

SEQUE

parallel R
in the cloud
two lines of code

no kidding!

SEQUEL



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close

SEGUE

syntax...

```
require(segue)
```

```
myCluster <- createCluster()
```

contratulations. we've built a hadoop
cluster!

more syntax...

parallel apply() on lists:

base R:

`lapply(X, FUN, ...)`

segue:

`emrlapply(myCluster, X, FUN, ...)`

SE
GUE
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howzit work?

emrapply()

SEGUE

list is serialized to CSV and uploaded to S3 – streaming input file

function, arguments, r objects, etc are saved & uploaded

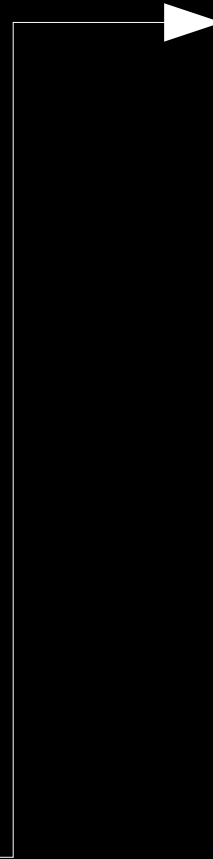
EMR copies files to nodes – mapper.R picks them up

CSV is input to mapper.R applies function to each list element

output is serialized into emr part-xxxxx files on s3

part files are downloaded to R and deserialized

deserialized results are reordered and put into a list object



when to use segue...

SEGUE

embarrassingly parallel

cpu bound

apply on lists with many items

object size: to / from s3 roundtrip

SEQUE

segue project page

<http://code.google.com/p/segue/>

google groups

<http://groups.google.com/group/segue-r>