



SimTK 1.5 Workshop Installation and Components

Jack Middleton
September 25, 2008



SimTKcore

- Webpage overview
- Overview of download contents
- Help with installs and compiling examples during break



SimTKcore Webpage

- Documents Link
 - Doxygen docs (also in download)
 - User guides (also in download)
 - Workshop slides
- Wiki : tips, know issues
- Bug tracking, forums
- News : announcements

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SimTKcore

Overview

The SimTK Core biosimulation toolkit brings together in a robust, convenient, open source form the collection of highly-specialized technologies necessary to building successful physics-based simulations of biological structures. These include: strict adherence to an important set of abstractions and guiding principles, robust, high-performance numerical methods, support for developing and sharing physics-based models, and careful software engineering.



We released SimTK 1.5 in early August and will be hosting a 2-day workshop on it on Sept. 25-26 (see <http://simbios.stanford.edu/EventsOfInterest/WorkshopSimTK1.5.htm>).

Accessible High Performance Computing

We believe that a primary concern of simulation scientists is performance, that is, speed of computation. We seek to build valid, approximate models using classical physics in order to achieve reasonable run times for our computational studies, so that we can hope to learn something interesting before retirement. In the choice of SimTK technologies, we are focused on achieving the best possible

Project Lead

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SimTKcore

Documents

[Collapse all](#)

[Current SimTK 1.5 Programming APIs](#)

[Advanced Programming Guide](#)

Guide to writing your own code to extend SimTK Core
Last Update: Aug 28, 2008

[Doxygen Docs, v 1.5](#)

Online SimTK reference manual
Last Update: Aug 08, 2008

[Simbody User Guide](#)

Theory and algorithms used in SimTK multi-body dynamics engine
Last Update: Aug 28, 2008

[Simmatrix User's Guide](#)

Introduction to using SimTK's Vector and Matrix classes
Last Update: Aug 28, 2008

[SimTK 1.5 Tutorial](#)

Introduction on creating multibody dynamics simulations in SimTK
Last Update: Aug 28, 2008

[SimTKlapack Overview](#)

Introduction to using SimTK's LAPACK implementation
Last Update: Aug 28, 2008

[SimTKmath User's Guide](#)

User guide for SimTK's numerical methods library
Last Update: Aug 28, 2008

[SimTK Molecular Modeling Guide](#)

User guide for SimTK's molecular modeling library: Molmodel
Last Update: Aug 28, 2008

SimTKcore - Windows Internet Explorer

https://simtk.org/api_docs/api_docs15/


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SimTKcore

Main Page Namespaces Classes Files Directories

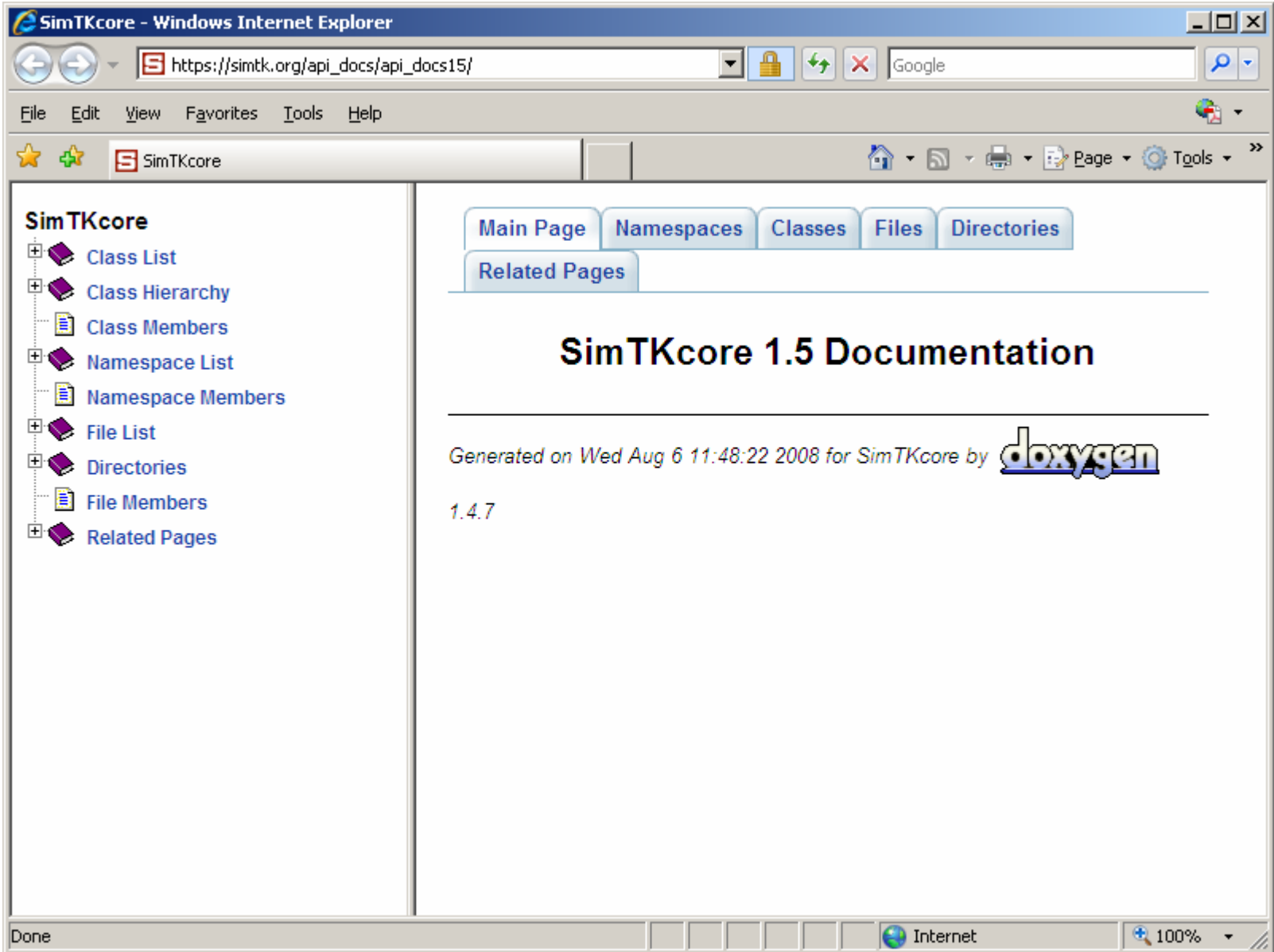
Related Pages

SimTKcore 1.5 Documentation

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SimTKcore

- [Class List](#)
- [Class Hierarchy](#)
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SimTKcore Class List

Here are the classes, structs, unions and interfaces with brief descriptions:

_DisMat	
_generic_N_Vector	
_generic_N_Vector_Ops	
_N_VectorContent_Parallel	
_N_VectorContent_Serial	
_SpgmrMemRec	
AbstractValue	<i>Abstract base class representing an arbitrary value of self-describing type</i>
AcetylResidue	<i>Widely used acetyl protein N-terminal end cap</i>
AdenineBase	
AlcoholOHGroup	<i>AlcoholOHGroup is OH group for attachment to aliphatic carbon</i>
AliphaticCarbon	<i>AliphaticCarbon is a tetrahedral sp^3 carbon atom for bonding to four other things</i>
AliphaticHydrogen	<i>AliphaticHydrogen is a hydrogen atom for bonding to AliphaticCarbon atoms (see below)</i>
Amber99ForceSubsystem	
AminoAcidResidue	<i>Amino acid residue building block for protein polypeptide chain molecules</i>
AminoAcidResidue::Alanine	

- Simbody
 - Class List
 - Class Hierarchy
 - Class Members
 - Namespace List
 - Namespace Members
 - File List
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 - File Members

Constraint::Ball Class Reference

```
#include <Constraint.h>
```

[List of all members.](#)

Detailed Description

Three constraint equations.

This constraint enforces coincident location between a point on one body and a point on another body.

The constraint is enforced by an internal (non-working) force applied at the spatial location of the point on body 2, on material points of each body that are coincident with that spatial location. Note that this is somewhat asymmetric when the ball is not properly assembled -- it acts as though the contact occurs at the point on body 2, *not* at the point on body 1.

The assembly condition is the same as the runtime constraint -- the two points can be brought together by driving the perr to zero.

Public Member Functions

```
Ball (MobilizedBody &body1, MobilizedBody &body2)  
Ball (MobilizedBody &body1, const Vec3 &defaultPoint1, MobilizedBody
```


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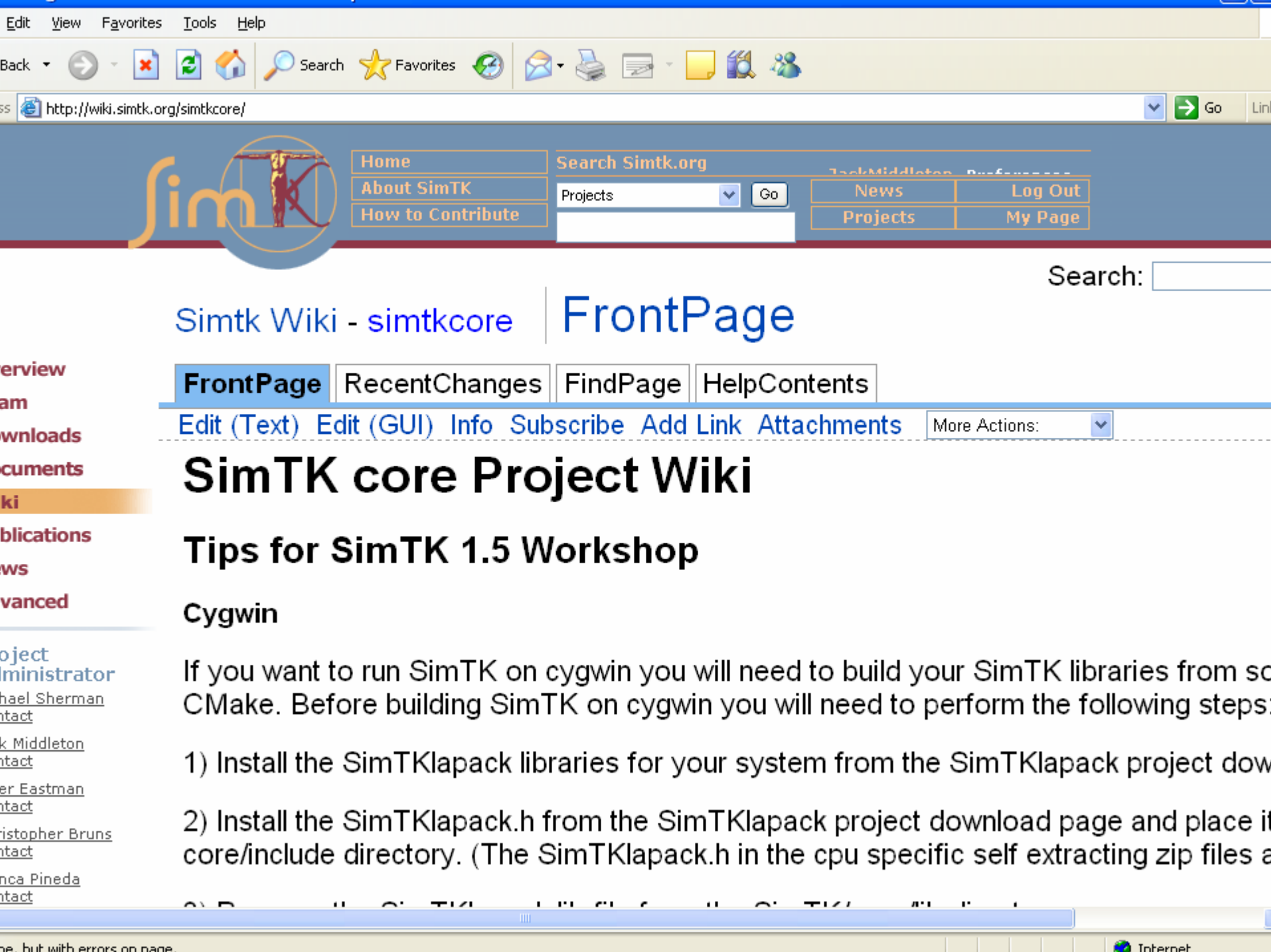
SimTKcore

Public Forums

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Topic	Topic Starter	Replies	Last Post
using SimTK with XCode (Mac)	David Wagner	5	2008-08-28 15:40
Fortran compiler optional?	Ralf Grosse-Kunstleve	3	2008-07-17 13:05
how to avoid leaks?	Ralf Grosse-Kunstleve	5	2008-07-16 17:17
Getting started with SimTK Core	Joy Ku	0	2008-07-10 10:57
simtkcore win32 mingw port	rob rein	1	2008-07-07 14:43
possibilities of the simtk core?	Pablo Ortega	1	2008-06-30 17:49
Porting SimTK Core	Marko Havu	0	2008-01-07 22:55
Welcome to Help	SimTK.org Admin	0	2006-04-04 13:26

Search Forums



Simtk Wiki - simtkcore | FrontPage

SimTK core Project Wiki

Tips for SimTK 1.5 Workshop

Cygwin

If you want to run SimTK on cygwin you will need to build your SimTK libraries from source using CMake. Before building SimTK on cygwin you will need to perform the following steps:

- 1) Install the SimTKlapack libraries for your system from the SimTKlapack project download page.
- 2) Install the SimTKlapack.h from the SimTKlapack project download page and place it in the core/include directory. (The SimTKlapack.h in the cpu specific self extracting zip files are not the same as the one in the core/include directory.)



SimTKcore Downloads

- Supported Platforms:
 - Windows XP
 - Mac (Intel)
 - Linux
- Temporarily supported Platforms
 - Mac PPC



Debug vs. Release

Workshop can be done with either

Release is faster

Windows: Release vs. RelWithDebInfo

Linux

Linux

SimTK core software for Linux



1.5.1

August 28, 2008

Patch to SimTK 1.5. Please download and read [SimTKcore_README_Linux.pdf](#) for installation instructions.

[Release Notes](#)

Download Links

Name	File Type	Platform	Updated
SimTKcore_Debug.sh (99948 kB) <i>Description:</i> Compiled with symbols for debugging.	binary	Linux	Aug 28, 2008
SimTKcore_Release.sh (55895 kB) <i>Description:</i> Compiled with full optimization.	binary	Linux	Aug 28, 2008

Documentation Links

[SimTKcore_README_Linux.pdf](#) (17 kB)

Description: Overview and installation instructions.

Sep 02, 2008

Please cite these papers

- Jeanette P. Schmidt, Scott L. Delp, Michael A. Sherman, Charles A. Taylor, Vijay S. Pande, Russ B. Altman, "The Simbios National Center: Systems Biology in Motion," IEEE Transactions on Biomedical Engineering, in press (2008)

■ Built on 32bit with gcc 4.1.2

Mac OSX (Intel)

Mac OS X (intel)

SimTK core software for intel based Max OS X. Please see Other Platforms for other MACs/OS.



1.5.1
August 28, 2008

Patch for SimTK 1.5. Please download and read [SimTKcore_README_Mac.pdf](#) for installation instructions.

[Release Notes](#)

Download Links

Name	File Type	Platform	Updated
SimTKcore_Debug.dmg (96251 kB)	binary	Mac	Aug 28, 2008
<i>Description:</i> Compiled with symbols for debugging.			
SimTKcore_Release.dmg (71276 kB)	binary	Mac	Aug 28, 2008
<i>Description:</i> Compiled with full optimization.			

Documentation Links

[SimTKcore_README_Mac.pdf](#) (16 kB)

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Description: Overview and installation instructions.

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- Xcode 3.0
- Add `#include <typeinfo>`

Windows XP and Vista

Windows

SimTK core software for Windows



1.5.1

August 28, 2008

Patch for SimTK 1.5. Please download and read [SimTKcore_README_Windows.pdf](#) for installation instructions.

[Release Notes](#)

Download Links

Name	File Type	Platform	Updated
SimTKcore_Release.exe (47012 kB)	binary	Windows	Aug 28, 2008
<i>Description:</i> Compiled with full optimization.			
SimTKcore_RelWithDebInfo.exe (50737 kB)	binary	Windows	Aug 28, 2008
<i>Description:</i> Compiled with optimization and symbols for debugging			

Documentation Links

[SimTKcore_README_Win.pdf](#) (16 kB)

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Description: Overview and installation instructions.

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- Jeanette P. Schmidt, Scott L. Delp, Michael A. Sherman, Charles A. Taylor, Vijay S. Pande, Russ B. Altman, "The Simbios National Center: Systems Biology in Motion," IEEE Transactions on Biomedical Engineering, in press (2008)

■ RelWithDebInfo for debugging

Temporarily Supported Platforms

Z Other Platforms-unsupported

As a courtesy to our users these downloads are provided for platforms which are not officially supported.



1.5.1

August 28, 2008

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[Release Notes](#)

Download Links

Name	File Type	Platform	Updated
SimTKcore_PPC_Debug.dmg (113051 kB)	binary	Mac	Aug 28, 2008
<i>Description:</i> Compiled with symbols for debugging.			
SimTKcore_PPC_Release.dmg (70722 kB)	binary	Mac	Aug 28, 2008
<i>Description:</i> Compiled with full optimization.			

Documentation Links

[SimTKcore_README_Mac_PPC.pdf](#) (16 kB)

Description: Overview and installation instructions.

Sep 02, 2008

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- Jeanette P. Schmidt, Scott L. Delp, Michael A. Sherman, Charles A. Taylor, Vijay S. Pande, Russ B. Altman, "The Simbios National Center: SystemsBiology in Motion," IEEE Transactions on Biomedical Engineering, in press (2008)

■ PPC needs libg2c

SimTK and VTK

- Files or directories with “SimTK” in name developed at Stanford (SimTKsimbody.dll)
- Files or directories with “vtk” part of Visualization Tool Kit <http://vtk.org> (vtkCommon.dll)
- SimTK examples use VTK for graphics but SimTK does not require VTK for simulation



Default Install Locations

Mac: /Developer/SimTK

Windows: C:/Program Files/SimTK

Linux: /usr/local/SimTK



SimTKcore Directory Structure

SimTK

core

bin

doc

examples

include

lib

Library Naming Conventions

- Debug have “_d” suffix
 - SimTKsimbody_d.dll
- Release and RelWithDebInfo no suffix:
 - SimTKsimbody.dll
- Static linkable libraries have “_static”
 - SimTKsimbody_static_d.lib

SimTK Libraries

- SimTKmolmodel Molecular modeling
- SimTKsimbody_aux VTK interface
- SimTKsimbody Multibody dynamics
- SimTKmath Numerical methods
- SimTKcommon Vector Matrix classes
- SimTKcpodes ODE, DAE solvers
- SimTKlapack Linear Algebra

SimTKlapack

- High speed implementation of LAPACK
 - Threaded for number of cpu's
 - Blocked algorithms for best cache reuse
- Uses ATLAS to generate optimized BLAS
- FORTRAN interface can be called from C/C++.
- `#include SimTKlapack.h`
- Link to SimTKlapack library

SimTKcpodes

- Multistep, error controlled numerical integrator for multibody dynamics problems
- Advances the ODE, and then performs coordinate projection back to the constraint manifold to exactly solve the DAE
- Joint project between Simbios and LLNL
- Other Integrators available in SimTK (verlet, Runge-Kutta)



SimTKcommon

- System, Subsystem, State
- Vector
- Matrix
- Random (random number generation)
- Rotation, Quaternion
- Event handling
- PolynomialRootFinder

SimTKmath: Nonlinear Optimization

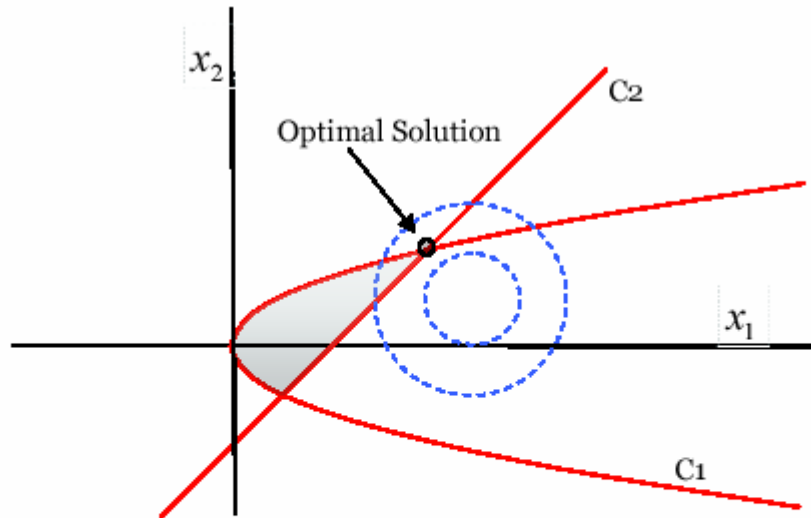


Figure 3: Graph of problem to finding optimal solution to the problem described above. The two constraints are shown in red and the feasible region is shown in grey. The contours of the objective function are shown as dashed blue circles.

SimTKmath: Numerical Integration

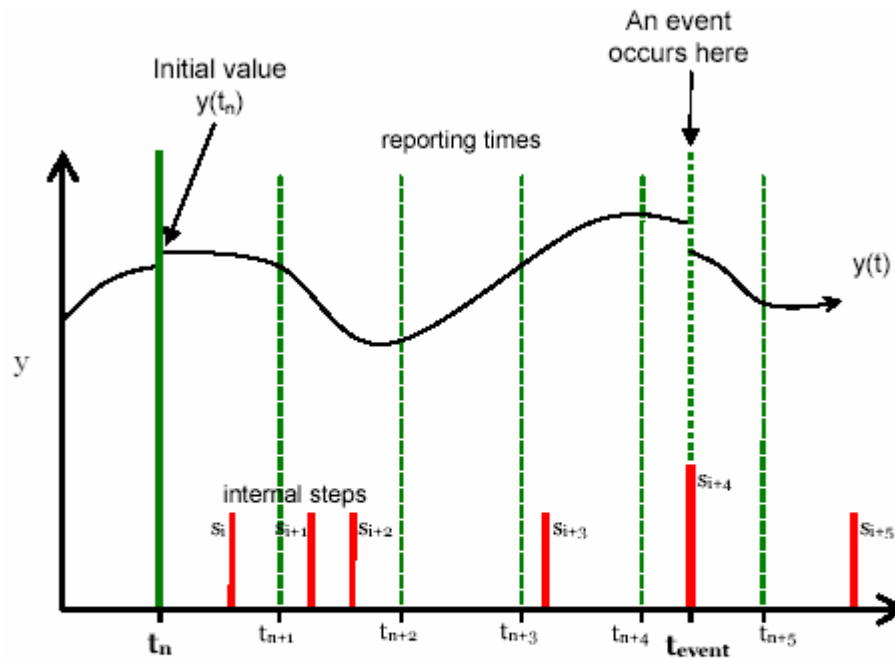


Figure 1: Numerical integration over a continuous segment of a time-stepping trajectory. The continuous region starts at t_n and terminates at t_{event} . Each tall green line represents a return of control to the time stepper; thin dotted ones are caller-requested reporting times; the thick dotted green line is an unrequested return of control just prior to event occurrence and just after event handling.

FactorLU: solve linear systems


$$\begin{aligned}x + 2y - z &= 2 \\4x + 3y + z &= 3 \\2x + 2y + 3z &= 5\end{aligned}$$

$$\begin{bmatrix} 1 & 2 & -1 \\ 4 & 3 & 1 \\ 2 & 2 & 3 \end{bmatrix} \begin{bmatrix} x \\ y \\ z \end{bmatrix} = \begin{bmatrix} 2 \\ 3 \\ 5 \end{bmatrix}$$

$$A \quad x \quad = \quad b$$

FactorQTZ: Linear Least Squares Solution

- Underdetermined: equations $<$ unknowns
infinite number of solutions
find: solution with minimum $\|x\|_2$
- Over determined: equations $>$ unknowns
No solution exists
find: minimum $\|Ax - b\|_2$



SimTKmath (continued)

Differentiator: Numerical Differentiation

Eigen: Eigen values and Eigen vectors

FactorSVD: Singular value
decomposition



SimTKsimbody

- Algorithms and data structures for modeling rigid multibody systems in internal coordinates.
- Order(n) algorithm
- Useful for internal coordinate and course grained molecular modeling
- Also for large scale mechanical models like skeletons



SimTKmolmodel

- Molecular modeling layer for Simbody
- Tools for modeling biological macromolecules such as proteins and nucleic acids.
- Dynamics of macromolecules can be simulated using Simbody

Verifying Your Install

- Install checkers in: core/bin
- CoreInstallCheck
 - “Success SimTK has been successfully installed”
- AuxInstallCheck
 - Displays 2 bouncing balls and pendulum

Examples

- Download examples
- Read instructions
- Compile, run

Linux

SimTK core software for Linux



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Tips for Compiling Examples

- Check Wiki
- Windows: release download, MVS configured Release
- Mac: Upgrade Xcode tools
- Linux: gcc 4.1



During Break

- Help with installs
- Help compiling and running examples`