

# Harnessing cloud computing for high capacity analysis of neuroimaging data from NDAR

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## Introduction

- The National Database for Autism Research (NDAR)<sup>1</sup> hosts a vast collection of neuroimaging datasets that can be processed and utilized to yield significant scientific discoveries.
- This amount of resources necessitates a high-performance computing (HPC) infrastructure, which is not always readily available for researchers in-house.
- Amazon Web Services (AWS) Elastic Compute Cloud (EC2) computing service offers a “pay as you go” model that allows researchers to utilize HPC performance without the up-front capital costs and maintenance of an in-house solution.
- The developers of the Laboratory of Neuro Imaging (LONI) Pipeline, the Neuroimaging Informatics Tools and Resources Clearinghouse (NITRC) Computational Environment (CE) and the Configurable Pipeline for the Analysis of Connectomes (C-PAC) have implemented pipelines in EC2 that interface with NDAR

## Methods

### LONI Pipeline

- The LONI Pipeline software was extended to include new pipeline modules to access data from the NDAR database, transfer input data out of Amazon S3 (Simple Storage Service), and to load results back into S3<sup>2</sup>
- A pipeline was constructed to extract cortical thickness and subcortical region volume data from structural MRI images in the NDAR database, which included:
  1. Reorient images to standard orientation using FSL's reorient2std module
  2. Extract cortical thickness using FreeSurfer recon-all
  3. Calculate volumes of subcortical regions using FSL's BET and FIRST all
- The resulting pipeline was used to process 780 T1-weighted structural images and return the results to NDAR

### Configurable Pipeline For the Analysis of Connectomes (C-PAC)<sup>3</sup>

- C-PAC modules were written in Python to build input data lists by querying NDAR, read input data from S3, write processed results to S3 and write values back to the NDAR database
- New pipelines were created to perform the ANTS cortical thickness<sup>4</sup> procedure and the Preprocessed Connectomes Project's Quality Assessment Protocol (<http://preprocessed-connectomes-project.github.io/quality-assessment-pipeline>)<sup>5</sup>
- The resulting pipelines and modules were used to process several datasets and return the results to NDAR
  1. Cortical extraction from 3,197 T1-weighted structural images
  2. Structural and functional processing for 1,112 datasets from ABIDE
  3. Automated quality assessment of 1,112 datasets from ABIDE

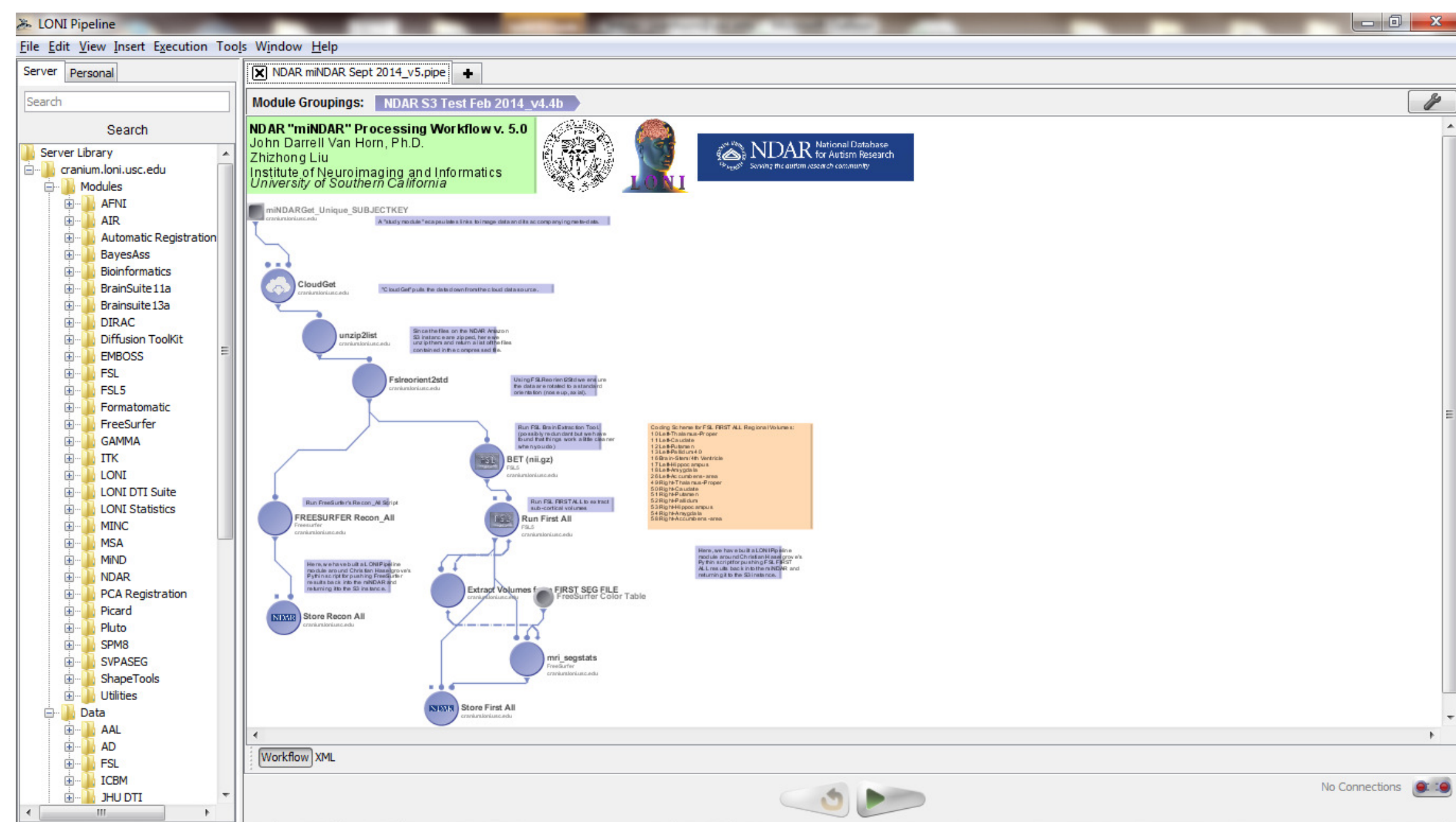


Figure 1 : Graphical layout of the constructed pipeline

Figure 2 : miNDAR database

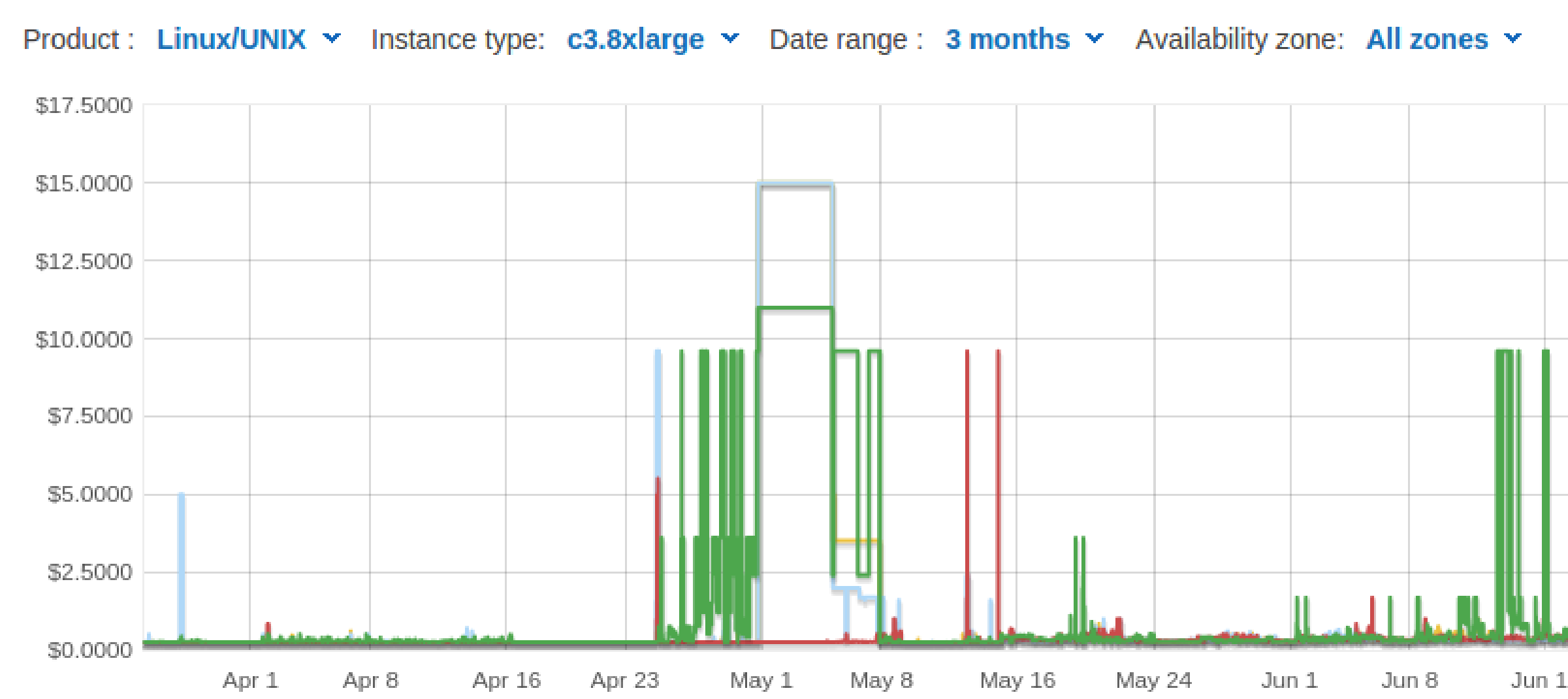


Figure 3 : Spot price history for the c3.8xlarge instance in the us-east-1 region on AWS across the four different availability zones for the past three months

## Results

## Results cont.

### Costs and run time models

- Models assume that users upload their input data to the cloud, run their pipelines and (optionally) downloads their outputs as they become available.
- Outputs are stored on an AWS Elastic Block Store (EBS) hard drive mounted to the master node and is NFS-shared across all of the slave nodes.
- Users can also upload their processed outputs to a cloud storage solution, like AWS Simple Storage Service (S3) directly from the cluster - this is the approach that was taken in processing data for NDAR.
- Storing results in AWS S3 avoids costly download time and provides for a viable solution for backing up data; however this does incur additional storage costs.

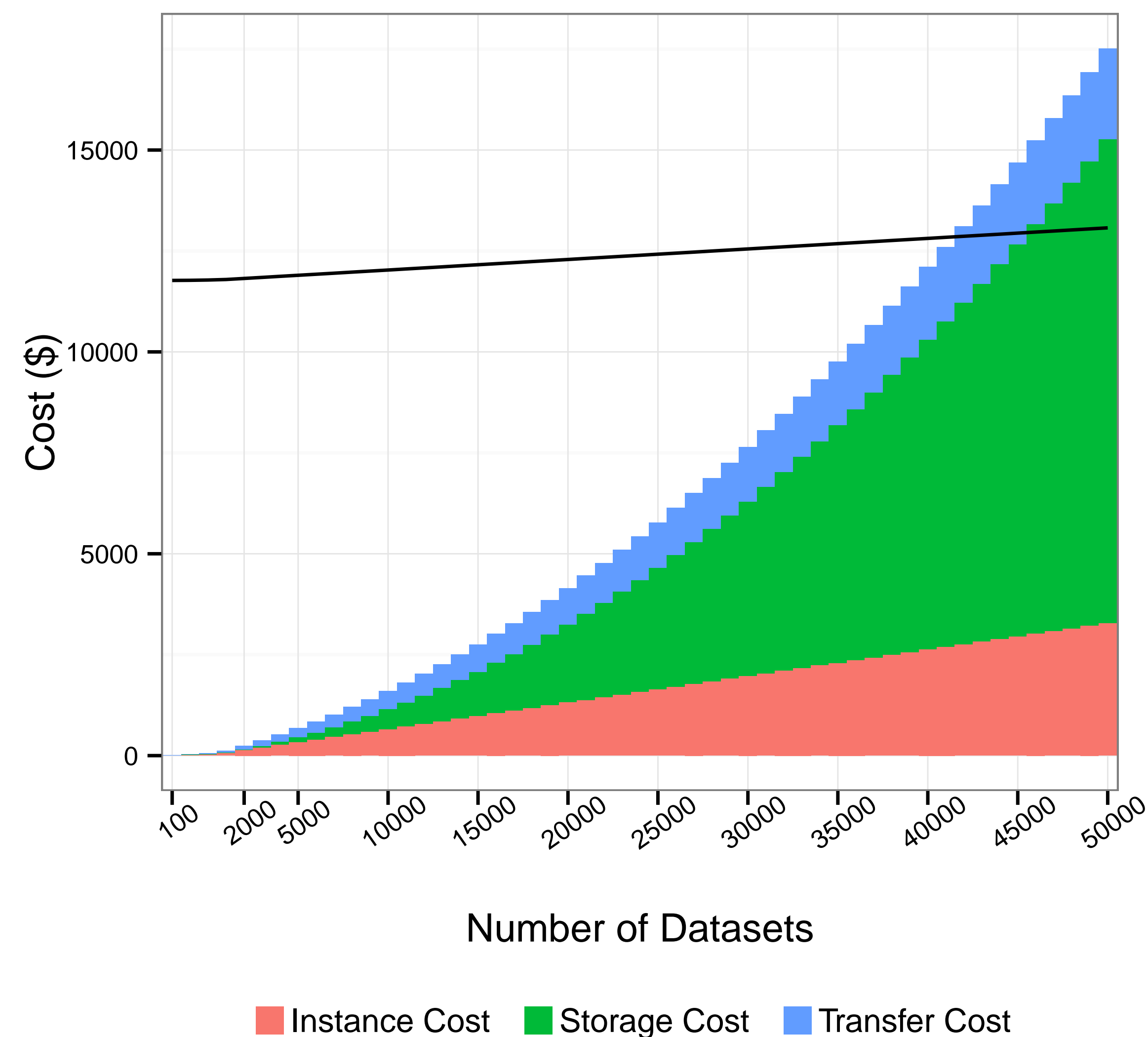


Figure 4 : AWS EC2 costs, grouped by cost type, for a typical C-PAC pipeline for different sized datasets versus owning and maintaining own server; costs are based on using the cluster configuration shown in Table 1 for the cloud, and a single compute node locally

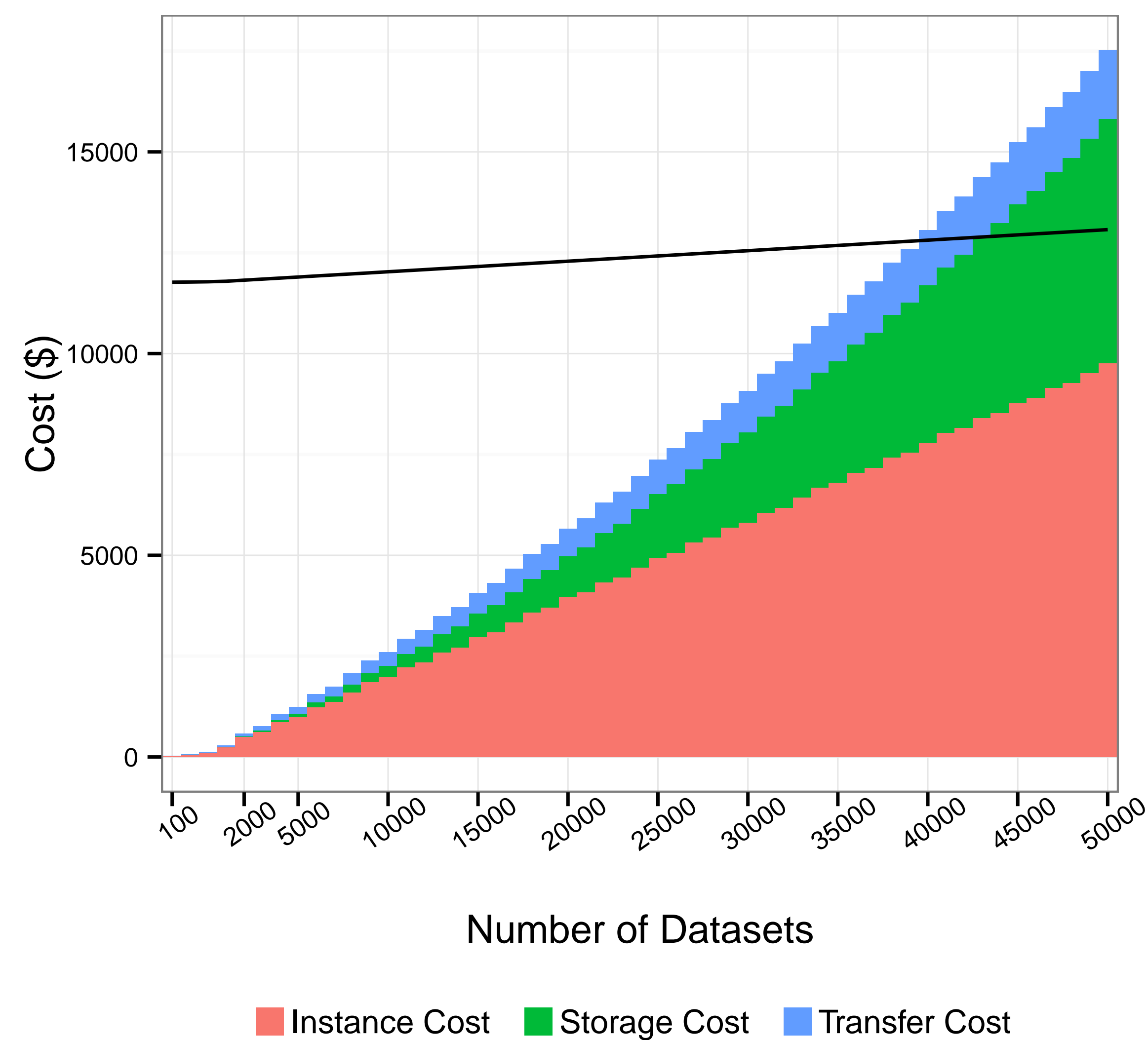


Figure 5 : AWS EC2 costs, grouped by cost type, for a typical FreeSurfer pipeline for different sized datasets versus owning and maintaining own server; costs are based on using the cluster configuration shown in Table 1 for the cloud, and a single compute node locally

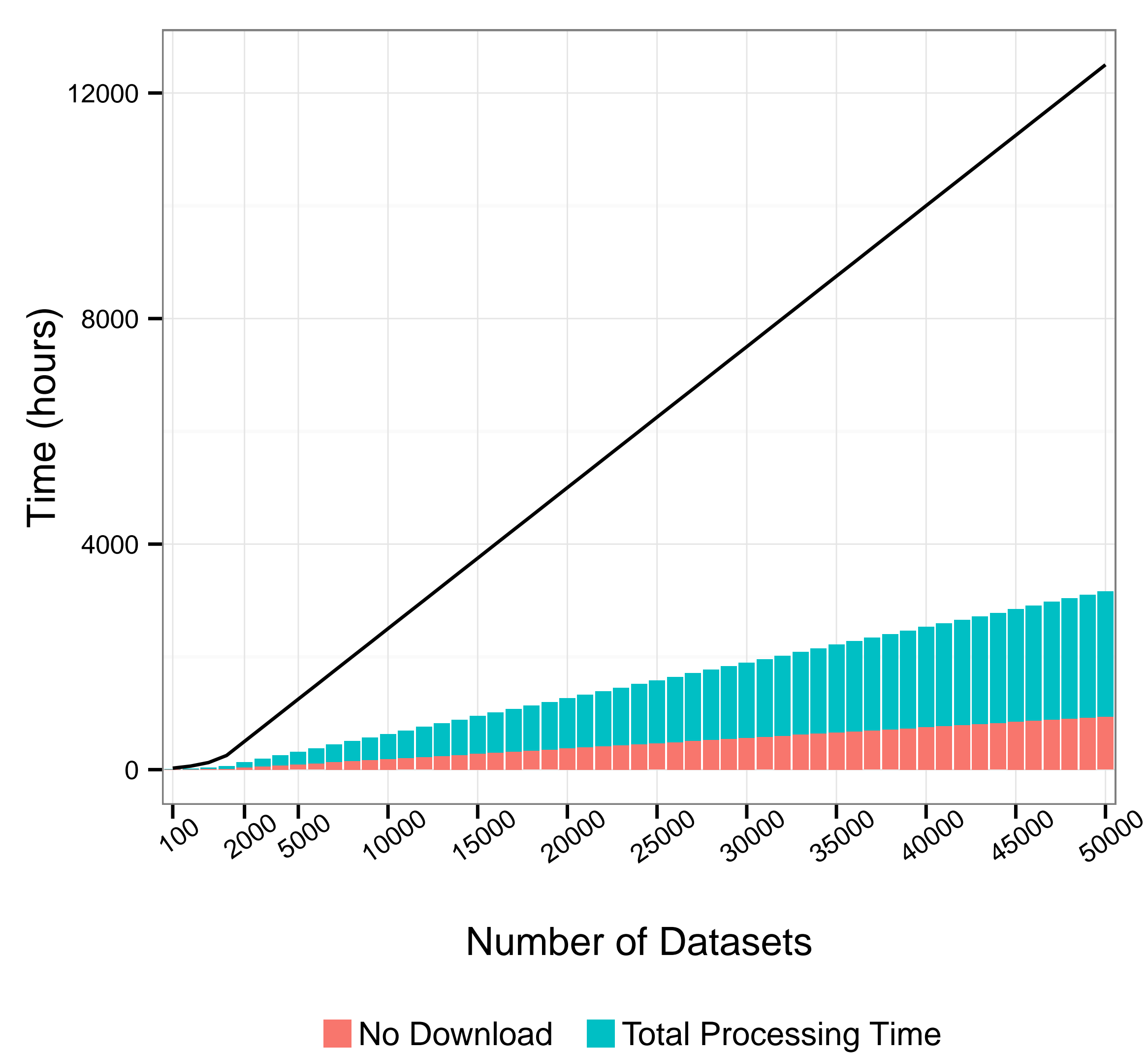


Figure 6 : Run times using the C-PAC pipeline, grouped by downloading vs non-downloading output data, for different sized datasets versus running locally; times are based on using the cluster configuration shown in Table 1 for the cloud, and a single compute node locally

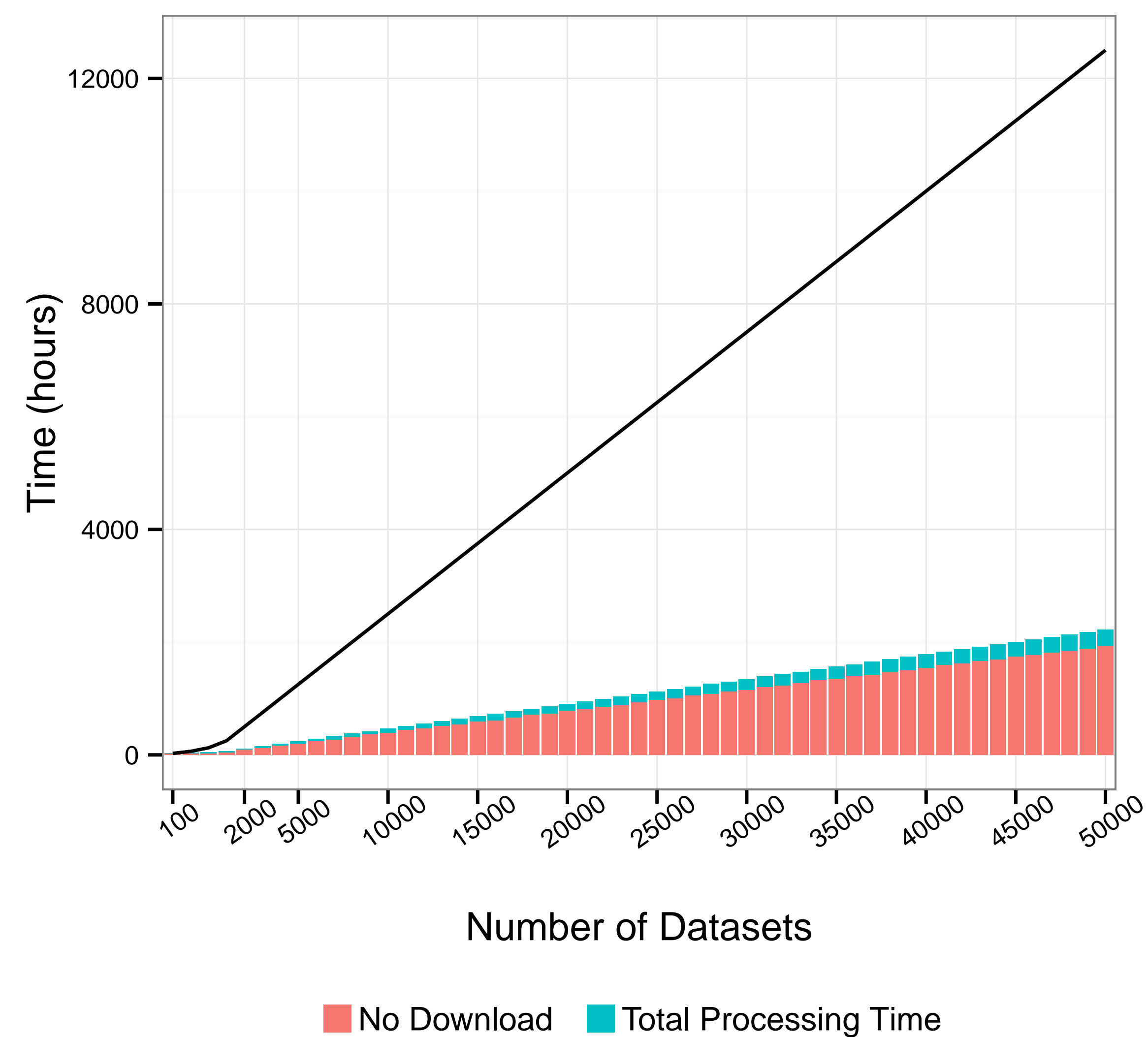


Figure 7 : Run times using the FreeSurfer pipeline, grouped by downloading vs non-downloading output data, for different sized datasets versus running locally