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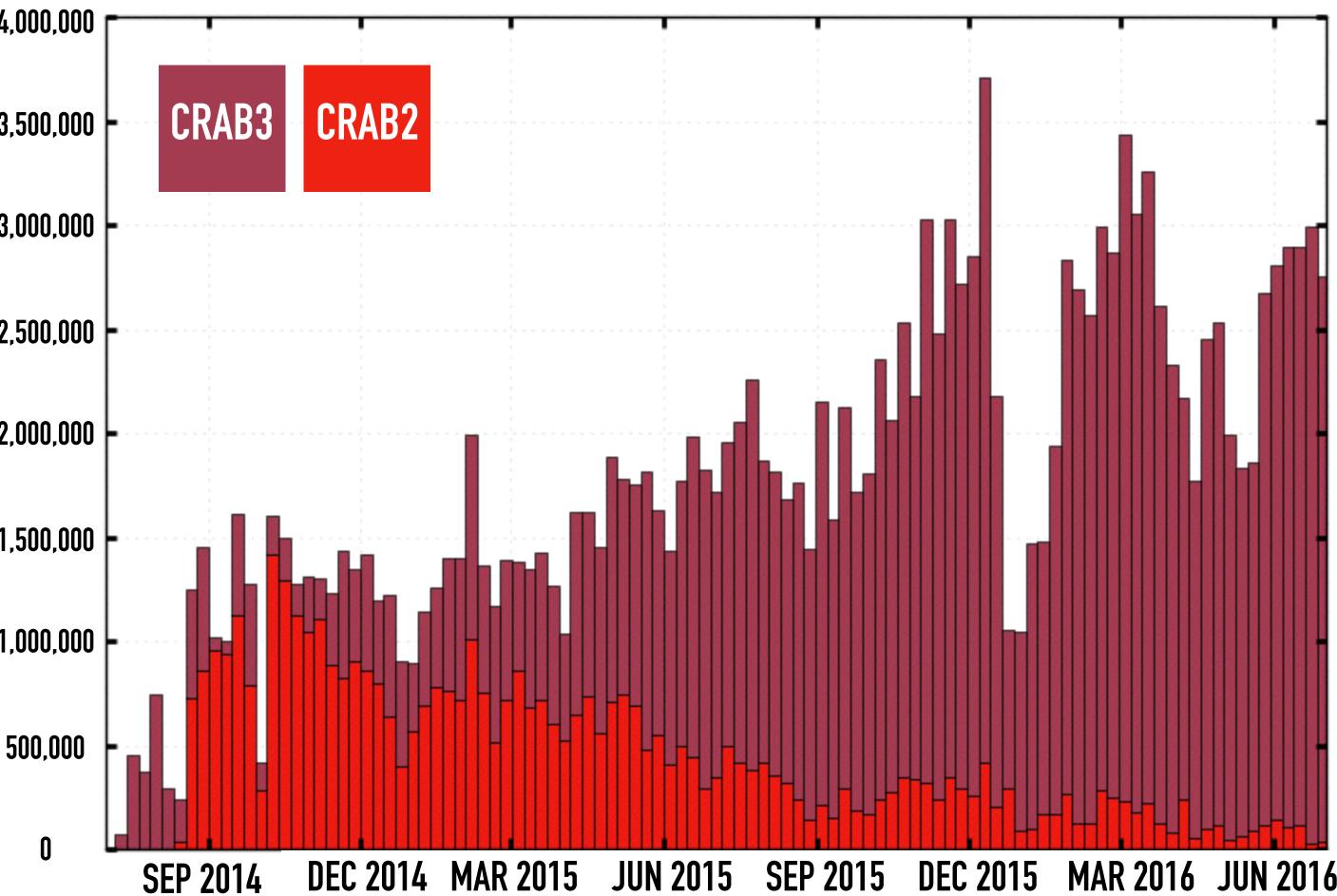
# USING DAGMAN IN CRAB3 TO IMPROVE TASK SPLITTING FOR CMS USERS

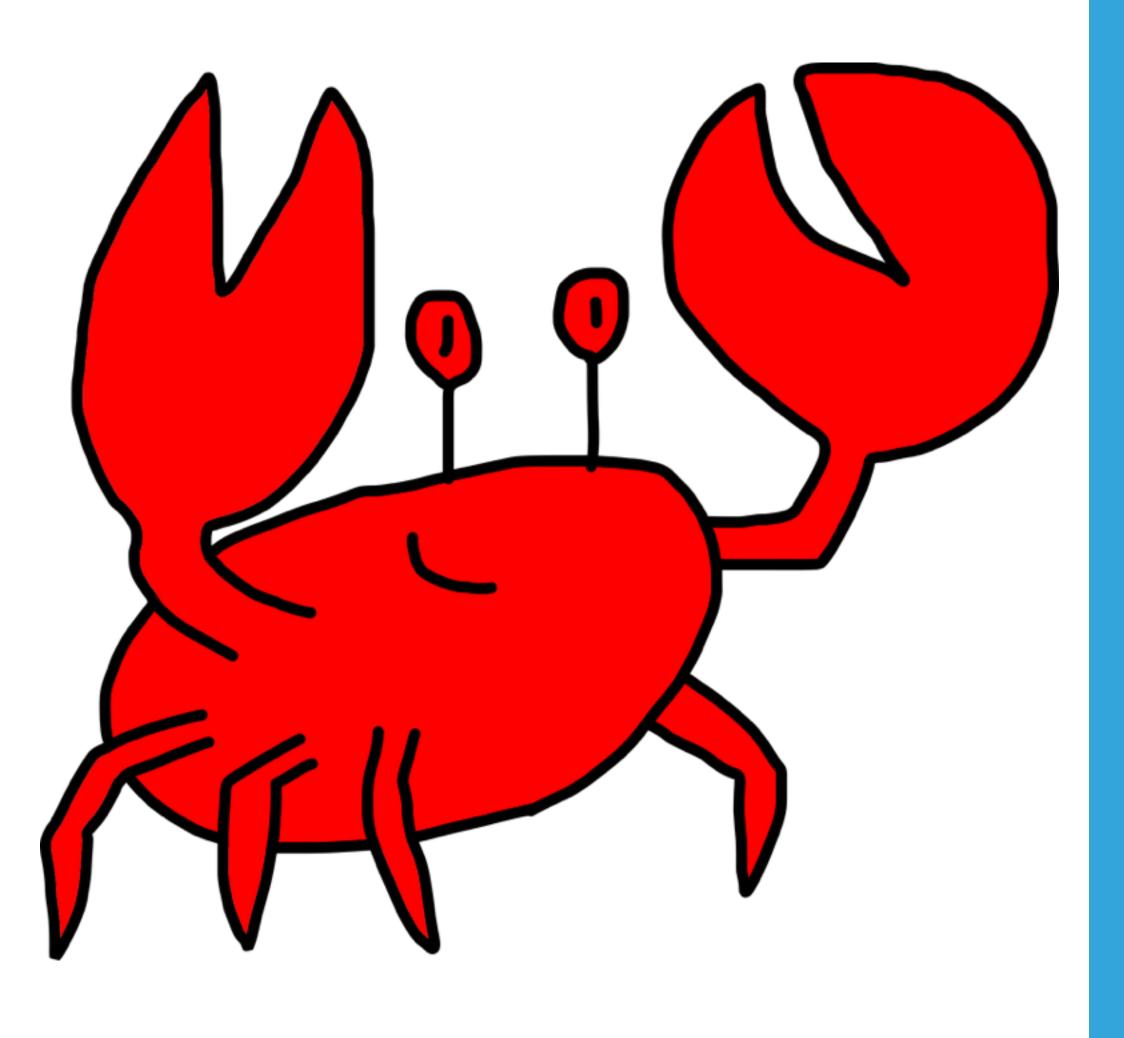


## WHAT IS CRAB3?

- Hundreds of physicists regularly submit analysis jobs to the Grid using a tool called CRAB3
- Architecture: lightweight user client and CRAB3 server which accepts user requests ("tasks")
- CRAB3 manages ~3 million jobs per week
- 4,000,000 3,500,000 3,000,000 2,500,000 2,000,000 1,500,000 1,000,000

### **COMPLETED JOBS**





# CONVENTIONAL

## SPECIFICATION OF A CRAB3 ANALYSIS TASK

• User responsible for specifying:

- Dataset (group of files) to process
- Code to run
- should process
  - Units = files, lumi sections, or events

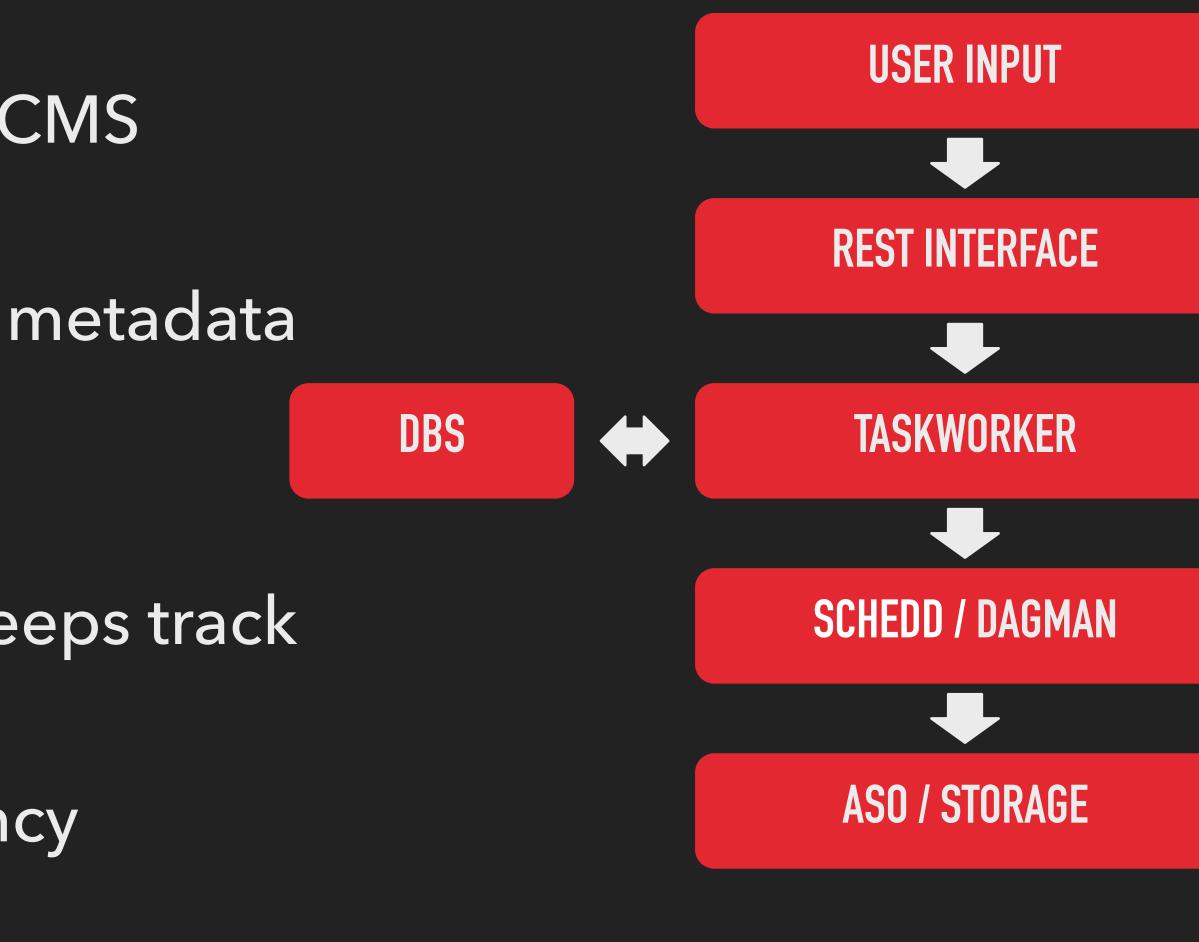
### • "Splitting parameters", i.e., how many units each job

## **CRAB3 SERVER'S HANDLING OF THE TASK**

Server initializes task and:

- Pulls dataset metadata in from the CMS dataset bookkeeping catalog (DBS)
- Splits input dataset into jobs using metadata + splitting parameters
- Creates a single-depth DAG:
  - DAG=Directed Acyclic Graph, keeps track of dependencies between jobs
  - Single depth=no interdependency between jobs

• DAG submitted to scheduler for processing







### SPLITING IS HARD FOR USERS

UHM.

### WHY DID YOU SUBMIT 10,000 FIVE MINUTE JOBS?!?









### HOW LONG DOES YOUR CODE TAKE TO RUN **PER EVENT? HOW BIG IS YOUR DATASET?**

ANNA WOODARD

**CRAB3 OPERATOR** 



## SPLITTING IS HARD FOR USERS...

- Difficult for users to provide optimal task splitting parameters • Code runtime may differ for each:
  - - Code iteration
    - Dataset (different event complexity)
- Tendency for user to choose parameters which create many short jobs
  - Hard limit for jobs per task is quite high (10,000 jobs)
  - Users can get away with memory leaks with short jobs

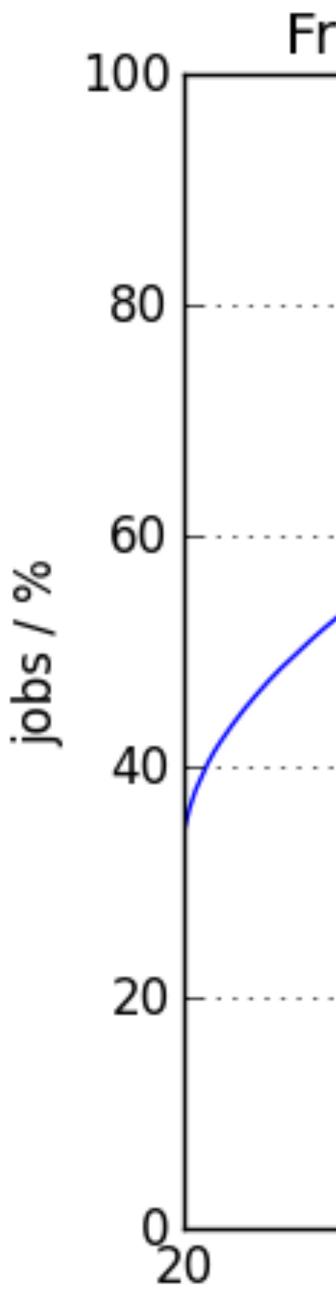
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## **TOO MANY SHORT** JOBS

- Almost 50% of the jobs have a runtime of ~40 minutes!
- Large number of small jobs causes excessive load on the schedulers and other central components.

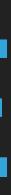


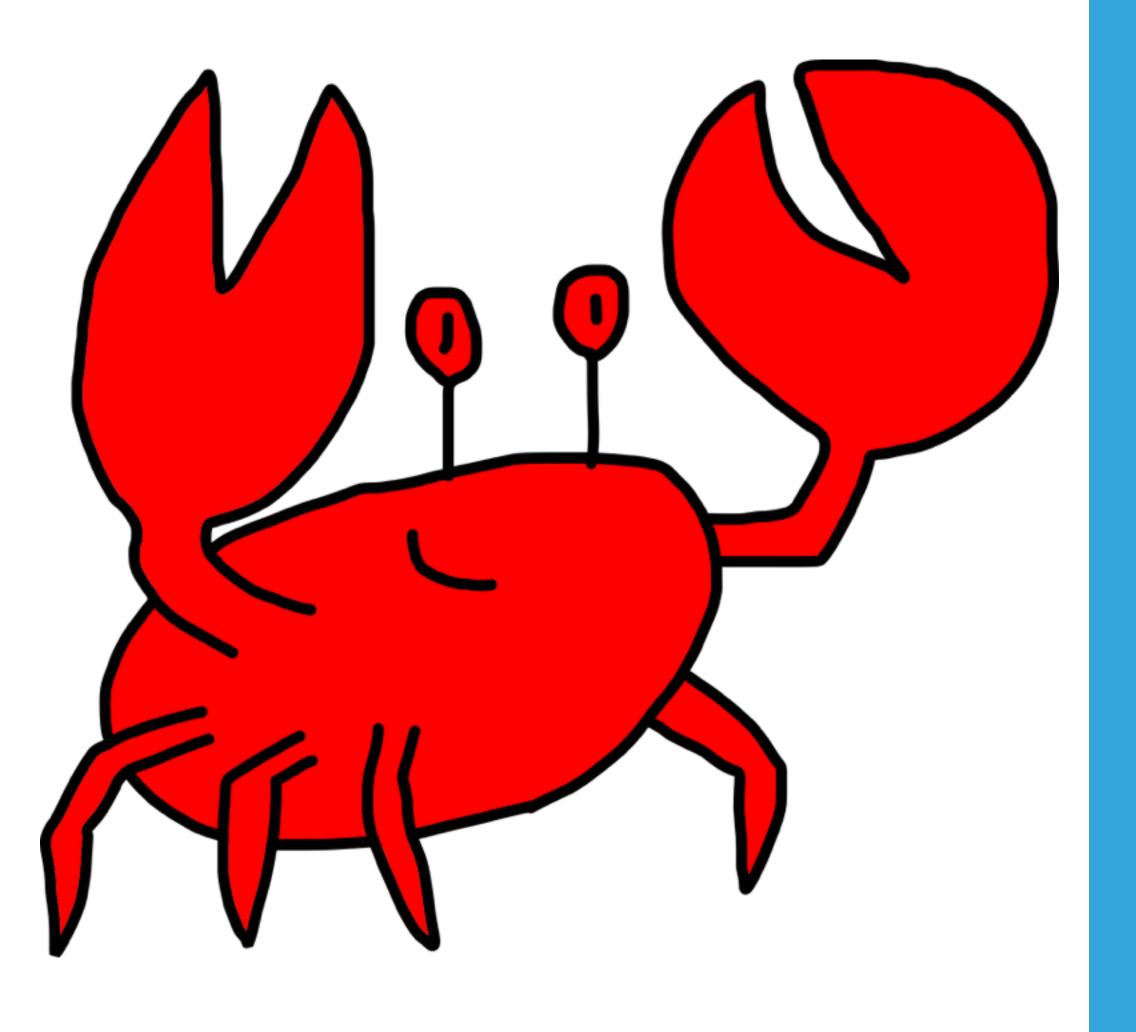
# Fraction of jobs with maximum runtime 240 120 180 60 max job runtime / m



# CAN CRAB3 DO A BETTER JOB THAN THE USERS?

### THERE'S NO ADVANCED PHYSICS IN SPLITTING.





# DRV-RUN FOR

### **MPROVING THE SPLITING EXPERIENCE: DRY-RUN**

lxplus066 @ ~/work/ttH/CMSSW\_8\_0\_15/src/ttH/TauRoast/test (ssh)

Creating temporary directory for dry run sandbox in /tmp/matze/tmproSmmL Executing test, please wait...

Using LumiBased splitting Task consists of 16 jobs to process 30881 lumis The longest job will process 2000 lumis, with an estimated processing time of 642 minutes The average job will process 1930 lumis, with an estimated processing time of 541 minutes The shortest job will process 1191 lumis, with an estimated processing time of 300 minutes The estimated memory requirement is 598 MB

Timing quantities given below are ESTIMATES. Keep in mind that external factors such as transient file-access delays can reduce estimate reliability.

For ~480 minute jobs, use: Data.unitsPerJob = 1710 You will need to submit a new task

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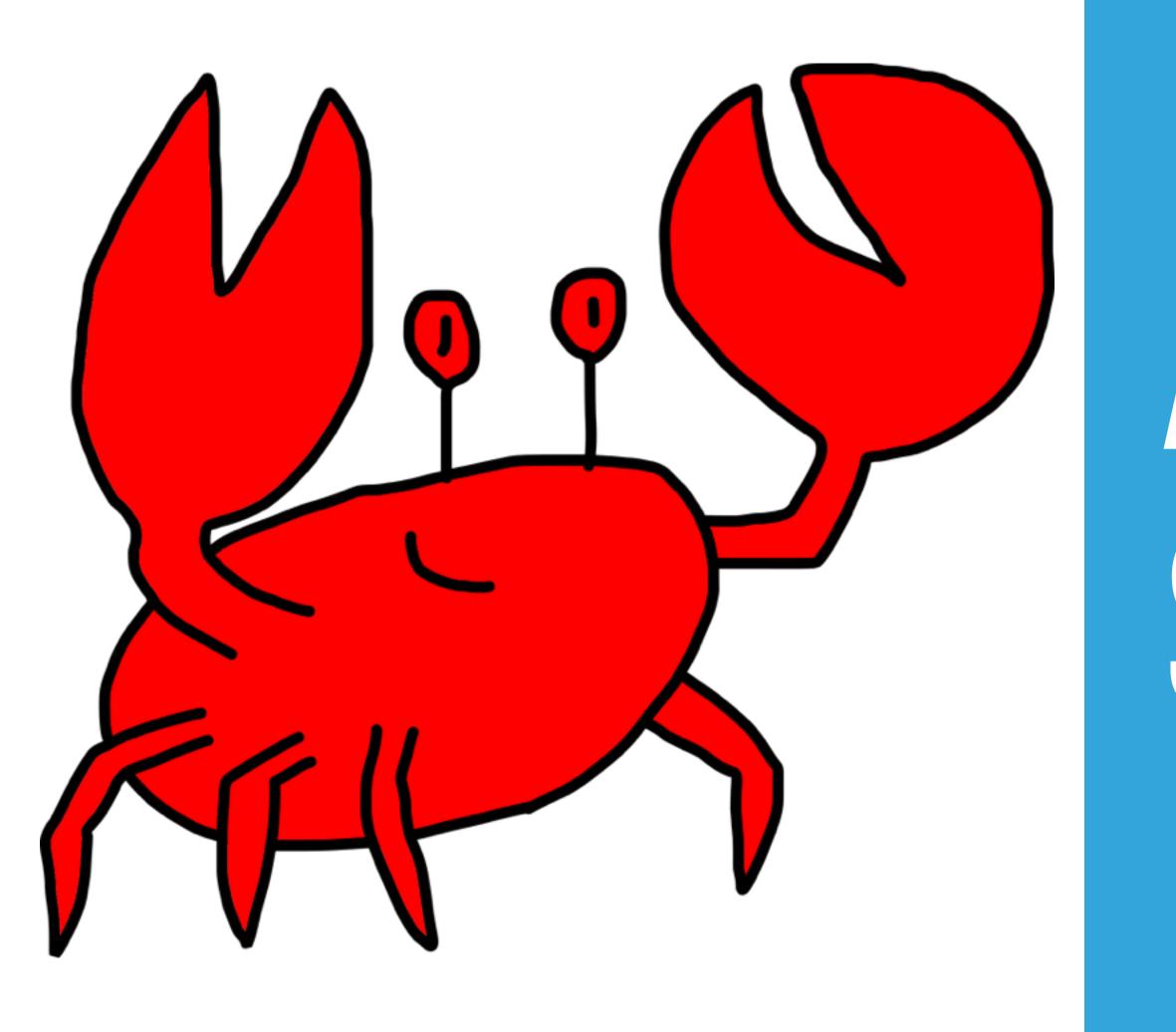
Dry run requested: task paused To continue processing, use 'crab proceed'

Run a small test job locally to provide timing estimate and recommend splitting parameters!









# AUTOMATIC Splitting

## **USING DAGMAN TO DO THE SPLITTING AUTOMATICALLY**

 The probe's post processing step then: Performs splitting Creates a SubDAG with processing jobs



# TaskWorker sends a single node DAG ("probe" for timing)

### **PROCESSING SUBDAG**

PRE	JOB	POST
PRE	JOB	POST
	• • •	



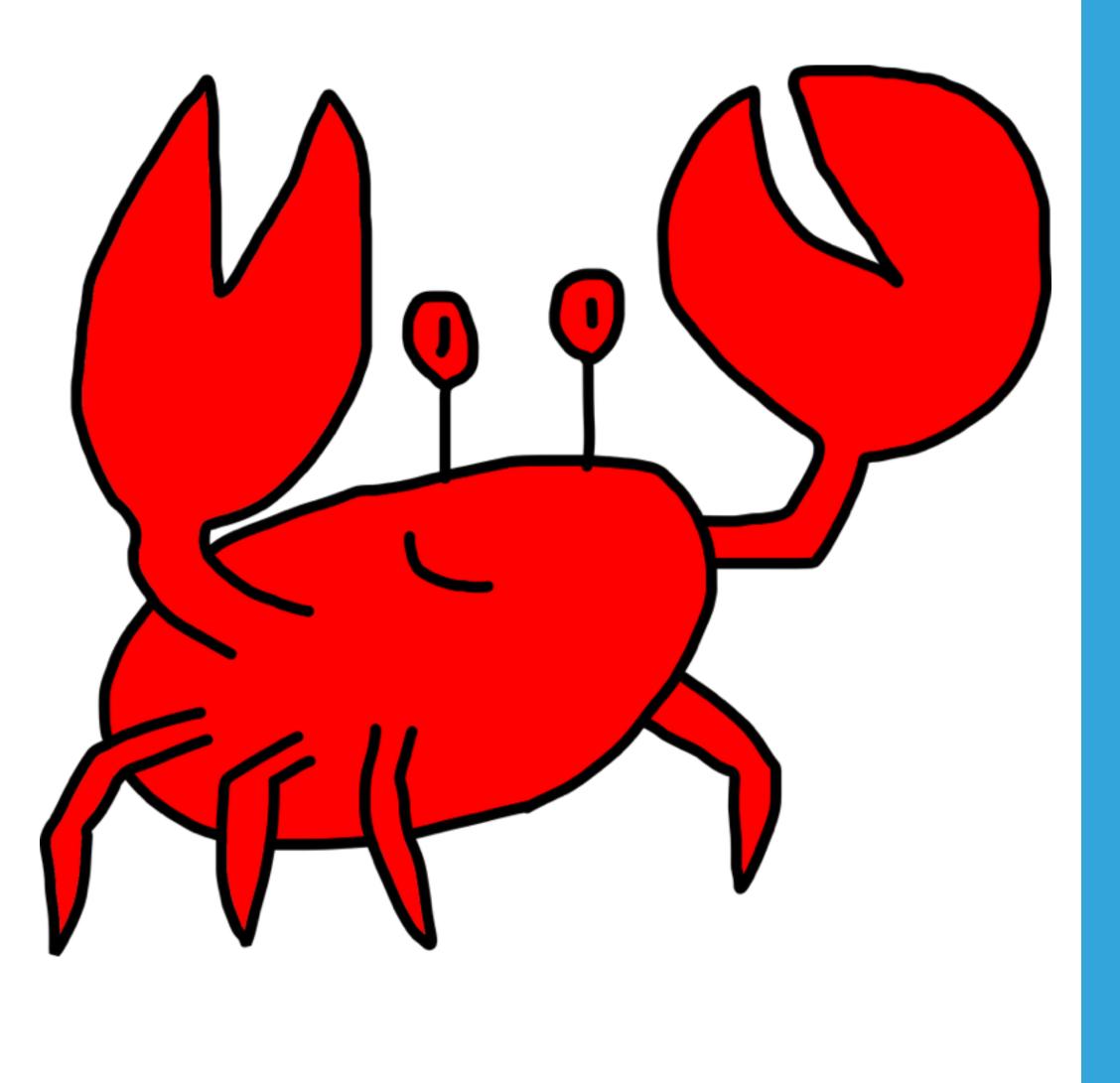
### BENEFITS

### For users- removes guesswork: only need to configure desired runtime

### For CRAB3 operations— allows enforcement of a minimum job runtime







# FOR FASTER



## **CONSTRAINING TASK RUNTIME**

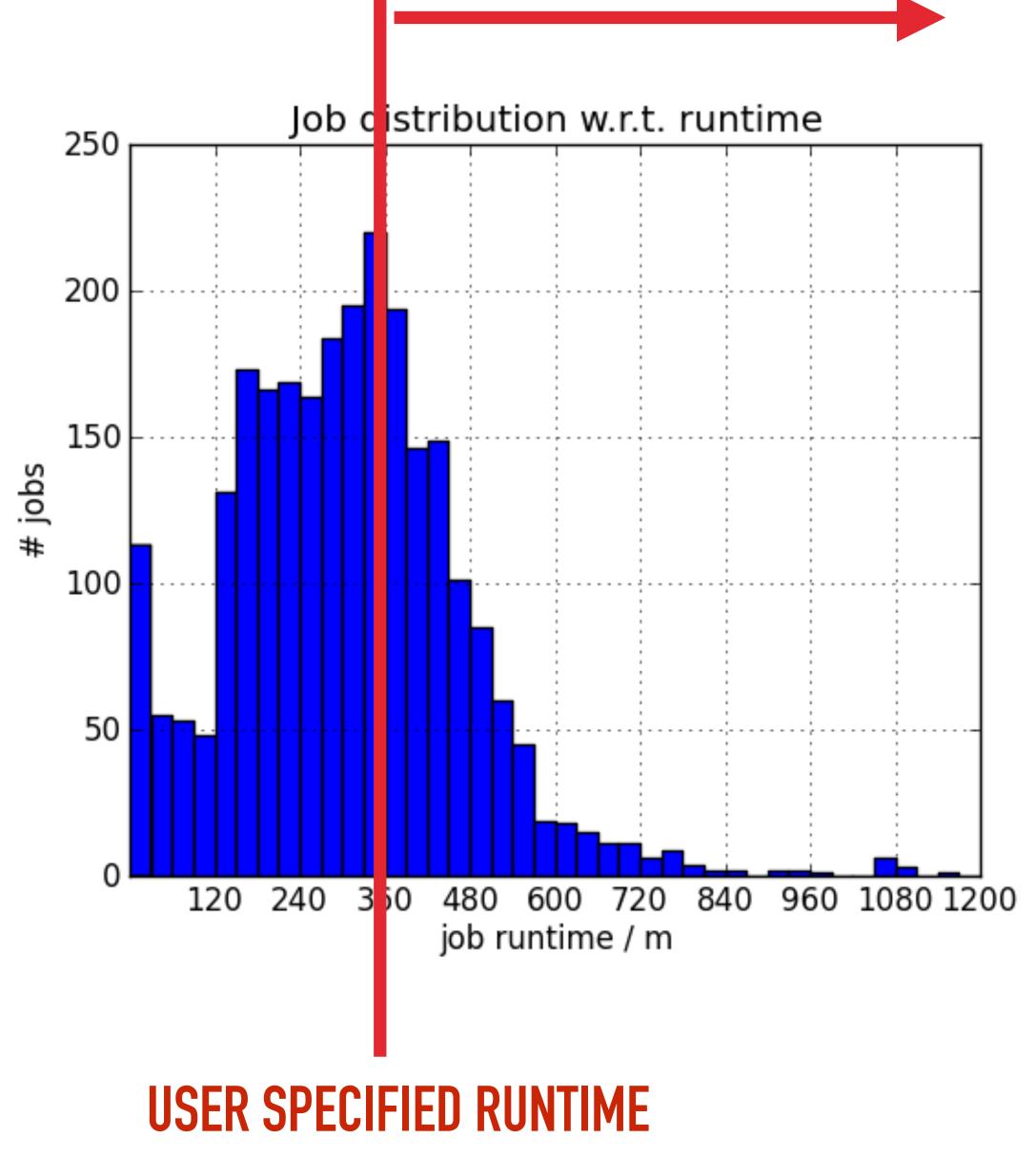
Not all machines/sites equal: some jobs still take more time than expected.

• Since CMSSW\_7\_2\_0: Can configure runtime limitations

New possibility:

- Use this to cut off tails in the runtime distribution!
- Resplit unfinished work to create subDAGs with small jobs which run in parallel
- Faster task completion!

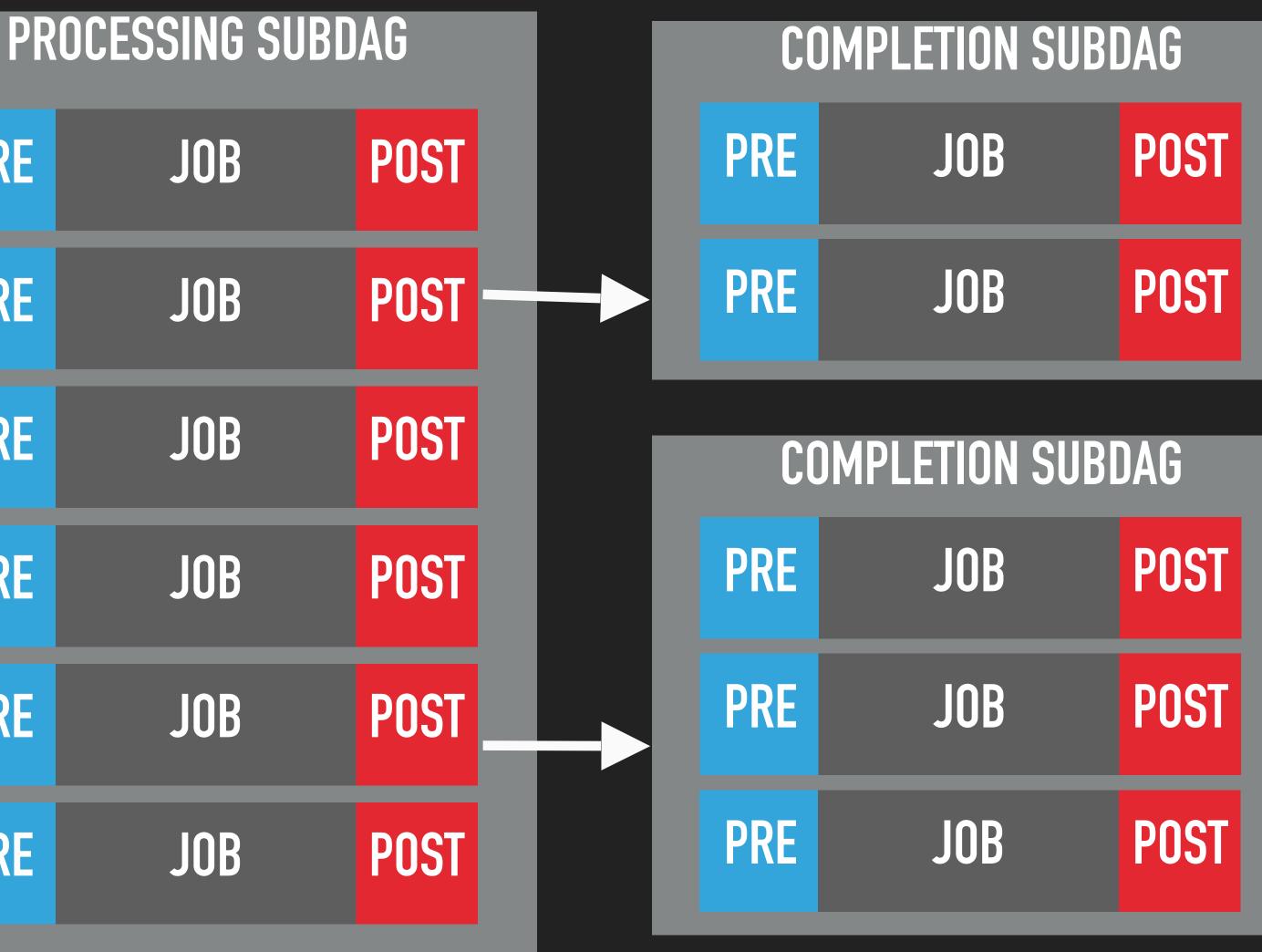
TAIL



## TAIL-SPLITTING FOR FASTER TASK COMPLETION

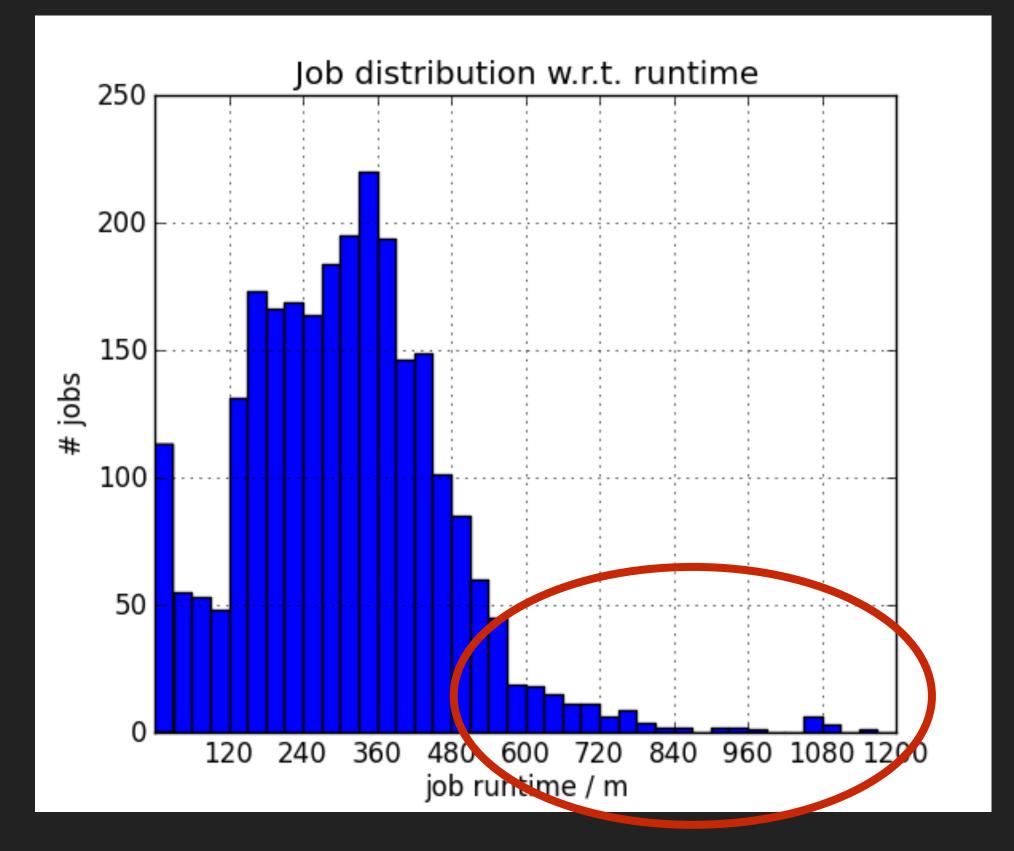


PRE	JO
PRE	JO



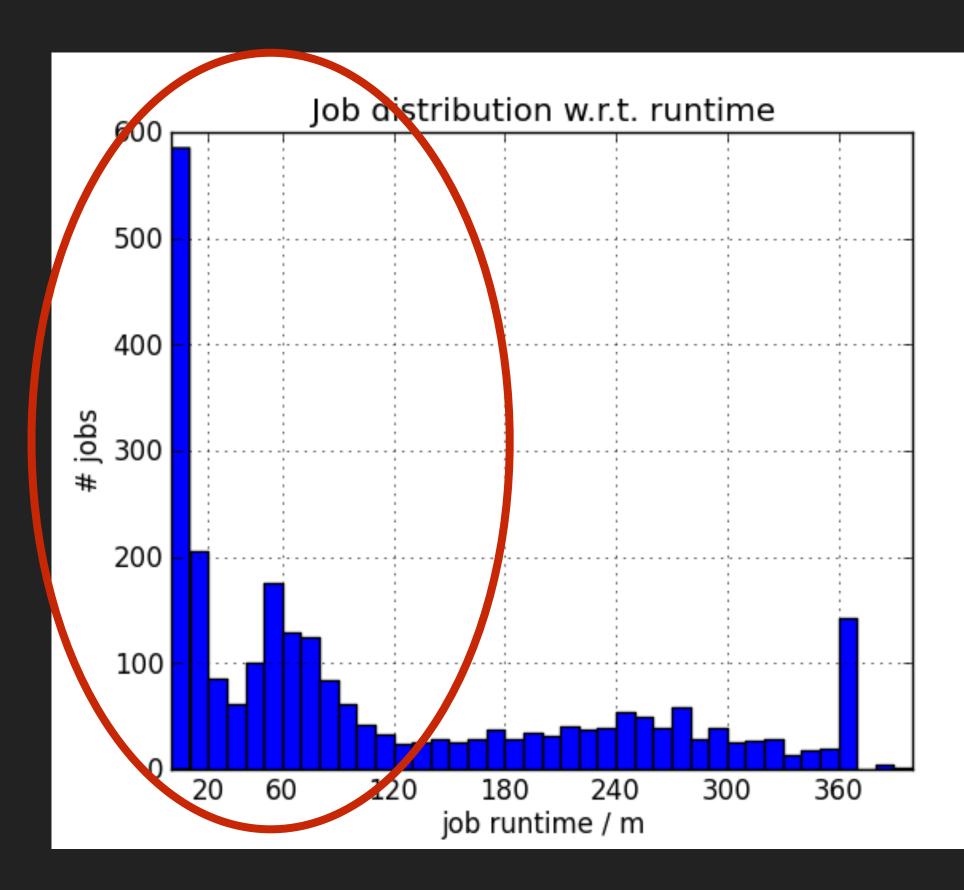


### **TAIL-SPLITTING FOR FASTER TASK COMPLETION**



### **TWENTY HOURS TO COMPLETE!**

### **CONVERT TAIL INTO SMALLER JOBS**





### CHALLENGES

SubDAGs for the tail-splitting jobs increase resource consumption

### CONCLUSIONS

- We have implemented automatic task splitting
  - Will be included in the October CRAB3 release
  - Tail-splitting feature is currently disabled, will be enabled pending more tweaks to address resource consumption

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