1	Accelerating the Lagrangian particle tracking of residence time
2	distributions and source water mixing towards large scales
3	Chen Yang <sup>a</sup> , You-Kuan Zhang <sup>b,c</sup> , Xiuyu Liang <sup>b,c</sup> , Catherine Olschanowsky <sup>d</sup>
4	Xiaofan Yang <sup>e</sup> *, Reed Maxwell <sup>a</sup> *
5	
6	<sup>a</sup> Department of Civil and Environmental Engineering and the Princeton Environmental Institute, Princeton University,
7	Princeton, NJ 08544, USA. e-mail: reedmaxwell@princeton.edu
8	<sup>b</sup> Guangdong Provincial Key Laboratory of Soil and Groundwater Pollution Control, School of Environmental Science
9	and Engineering, Southern University of Science and Technology, Shenzhen 518055, China.
10	°State Environmental Protection Key Laboratory of Integrated Surface Water-Groundwater Pollution Control, School
11	of Environmental Science and Engineering, Southern University of Science and Technology, Shenzhen 518055, China.
12	<sup>d</sup> Computer science department, College of Engineering, Boise State University, Boise, ID 83704, USA.
13	°State Key Laboratory of Earth Surface Processes and Resource Ecology, Faculty of Geographical Science, Beijing
14	Normal University, Beijing 100875, China. e-mail: xfyang@bnu.edu.cn
15	*Corresponding author
16	
17	All codes for this work are publicly available on GitHub along with the Hillslope model used for the simulations:
18	https://github.com/reedmaxwell/EcoSLIM
19	
20	This manuscript has been submitted for publication in Computers & Geosciences. Please note that, despite
21	having undergone peer-review, the manuscript has yet to be formally accepted for publication. Subsequent
22	versions of this manuscript may have slightly different content. If accepted, the final version of this manuscript
23	will be available via the 'Peer-reviewed Publication DOI' link on the right-hand side of this webpage. Please
24	feel free to contact any of the authors; we welcome feedback.

Dr. Chen Yang was responsible for code development, code tests, and manuscript preparation. Dr. Zhang and Liang assisted with the computing resources. Dr. Olschanowsky assisted with the code development. Dr. Xiaofan Yang assisted with the code development, computing resources, and manuscript preparation. Dr. Maxwell assisted with the code development, code tests, and manuscript preparation.

### 25 Abstract

26 Travel/residence time distributions (TTDs/RTDs) are important tools to evaluate the vulnerability of catchments 27 to contamination and understand many aspects of catchment function and behavior. In recent years, the calculation of 28 TTDs/RTDs based on the Lagrangian particle tracking approach together with the integrated hydrologic modeling has 29 become a popular counterpart to analytical approaches and lumped numerical models. As global water availability 30 becomes more stressed due to anthropogenic disturbance and climate change, the requirement of large-scale and long-31 term simulations for TTDs/RTDs further pushes the high computational costs of Lagrangian particle tracking. Hence, 32 speeding up the Lagrangian particle tracking approach becomes an important barrier to advancement. In this study, 33 we accelerate the Lagrangian particle tracking program EcoSLIM, using a combination of distributed (e.g. MPI) and 34 multi-core accelerator (CUDA) approaches for large-scale and long-term simulations. EcoSLIM was developed to be 35 seamlessly paired with the integrated ParFlow.CLM model for calculations of transient RTDs and source water mixing 36 and was originally developed using threaded OpenMP. This work extends this implementation to compare 37 combinations of MPI, CUDA and OpenMP. Of these combinations, the OpenMP-CUDA parallelism performed the 38 best moving from single-GPU to multi-GPU. The multi-GPU shows strong scalability which becomes increasingly 39 efficient with more particles, demonstrating a potential feasibility for regional-scale, transient residence time 40 simulations. This work largely improves the computational capability of EcoSLIM, and results also show the 41 advantages of using GPU-parallel to traditional parallel-APIs (application programming interfaces) and its potential 42 to widely accelerate the next generation programs in subsurface environment modeling.

Keywords: Lagrangian particle tracking, Integrated modeling, Travel/residence time distributions, MPI, Multi GPU

# 46 **1. Introduction**

Travel time distribution (TTD) and residence time distribution (RTD) are probability density 47 functions of age for water parcels leaving from and storing in the catchment subsurface (Yang et 48 al., 2018). Both TTD and RTD are important tools to reflect how catchments retain and release 49 water which in turn indicate the storage and turnover of the contaminants (Jing et al., 2019; 50 Kirchner et al., 2000; McDonnell et al., 2010; van der Velde et al., 2010; Yang et al., 2018). 51 Traditionally, analytical approaches and lumped numerical models are common tools to determine 52 the TTDs/RTDs (Basu et al., 2012; Benettin et al., 2015; Benettin et al., 2013a; Benettin et al., 53 2013b; Botter et al., 2011; Engdahl et al., 2016; Kirchner et al., 2001; Małoszewski and Zuber, 54 1982; McDonnell et al., 2010). 55

However, in recent years, Lagrangian particle tracking together with physically-based, 56 integrated hydrological models has emerged as a promising tool for estimating TTD/RTD 57 (Danesh-Yazdi et al., 2018; de Rooij et al., 2013; Engdahl and Maxwell, 2015; Engdahl et al., 58 2016; Jing et al., 2019; Jing et al., 2020; Wilusz et al., 2019; Yang et al., 2018). Though particle 59 tracking generally is more computationally efficient than most Eulerian approaches (Tompson and 60 Gelhar, 1990; Yang et al., 2018), the computation burden can still be quite high if fine spatial and 61 temporal resolutions or large-scale and long-term simulations are considered. As such, most of the 62 current studies are limited by the total number of particles thus constraining the problem size or 63 simulation duration. For example, the study areas of some recent applications, e.g. Yang et al. 64 (2018), Weill et al. (2019), and Wilusz et al. (2019) are quite small, only 1.44-, 0.8-, and 0.4-km<sup>2</sup> 65 respectively. Wilusz et al. (2019) adopted an hourly timestep while the total simulation time was 66 3 years. Though a longer, 10-year simulation was conducted by Yang et al. (2018), a daily timestep 67 68 was used. Similar studies with larger scale study area but coarse timestep include but not limited to the Nägelstedt catchment of 850 km<sup>2</sup> in Jing *et al.* (2019) and Jing *et al.* (2020) and the Little Washita watershed of about 600 km<sup>2</sup> in Kollet and Maxwell (2008b). While the continental US of  $6.3 \times 10^6$  km<sup>2</sup> was simulated in Maxwell *et al.* (2016), this simulation was steady-state, and the particles were run separately in several subsets of the domain.

More importantly, the objectives of these large-scale studies are all TTDs of outflow 73 (streamflow) instead of RTDs of groundwater. Simulation of TTDs is conducted by injecting a 74 fixed number of particles and tracking them until all of them leave the modeling domain. Whereas, 75 76 the simulation of RTDs has to be spun-up; running the system to a dynamic equilibrium when particles filling up the subsurface (Maxwell et al., 2019; Yang et al., 2018). The number of total 77 particles required for RTDs cannot be determined a priori, which results in a much higher 78 79 computational requirement for RTDs than TTDs. With the recent advances in TTDs/RTDs, the travel time of evapotranspiration (i.e., the evapotranspiration time) (Botter, 2012; Maxwell et al., 80 2019) and the age of pumped groundwater under anthropogenic disturbance, have the same 81 82 requirement of reaching a dynamic equilibrium. In addition, under the climate change, long-term 83 simulations are necessary for in-depth understanding of the RTDs (Jing et al., 2020; Manning et al., 2012; Yang et al., 2018). Last but not least, transport modeling at larger scales also motivates 84 85 further development of Lagrangian particle tracking (Hartmann et al., 2020; Jing et al., 2019; 86 LaBolle et al., 1996; Tompson, 1993; Tompson and Gelhar, 1990). As a result, it is critical to accelerate the computation speed of the Lagrangian particle tracking. 87

Modeling platforms including both hardware and software are required to meet the increasing demand of memory and speed, which can benefit the continuous development of subsurface environmental modeling, and, otherwise, would be a bottleneck (Jacobsen *et al.*, 2010; Scheibe and Smith, 2015). CPUs (Central Processing Units)-based high performance clusters are generally

used to solve computationally expensive problems. MPI (Message Passing Interface)-based 92 ParFlow.CLM (Kollet and Maxwell, 2008a), PFLOTRAN (Hammond et al., 2014), ParMIN3P-93 THCm (Su et al., 2017), and eSTOMP (Freedman et al., 2016) are representatives in massively-94 parallel groundwater and/or reactive transport models (Steefel et al., 2015). Particularly, Engdahl 95 et al. (2019) parallelized the Lagrangian simulations of mixing-limited reactive transport using 96 MPI based on a novel parallel particle scheme. OpenMP (Open Multi-Processing), supporting 97 shared memory multi-threading programming on CPU, is also widely used, which, however, when 98 99 used as the only parallelization strategy is more suitable for medium-size computation or on personal workstations. OpenMP can also be nested within MPI for further intra-node speedup on 100 clusters. Typical examples are the current version of EcoSLIM (Maxwell et al., 2019) using 101 102 OpenMP and the TOUGHREACT using hybrid MPI-OpenMP (Steefel et al., 2015).

Since 2005, the application of GPU (Graphics Processing Unit) to scientific computing has 103 attracted more attention due to the large number of cores contained within each GPU. With the 104 105 appearance of NVIDIA's CUDA (Compute Unified Device Architecture) in 2007, GPU-based 106 parallel programing has been increasingly utilized in scientific computing (Ruetsch and Fatica, 2014). In geoscientific modeling, GPU-based acceleration mainly focus on earth system models 107 such as the atmosphere and ocean models (Porter et al., 2018; Xu et al., 2015). MIKE 21 GPU is 108 109 the GPU-parallel version of MIKE 21 which is for coastal modeling (DHI, 2019). GPUs have also been widely used in particle/lattice-based computational fluid dynamics (CFD) methods and 110 simulations (Cheng et al., 2015; Jacobsen et al., 2010; Molinero et al., 2019). Whereas, the GPU-111 parallelism is still in its infancy in modeling the subsurface environment and related integrated 112 113 modeling. Ji and her collaborators conducted GPU-parallel on MODFLOW and MODPATH (Ji et al., 2010; Ji et al., 2014; Ji et al., 2019; Sun et al., 2019). The reported speedup of MODFLOW 114

using NVIDIA Tesla C1060 is 1.6- to 10.6-fold for models with more than 10<sup>5</sup> cells while the
speedup of MODPATH on simulating more than 5 million streamlines is over 1000 by using 8
NVIDIA GPUs. However, for the next generation code/program handling coupled processes at
large scales, the CUDA-version ParFlow is expected to become a powerful tool (Kuffour *et al.*,
2020).

EcoSLIM (Maxwell et al., 2019), based on a Lagrangian particle tracking approach, works 120 seamlessly with ParFlow.CLM, an integrated hydrologic model (IHM) coupling the groundwater 121 122 model and land surface model (Kuffour et al., 2020; Maxwell and Miller, 2005). The current EcoSLIM focuses on transient RTDs of different hydrological components at the watershed scale, 123 such as groundwater in variably-saturated zone, evapotranspiration (ET), and outflow. The latter 124 125 two are the same to travel times of ET and outflow mentioned above. Another feature of EcoSLIM is to identify the source water mixing of hydrological components (Maxwell et al., 2019). For 126 example, ET particles that exit the modeling domain can be further characterized by source water 127 128 composition: for example, the fractions of groundwater, rain, or snow. Particularly, the RTD of 129 ET and the source water mixing are rarely addressed in previous modeling approaches (Maxwell et al., 2019), which makes EcoSLIM as a promising tool to model RTDs by IHM and particle 130 tracking, advancing the work of lumped models and more simple analytical solutions. Nested 131 132 within ParFlow.CLM, EcoSLIM also has a possible extension to large-scale contaminant transport modeling, which can trace the sources and traveling trajectories of the contaminants as well as 133 calculate the RTDs that are important for remediation. 134

The objective of this study is to accelerate the EcoSLIM code for large-scale and long-term simulations. The EcoSLIM code was extended to include MPI and GPU parallelism. This was constructed using both the CPU-based MPI and the GPU-based CUDA Fortran as extensions to

its original OpenMP parallel formulation. More importantly, different approaches from single 138 GPU to multi-GPU were inter-compared and evaluated. The rest of this article is organized as 139 follows. In the second section, the parallelism currently in EcoSLIM (i.e., the particle loops) is 140 identified and profiled; the implementation of new parallelism with different APIs (application 141 programming interfaces) is described. In the third section, the model setup and hardware for a suite 142 of test cases are introduced. Then the simulation results from MPI- and CUDA-parallel are verified 143 against those from the original OpenMP-parallel, which are presented to demonstrate the accuracy 144 145 of new parallel approaches. In the fourth section, the computation performance of three parallel-APIs are intercompared. The computation capability based on multi-GPU is compared to previous 146 147 studies, which to the best of our best knowledge, is the first time. Finally, conclusions are drawn 148 for this study.

# 149 **2. Methodology**

### 150 2.1. The parallel particle loop in EcoSLIM

We first evaluated the computational cost of EcoSLIM using the hillslope model introduced in Maxwell *et al.* (2019). Details of the hillslope model are presented later in section 3.1. Using one CPU thread based on OpenMP, results show that the time used by the particle loop in EcoSLIM is over 99.8% of the total simulation time (Table 1). Hence, the parallelization in this study is conducted only for the particle loop in EcoSLIM. A flow chart of the particle loop is shown in Fig. 1, with key variables summarized in Table 2. For each active particle in a global timestep (i.e., the timestep in ParFlow), the execution proceeds as follows:

a) It is first looped by the local particle timestep. This loop continues until the total
accumulated time is larger than the current global timestep. Then the adjacent cell of this
particle (#9) and the grid-cell aggregated information [C(1:5, :, :, :)] (#10) are updated.

b) In the local time loop, the adjacent cell of a particle is determined by its coordinates
[P(:,1:3)] (#1). Next, cell fractions in x, y, and z directions of a particle are determined (#3)
before its advection velocities being linearly interpolated from the velocities (variables 12–
14 in Table 2) distributed on cell-interfaces (#4). Thus, the local particle timestep can be
calculated (#5).

- c) The location of the particle is updated together with the deviation caused by molecular
  diffusion (#7). Finally, the particle's trajectory is amended with the reflection principle
  (LaBolle et al., 1996) if the particle is out of the domain (#8).
- d) In the local time loop of each particle, whether the particle is out of the domain either through outflow or ET is evaluated (#2 and #6). If a particle leaves as outflow or ET, its status is switched to inactive. Meanwhile, the age, mass, composition, and particle-number of the outflow or ET are updated (variables 3–10 in Table 2). For ET, *C* array [*C*(6:9, :, :, :)] is also updated for grid-cell aggregated information of ET.

174 Table 1. Baseline simulation time for EcoSLIM using one CPU thread based on OpenMP

Case name	Particle loop (hour)	Total time (hour)	Percentage (Loop/Total)
ER_Shrubs	33.81	33.87	99.83%
ER_Trees	70.21	70.33	99.83%
LW_Shrubs	111.02	111.21	99.83%
LW_Trees	84.54	84.70	99.81%

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#### Table 2. Key variables in the particle loop

Index	Array	Function	GPU memory
1	P(1:np,1:10) np is the maximum permitted number of particles in the simulation. For each particle, from 1 to 10, the array includes: xyz coordinates, residence time (RT), saturated RT, mass, source, active/inactive status, concentration, and exit status.		Global/Shared. Copied to shared memory before computation and copied back to global memory after computation.
2	C(9,nx,ny,nz)	nx, ny, and nz are the grid dimension of the modeling domain. 1 to 9 is the grid-cell aggregated information: concentration, age, mass, source, P(:,9), ET particles, ET mass, ET age, and ET source.	Global. Atomic operations.
3	ET_age(pfnt,5)	pfnt is the total timesteps in the simulation. ET_age(:,1) is the mass weighted age of ET.	Global/Register

4	ET comp(pfnt,3)	ET_comp(:,1:3) is the source water composition of ET_mass. From 1 to 3: initial water, rain, and	Global/Register
4		snowmelt.	
5	ET mass(pfnt)	The mass of ET.	Global/Register
6	ET np(pfnt)	The number of particles exit as ET.	Global/Register
7	Out_age(pfnt,5)	The mass weighted age of outflow.	Global/Register
8	Out_comp(pfnt,3)	Out_comp(:,1:3) is the source water composition of outflow. From 1 to 3: initial water, rain, and snowmelt.	Global/Register
9	Out mass(pfnt,5)	Out_mass(:,1) is the mass of outflow.	Global/Register
10	Out_np(pfnt)	Number of particles exit as outflow.	Global/Register. Variables 3–10 have private copies on registers. Atomic summation of threads on global memory is conducted.
11	dz(nz)	Vertical thickness of layers	Texture
12	Vx(nx+1,ny,nz)	Transient velocity in x direction.	Texture
13	Vy(nx,ny+1,nz)	Transient velocity in y direction.	Texture
14	Vz(nx,ny,nz+1)	Transient velocity in z direction.	Texture
15	Saturation(nx,ny,nz)	Saturation.	Texture
16	Porosity(nx,ny,nz)	Porosity.	Texture
17	EvapTrans(nx,ny,nz)	Evapotranspiration (ET). 12–15 and 17 change for each global timestep. Porosity keep constant in the whole simulation.	Texture

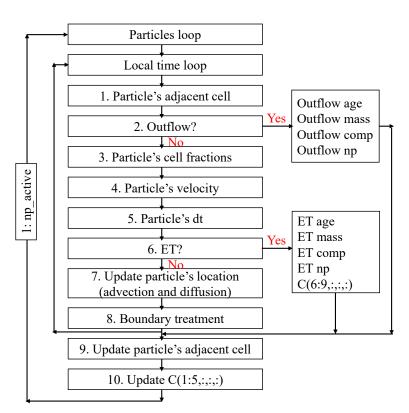


Fig. 1. Flow chart of the particle loop in a global timestep in EcoSLIM.

#### 181 **2.2.** Memory management via CUDA-capable GPU

Memory management via CUDA-capable GPU is shown in Fig. 2. The shared memory, L1 182 cache, local memory, and registers are on-chip memory; while global, constant, and texture 183 memories are all in DRAM (dynamic random-access memory) of the GPU device. There are also 184 on-chip caches for constant and texture memories. Global memory can be read and written by both 185 CPU and GPU, and it is available by all threads launched on GPU. Constant and texture memories 186 can be read and written by CPU while they are read-only for threads on GPU. For each thread 187 188 block, shared memory can be read and written by all threads in that block. Shared memory and L1 cache are configurable memories of 64 KB in total. Variables stored in registers are private for 189 each thread. If the memory of registers is insufficient, data will be stored in local memory which 190 191 is also thread private. CUDA Fortran, which was first developed in late 2009, is adopted for GPU programming in this study. CUDA Fortran can be implemented using The Portland Group<sup>®</sup> (PGI<sup>®</sup>) 192 Fortran compiler which has the NVIDIA's CUDA architecture. 193

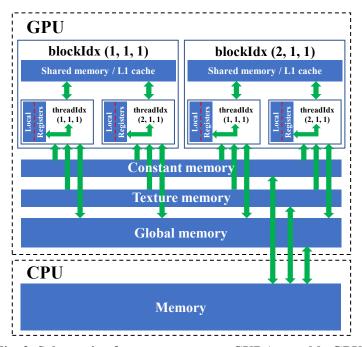


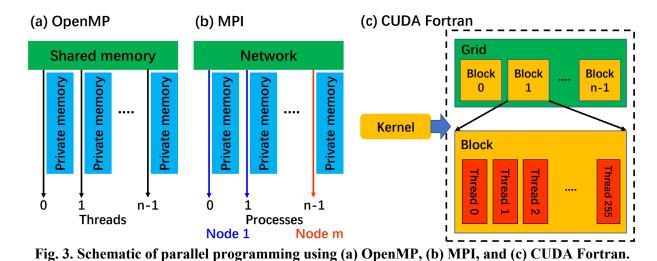


Fig. 2. Schematic of memory types on CUDA-capable GPU.

#### 2.3. Parallel programing 196

#### 2.3.1. OpenMP 197

The original EcoSLIM (Maxwell et al., 2019) code was implemented in OpenMP and is 198 described briefly here for comparison among different parallel APIs. The total np active particles 199 are implicitly decomposed by OpenMP and assigned to *n* threads (Fig. 3a). Threads number *n* can 200 be set by the environment variable OMP NUM THREADS before starting the run. All variables 201 in Table 2 and other public information (e.g., np active, dimensions of modeling domain, global 202 203 timestep) are shared by all threads while the particle-dependent information (e.g., particle's adjacent cell and cell fractions, random numbers for ET and diffusion, particle's velocities, etc.) 204 are thread private. For calculations related to outflow, ET, and C in the particle loop (#2, #6, and 205 206 #10 in Fig. 1), atomic operations are used to avoid the concurrent writing of these arrays by more than one thread. 207



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211
      2.3.2. MPI
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In each global timestep, the loop of *np* active particles are evenly decomposed to *n* sub-loops. 212 213 *n* is the number of processes launched in the run (Fig. 3b). Hence, *P*(1:*np* active,:) is divided into

n segments. If  $np\_active$  is not divisible by *n*, the remainder *m* particles are distributed to the first *m* processes. *P* segments, velocities, saturation, ET, and porosity are then distributed from the root process to others. New variables corresponding to variables 2–10 are declared and set as zero in each global timestep before the sub-loop to gather the information of outflow, ET, and *C* on each process. Once finished, the *P* array on the root process is updated by segments from other MPI processes. ET, outflow, and *C* from each process represented by new variables are summed to update variables 2–10 in Table 2.

#### 221 **2.3.3. CUDA Fortran**

When implementing EcoSLIM on the GPU, one kernel is defined (Fig. 3c) and launched once in each global timestep. Size of the thread block (*block\_size*) is 256, and the number of thread blocks is determined by *ceiling(real(np\_active)/block\_size)*. Taking the 1080 Ti GPU card as an example, the maximum number of thread blocks permitted is 2,147,483,647; which is 549,755,813,632 threads in total if the block size is 256. This number is beyond the number of particles in all test cases in this study, so each thread is responsible for only one particle.

228 Arrays of P, C, and ET and outflow (variables 1-10 in Table 2) are both read and written in the particle loop and thus stored in the global memory. Input variables such as layer thickness, 229 porosity, velocities, saturation, and ET for each global timestep (variables 11-17 in Table 2) use 230 231 the read-only texture memory. Texture memory has caches on chip and can be accessed more rapidly compared to global memory. It is also advantageous for noncontiguous access to these 232 arrays by threads because variables 12-17 in Table 2 are stored with the sequence by index of grid-233 cells while threads are indexed by the sequence of particles in P. After P is copied from CPU to 234 235 GPU, it is further copied from global memory to shared memory of each thread block to improve the efficiency of thread access. 236

In the GPU kernel, a group of thread-private variables for outflow and ET are declared using 237 registers. Outflow and ET information are first obtained on each thread without waiting. After the 238 particle loop, the summation of these variables (variables 3–10 in Table 2) on each thread are 239 computed by the reduction method used in the  $\pi$  computation with Monte Carlo method (Ruetsch 240 and Fatica, 2014). For the C array, atomic operation is directly performed in the particle loop. For 241 molecular diffusion and ET processes, random numbers are generated by a GPU function called 242 in the kernel function. After the execution, arrays of P, C, and ET and outflow (variables 1–10 in 243 244 Table 1) are copied back to CPU to update the information.

#### 245 **2.3.4. Multi-GPU**

Three multi-GPU approaches were considered for EcoSLIM (Fig. 4): (1) Multi-GPU via 246 247 asynchronous data transfer with CUDA stream, (2) Multi-GPU with MPI, and (3) Multi-GPU with OpenMP. For CUDA stream (Fig. 4a), the particle loop is equally divided into sub-loops, and each 248 GPU is assigned to one sub-loop. The total number of sub-loops is determined by the total number 249 250 of GPUs. Computation of sub-loops on GPUs use the same kernel explained in section 2.3.3. A single CPU thread is used while the switch between different GPUs is through asynchronous data 251 transfers with specific CUDA stream for each GPU. Communication with MPI (Fig. 4b)/OpenMP 252 (Fig. 4c) is similar to that with CUDA stream (Fig. 4a), only that different GPUs are assigned to 253 254 different CPU processes/threads. The main difference between MPI and OpenMP is that processes can belong to different computation nodes while all threads belong to the same node (Fig. 3a and 255 3b). 256

Obviously, for all three approaches using multi-GPU, the overhead on CPU is necessary to decompose the loop of particles to sub-loops, distribute necessary information to different streams/processes/threads, and combine results from different streams/processes/threads for updates. For CUDA stream, this is realized by *do loops* while for MPI, this is realized by MPI communicators. However, for OpenMP, variables are simply set with shared or private attributes at the beginning of the parallel region. Through tests in this study, this overhead for CUDA stream and MPI is over the GPU time decreased by multi-GPU relative to the single GPU code in section 2.3.3. Therefore, CUDA stream and MPI are not suitable to parallelize EcoSLIM with current structure using multi-GPU and hence will not be discussed any more.

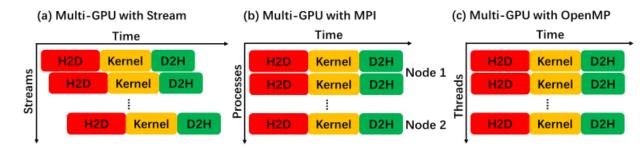


Fig. 4. Schematic of parallel programming using multi-GPU. H2D represents data transfer from CPU
to GPU while D2H represents GPU to CPU.

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# 270 3. Model setup, hardware, and validity of parallels

### 271 **3.1. Hillslope model**

The Hillslope model (Maxwell et al., 2019) was adopted to test the parallel performance of 272 273 EcoSLIM. Geometry of the modeling domain has the length of 100-, 1-, and 9.4-m in x, y, and zdirections, respectively. It was divided into 20 columns, 5 rows, and 20 layers with constant 274 275 resolutions of 5- and 0.2-m in x and y directions while, in z direction upward, various thickness of layers was set as: 0.5 m for the bottom 18 layers and 0.3- and 0.1-m for the top 2 layers. The soil 276 properties are homogeneous, including saturated hydraulic conductivity of 0.05 m/h, Manning's 277 N of 10<sup>-6</sup> m<sup>1/3</sup>h<sup>-1</sup>, porosity of 0.2, and van Genuchten parameters with  $\alpha$  of 1.0 m<sup>-1</sup> and exponent n 278 of 2.0. Two real meteorological-forcings were considered to drive ParFlow.CLM, representing a 279 high elevation, snow dominated mountain headwaters (ER) and a semiarid, rain-dominated plains 280

system (LW). Two homogeneous land-cover types were used which are the Evergreen Needleleaf
plant functional type (Trees) and the Shrub plant functional type (Shrubs). Thus, four cases were
tested with combination of the meteorological forcings and the land-cover types named as: ERShrubs, ER-Trees, LW-Shrubs, and LW-Trees.

For both ParFlow.CLM and EcoSLIM, no flux boundaries were used except the top boundary 285 which is open for precipitation and exit of outflow and ET. ParFlow.CLM simulations of 5 286 consecutive years with hourly timestep were conducted. One-year forcing data were repeatedly 287 288 used in all the simulations. Dynamic equilibrium of the flow field was achieved at the end of the simulation for all four cases. Thus, the transient flow field in the last year of the 5-year simulation 289 is repeatedly used in the EcoSLIM simulation of 20 years with hourly timestep. At the end of the 290 291 simulation, EcoSLIM system achieved the dynamic equilibrium. By injecting 2 particles into the modeling domain per precipitation event, the mean particle-numbers in the last year of the 292 simulation for four cases are 0.39, 0.83, 1.30, and 1.03 million, respectively. For the test in section 293 294 4.3, it was increased to 2000 particles per precipitation event to test the capability using multi-295 GPU toward large scales.

### 296 **3.2. Hardware**

The following platforms were used to test the OpenMP-, MPI-, and CUDA-version EcoSLIM.
Platform (1) is to test OpenMP and MPI while (1), (2), and (3) are set for CUDA.

(1) A workstation. It is equipped with GeForce GTX 1080 Ti (GPU) and Intel<sup>®</sup> Xeon<sup>®</sup> CPU
E5-2683 v3 @ 2.00GHz. There are 28 physical CPU cores with hyper-threading technology. The
maximum available threads for OpenMP is 56 while the maximum available processes for MPI is
28 with the PGI compiler. The driver version for NVIDIA GPU is 440.33.01, CUDA version is
10.2, and PGI version is 19.10 with Open MPI of 3.1.3. All OpenMP, MPI, and CUDA version
codes were compiled and built by the PGI compiler.

Tianhe-2 (TH-2) supercomputer in the National Supercomputer Center in Guangzhou, China, *gpu\_v100 partition*. It has 4 GPU cards of Tesla V100 (SXM2, 16GB) per computation node and the CPU processor of Intel<sup>®</sup> Xeon<sup>®</sup> Gold 6132 CPU @ 2.60GHz. Driver version for NVIDIA GPU is 418.67, CUDA version is 10.1, and PGI version is 19.10.

(2) Tianhe-2 (TH-2) supercomputer in the National Supercomputer Center in Guangzhou,
China, gpu partition. It has the GPU card of Tesla K80 (11 GB) and the CPU processor of Intel®
Xeon® CPU E5-2660 v3 @ 2.60GHz. Driver version for NVIDIA GPU is 390.30, CUDA version
is 8.0, and PGI version is 17.1.

#### 313 **3.3.** Code-to-code verification: MPI- and CUDA-parallel of EcoSLIM

Taking outflow as an example, daily flow rate and residence time for four cases via MPI- and 314 OMP were plotted in Fig. 5; while those using CUDA- and OMP were plotted in Fig. 6. OpenMP 315 316 is abbreviated as OMP here. Only results for the last year of the 20-year simulation are shown due to the achievement of dynamic equilibrium in all simulations as mentioned in section 3.1. Results 317 from MPI and CUDA are highly consistent with those from OMP for all cases, which demonstrated 318 319 the accuracy of the parallel implementation. The little discrepancy between MPI/CUDA and OMP should be attributed to the generation of random numbers for initial position of particles, particles 320 captured as ET, and random walk induced by molecular diffusion. 321

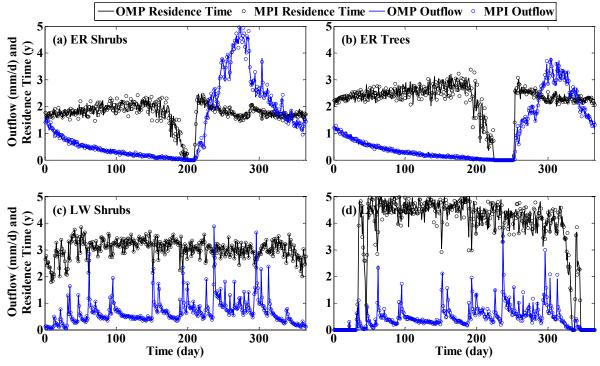
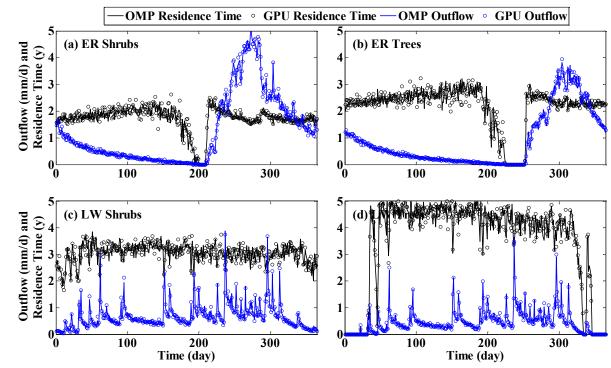
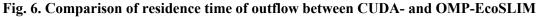


Fig. 5. Comparison of residence time of outflow between MPI- and OMP-EcoSLIM



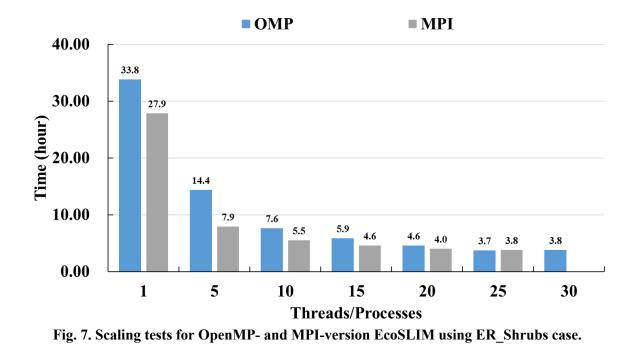


#### 328 **4. Parallel performance**

## 329 4.1. Speedup by OMP and MPI

Scaling tests for OMP and MPI conducted on the workstation based on ER Shrubs are shown 330 in Fig. 7. For MPI using PGI compiler, a maximum of 28 processes can be launched since there 331 are 28 CPU cores on the workstation. In general, MPI has the advantage that more processes can 332 be used on the distributed architecture while the speedup by OMP may be limited by the number 333 of CPU cores on one computational node. However, through the results from the scaling tests 334 shown in Fig. 7, both OMP and MPI cannot unlimitedly speedup the computation with high 335 efficiency even though CPU threads/processes are continuously increasing. For OMP, the time 336 337 consumption decreases with increasing number of CPU threads to a maximum of 25 (3.74 h). The time used for 30 threads (3.78 h) is even more than that with 25 threads. Limited speedup can also 338 be observed for MPI from 20 to 25 processes. We further tested all four cases on a distributed 339 architecture using MPI with 20, 25, 30, 35, and 40 processes. For all four cases, speedup plateaus 340 using more than 25 processes. However, it is noted that the upper limit of the processes should be 341 case and platform dependent for both OMP and MPI. 342

Additionally, MPI is generally faster than OMP for the same number of threads/processes (Fig. 7). This difference becomes smaller when the number of threads/processes is increased. As an exception, MPI (3.78 h) becomes slower than OMP (3.74 h) with 25 threads/processes. This can be explained by the fact that computation tasks are assigned to different threads implicitly by OMP while they are explicitly decomposed and distributed to different processes via MPI. However, the quick access to shared memory by different threads using OMP may be more advantageous than the message passing among different processes when a larger number of threads/processes are used, which is why MPI becomes slower when the number of threads/processes is beyond athreshold such as 25 in Fig. 7.





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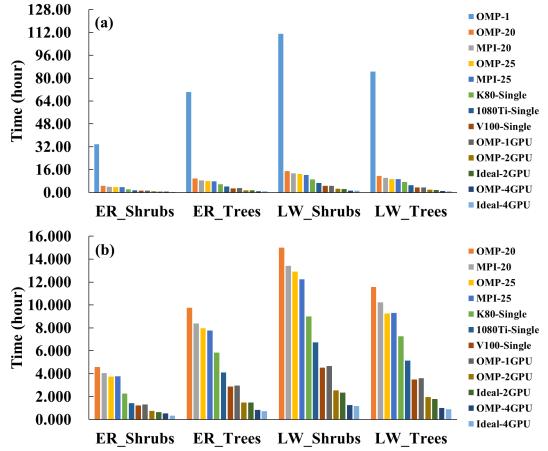
353



Parallel performance for different APIs were plotted in Fig. 8. The original data and speedup 356 were summarized in Tables 3 and 4. Results marked with OMP-number and MPI-number were 357 obtained from the workstation where number represents number of threads/processes. Only those 358 for 20 and 25 threads/processes are shown since no obvious speedup can be achieved when using 359 more than 25 processes for all four cases. GPU-single represents results by using different GPU 360 cards for the parallel with single-GPU described in section 2.3.3. OMP-nGPU represents results 361 for multi-GPU with OpenMP using Tesla V100 cards where n is the number of GPU cards used. 362 Ideal-*n*GPU represents results using multi-GPU which are scaled from the time of OMP-1GPU 363 with the assumption of rigorous scaling. 364

For CUDA parallel using all three kinds of GPU cards, further speedup is observed over the 365 upper limit of the CPU, i.e., MPI-25 here (Fig. 8b). Speedup relative to MPI-25 using single card 366 is 1-  $\sim$  2-fold for K80, 1-  $\sim$  3-fold for 1080 Ti, and about 3-fold for V100 while that relative to 367 OMP-1 is over 10-fold for all GPU cards with a maximum of 27.5 for V100 (Table 4). For OMP-368 4GPU, i.e., using 4 GPU cards with OpenMP, speedup achieves a maximum of 9.69-fold relative 369 to MPI-25 and 87.82-fold relative to OMP-1 (Table 4). In addition, speedup by multi-GPU is 370 higher with more particles. For example, it is 65.35-, 84.50-, 84.81-, and 87.82-fold speedup 371 relative to OMP-1 for ER Shrubs, ER Trees, LW Trees, and LW Shrubs, with increasing 372 particles number. 373

Difference between V100-Single and OMP-1GPU, also that between OMP-2GPU and Ideal-374 375 2GPU, and that between OMP-4GPU and Ideal-4GPU is the overhead of OpenMP (Fig. 8b and Table 3), which is minimal. Therefore, we can conclude that the current codes exhibit superior 376 scaling performance with increasing GPUs for multi-GPU with OpenMP (Table 4 and Fig. 8b). It 377 378 is also demonstrated better scaling if using more particles. For example, it is 3.69 and 2.01 (even beyond 2) for OMP-4GPU/OMP-1GPU and OMP-4GPU/OMP-2GPU for LW Shrubs which has 379 the maximum particles. Hence, for LW Shrubs using 16 GPUs which is the maximum extension 380 for multi-GPU with OpenMP (NVIDIA, 2019), the speedup is expected to be 351.82-fold (87.82×4) 381 382 if converted from OMP-1. However, due to the scaling performance of MPI discussed in section 4.1, speedup could achieve 969-fold  $(9.69 \times 25 \times 4)$  if converted from MPI-25. 383



384 ER\_Shrubs ER\_Trees LW\_Shrubs LW\_Trees
 385 Fig. 8. Computation performance for different parallels. (b) is re-scaled from (a) without OMP-1.

Table 3. Time used by particle loop for different APIs (Unit: hour)

	ER_Shrubs	<b>ER_Trees</b>	LW_Shrubs	LW_Trees
OMP-1	33.8099	70.2118	111.0170	84.5417
OMP-20	4.5739	9.7658	15.0169	11.5777
OMP-25	3.7404	7.9578	12.9045	9.2516
MPI-20	4.0381	8.3705	13.4108	10.2140
MPI-25	3.7808	7.7594	12.2478	9.3124
K80-Single	2.2570	5.8318	8.9997	7.2757
1080Ti-Single	1.4119	4.0957	6.7310	5.1485
V100-Single	1.2310	2.8758	4.5249	3.4911
OMP-1GPU	1.3056	2.9561	4.6690	3.5950
OMP-2GPU	0.7546	1.4914	2.5448	1.9685
Ideal-2GPU	0.6528	1.4781	2.3345	1.7975
OMP-4GPU	0.5174	0.8309	1.2642	0.9968
Ideal-4GPU	0.3264	0.7390	1.1673	0.8988

		1		1
	ER_Shrubs	ER_Trees	LW_Shrubs	LW_Trees
K80-Single/OMP-1	14.9800	12.0395	12.3356	11.6197
1080Ti-Single/OMP-1	23.9464	17.1428	16.4934	16.4206
V100-Single/OMP-1	27.4654	24.4147	24.5347	24.2163
OMP-1GPU/OMP-1	25.8961	23.7515	23.7775	23.5165
OMP-2GPU/OMP-1	44.8051	47.0778	43.6250	42.9473
OMP-4GPU/OMP-1	65.3458	84.5009	87.8160	84.8131
K80-Single/MPI-25	1.6751	1.3305	1.3609	1.2799
1080Ti-Single/MPI-25	2.6778	1.8945	1.8196	1.8088
V100-Single/MPI-25	3.0713	2.6982	2.7068	2.6675
OMP-1GPU/MPI-25	2.8958	2.6249	2.6232	2.5904
OMP-2GPU/MPI-25	5.0103	5.2028	4.8129	4.7307
OMP-4GPU/MPI-25	7.3073	9.3385	9.6882	9.3423
OMP-2GPU/OMP-1GPU	1.7302	1.9821	1.8347	1.8263
OMP-4GPU/OMP-1GPU	2.5234	3.5577	3.6932	3.6065
OMP-4GPU/OMP-2GPU	1.4584	1.7949	2.0130	1.9748

Table 4. Speedup for different parallel APIs (A/B: A relative to B)

390

## **4.3.** Toward large-scale simulations using multi-GPU

To test the capability of multi-GPU with OpenMP version of EcoSLIM at large scales, up to 392 2000 particles were added into the modeling domain per precipitation event. The maximum 393 permitted number of active particles is also increased from  $3 \times 10^7$  to  $3 \times 10^8$ . Numerical experiment 394 was conducted based on ER Shrubs with 4 Tesla V100. The maximum number of active particles 395 was achieved after about 2 years of simulation (Fig. 9a). The ratio of time used per timestep to the 396 number of active particles was plotted in Fig. 9b, which has a constant value of 0.023 in average. 397 We also tested that the maximum number of active particles on one V100 (16 GB) is about  $2 \times 10^8$ 398 due to the size of the memory. Hence, with the maximum extension of 16 V100 cards for OpenMP 399 (NVIDIA, 2019), the estimated computational time used for one-year simulation with hourly 400 resolution using 3.2 billion (0.2 billion×16) active particles would be only 44.77 hours. To our best 401 knowledge, this has been so far the largest application of Lagrangian particle tracking in subsurface 402 environment modeling, which shows a promising future of the EcoSLIM with multi-GPU. 403

For example, the particles number is 50,000 in Danesh-Yazdi et al. (2018), 80,000 in Jing et 404 al. (2019), 100,000 in Jing et al. (2020), 765,000 in Kollet and Maxwell (2008b), 1 million in 405 Engdahl and Maxwell (2015), and 4.14 million in Maxwell et al. (2016). In Wilusz et al. (2019) 406 and Weill et al. (2019), which are the latest applications using IHM and particle tracking on RTDs, 407 408 the number of particles injected into the modeling domain is 4,506,720 and 6.8 million in the whole simulation, respectively. More specifically, the number of active particles in this study (if using 409 16 V100 cards) is about 710-fold and 470-fold to those total injected in Wilusz et al. (2019) and 410 411 Weill et al. (2019), respectively. If we count the total number of particles injected in the modeling domain, it is  $1.11 \times 10^{12}$  for the specific simulation in Fig. 9, which is evidently a huge step forward 412 413 for improving the computational capability of EcoSLIM at exa-scales (e.g., continental).

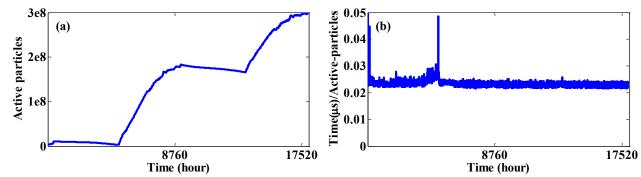


Fig. 9. Number of active particles (a) and the ratio of time used per timestep to number of active particles at that timestep. It is based on ER\_Shrubs with 2000 particles added into the modeling domain per precipitation event and a maximum permitted number of active particles of 3×10<sup>8</sup>. Test was conducted using 4 Tesla V100 with OpenMP.

419

## 420 5. Conclusions and perspectives

In this work, the EcoSLIM, a Lagrangian particle tracking program for calculating
travel/residence time distributions (TTDs/RTDs), is accelerated via both MPI- and CUDAparallelism. The latter is conducted through both single GPU and multi-GPU with OpenMP. The

MPI- and CUDA-parallel were verified by comparing the results to those from the original 424 OpenMP parallel, which shows excellent agreement. The computation capability of EcoSLIM is 425 drastically improved. With a maximum extension of 16 Tesla V100 cards and OpenMP, the wall 426 clock time used for one-year simulation at hourly timesteps with 3.2 billion active particles is 427 428 expected to be only 44.77 hours. The multi-GPU parallelism has demonstrated strong scalability especially using a very large number of particles. The accelerated EcoSLIM is expected to be a 429 powerful tool for large-scale and long-term simulations with high spatiotemporal resolutions, 430 431 which will advance our understanding of TTDs/RTDs under the evident climate change and intensified human disturbance. Although the current EcoSLIM is primarily developed to provide 432 an efficient tool for simulating TTDs/RTDs, it can be extended to large scale contaminant transport 433 434 modeling. In addition, the excellent performance of GPU parallel demonstrated in this study is expected to widely accelerate the next generation programs in subsurface environment modeling. 435 Finally, it has to be noted that the tests based on ideal cases in this study are still limited. We expect 436 437 more practical applications in the following work to further test and improve the computational capability of EcoSLIM. 438

# 439 **Computer code availability**

All codes for this work are publicly available on GitHub along with the Hillslope model used
for the simulations: <u>https://github.com/reedmaxwell/EcoSLIM</u>

# 442 Acknowledgements

This study was supported by the National Natural Science Foundation of China (NSFC-444 41807198), by the Strategic Priority Research Program of Chinese Academy of Sciences (Grant 445 No. XDA20100104), the Special Program for Applied Research on Super Computation of the 446 NSFC-Guangdong Joint Fund (the second phase), by the U.S. Department of Energy Office of

Science, Offices of Advanced Scientific Computing Research and Biological and Environmental 447 Sciences **IDEAS-Watersheds** project, by the U.S. National Science Foundation 448 CyberInfrastructure project, HydroFrame (NSF-OAC 1835903), and by the U.S. National Science 449 Foundation INFEWS-China (NSF-1805160). This study was also supported by the Center for 450 Computational Science and Engineering of Southern University of Science and Technology and 451 the National Supercomputer Center in Guangzhou, China. 452

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