

Handling Domain Decomposition in Massively Parallel Implementations of Stochastic Lattice Models

Jeffrey Kelling¹, Géza Ódor², Sibylle Gemming^{1,3}

¹Institute of Ion Beam Physics and Materials Research, Helmholtz-Zentrum Dresden-Rossendorf, Germany

²MTA TTK MFA Research Institute for Natural Sciences, Budapest, Hungary

³Institute of Physics, TU-Chemnitz, Germany

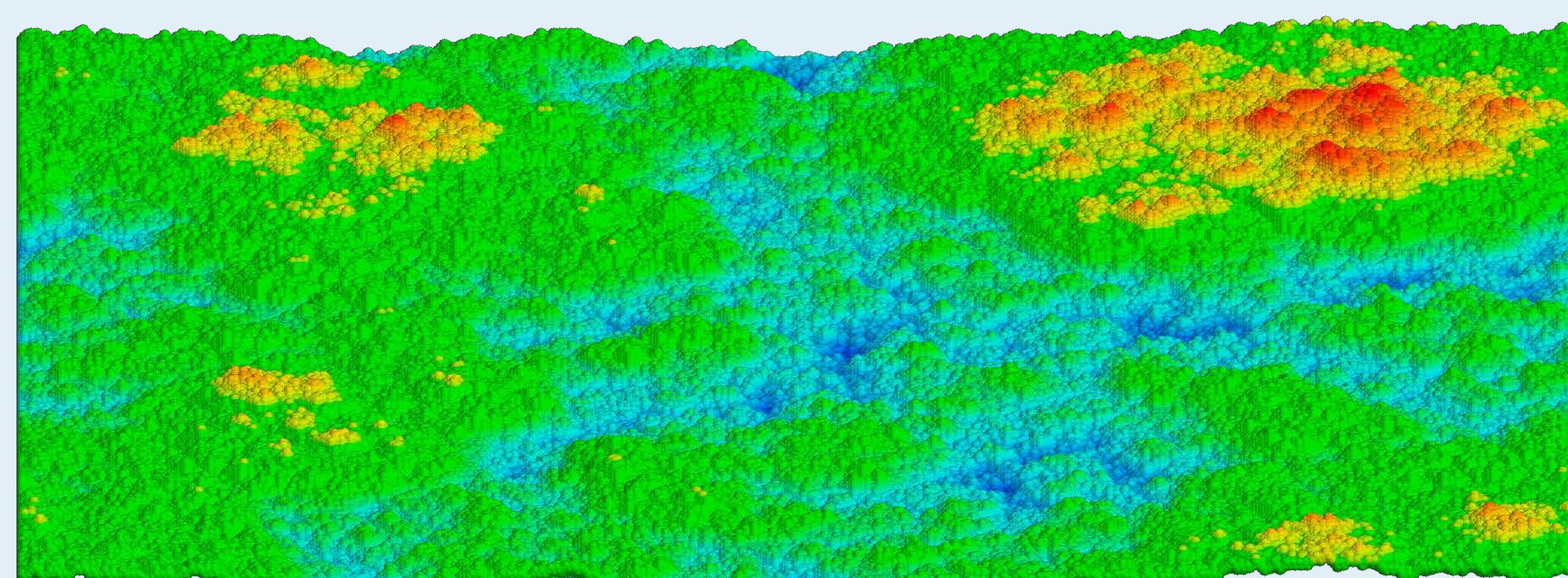


DRESDEN
concept



HELMHOLTZ
ZENTRUM DRESDEN
ROSSENDORF

- nanopatterning of materials from molecular electronics to photovoltaics
- need large scale atomistic simulations to understand self-organization
- only stochastic models can bridge the gap from nano to micro
- GPUGPU enables simulations of micron-sized volumes, billions of atoms and studies of the long-time evolution of systems
- random site-selection is essential but can be harmed by domain decomposition



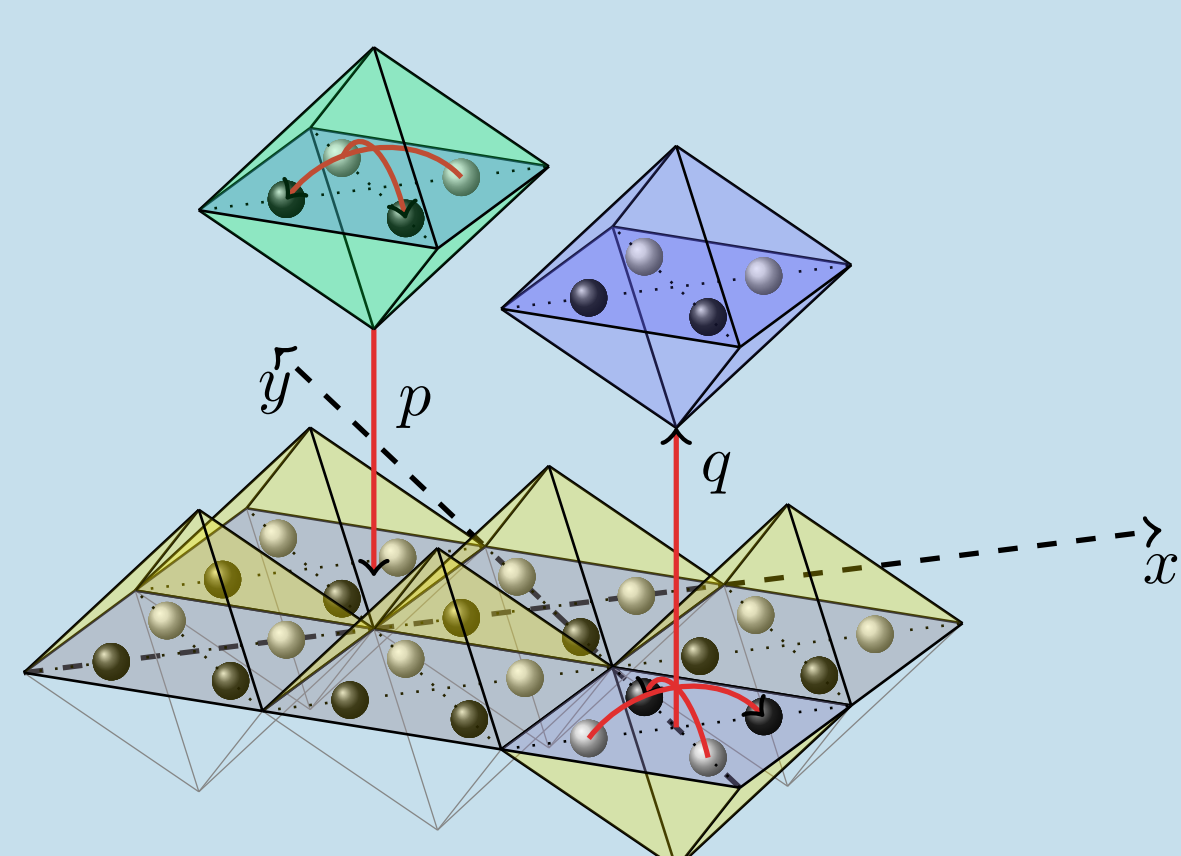
KPZ surface in the steady state

- Kardar-Parisi-Zhang (KPZ) universality class [1]
- can describe *growth processes*, directed polymers in random media, randomly stirred fluids, dissipative transport and magnetic flux lines in superconductors.

$$d_t h(\mathbf{x}, t) = \underbrace{v}_{\text{mean growth vel.}} + \underbrace{\sigma_2 \nabla^2 h(\mathbf{x}, t)}_{\text{surface tension}} + \underbrace{\lambda [\nabla h(\mathbf{x}, t)]^2}_{\text{local growth vel.}} + \underbrace{\eta(\mathbf{x}, t)}_{\text{noise}}$$

KPZ stochastic differential equation

Roof-Top Model



2 + 1D roof-top model—octahedron model

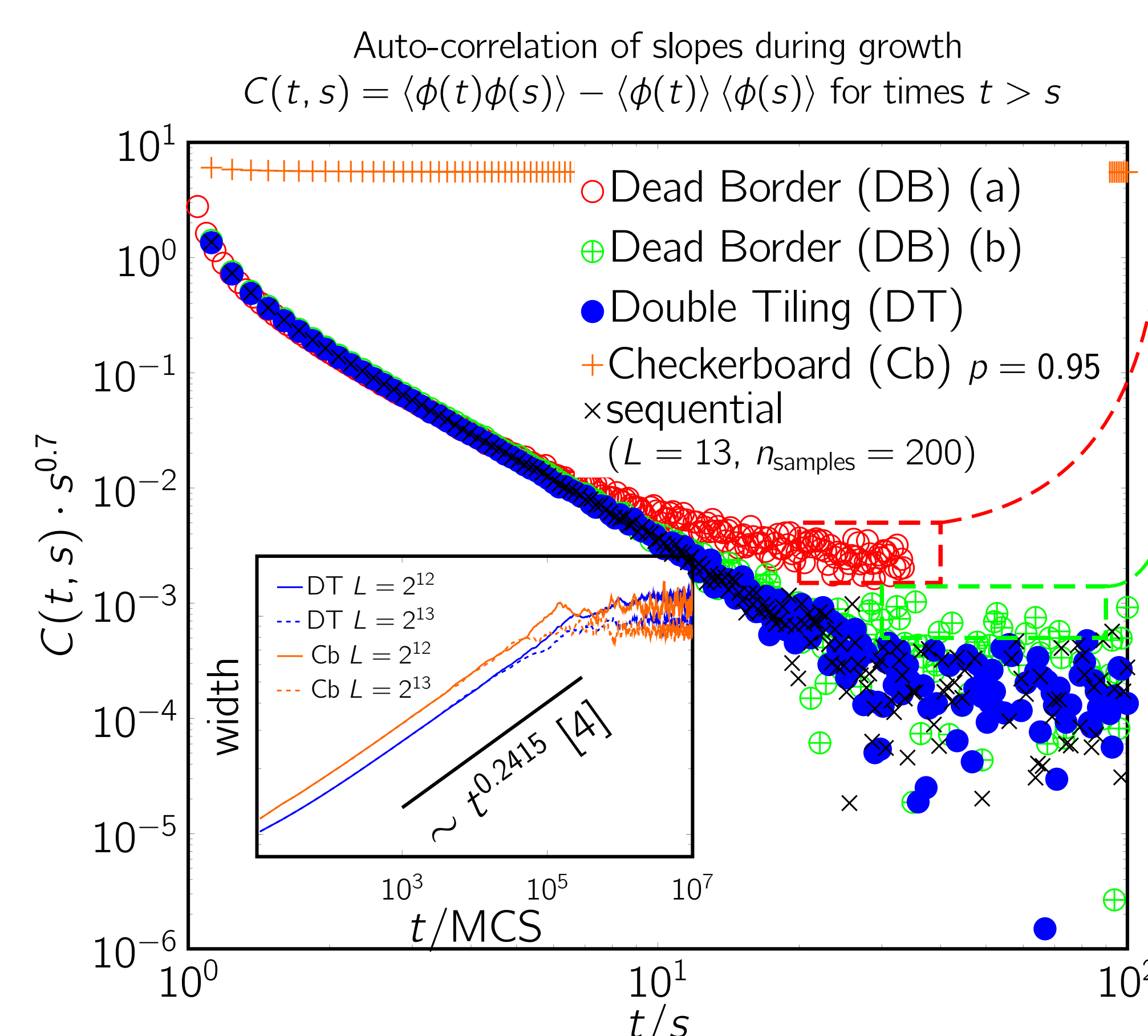
- stochastic model for surface growth [2, 3]
- 2D square lattice (octahedron model)
- carry out *random* depositions (probability p) or removals (prob. q)
- time t measured in sweeps of the lattice—Monte-Carlo-Steps (MCS)
- shows KPZ universality for $q = 0$
- bit-coded up- or down-slopes in between lattice-sites

Parallel Implementation

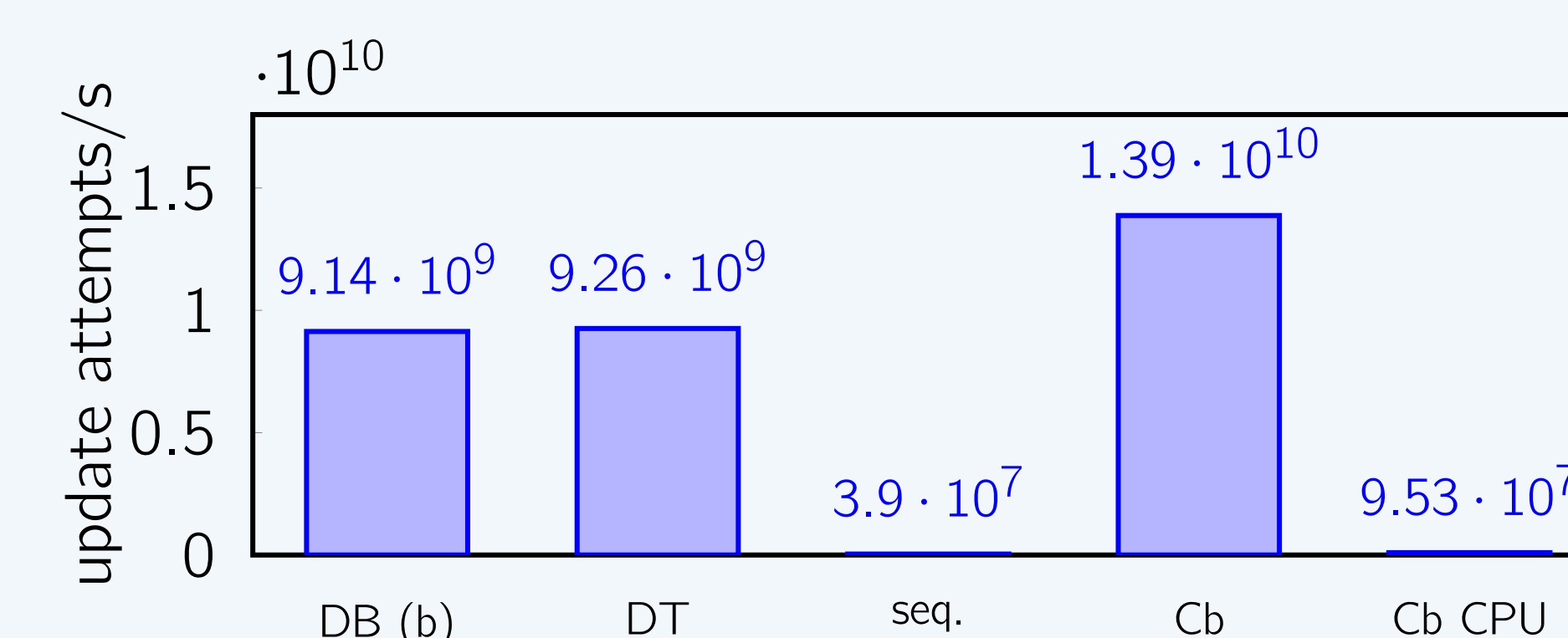
- CUDA implementation enables runs with $2^{17} \times 2^{17} \approx 16 \times 10^9$ lattice-sites [4] + self-averaging system \Rightarrow low noise in observables
- aim*: GPU implementation should preserve random site-selection \Rightarrow *domain decomposition* to distribute work among multiprocessors (blocks) see right-hand side \nearrow
- cells loaded into shared memory, updated collectively by threads
 - single-hit double tiling scheme with fixed borders
 - each thread maintains independent RNG state (TinyMT [5])

Domain Decomposition

- simulation-cell is split into non-interacting cells
- cells are treated independently for short periods of time, effectively with fixed boundary conditions \Rightarrow introducing small errors that must be attenuated
 - larger active domains give better results
- Dead Border Decomposition*: tiles, sites interacting with neighboring cells are inactive, origin randomized in intervals
- Double Tiling Decomposition*: tiles split into 4 subtiles, only one set of subtiles active at a time, origin randomized



Benchmarks



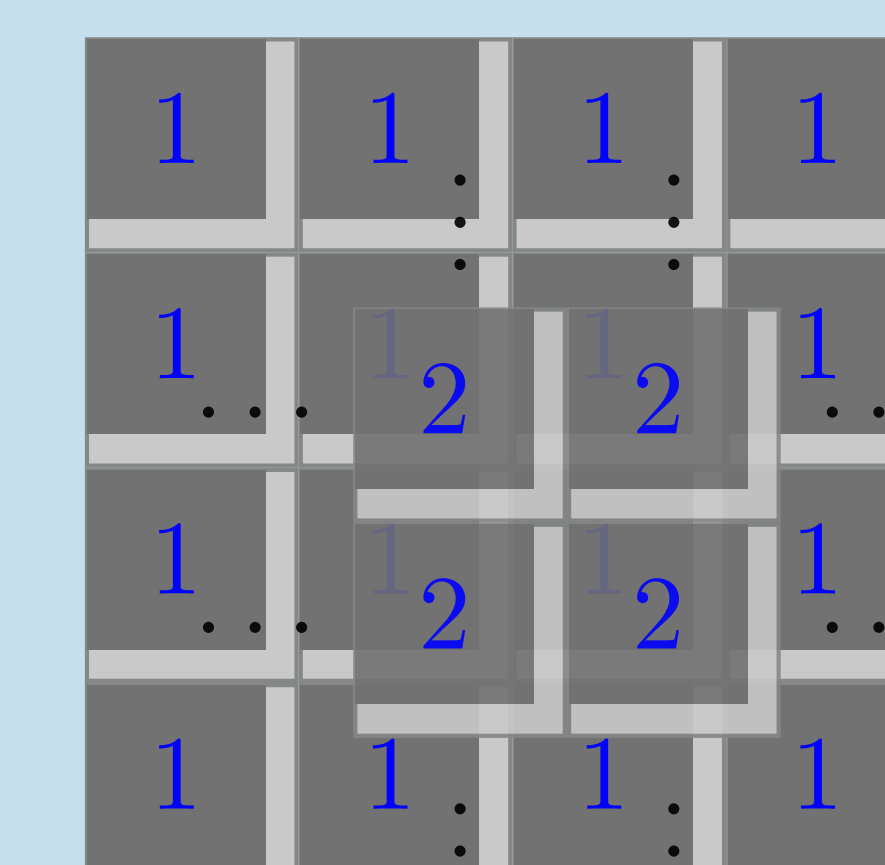
- sequential code: Intel Core i7-4930K @3.4 GHz
- CUDA code: NVIDIA GTX Titan Black

Conclusions

- \Rightarrow enabling long-time studies and large-scale simulations
- no correlations using double tiling with randomized origin
 - speedup $\sim 238\times$ vs. sequential code
- straightforwardly adaptable for multi-GPU

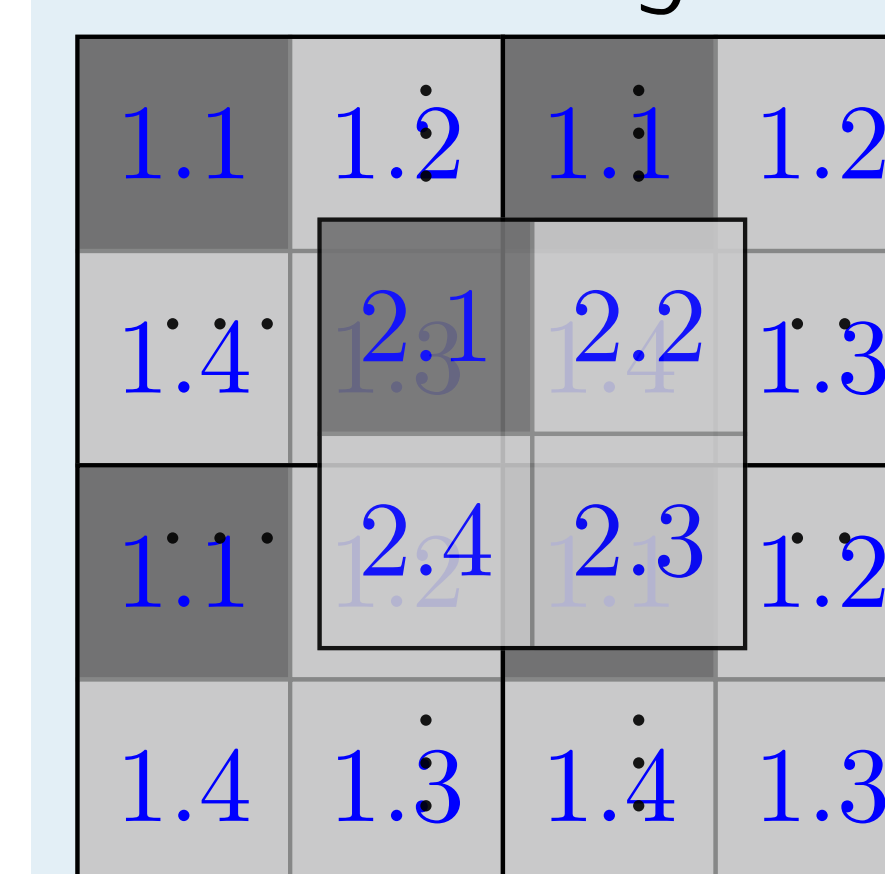
Dead Border Decomposition

- (a) smallest possible border for the Roof-Top Model: one lattice site to the right and bottom (see figure)
- borders can only be moved to word boundaries for encoding/performance reasons (chunks of 4×4 lattice sites)
 - + fast, good scaling results
 - restricted border-movement leads to correlations
- (b) remove restriction for borders by using wider borders (four sites)
- + removes correlation
 - bad signal-to-noise-ratio compared to sequential code ... due to added disorder from crossing of wide borders



Dead-border decomposition: Light-gray areas are domain borders with inactive sites.

Double Tiling Decomposition

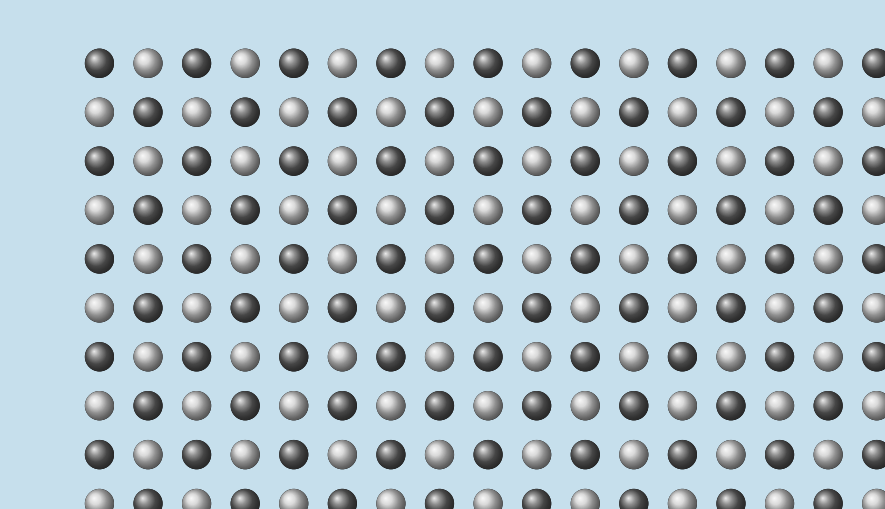


Double-tiling decomposition: Light-gray sub-cells are inactive.

- four non-interacting sets of sub-cells are updated in random order
- active regions overlap with inactive cells \Rightarrow works without shifting the origin for different models like Kinetic Metropolis Lattice Monte-Carlo [6]
- randomly shift origin of decomposition after every sweep to avoid auto-correlation error
- + noise en par with sequential code \Rightarrow *capability to perform large-scale aging studies*

Even/Odd site Checkerboard

- even sites do not interact with each other in single update \Rightarrow update even sites, then odd sites—often used for Ising model [7]
- replacing random site selection by finite update probability $p < 1$
- + correct growth results, very fast
- updates correlated, correlation depends on p



Lattice sites colored by parity.

Acknowledgements

We thank M. Bussmann for providing access to additional GPUs, the computing center at HZDR for support and the CCoE Dresden and the Helmholtz Initiative and Networking Funds for funding.

- Kardar, M., Parisi, G. & Zhang, Y.-C. *Phys. Rev. Lett.* **56**, 889–892 (1986).
- Plischke, M., Rácz, Z. & Liu, D. *Phys. Rev. B* **35**, 3485–3495 (1987).
- Ódor, G., Liedke, B. & Heinig, K.-H. *Phys. Rev. E* **79** (2009).
- Kelling, J. & Ódor, G. *Phys. Rev. E* **84**, 061150 (2011).
- TinyMT. URL <http://www.math.sci.hiroshima-u.ac.jp/~%20m-mat/MT/TINYMT/index.html>.
- Kelling, J., Ódor, G., Nagy, M. F., Schulz, H. & Heinig, K. *The European Physical Journal - Special Topics* **210**, 175–187 (2012).
- Preis, T. *et al. J. Comp. Phys.* **228**, 4468–4477 (2009).
- Kelling, J., Ódor, G. & Gemming, S. *in preparation* (2015).