NAMDのインストール(1)

<u>https://www.ks.uiuc.edu/Research/namd/</u>にアクセス 「Software」→「NAMD」→「Download」を選ぶ

NIH RESOURCE FOR MACROMOLECULAR MODELING & VISUALIZATION UNIVERSITY OF ILLINOIS AT URBANA-CHAMPAIGN Type Keywords SEARCH												
THEORETICAL and COMPUTATIONAL BIOPHYSICS GROUP									2			
Home	Researc	h Publications	Software	Instruction	News	Galleries	Facilities	About Us				
			NAMD	Dov	vnload							
Home		NIAME	VMD	Use	User's Guide							
Overview		Scalable	GPU Computin	g Mai	ling List							
Overview		NAMD recipient of a 20	Cloud Computi	ng	240 Olden. Erstende Angelende 2000 Onder Bull Dies is a sealled webendende demonies ande designed for high auformance							
Publications		simulation of large bion	Lattice Microbe	s Cha	Charm++ parallel objects, NAMD scales to hundreds of cores for typical simulations and beyond 500,000 cores for the largest						gest	
Research		simulations. NAMD use	Atomic Resolut	tion ohics	program VMD	for simulation	ation setup and trajectory analysis, but is also file-compatible with AMBER, CHARMM, and X-PLOR. NAMD is					OR. NAMD is
Software		biomolecular modeling.			an build wave yoursel or download binaries for a wide variety of platforms. Our totomars show you now to use NAIMD an							
NAMD		Search all NAMD reso	QwikMD	site a	and tutorials		Google					
VMD		Dracking News	VND (Neuronal)									
GPU Comput	ting	Dreaking News	Other									
Atomic Resc	olution	NAMD 3.0 New Features - webpage is posted										

NAMDのインストール(2)

3. Version 2.14から「MacOSX-x86_64」または「Win64」を選ぶ

Version 2.14 (2020-08-05) Platforms:

- Linux-x86_64-multicore (64-bit Intel/AMD single node)
- Linux-x86_64-multicore-CUDA (NVIDIA CUDA acceleration)
- Linux-x86_64-netIrts (Multi-copy algorithms, single host or ethernet)
- Linux-x86_64-netIrts-smp-CUDA (Multi-copy algorithms, single process per copy)
- Linux-x86_64-verbs (InfiniBand, no MPI needed, supports multi-copy algorithms)
- Linux-x86_64-verbs-smp (InfiniBand, no MPI needed, supports multi-copy algorithms)
- Linux-x86_64-verbs-smp-CUDA (InfiniBand, no MPI needed, supports multi-copy algorithms)
- Linux-KNL-multicore (Intel Xeon Phi KNL processor single node)
- MacOSX-x86 64 (Mac OS X for 64-bit Intel processors)
- MacOSX-x86_64-CUDA (NVIDIA CUDA acceleration)
- MacOSX-x86_64-netIrts (Multi-copy algorithms)
- Win64 (Windows 7, 8, 10, etc.)
- Win64-CUDA (NVIDIA CUDA acceleration)
- Win64-MPI (Windows HPC Server, multi-copy algorithms)
- Win64-MPI-smp-CUDA (HPC Server with CUDA)
- Source Code

NAMDのインストール(3)

4. 適当なユーザー名とパス ワードを入力して 「Continue …」

Registration/Login

You will need a username and password to download software.

If this is your first download, please choose a username and password to register. Current NAMD or VMD users, please enter your existing username and password.

Username:		
Password:		
Continue v	vith registration or download	

Your download will continue after you have registered or logged in.

5. 右の画面で情報を埋めて 「Register」

New User Registration

New User Registration for 'Tohru.Terada':

First and La	st Name:]	
Email Addro	SS:]	
Affiliation:]	
Academ	ic 🔍 Government	Industrial	Other (specify)	
The numbe	of people using T 4 ○ 5-10 ○ 11-2	CBG software at	my site is:	
luse ICBG	software primarily	tor:		
Resear The work I	h	Commerce Second For Commerce (a vare is funded (a	^o ersonal t least partially)	by NIH:
◯ Yes ◯	No			
Re-enter pa	ssword for confirm	ation:		

Register

NAMDのインストール(4)

- 6. 右の画面で、「I am ... I をクリック
- 7. ダウンロードが完了し たら「開く」
 - Windowsはスライド5へ
 - macOSはスライド6へ \succ

Software Downloads

NAMD 2.14 for Win64

To download this software you must agree to abide by the terms of the following license:

(pre> Upon execution of this Agreement by the party identified below ("Licensee"). The Board of Trustees of the University of Illinois ("Illinois"), on behalf of The Theoretical Biophysics Group ("TBG") in the Beckman Institute, will provide the NAMD molecular dynamics software ("NAMD") in Executable Code and/or Source Code form ("Software") to Licensee, subject to the following terms and conditions. For purposes of this Agreement, Executable Code is the compiled code, which is ready to run on Licensee's computer. Source code consists of a set of files which contain the actual program commands that are compiled to form the Executable Code compiled to form the Executable Code.

I am Tohru Terada and I agree to the terms of this License Printable version

NAMDのインストール(5)

8. 解凍して生成されたフォルダ 「NAMD_2.14_Win64-multicore」内の 「namd2.exe」と「tcl85t.dll」をデスクトップに コピー

NAMDのインストール(5)

- 8. 解凍して生成されたフォルダ
 - 「NAMD_2.14_MacOSX-x86_64-multicore」内の「namd2」をデスクトップにコピー

NAMD 実行時の 注意

1. NAMD実行時に右のウ インドウが現れたら 「キャンセル」

macOS Sonoma

- Dockから「システム環境
 設定」を選ぶ
- 3. 「セキュリティとプライバ シー」を開き「このまま 許可」
- 4. 再度実行し、右のウイン ドウで「開く」

