysis and processing of clinical or preclinical image data, nuclear data, optical or electron microscopy image data, and all types of multidimensional data. Several modules are also provided to prepare presentations, demos and movies. The basic package contains the powerful Segmentation Editor and the Multi Planar Viewer.

#### Analysis

- Material statistics
- Volume and surface measurement
- 2D/3D linear measurement and annotation
- Co-localization analysis
- Arithmetic operations
- Densitometry
- Surface analysis
- Integration of MATLAB® engine
- Network statistics
- Cluster density

#### Visualization

- Orthogonal, oblique, and curved slicing
- Shaded multi-resolution volume rendering
- Surface and isosurface rendering
- Multi-channel image visualization
- Image fusion
- Height field and bump slicing
- Orthogonal projections
- Visualization of traced filaments
- 3D ROI selection
- Clipping and cutting
- Visualization of time lapse data
- 3D + Multi-Planar Viewer to explore complex data
- Illuminated line visualization
- Utilization of modern graphics hardware

#### Processing

- Image filtering
- Surface generation
- Image segmentation
- Slice alignment
- Automatic and manual registration
- Volume and image sampling
- Volume cropping
- Direct volume edit
- Surface editing
- Data type conversion
- Scripting
- Fusion in 3D Multi-Planar Viewer
- Automatic and manual registration in 3D Multi-Planar Viewer

#### Presentation

- DemoDirector for authoring complex animations
- High quality demos
- Movie generation
- Various 3D stereo display modes
- Auto stereoscopic display support
- 2D and 3D annotation

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### Editors

- **Segmentation Editor** interactive and semi-automatic tools for 3D image segmentation
- **Multi-Planar Viewer** to view up to two data sets simultaneously in a 3+1 Multi-Planar Viewer
- **Camera Path Editor** to create a camera path using keyframes for animations and movies
- **Color Dialog** graphical interface for defining a color value
- **2 Colormap Editors** to modify the RGBA values of a discrete colormap
- **Curve Editor** creates and edits curves
- **Digital Image Filters** to apply standard image processing filters
- **Image Crop Editor** to crop 3D images, change bounding box, and voxel size
- **Landmark Editor** to add, move, or delete markers in a landmark set
- **LineSet Editor** to select, create, modify, and delete polylines
- **Parameter Editor** to add, change, or delete attributes of a data object
- **Surface Simplification Editor** to reduce the number of triangles of a surface
- **Surface Editor** to modify triangles, remove intersections, assign boundary IDs
- **Transform Editor** to translate, rotate, or scale any 3D data object
- **Plot Tool** graphical interface for displaying 2D plots
- **Demo Manager** graphical interface to manage and control demos

### Customization

- **Amira Script Object** Amira script objects are custom modules with user-defined GUI elements (ports) written in the Tcl command language.

### Import/Export

File Format	Extension	Read/Write
<b>Amira Script</b> Amira Tcl script	hx	rw
<b>AmiraMesh Format</b> Amira's native general purpose format	am	rw
<b>AmiraMesh, StackedSlices and Raw data as LargeDiskData</b>	ldd, ldm	rw
<b>Analyze 7.5</b> 3D image data with separate header file	hdr	rw
<b>AnalyzeAVW</b> contains 2D and 3D medical image data	avw	rw
<b>BMP</b> uncompressed Windows bitmap format	bmp	rw
<b>DXF</b> Drawing Interchange Format for AutoCAD 3D models	dxf	rw
<b>EPS</b> (Encapsulated PostScript) for 2D raster images only	eps	w
<b>HTML</b> Hypertext document format	html	r
<b>Hoc</b> Hoc file reader of morphometric models for NEURON environment	hoc	rw
<b>HxSurface</b> Amira's native format for triangular surfaces	surf	rw
<b>Icol</b> ASCII format for colormaps with alpha channel	icol	rw
<b>Interfile</b> format used in nuclear medicine, based on Analyze 7.5	img/hdr	r
<b>JPEG</b> 2D image format with lossy compression	jpg	rw
<b>LargeDiskData</b> access image data blockwise	ldd	rw
<b>MATLAB Binary</b> MATLAB matrices	mat	rw
<b>MATLAB M-files</b> MATLAB script	m	rw
<b>Nifti</b> successor of the popular Analyze 7.5 format	nii/img/hdr	rw
<b>Open Inventor</b> standard file format for 3D models	iv	rw

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### Import/Export (cont'd)

File Format	Extension	Read/Write
<b>PNG</b> portable network graphics format for 2D images	png	tw
<b>PNM</b> simple uncompressed 2D image format	pnm	rw
<b>PSI</b> ASCII format for 3D points and associated data values	psi	rw
<b>Ply</b> Stanford triangle format for points and surfaces	ply	rw
<b>Raw Data</b> binary data as a 3D uniform field	raw	rw
<b>SGI-RGB</b> 2D image format with run-length encoding	sgi/bw/rgb	rw
<b>STL</b> simple format for triangular surfaces, no connectivity	stl	rw
<b>Stacked-Slices</b> info file grouping together 2D images	info	rw
<b>SWC</b> interchange format for neuroscience	swc	rw
<b>TIFF</b> standard format for 2D and 3D image data	tif/tiff	rw
<b>Tecplot</b> Tecplot ASCII and Binary file reader	tec/plt	rw
<b>VRML</b> virtual reality markup language for 3D models	wrl	rw
<b>Vevo Raw Images</b> 2D and 3D ultrasound images from Visual Sonics' Vevo 770	rdb/rdi	r

### Supported Platforms

- **Microsoft Windows** – Microsoft Windows XP/Vista/7, 32-bit and 64-bit editions
- **Mac OS X** – Mac OS X 10.5/10.6/10.7, Amira runs as 32-bit application
- **Linux** - Red Hat Enterprise Linux 5.5 for x86\_64 or compatible

### System Requirements

A processor (CPU) supporting SSE2 instruction set is required.

A graphics card with hardware accelerated texture mapping capabilities and OpenGL support is required. Some visualization modules, e.g., the highest quality volume rendering, require graphics hardware with vertex and fragment shader support, as well as installation of the most recent graphics drivers.

**Note:** Amira and all its extension options are intended for research only and not for diagnosis or treatment of patients.

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### Mesh Option – Surfaces, 3D grids, and vector fields

The Mesh Option offers advanced features for scientists working with complex surface analysis, volume reconstruction using tetrahedral grids, vector field visualization, and complex tensor computation. This package is designed especially for Finite Element Modeling (FEM), for computational analysis of 3D regular fields, tensors, and 3D fluid dynamics simulations.

#### Main features

- Surface editing
- Tetrahedral and hexahedral grid visualization
- Tetrahedral grid generation
- Grid analysis and refinement
- Grid parameterization
- Conversion from tetrahedral to hexahedral grids and vice versa
- Boundary condition visualization
- Vector field visualization
- Planar and surface LIC (line integral convolution)
- Tensor computation
- Tensor visualization
- Out-of-core tensor computation
- Animated illuminated streamline visualization
- Animated cone plot visualization
- Ribbon visualization
- Stream surface

#### Editors

- **Grid Editor** to edit and simplify tetrahedral grids

#### Import/Export

File Format	Extension	Read/Write
<b>Abaqus</b> stores geometry and data values of unstructured grids	inp	w
<b>AVS Field</b> stores data defined on regular grid	fld	raw
<b>AVS UCD Format</b> stores unstructured cell data	inp	raw
<b>FIDAP NEUTRAL</b> stores FEM meshes and solution data	ascii fmt	r
<b>Fluent-UNS</b> contains FEM meshes, boundary ids, solution data	hdr	raw
<b>Hypermesh</b> used by Altair HyperWorks FEM software	hm/hmascii	raw
<b>IDEAS universal format</b> describes FEM grids and simulation data	unv	raw
<b>Plot 3D Single Structured</b> stores curvilinear grids and associated data	p3d	raw

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

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### Microscopy Option – Readers and tools for microscopy

The Microscopy Option is the comprehensive option for visualization of microscopy data. It includes specific readers for microscopy data as well as a powerful 3D image deconvolution suite. The Microscopy Option comes with several examples and step-by-step tutorials to easily learn how to enhance images with filters and deconvolution process. This option includes the Filament Editor.

#### Main features

- Microscopy file readers
- Deconvolution
- Z-Drop correction
- Tool for bead measurements
- Background image and flatfield correction
- Fourier transform
- Automatic and interactive filament tracing
- Filament network labeling and editing
- Filament network automatic hierarchy and loop detection
- Watershed separation
- Multi-component analysis

#### Editors

- **Filament Editor** to extract, modify and label neuron and vessel networks (including features of the Skeletonization Option)

#### Import/Export

File Format	Extension	Read/Write
<b>Bio-Rad Confocal</b> simple uncompressed format for 3D image stacks	pic	r
<b>Leica 3D TIFF</b> contains 3D image data and meta information	tif	r
<b>Leica Binary</b> 3D image stacks, time series, and meta information	lei	r
<b>Leica Image</b> 3D image stacks, time series, and meta information	lif	r
<b>Leica Slice Series</b> contains list of 2D TIFF files, raw data and meta info.	info	r
<b>MRC</b> MRC file format for electron microscopy	mrc/rec	rw
<b>Metamorph STK</b> special TIFF variant for 3D image stacks	stk	r
<b>Olympus OIB/OIF</b> binary file format (Windows only)	oib/oif	r
<b>Zeiss LSM</b> 3D raster image format	lsm	r

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

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### Neuro Option – Medical image analysis for DTI and Brain Perfusion

This option offers specialized modules to analyze medical images obtained using CT or MRI. Diffusion weighted MR imaging (DWI) is used to image the rate of water diffusion in tissues like brain or muscle. This option allows the user to compute the apparent diffusion coefficient (ADC), fractional anisotropy (FA), and directionally encoded color (DEC) maps. Diffusion tensor imaging (DTI) aims at localizing and visualizing fiber tracts in the human or primate brain and is supported by several modules. Brain perfusion analysis is supported based on tracking a contrast bolus in a time series of MR or CT images.

#### Main features

- Fiber tracking supporting several stream-line based algorithms
- Fiber separation into fiber bundles based on user defined source and destination regions
- Computation of tensor fields from an arbitrary number of diffusion weighted MR images
- Generation of test data for tensor field and gradient weighted image stack analysis
- Computation of diffusion weighted maps like ADC, FA, relative anisotropy (RA), and DEC
- Eigenvalue decomposition of tensor fields
- Supports both CT and MR bolus tracked images
- Computes mean transit time (MTT), cerebral blood flow (CBF), and cerebral blood volume (CBV)
- Manual selection and export of arterial/venous functions from perfusion time series
- Tracer delay time corrected
- Computation based on deconvolution
- Automatic brain segmentation

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

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### Very Large Data Option – Explore data sets larger than 8 GByte

The Very Large Data Option allows Amira to display large data sets that exceed the available main memory, e.g., on computers with only a small amount of RAM or when using a 32-bit operating system. Using volumes of interest and multi-resolution techniques, the Very Large Data Option overcomes these limitations and allows interactive visualization and processing of large data volumes. Depending on the available resources, this option allows exploration of image volumes greater than 8 GByte, and has been successfully tested with data sets larger than 300 GByte.

#### Main features

- Interactive visualization with orthogonal/oblique slicing, volume- and isosurface rendering
- Fast disk access to sub-regions and sub-samples
- Optimal image quality based on available hardware resources
- Automatic refinement of displayed images
- Import/export based on native file format (LDA VolumeViz)

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

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### Molecular Option – Visualize complex molecular structures and properties

This option combines Amira's strong capabilities for 3D data visualization, such as hardware-accelerated volume rendering, with specific tools for molecular visualization and analysis, such as molecular surfaces, sequence alignment, configuration density computation, molecule trajectories, and more. The Amira Molecular Option comes with self-running demos and step-by-step tutorials for the most common molecular visualization tasks.

#### Main features

- Secondary structure extraction and visualization
- Molecular trajectory computation and visualization
- Potential field computation
- Pseudo-electron density computation
- Molecule labeling
- Molecule surface computation
- Align molecule and sequences
- Pareto-optimal set computation
- H bond computation and visualization
- Ribbon, tube, cartoon chain visualization
- Dynamic visualization
- Configuration density evaluation
- Mean-molecule computation

#### Editors

- **Molecule Attribute Editor** allows editing of attributes of groups contained in a molecule
- **Molecule Editor** allows you to change geometry and topology of a molecule

#### Import/Export

File Format	Extension	Read/Write
<b>AMBER</b> Assisted Model Building with Energy Refinement format	top/crd/vel	r
<b>AMF</b> Amira Molecule Format	amf	rw
<b>DX</b> APBS DX electrostatic field file	dx/apbs	r
<b>GROMACS</b> Groningen Machine for Chemical Simulations format	gro/top/trr	rw
<b>MDL</b> MDL file format for chemical structures	sdf	rw
<b>PDB</b> protein data base file format	pdb	rw
<b>PHI</b> Congen PHI Electrostatic field file	phi	r
<b>PSF/DCD (CHARMM)</b> file format used by CHARMM	psf/dcd	r
<b>Tripes</b> file format used to save Tripes Sybyl mol2 molecules	mol2	rw
<b>UniChem</b> file format used by the UniChem molecular software	uni	rw
<b>ZIB Molecular</b> format developed as molecular exchange	zmf	rw

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

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## Specification Sheet

### Skeletonization Option – Reconstruct and analyze networks

This free-of-charge option supports reconstruction and analysis of vascular, dendritic, and fracture networks. It combines specific micro-detailed image mosaic management with advanced automatic and semi-automatic tools for reconstruction of a 3D vessel network from confocal microscopy or synchrotron image acquisition. The Skeletonization Option delivers a unique set of features for the researcher who works at studying and analyzing neurons or micro-vascular networks.

#### Main features

- Automatic skeletonization tools
- Distance mapping
- Thinning
- Smoothing
- Support for bricked data (mosaic)
- Radius histogram

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

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### Developer Option – Extend Amira through C++ development

The Developer Option allows you to extend Amira by developing custom data types, visualization or processing modules, and input/output routines through a well-documented C++ API. The object-oriented language together with the open design makes this option the ideal answer to specific technical requirements.

#### Main features

- Provides the C++ header files needed to compile and develop custom extensions
- Offers a development wizard (assistant) to simplify code development
- Build and test new modules for computation, I/O, and visualization
- Based on a number of industry standard libraries (OpenGL®, Open Inventor™, Qt®)
- Comes with the Open Inventor graphics libraries for creating new Amira visualization modules
- The unique object-oriented interface provides maximum flexibility also for developers with basic experience in C++ coding
- Expert programmers can easily include new graphical user interface (GUI) elements based on Qt

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux

#### Supported Compilers

In order to add custom extensions to Amira using the Developer Option, a C++ compiler is required.

The following C++ compilers are supported:

- **Windows:** 32-bit: Microsoft Visual Studio 2005 (VC++ 8)  
64-bit: Microsoft Visual Studio 2008 (VC++ 9)
- **Mac OS X:** GCC 4.2.x
- **Linux:** GCC 4.1.x

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### Virtual Reality (VR) Option – Display 3D data in immersive environments

The Amira VR Option enables advanced data visualization and analysis with large tiled displays as well as immersive VR configurations like CAVEs. The VR Option supports efficient multi-threaded rendering on multi-pipe systems and distributed rendering on cluster systems at optimal performance with minimal bandwidth requirements. It also supports active or passive stereo, edge blending, head tracking, and 3D interaction for using Amira capabilities with a true immersive experience.

#### Main features

- Full immersive 3D interaction
- Support for single and multiple screens
- Support for single and multi-pipe rendering
- Support for graphic clusters
- Flexible customization for specific display geometry
- Supports passive and active stereoscopic displays
- Supports auto-stereoscopic displays
- Head-tracking
- Support for tracked hand-held 3D input devices
- Control functionality via user interface
- Control of 2D user interfaces with 3D devices (virtual mouse)
- Scriptable using Tcl

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, 64-Bit Linux

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### DICOM Option – Read and write DICOM files

Beside the import and export potentials of Amira, this free-of-charge option offers an extended capability to read and write DICOM images using a clear user interface. This option includes also a module to “push” (save) the DICOM data set on a remote server. These features are especially useful in a medical research environment.

#### Main features

- **DICOM Push** to save data on a remote DICOM node

#### Import/Export

File Format	Read/Write
ACR-NEMA predecessor of the DICOM format for medical images	r
DICOM export medical image export	w
DICOM import standard file format for medical images	r

#### Platforms

Microsoft Windows 32-bit, Microsoft Windows 64-bit, Mac OS X, 64-Bit Linux