# Accurate load balancing accelerates Lagrangian simulation of water ages on distributed, multi-GPU platforms

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November 22, 2022

#### Abstract

Water age is a fundamental descriptor of source, storage, and mixing of water parcels in a watershed. The Lagrangian, particle tracking, approach is a powerful tool for physically-based modeling of water age distributions, but its application has been hampered since it is computationally demanding. In this study, we present a parallel approach for particle tracking simulations. This approach uses multi-GPU with MPI parallelism based on domain decomposition. An inherent challenge of distributed parallelization of Lagrangian approaches is the disparity in computational work or load imbalance (LIB) among different processing elements (PEs). Here, load balancing (LB) schemes were proposed to dynamically balance the distribution of particles across PEs during runtime. In the followed hillslope simulations, LIB was observed in all LB-disabled runs, e.g., with a load ratio of 423.62% by using 2-GPU in LW\_Shrub case. LB schemes then accurately balanced the load distribution and improved the parallel scaling. Additionally, the parallel approach showed excellent overall speedup: a 60-fold improvement using 4-GPU relative to the serial run. A regional scale application further demonstrated the LB performance. The parallel time used by 8-GPU without LB was 31.33% reduced after LB was activated. When increasing 8-GPU with LB to 16-GPU with LB, it showed parallel scalability by reducing the parallel time of ~50%. This work shows how massively parallel computing can be applied to particle tracking in water age simulations. It also demonstrates the practical importance of load balancing in this context, which enables the large-scale simulations with an increased complexity of flow paths.

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2	ages on distributed, multi-GPU platforms
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<ol> <li>23</li> <li>24</li> <li>25</li> <li>26</li> <li>27</li> <li>28</li> <li>29</li> <li>30</li> <li>31</li> <li>32</li> <li>33</li> <li>34</li> <li>35</li> <li>36</li> </ol>	<ul> <li>Massively parallel computing of Lagrangian water-age simulations is realized</li> <li>Mechanisms of load imbalance are identified and LB schemes are proposed</li> <li>Parallel performance of the approach is demonstrated at the regional scale</li> </ul>

# 37 Abstract

38 Water age is a fundamental descriptor of source, storage, and mixing of water parcels in a 39 watershed. The Lagrangian, particle tracking, approach is a powerful tool for physically-based 40 modeling of water age distributions, but its application has been hampered since it is 41 computationally demanding. In this study, we present a parallel approach for particle tracking 42 simulations. This approach uses multi-GPU with MPI parallelism based on domain 43 decomposition. An inherent challenge of distributed parallelization of Lagrangian approaches is 44 the disparity in computational work or load imbalance (LIB) among different processing 45 elements (PEs). Here, load balancing (LB) schemes were proposed to dynamically balance the 46 distribution of particles across PEs during runtime. In the followed hillslope simulations, LIB 47 was observed in all LB-disabled runs, e.g., with a load ratio of 423.62% by using 2-GPU in 48 LW Shrub case. LB schemes then accurately balanced the load distribution and improved the 49 parallel scaling. Additionally, the parallel approach showed excellent overall speedup: a 60-fold 50 improvement using 4-GPU relative to the serial run. A regional scale application further 51 demonstrated the LB performance. The parallel time used by 8-GPU without LB was 31.33% 52 reduced after LB was activated. When increasing 8-GPU with LB to 16-GPU with LB, it showed 53 parallel scalability by reducing the parallel time of  $\sim 50\%$ . This work shows how massively 54 parallel computing can be applied to particle tracking in water age simulations. It also 55 demonstrates the practical importance of load balancing in this context, which enables the large-56 scale simulations with an increased complexity of flow paths.

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# 60 Keywords

61 Water age, Particle tracking, Multi-GPU with MPI, Domain decomposition, Load balancing

# 62 **1. Introduction**

63 Water age is an important metric that can unravel the journey that water-parcels and 64 pollutants take while traveling through a watershed (Botter, Bertuzzo, & Rinaldo, 2011). 65 Methods used to quantify water ages mainly include tracer data, lumped analytical models, and 66 distributed numerical models (Nicholas B. Engdahl & Maxwell, 2014). Tracer data are direct 67 observations of system behavior, however they have technical challenges such as tracer selection 68 and data interpretation (Sprenger et al., 2019). Moreover, the limited sampling cannot provide a 69 full view of the spatiotemporal variations of water ages (Nicholas B. Engdahl, McCallum, & 70 Massoudieh, 2016; McCallum, Engdahl, Ginn, & Cook, 2014). Analytical solutions can also 71 provide an understanding of the real system, although the limitations of this approach due to 72 simplifications have been well acknowledged by the community (Basu, Jindal, Schilling, Wolter, 73 & Takle, 2012). For distributed numerical models, the application of Eulerian framework is 74 hampered by the complications to solve the high-dimensional governing equation of water age 75 (Gomez & Wilson, 2013). Therefore, Lagrangian approach based on integrated hydrologic 76 modeling has become a promising tool to simulate transient age distributions (Nicholas B. 77 Engdahl & Maxwell, 2014; Jing et al., 2019; Wilusz, Harman, Ball, Maxwell, & Buda, 2019).

78 The Lagrangian approach is computationally demanding which limits widespread application 79 (Sprenger et al., 2019; Wilusz et al., 2019). Current studies of water ages using particle tracking 80 are limited to either catchments at small scales (Wilusz et al., 2019; J. Yang, Heidbüchel, 81 Musolff, Reinstorf, & Fleckenstein, 2018) or larger scales with a limited number of particles or 82 for steady-state conditions (Jing et al., 2020; Maxwell et al., 2016). Recently, as global water 83 security and climate change become increasing concerns, a growing number of studies have 84 proposed simulating water age at larger scales over long time-periods at high resolution 85 (Maxwell et al., 2016; McGuire et al., 2005; Starn, Kauffman, Carlson, Reddy, & Fienen, 2021). 86 Accomplishing these goals will significantly increase the computational burden of particle 87 tracking; massively parallel computing represents a promising solution.

Currently, there are few studies documenting parallelization of particle tracking approaches
for water age (Jing et al., 2020; Wilusz et al., 2019; J. Yang et al., 2018). Maxwell, Condon,
Danesh-Yazdi, and Bearup (2019) developed EcoSLIM, a particle tracking code, using OpenMP
(Open Multi-Processing) on CPU (Central Processing Unit). Yang and coauthors (C. Yang et al.,
2021), added CPU-based MPI (Message Passing Interface) and multi-GPU (Graphics Processing

93 Unit) with OpenMP into EcoSLIM. Ji, Luo, and Wang (2019) sped up MODPATH (Pollock, 2016) through multi-GPU with OpenMP/MPI based on domain decomposition (DDC). However, 94 95 their parallelized codes are limited to steady state simulations, and MODPATH is unable to 96 simulate evapotranspiration (ET) age and source water composition. N. B. Engdahl, Schmidt, 97 and Benson (2019) proposed two schemes for speeding up particle tracking simulations with 98 mass transfer to represent chemical reactions. One of the schemes also implemented MPI 99 parallelism through DDC. In DDC-based MPI parallelism, a typical problem encountered is load 100 imbalance among processing elements (MPI processes and/or GPUs), which can present a 101 challenge for good parallel efficiency. However, load imbalance has not been quantified in water 102 age simulations and its effects on parallel performance are unclear.

103 Load balancing (LB) schemes have been used to overcome these parallelization challenges 104 discussed above. Other disciplines using particle tracking, such as the molecular dynamics (MD) 105 and the smoothed particle hydrodynamics (SPH) (Boulmier, Raynaud, Abdennadher, & Chopard, 106 2019; Egorova, Dyachkov, Parshikov, & Zhakhovsky, 2019; Eibl & Rüde, 2019; Fattebert, 107 Richards, & Glosli, 2012; Furuichi & Nishiura, 2017; Kunaseth et al., 2013) have presented LB 108 schemes that greatly improved parallel simulation performance. However, LB has not been 109 applied to hydrologic modeling based on particle tracking; even in the studies using MPI through 110 DDC mentioned above. Additionally, when applying the particle tracking at larger scales in 111 long-term simulations, spatiotemporal variations of water age drivers in the real-world 112 applications can increase the heterogeneity of flow paths. This heterogeneity causes uneven 113 particle distributions and velocities within the domain, presenting new challenges to efficiency. 114 This will further complicate the distribution of particles across different subdomains and thus the 115 processing elements (which we are using here as a generic term for compute resources such as 116 CPU cores or a GPU). Furthermore, it becomes challenging to implement LB when considering 117 the increasing complexity of code structure of particle tracking due to the growing capabilities 118 such as simulating ET age and source-water composition at transient state in EcoSLIM.

In this study, we present new LB approaches implemented in the EcoSLIM code which is a particle tracking code simulating water age (ET, outflow, and groundwater) and source water mixing (initial subsurface water, rainfall, and snow). EcoSLIM is a grid-based approach which is different from the mesh-free particle tracking in other disciplines. EcoSLIM works seamlessly with ParFlow.CLM (Kollet & Maxwell, 2008a), which is an integrated hydrologic model simulating the coupled land-surface and subsurface water- and energy-processes at transient state.
ParFlow.CLM provides the temporally variant hydrodynamics and spatially variable subsurfaceproperties for EcoSLIM as input files, such as saturation, precipitation minus ET, threedimensional velocity fields, and IDs and porosities of the subsurface units.

128 Objectives of this study are twofold. Firstly, we use MPI to manage multi-GPU instead of 129 OpenMP in our previous work (C. Yang et al., 2021). This changes the target of the 130 decomposition from computational load to modeling domain. MPI parallelism removes the 131 barrier of a limited number of GPUs on a single computational node by created when using 132 OpenMP and potentially extends the particle tracking applications to massively parallel 133 computing. Secondly, three schemes with increasing physical representation are proposed to 134 dynamically balance the load among different MPI-processes/GPUs during runtime, which is 135 crucial for parallel efficiency. In following sections, the EcoSLIM code, the implementation of 136 multi-GPU with MPI, and the load balancing schemes are introduced in section 2. The setup of 137 test cases and platforms are followed in section 3. In section 4, validation of the new code is 138 verified and parallel performances of the code with/without the LB schemes are illustrated. 139 Specifically, application of the code for a 40-year simulation at regional scale in the North China 140 Plain was shown. Finally, contributions and implications of this work to hydrologic modeling 141 using Lagrangian approach are concluded in section 5.

# 142 **2. Methodology**

## 143 **2.1. EcoSLIM code**

EcoSLIM is originally implemented in Fortran and further accelerated by GPUs using CUDA (Compute Unified Device Architecture) Fortran (C. Yang et al., 2021). Its original structure is briefly introduced here to understand the following implementations of multi-GPU with MPI and LB schemes. For more details, please refer to our previous work (Maxwell et al., 2019; C. Yang et al., 2021). In each timestep of a transient simulation, key steps are as follows:

- 149 (1) New particles are added into grid-cells where precipitation minus ET (PME) is positive.
- 150 (2) Mass balance of precipitation and ET is calculated based on PME.
- (3) Advancing each active particle by a do-loop. Each particle either moves forward with anincrease of age or exits the modeling domain through outflow or ET.

(4) After the particle loop, inactive particles that have left the modeling domain via outflow
or ET are sorted out of the array of particles, the number of active particles is updated and
space at the end of this array is made available for new particles.

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(5) The time loop moves to next timestep.

157 In the particle loop, other attributes of each particle are also updated, such as the 158 saturated/unsaturated travel time, the saturated/unsaturated travel length, and the travel 159 time/length in some specified subsurface units. Statistics of outflow and ET of the whole 160 modeling domain, such as mass, mass weighted age, and source water composition, are included. 161 Additionally, gridding information is recorded, such as mass, cell-averaged age, and source 162 water composition for each grid-cell. Particle loop is the main computational load in EcoSLIM 163 which occupies more than 99% of the total simulation time when serially executing the code (C. 164 Yang et al., 2021). Therefore, our previous work (C. Yang et al., 2021) and also this study 165 focused on parallelizing the particle loop.

## 166 2.2. Multi-GPU with MPI

167 Instead of the load decomposition in our previous work (C. Yang et al., 2021), MPI 168 parallelism is applied here based on DDC. The modeling domain is split into P and Q parts in x 169 and y directions respectively, which generates a computing topology using  $P \times Q$  MPI processes. 170 The quantity of GPUs utilized in a simulation equals that of MPI processes. We use the method 171 in Ruetsch and Fatica (2014) to assign a unique GPU to each MPI rank. GPUs, MPI ranks, and 172 subdomains are all numbered from 0 to  $P \times Q$  - 1. Each GPU is responsible for a subdomain of 173 the same number. Source of particles on each GPU is from the assigned subdomain while 174 transport of these particles is in the whole modeling domain. For the particle loop, we adopt the 175 same GPU kernel in our previous work (C. Yang et al., 2021) with a few modifications for 176 tracking particles in specified subsurface units. After the mapping to subdomains, each MPI-177 process/GPU works almost independently with a limited MPI collective communications. They 178 are the calculation of mass balance mentioned in item (2) in section 2.1 and the ET/outflow 179 statistics of the whole modeling domain also mentioned in section 2.1, which are performed 180 before and after the execution of the kernel respectively.

181 The transport of particles in the whole domain, instead of in the subdomain where they were 182 added, avoids the MPI communications of exchanging particles between subdomains when 183 particles move out of a subdomain. The disadvantage of such a design is the redundant copies of 184 the global information (e.g., velocities, porosities, saturations, and PME) for all MPI processes. 185 This is CPU-memory expensive. However, for present clusters which are commonly equipped 186 with 2/4/8 GPUs per node, the 2/4/8 copies of necessary information on one node are not a 187 bottleneck for regional modeling with scales of Tran, Zhang, Cohard, Condon, and Maxwell 188 (2020) and C. Yang et al. (2020). The multi-GPU with MPI was also conducted in our previous 189 work based on load decomposition (C. Yang et al., 2021). However, in each timestep, the 190 overhead of distributing load from rank 0 to others by MPI communication is over the speedup 191 by extending single GPU to multi-GPU. Though the DDC in this study avoids such a problem, 192 the load imbalance mentioned in section 1 becomes a new issue. Therefore, three schemes with 193 increasing physical representation are proposed to balance the load among GPUs/MPI-processes 194 during runtime in next section.

## 195 **2.3. Schemes of load balancing**

## 196 **2.3.1** Direct transfer (S1)

197 The movement of particles in current EcoSLIM are independent, so the loads on GPUs can 198 be redistributed by directly transferring particles between MPI processes with a user specified 199 frequency. It is implemented by communications on CPU using the MPI functions of MPI\_Send 200 and MPI\_Recv after the sort of particles. At a given time, the number of active particles on each 201 MPI process is gathered. Thus, the old numbers of the starting and ending particles (*np lo* and 202 np\_ro) on each process ranked in a global queue are obtained. Then the starting and ending 203 numbers are updated to new ones (np\_ln and np\_rn) based on an even division of the global 204 queue. Then the transfer is accomplished on each MPI process by the following four steps: (1) 205 sending particles to the upstream process if  $np\_lo$  is smaller than  $np\_ln$ , (2) receiving particles 206 from the downstream process if  $np \ ro$  is smaller than  $np \ rn$ , (3) sending particles to the 207 downstream process if *np\_ro* is larger than *np\_rn*, and (4) receiving particles from the upstream 208 process if *np\_lo* is larger than *np\_ln*. The number of active particles is updated after each transfer. 209 The overhead of this scheme is determined by the quantity of particles transferred and the 210 bandwidth of the computing platform.

# 211 2.3.2 Cyclic mapping (S2)

In this scheme, the DDC is static which is determined after initialization of the simulation. In a simulation using *n* GPUs/MPI-processes, the mapping between subdomains and GPUs (both numbered from 0 to n-1) is continuously shifted with a user specified frequency. For instance, at a given time  $t_1$ , the mapping is shifted from 'the *m*th subdomain  $\rightarrow$  the *m*th GPU' to 'the (m+1)th subdomain  $\rightarrow$  the *m*th GPU' where *m* ranges from 0 to *n*-1. If *m*+1 is larger than *n*-1, the subdomain numbered with the reminder of *m*+1 and *n* will be mapped to the *m*th GPU. Table 1 showed the cyclic mapping between subdomains and GPUs in a simulation using four MPI processes. Using this scheme, each GPU traverses the loads of all subdomains periodically, and thus the load distribution is dynamically balanced among GPUs. The overhead of shifting the mapping is almost negligible.

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 Table 1. Cyclic mapping in a simulation using four GPUs/MPI-processes

$t_0$	Subdomain number	0	1	2	3
(Initial)	GPU number	0	1	2	3
4	Subdomain number	1	2	3	0
$l_1$	GPU number	0	1	2	3
4	Subdomain number	2	3	0	1
$l_2$	GPU number	0	1	2	3

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### 224 2.3.3 Dynamic DDC (S3)

225 In the initialization of a simulation, particles are evenly distributed in space, so we 226 decompose the modeling domain into subdomains of an equal size. PME is spatiotemporally 227 variable, so the number of particles added into each subdomain is different at a given timestep 228 and such a difference varies with time. More importantly, particles added into different 229 subdomains have different exits from the whole modeling domain. Both source and exit are 230 responsible for heterogeneity of the flow paths. A subdomain of more source particles and longer flow paths imposes heavier load on its corresponding GPU. As a result, after the initial even-231 232 decomposition, we dynamically update the decomposition during runtime based on flow paths of 233 particles. At a given timestep, the initial location of an active particle is identified and the 234 corresponding grid-cell in the top layer get one score. After traversing all the active particles, we 235 get the accumulated scores for each grid-cell in the top layer. This was implemented through 236 atomic operations on a two-dimensional matrix in the GPU kernel mentioned in section 2.2. 237 Then we conduct DDC based on this weight matrix. The frequency of such a dynamic DDC can 238 be specified by users.

The orthogonal recursive bisection (ORB) method is used for DDC, which is popular in MD and SPH (Egorova et al., 2019; Fattebert et al., 2012). The domain or each subdomain is divided into two in a direction at one time. By switching the direction, the whole domain is recursively 242 divided into the scheduled number of subdomains  $(P \times Q)$ . DDC is only implemented in x and y 243 directions in this study. It starts in a direction which will be divided into more pieces. For 244 example, if Q is larger than P, DDC will start in y direction, otherwise, it starts in x direction. 245 Two algorithms are provided in the code to determine the dividing line. One calculates the 246 accumulated particles of columns (rows) in x (y) direction. Once the number of particles of n-1247 columns (rows) is less than half the total particles in this subdomain while that of n columns 248 (rows) is more than half the total particles in this subdomain, the dividing line is found as column 249 (row) n. The other algorithm to find the dividing line with higher efficiency is the typical 250 dichotomizing search.

# **3. Test setup**

## **3.1. Hillslope model**

253 Tests were conducted based on a hillslope model (Maxwell et al., 2019; C. Yang et al., 2021). 254 The modeling domain has the length of 100-, 1-, and 9.4-m in x, y, and z directions, respectively. 255 It was divided into 20 columns, 5 rows, and 20 layers with constant resolutions in x and y 256 directions. In vertical direction, the layer-thickness was variable: 0.5 m for the bottom 18 layers 257 while 0.3- and 0.1-m for the top 2 layers. Soil has the homogeneous properties: saturated hydraulic conductivity of 0.05 m/h, Manning's N of 10<sup>-6</sup> m<sup>1/3</sup>h<sup>-1</sup>, porosity of 0.2, and van 258 Genuchten parameters with  $\alpha$  of 1.0 m<sup>-1</sup> and exponent *n* of 2.0. Two real meteorological-forcings 259 260 were used to drive ParFlow.CLM, representing a high elevation, snow dominated mountain 261 headwaters (ER) and a semiarid, rain-dominated plains system (LW). Two homogeneous land-262 cover types were used which are the Shrub plant functional type (Shrub) and the Evergreen 263 Needleleaf plant functional type (Trees). Thus, four cases were tested with a combination of the 264 meteorological forcings and the land-cover types, which were named as: ER\_Shrub, ER\_Trees, 265 LW\_Shrub, and LW\_Trees. For simulations of both ParFlow.CLM and EcoSLIM, no flux 266 boundaries were adopted except the land surface which was open for precipitation, outflow and 267 ET. For each case, ParFlow.CLM simulation of 5 years was conducted using hourly timestep. 268 One-year forcing data were repeatedly used in the whole simulation. Dynamic equilibrium of the 269 flow field was approached at the end of simulation. Hence, the transient flow field of the last 270 year in ParFlow.CLM simulation was repeatedly used in EcoSLIM simulation of 20 years with 271 hourly timestep. At the end of each simulation, EcoSLIM system achieved the dynamic 272 equilibrium.

273 By injecting 2 particles into the modeling domain per precipitation event, the average 274 particle-numbers in the last year of the simulations for four cases are 0.39-, 0.83-, 1.30-, and 275 1.03-million, respectively. Such quantities of particles are comparable to those in most of the 276 previous studies (Danesh-Yazdi, Klaus, Condon, & Maxwell, 2018; Nicholas B. Engdahl & 277 Maxwell, 2015; Jing et al., 2019; Jing et al., 2020; Kollet & Maxwell, 2008b; Maxwell et al., 278 2016; Weill, Lesparre, Jeannot, & Delay, 2019; Wilusz et al., 2019). It has a maximum of 6.8 279 million in Weill et al. (2019) to the best of our knowledge. In fact, the particle-number in most 280 previous studies is the total injected particles while that during runtime is a fewer quantity. 281 However, the number in this study is the active particles in the modeling domain and the 282 particles out of the domain through ET and outflow are not included. Thus, the particle-283 quantities are much more than those in previous studies.

#### 284 **3.2. Test platform**

285 Tests were conducted on the Casper cluster in the Computational and Information Systems 286 Laboratory at the National Center for Atmospheric Research. The computational node used for the following simulations is equipped with 2.3-GHz Intel<sup>®</sup> Xeon<sup>®</sup> Gold 6140 processors and 287 288 NVIDIA Tesla V100 32GB SXM2 GPUs with NVLink. The compiler is NVIDIA HPC SDK of 289 version 20.11, the MPI is implemented using Open MPI of version 4.0.5, the GPU driver version 290 is 450.51.06, and the CUDA version is 11.0.3. Tests were also repeated on a personal workstation (WS). The WS is equipped with 2.00-GHz Intel<sup>®</sup> Xeon<sup>®</sup> E5-2683 v3 processors 291 292 together with four GPUs of 12 GB GeForce GTX 1080 Ti. Other necessary setups of the WS environment are NVIDIA HPC SDK 20.11, Open MPI 3.1.5, GPU driver 440.118.02, and 293 294 CUDA 10.2.

# **4. Results and discussion**

## 296 **4.1. Code-to-code verification**

To verify the availability of the new code, simulation results using the new code were compared to those of the original OpenMP version (Maxwell et al., 2019). Comparisons were performed for all four cases while that of the ER\_Shrub case was shown in Figures 1 (outflow) and 2 (ET). Results of other test cases had performances as good as that of ER\_Shrub. Tests in this study were conducted using one, two, and four GPUs successively while the results using four GPUs were illustrated in Figures 1 and 2. Subplots in Figures 1 and 2 were for results without LB and with each LB scheme. The water-age and -mass for both outflow and ET 304 simulated by the new code well fitted those generated by the original code. The deviations 305 between them were attributed to the generation of pseudo-random numbers (PRNs). Though the 306 ensembles of the PRNs were statistically the same for each run, the PRN for a specific particle 307 probably changed due to the invoking sequence of the generation-function which was dependent 308 on the parallelism, i.e., the OpenMP or the multi-GPU with MPI. For the same parallelism, if 309 different numbers of CPU-threads/GPUs were used, there were also such deviations during our 310 tests. The fitness of outflow was better than that of ET because ET in EcoSLIM were directly 311 dependent on PRNs.



312Time (day)Time (day)313Figure 1. Comparisons for age and mass of outflow based on ER\_Shrub case between the original

- 314 **EcoSLIM code and that parallelized in this study.**
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316Time (day)Time (day)317Figure 2. Comparisons for age and mass of ET based on ER\_shrub case between the original318EcoSLIM code and that parallelized in this study.

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320 4.2. Parallel performance





322 Figure 3. Wall-clock time consumption of each test. *n*-GPU represents the number of GPUs used in

simulations. Sn represents different LB schemes. WS indicates tests conducted on workstation
 while others on Casper. 720 indicates S3 worked every 720-hour while others worked every 8760-

- 324 while 325 hour.
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- 327





Figure 4. Load distribution for LW Shrub by using 2-GPU. Sn represents LB schemes. 720h represents that S3 worked every 720-hour while others were 8760-hour.



333 Figure 5. Load distribution for LW\_Shrub by using 4-GPU. Sn represents LB schemes. 720h represents that S3 worked every 720-hour while others were 8760-hour.

337 Figure 3 shows wall-clock time consumption of each test on both Casper and WS. Load 338 distributions for tests using 2- and 4-GPU were shown in Figures 4 and 5, respectively. Tests 339 using 2-GPU (P = 2 and Q = 1) were performed by evenly dividing the domain in x direction while those using 4-GPU (P = 2 and Q = 2) had an additional division in y direction. All LB 340 341 schemes worked every 8760-hour except that in Figures 4d and 5d were 720-hour for S3. 342 Though only LW\_Shrub was taken as an example in Figures 4 and 5, other cases had similar 343 performances. Time used for four cases by one CPU-thread based on the original code were 344 tested on WS, which were 36.41-, 76.34-, 121.56-, and 92.40-hour for ER Shrub, ER Trees, 345 LW\_Shrub, and LW\_Trees, respectively. Speedup of each case was then calculated and listed in 346 Table 2.

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Table 2. Speedup of each test relative to the serial run using one CPU-thread

	Case name	Speedup								
Platform		Without LB		Scheme 1		Scheme 2		Scheme 3		
		1-GPU	2-GPU	4-GPU	2-GPU	4-GPU	2-GPU	4-GPU	2-GPU	4-GPU
	ER_Shrub	20.5403	20.5693	33.5134	30.5237	54.8938	31.0469	52.7465	30.6935	46.3058
Common	ER_Trees	21.1316	20.1687	22.3125	23.5297	57.5619	23.5857	58.7581	24.8849	48.6518
Casper	LW_Shrub	21.1858	20.3505	34.5421	23.7138	51.5341	26.3410	47.6590	26.4183	41.5489
	LW_Trees	15.9127	18.8541	32.9764	25.3874	56.0544	24.1910	54.8824	26.9184	46.2856
	ER_Shrub	13.9637	17.2900	36.0809	34.4456	48.8535	34.4522	54.4261	33.4696	49.0443
WS	<b>ER_Trees</b>	13.8955	11.5971	26.0117	25.0581	47.8438	25.1381	60.4948	24.5480	51.0523
VV 0	LW_Shrub	13.5910	17.5129	27.0710	24.7589	37.1392	24.9531	43.0452	23.8150	37.1698
	LW_Trees	13.6440	17.7556	26.1808	24.6492	45.1304	24.8835	57.2562	24.7449	40.4952

348 In tests without LB, time used by 2-GPU was even more than that by 1-GPU for ER Trees 349 and LW\_Shrub on Casper (Figure 3). This performance degradation has two reasons. Firstly, 350 severe load imbalance can be observed in Figure 4 when using 2-GPU with a particle-number 351 ratio of 423.62%. The larger one (yellow lines in Figure 4) which determined the parallel 352 efficiency almost achieved the total load. It confirmed that load imbalance also exists in 353 Lagrangian hydrologic modeling and decreases the parallel performance. Secondly, collective 354 MPI communications mentioned in section 2.2 introduced overhead when increasing the GPU 355 number from one to two. When using 4-GPU without LB, the maximum load largely decreased 356 (Figure 5) due to the additional decomposition in y direction. The hillslope model is quasi-three-357 dimensional with a x slope of 0.1 and a y slope of 0. Hence, the movement of particles in y 358 direction can be neglected which formed the parallel flow paths along x direction. Along x359 direction, particles added upstream had much longer flow paths than those added downstream. 360 As a result, the division in y direction was much more effective than that in x direction to

improve the parallel performance. However, load imbalance was still significant (Figure 5) and
 parallel scalability was not shown by increasing 2-GPU to 4-GPU (Figure 3).

363 When S1 was activated, time used by 2- and 4-GPU were dramatically decreased relative to 364 those without LB. With S1, time used by 4-GPU was even less than half of that used by 2-GPU 365 on Casper (Figure 3). The overhead of particle transfer was small for all four cases, which was 366 less than 3 seconds in each 20-year simulation. S2 had performance as good as S1 (Figure 3). It 367 was mentioned in section 2.3.2 that the overhead of S2 was small enough to be neglected. For S3, 368 more time was used by 4-GPU when compared to that of S1 and S2 (Figure 3). In Figures 4c and 369 5c, the load distribution was not well balanced relative to that of S1 and S2. The flow paths of 370 particles were transient, so the weight matrix at a moment cannot effectively balance the load for 371 a long-period of simulation (i.e., 8760-hour). When S3 was activated with a higher frequency of 372 every 720-hour, the improved load balance can be observed in Figures 4d and 5d and the further 373 speedup was indicated in Figure 3. However, difference of the loads between GPUs 1-2 and 3-4 374 was still observed in Figure 5d. This is due to the model dimension which is five grid-cells in y 375 direction. Hence it cannot be evenly divided by ORB introduced in section 2.3.3.

376 S3, a physically-based scheme, not only aims to balance the load but also to understand the 377 load imbalance in Lagrangian hydrological modeling. To build S3, we also tried DDC based on 378 the source of particles (i.e., PME). The score/weight of a grid-cell is determined by the 379 accumulated particle-number added into it in a period. However, it didn't show good 380 performance. Current implementation actually integrates the effects of both the quantity of 381 source particles and the flow-path lengths. This trial and error indicates that the flow-path 382 lengths instead of the quantity of added particles dominate the load distribution. This has 383 important implications to efficiently build other physically-based LB schemes. Generally, with 384 LB, the new code showed excellent parallel performance in the tests on both Casper and WS 385 (Table 2). The speedup by one GPU is ~13-fold on WS with 1080 Ti while ~21-fold on Casper 386 with Tesla V100. The speedup by 4-GPU is over 50-fold on both Casper and WS and has a 387 maximum over 60-fold. With LB schemes, the code showed parallel scalability from 2-GPU to 388 4-GPU.

## 389 **4.3.** Application in the North China Plain

We applied the new parallel code on a North China Plain (NCP) domain to demonstrate its capacity for large-scale simulations. To the best of our knowledge, there have been no previous 392 studies based on particle tracking at such a regional scale for water ages of ET, groundwater 393 (GW), and outflow in a unified framework. The NCP ParFlow.CLM model was adopted from C. 394 Yang et al. (2020) with a few modifications. The model has 509 and 921 grid-cells in x and y 395 directions respectively while it is discretized into five layers in vertical direction. The horizontal 396 resolution is 1 km while the layer thickness from bottom to top is 100-, 1-, 0.6-, 0.3-, and 0.1-m. 397 Thus, the NCP model has a dimension of 509 km  $\times$  921 km  $\times$  102 m in total. We conducted an 398 EcoSLIM simulation of 40 years on Casper with hourly timestep, in which the hourly outputs of 399 one-year simulation from ParFlow.CLM were repeatedly used.

400 The EcoSLIM simulation was started using 8-GPU (P = 2 and Q = 4) without LB 401 (abbreviated as R1 hereafter). From the 155,928th hour, S1 was activated every 240-hour (R2) 402 while R1 was continued for the following 7.8 years. At the 247,032th hour, the load of R2 was 403 evenly divided into 16 portions and a new run (R3) was started using 16-GPU (P = 4 and Q = 4) 404 with S2 activated every 240-hour. The overlap between R2 and R3 is 3.4 years. We also tried 16-405 GPU without LB for the first five years of the simulation (R4). Figure 6 showed the active-406 particle-number and the wall-clock time consumption of each timestep during the latter 22 years 407 of the simulation (the 155,929th to the 350,400th hour). R1, R2, and R3 were indicated by green, 408 blue, and red in Figure 6 respectively. The active-particle-number is around 200 million during 409 this simulation time-interval (Figure 6a). The discrepancy of the particle number between 410 different runs in the overlaps are due to the generation of random numbers discussed in section 411 4.1.





412 413 Figure 6. Computational load (a) and wall-clock time consumption (b) for the EcoSLIM simulation 414 in the North China Plain. The time interval is from the 155929th to the 350400th hour in the 40-415 year simulation.

416 For the overlap between R1 and R2, parallel time of the particle loop was 98.390- and 417 65.987-hour for R1 and R2 respectively (Figure 6b). The overhead of S1 for transferring data 418 was 1.573-hour. S1 (overhead included) decreased 31.33% of the time used by R1, which demonstrated the high efficiency of S1. Figures 7a and 7b showed the well-balanced load by S1. 419 420 For the overlap between R2 and R3, the parallel time was 43.311- and 21.844-hour for R2 and 421 R3 respectively (Figure 6b). Though it showed 50% decrease of the parallel time, the obvious 422 jitters of the time in R2 has to be considered. Based on their baselines, the time used by R3 was a 423 little longer than half the time used by R2. This should be due to the better load balancing effect 424 of S1 than that of S2, which was shown in Figures 7b and 7d. However, when comparing the 425 load distribution between R3 and R4 for a time interval of the same length (4.94-year), the load 426 balancing effect of S2 was significant. The difference between the maximum- and minimum-427 load at the end of the comparing time-interval in R4 was 6.66 million (Figure 7d) while that in 428 R3 was 3.52 million (Figure 7c), which was 47.21% decrease of the load variance. Additionally, 429 based on the increasing trend in Figures 7c and 7d, the load variance in R3 with S2 gradually 430 achieved a steady state while that in R4 continued increasing.



431Timestep (×10<sup>3</sup>)Timestep (×10<sup>3</sup>)432Figure 7. Load distributions in the application in the North China Plain. Load distribution on 8-433GPU before and after using S1 in R2 (a), on 8-GPU with S1 in R2 (b), on 16-GPU without LB in R4434(c), and on 16-GPU with S2 in R3. (b) was magnified from (a).

435

# 436 **5. Conclusions**

437 Water age can reveal the source, storage, and mixing of water parcels in a watershed. Though 438 data- and model-driven methods have significantly advanced our understanding of water ages, 439 the quantification of water ages is still technically challenging. Lagrangian particle tracking is an 440 invaluable tool for physically-based transient modeling of water ages, but it is computationally 441 expensive. When considering climate change and global water security, it is essential to conduct 442 simulations of water ages at large scale with high resolution, which makes the implementation of 443 massively parallel computing in particle tracking for this purpose pressing. Though parallel 444 computing is widely implemented for Eulerian hydrological modeling, applications to 445 Lagrangian based simulations are developing. This is likely due to the inherent difficulties such 446 as load imbalance across computational resources which will become more challenging when 447 modeling a real hydrologic system with high spatiotemporal variability.

In this study, multi-GPU with MPI parallelism based on domain decomposition (DDC) was
 implemented in the Lagrangian, particle tracking code EcoSLIM, to accelerate simulations of

450 water age and source-water mixing. Three load balancing (LB) schemes with increasing physical 451 representation (i.e., direct transfer, cyclic mapping, and dynamic DDC) were built to 452 dynamically balance the quantity of particles across GPUs during runtime. With LB, the code 453 showed excellent parallel performance in the hillslope simulations on two different platforms, 454 e.g., a maximum of 60-fold speedup on 4-GPUs and the parallel scalability from 2-GPU to 4-455 GPU that is almost ideal. A 40-year simulation conducted in the North China Plain further 456 demonstrated the high parallel efficiency of LB for a large-scale application. Using 8-GPU with 457 LB, it reduced 31.33% of the parallel time using 8-GPU without LB. When increasing 8-GPU 458 with LB to 16-GPU with LB, ~50% reduction of the parallel time demonstrated the parallel 459 scalability.

460 More importantly, results confirmed the load imbalance in Lagrangian hydrologic modeling. 461 In LW\_Shrub case using 2-GPU, the particle-number ratio achieved 423.62%, which severely degraded the parallel performance without LB. For LB schemes, physically-based dynamic DDC 462 463 performed as well as other schemes in hillslope simulations. Trial and error of building this 464 scheme identified that the distribution of flow-path lengths in the domain instead of the quantity 465 of particles added into the domain dominates the load distribution. This illustrated both the 466 mechanisms of load imbalance and the directions to build efficient physically-based LB schemes 467 in this context. This study realized the massively parallel computing of particle tracking in water 468 age simulations which is lacking in hydrologic modeling. It also demonstrated that LB have 469 practical importance enabling its applications at large scales with increased heterogeneity of flow 470 paths. The LB schemes can be borrowed to other hydrologic models using Lagrangian approach 471 and the parallelized EcoSLIM is a promising tool to accelerate the scientific progress of water 472 age studies.

473

# 474 Acknowledgements

This work was supported by the U.S. Department of Energy Office of Science, Offices of Advanced Scientific Computing Research and Biological and Environmental Sciences IDEAS project and Watershed Function Scientific Focus Area under Award Number DE-AC02-05CH11231. The authors acknowledge high-performance computing support from Cheyenne (doi:10.5065/D6RX99HX) provided by NCAR's Computational and Information Systems Laboratory, sponsored by the National Science Foundation. Fortran/CUDA-Fortran code

- 481 (https://github.com/aureliayang/EcoSLIM/tree/multi-GPU) which can reproduce the results is
- 482 located on Github. Once the manuscript is accepted, this repository will be archived in Zenodo to
- 483 get a DOI for citation purpose.
- 484

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