SC13 Short lecture @ AICS

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Title of your	Supporting Workflow Management of Scientific Applications by MapReduce
Presentation	Programming Model
Abstract	There are many applications that run many tasks as a workflow, such as
	ensemble simulation and data processing, in scientific applications. We
	surveyed various applications that run on large scale parallel systems,
	extracted their workflow patterns and then investigated requirements for
	building the workflow on the MapReduce programming model. To fulfill the
	requirements, we implemented them to 'K MapReduce' MapReduce
	implementation. To illustrate the scientific applications' workflows on the
	MapReduce model, we implemented two application workflows on the K
	MapReduce; Replica exchange molecular dynamics and Genome analysis.
	As a result of comparison with implementations without MapReduce, we
	confirmed a performance benefit in the former case and a coding benefit in
	the latter case.