## Using Cloud Computing for Workshops

- Problem: Setting up the computing environment on each participant's computer takes time away from workshop content; versioning can cause problems
- Solution: Provide each participant with a cloud server with a preconfigured environment to access and use during the workshop

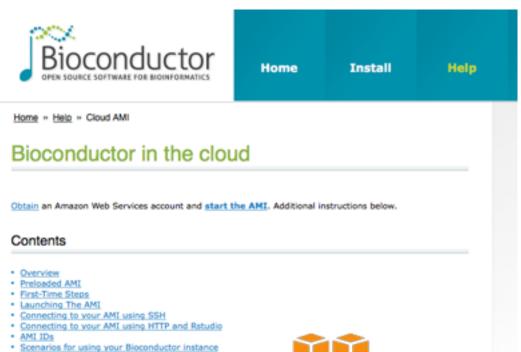




## **Bioconductor supports preconfigured AMIs**

amazon

webservices



- Pros:
  - Pre-loaded with latest R version, top 80 BioC software pkgs plus many annotation packages
  - Supported & updated

### Cons:

- Limited customizability of AMI properties
- Pricing structure can be confusing (e.g. bidding on Spot Instances)

## **Digital Ocean**

Droplets Images Networking API Support

### Create Droplets

#### Choose an image 👔

Distributions One-click apps Snapshots



- Fully customizable droplet properties
- Simple pricing structure



### Cons:

All setup of R/Rstudio Server, packages, etc done manually

### Steps to Create Rstudio Server Instances

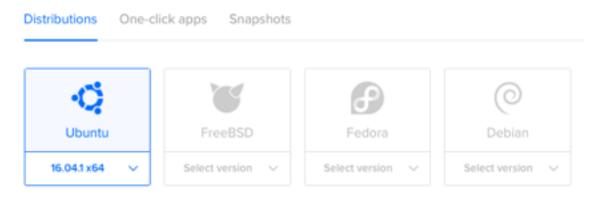
- 1. Launch empty droplet (instance)
  - Choose OS version, RAM, # cores, disk size, location, etc
- 2. Configure environment
  - Install R, Rstudio Server, packages, etc
- 3. Take a snapshot image
- 4. Launch many droplets based on snapshot



Droplets Images Networking API Support

### Create Droplets

#### Choose an image 💿



#### Choose a size

Standard High memory

\$ <b>5</b> /mo	\$ <b>10</b> /mo	\$20/mo	\$40/mo	\$ 80/mo	\$160/mo
\$0.007/hour	\$0.015/hour	\$0.030/hour	\$0.060/hour	\$0.119/hour	\$0.238/hour
512 MB / 1 CPU	1 GB / 1 CPU	2 GB / 2 CPUs	4 GB / 2 CPUs	8 GB / 4 CPUs	16 GB / 8 CPUs
20 GB SSD disk	30 GB SSD disk	40 GB SSD disk	60 GB SSD disk	80 GB SSD disk	160 GB SSD disk
1000 GB transfer	2 TB transfer	3 TB transfer	4 TB transfer	5 TB transfer	6 TB transfer
\$ <b>320</b> /mo \$0.476/hour	\$ <b>480</b> /mo \$0.714/hour	\$ <b>640</b> /mo \$0.952/hour			
32 GB / 12 CPUs 320 GB SSD disk 7 TB transfer	48 GB / 16 CPUs 480 GB SSD disk 8 TB transfer	64 GB / 20 CPUs 640 GB SSD disk 9 TB transfer			

#### Choose a size

Standard High memory

...

\$ <b>5</b> /mo	\$ <b>10</b> /mo	\$20/mo	\$40/mo	\$ 80/mo	\$160/mo
\$0.007/hour	\$0.015/hour	\$0.030/hour	\$0.060/hour	\$0.119/hour	\$0.238/hour
512 MB / 1 CPU	1 GB / 1 CPU	2 GB / 2 CPUs	4 GB / 2 CPUs	8 GB / 4 CPUs	16 GB / 8 CPUs
20 GB SSD disk	30 GB SSD disk	40 GB SSD disk	60 GB SSD disk	80 GB SSD disk	160 GB SSD disk
1000 GB transfer	2 TB transfer	3 TB transfer	4 TB transfer	5 TB transfer	6 TB transfer
\$ <b>320</b> /mo \$0.476/hour	\$ <b>480</b> /mo \$0.714/hour	\$640/mo \$0.952/hour			
32 GB / 12 CPUs 320 GB SSD disk 7 TB transfer	48 GB / 16 CPUs 480 GB SSD disk 8 TB transfer	64 GB / 20 CPUs 640 GB SSD disk 9 TB transfer			

#### Choose a size

Standard High memory

...

\$ <b>5</b> /mo	\$ <b>10</b> /mo	\$20/mo	\$ <b>40</b> /mo	\$ 80/mo	\$ <b>160</b> /
\$0.007/hour	\$0.015/hour	\$0.030/hour	\$0.060/hour	\$0.119/hour	\$0.238/h
12 MB / 1 CPU	1 GB / 1 CPU	2 GB / 2 CPUs	4 GB / 2 CPUs	8 GB / 4 CPUs	16 GB / 8 C
0 GB SSD disk	30 GB SSD disk	40 GB SSD disk	60 GB SSD disk	80 GB SSD disk	160 GB SSD
00 GB transfer	2 TB transfer	3 TB transfer	4 TB transfer	5 TB transfer	6 TB trans
\$ <b>320</b> /mo \$0.476/hour	\$ 480/mo \$0.714/hour	\$ 640/mo \$0.952/hour			
2 GB / 12 CPUs 20 GB SSD disk 7 TB transfer	48 GB / 16 CPUs 480 GB SSD disk 8 TB transfer	64 GB / 20 CPUs 640 GB SSD disk 9 TB transfer			

#### Choose a datacenter region

New York	San Francisco	Amsterdam	Singapore	London	Frankfurt
1 2 3	1 2	2 3	1	1	1
Torento	Bangalore				
1	1				

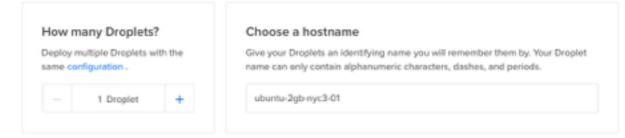
#### Choose a size

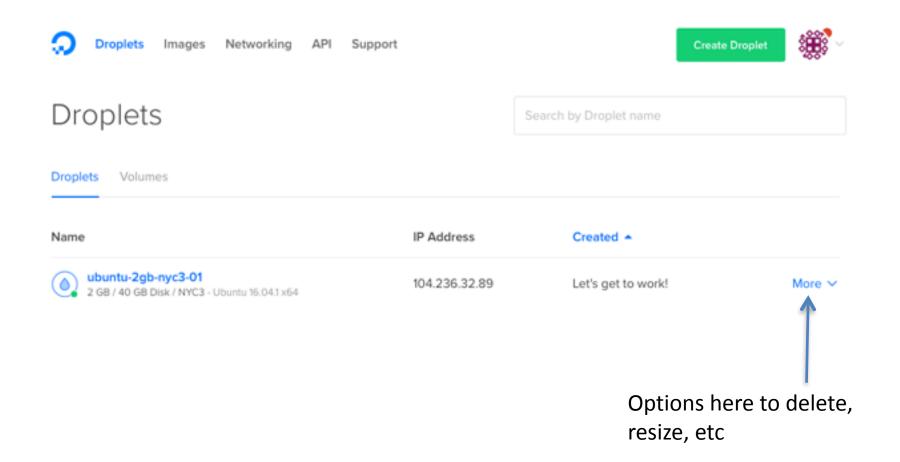
Standard High memory

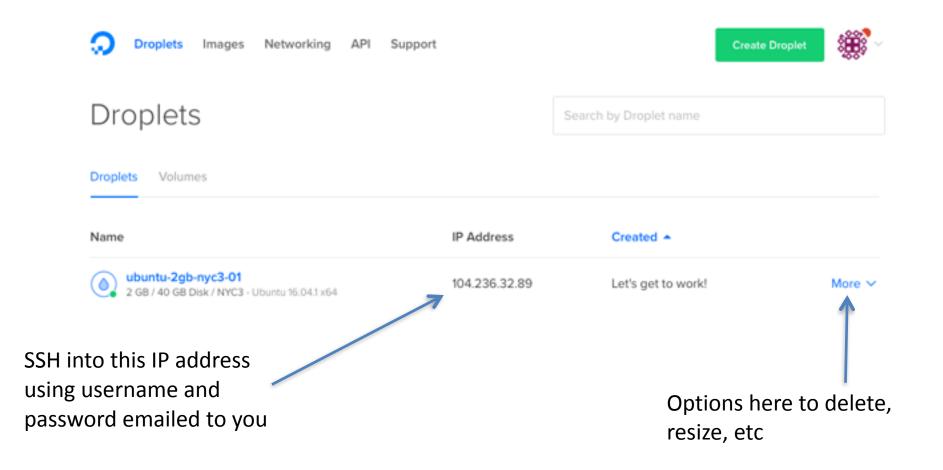
...

\$ <b>5</b> /mo \$0.007/hour	\$ <b>10</b> /mo \$0.015/hour	\$20/mo \$0.030/hour	\$ <b>40</b> /mo \$0.060/hour	\$ 80/mo \$0.119/hour	\$160/mo \$0.238/hour
512 MB / 1 CPU	1 GB / 1 CPU	2 GB / 2 CPUs	4 GB / 2 CPUs	8 GB / 4 CPUs	16 GB / 8 CPUs
20 GB SSD disk	30 GB SSD disk	40 GB SSD disk	60 GB SSD disk	80 GB SSD disk	160 GB SSD disk
000 GB transfer	2 TB transfer	3 TB transfer	4 TB transfer	5 TB transfer	6 TB transfer
\$ <b>320</b> /mo \$0.476/hour	\$480/mo \$0.714/hour	\$ <b>640</b> /mo \$0.952/hour			
32 GB / 12 CPUs	48 GB / 16 CPUs	64 GB / 20 CPUs			
320 GB SSD disk	480 GB SSD disk	640 GB SSD disk			
7 TB transfer	8 TB transfer	9 TB transfer			

#### Finalize and create







GitHubGist Search	All gists GitHub			Net	w gist	<b>3</b> -	
kdkorthauer / RstudioServerSetup.sh		🖋 Edit	1 Delete	★ Star	0	•	
<> Code Revisions 1	Embed - <sc< td=""><td>cript src="https://</td><td>gist. 🕃</td><td>٤</td><td>Downlo</td><td>ad ZIP</td><td></td></sc<>	cript src="https://	gist. 🕃	٤	Downlo	ad ZIP	

Bash script to set up R, install a few R packages, and get Rstudio Server running on ubuntu.

🕀 Rstu	dioServerSetup.sh	Raw
1	<pre>sudo sh -c 'echo "deb http://cran.rstudio.com/bin/linux/ubuntu trusty/" &gt;&gt; /etc/apt/sources.list'</pre>	
2	gpgkeyserver keyserver.ubuntu.comrecv-key E084DAB9	
3	gpg -aexport E084DA89   sudo apt-key add -	
- 4	sudo apt-get update	
5	sudo apt-get -y install r-base libapparmor1 libcurl4-gnutls-dev libxml2-dev libssl-dev gdebi-core	
6	sudo apt-get install libcairo2-dev	
7	sudo apt-get install libxt-dev	
8	sudo apt-get install git-core	
9		
10	sudo /bin/dd if=/dev/zero of=/var/swap.1 bs=1M count=1024	
11	sudo /sbin/mkswap /var/swap.1	
12	sudo /sbin/swapon /var/swap.1	
13	sudo sh -c 'echo "/var/swap.1 swap swap defaults 0 0 " >> /etc/fstab'	
14		
15	<pre>sudo suc "R -e \"install.packages('devtools', repos='http://cran.rstudio.com/')\""</pre>	
16	<pre>sudo suc "R -e \"install.packages('Rcpp', repos='http://cran.rstudio.com/')\""</pre>	
17	<pre>sudo suc "R -e \"install.packages('RcppEigen', repos='http://cran.rstudio.com/')\""</pre>	
18	<pre>sudo suc "R -e \"install.packages('ggplot2', repos='http://cran.rstudio.com/')\""</pre>	
19	<pre>sudo suc "R -e \"install.packages('Cairo', repos='http://cran.rstudio.com/')\""</pre>	

#### GitHub Gist: /kdkorthauer/RsudioServerSetup.sh

- Create a non-root account for yourself with sudo privileges, e.g. 'adduser keegan sudo'
- Set up an account for each user with a password and homedir using the 'adduser' command

▦	FoG_RstudioSe File Edit View In		Data Tools A	<b>dd-ons Help</b> La	ist edit was on Sept	ember 19
	8527 1	% .0 <u>_</u> .0 <u>0</u> 12	3 - Arial	· 10 ·	в <i>I</i> -5 <u>А</u> -	🌺 = 🖽 = 33
fx	IP address					
	A	в	с	D	E	F
1	IP address	Username	Password	Participant Name	Participant Email	
2	104.236.148.226:8787	festival1	hj5MQXTB			
3	45.55.20.73:8787	festival2	msG7PLn2			
4	45.55.20.98:8787	festival3	YT7QgH55			
5	45.55.21.29:8787	festival4	CfjbNPSs			
6	45.55.23.70:8787	festival5	Csc8mZaf			
7	45.55.24.97:8787	festival6	UdpgcqMU			
8	45.55.24.163:8787	festival7	Us59MXH8			
9	45.55.24.196:8787	festival8	4XPuqFD8			
10	45.55.25.50:8787	festival9	rGq2yzjH			
11	45.55.25.87:8787	festival10	nJqktFgT			
12	45.55.12.150:8787	festival11	s4NNFVN9			
13	45.55.30.170:8787	festival12	ZmAwLb3T			
14	45.55.30.192:8787	festival13	dbn2PfLh			
15	45.55.31.79:8787	festival14	B75eucWF			
16	159.203.240.96:8787	festival15	TELBZADB			

### 3. Take a snapshot image

Support



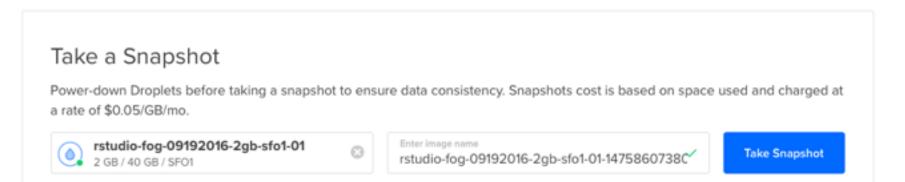
Images Networking API



### Images

Snapshots Backups

Droplets

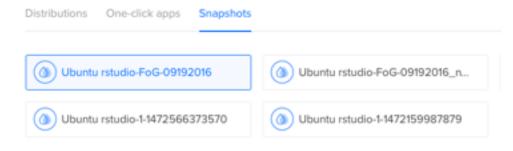


## 4. Launch many droplets based on snapshot

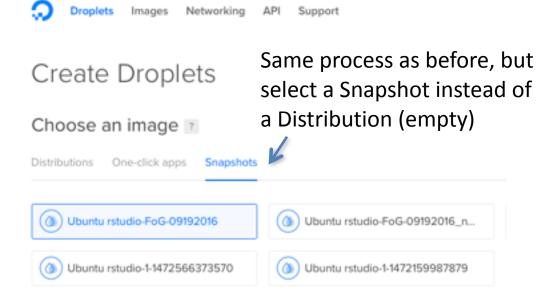
Droplets Images Networking API Support

### Create Droplets

#### Choose an image 📧



## 4. Launch many droplets based on snapshot



## 4. Launch many droplets based on snapshot

create Dropiets		t instead o		
() Ubuntu rstudio-1-1472566373570	Depl	y many Droplets? ay multiple Droplets with the configuration . 5 Droplets	•	Choose a hostname Give your Droplets an identifying name you will remember them by. Your Droplet name can only contain alphanumeric characters, dashes, and periods. rstudio-fog-09192016-2gb-sfo1-01 rstudio-fog-09192016-2gb-sfo1-02 rstudio-fog-09192016-2gb-sfo1-03 rstudio-fog-09192016-2gb-sfo1-04

# End Result

A SUBMIT AND A DEALER AND A SUBMIT AND A SUB			
••• <> 🗉 🛈 🔶 🚺	9 192.241.236.186:8787	Ċ	0 1 7 +
Studio	Sign in to RStudio Username: Password: Stay signed in Sign In		

# **End Result**

• • • < >	3,186 Č	0 ( 1 ( 1 ) (
R File Edit Code View Plots Session Build Debug Too	ls Help	keegan 🕞 🕘
💴 🍳 •   🔒 🔝   🚔   🏕 Co to file/function 🔄   🔯 •   Addi	is •	🖲 Project: (None) 🔹
SingleCellAnalyses.Rmd ×	Environment History	
🗇 🖓 🖳 🖓 💁 🚅 Knit HTML + 💿 + 🚱 🖓 🕀 🕀 🕀 🛪	🞯 🔒 📑 Import Dataset + 🥑	🗏 List + 🏼 🎯
25 - ```{r Check for data files, eval=TRUE, echo=TRUE} ◎ ≚ ▶	Global Environment -	Q,
<pre>27 file.exists("cell_metadata.csv") 28 file.exists("genes_counts.csv") 29 file.exists("genes_rpkm.csv") 30 file.exists("ercc_counts.csv")</pre>	Environment is empty	
<pre>31 file.exists("README.txt")</pre>	Files Plots Packages Help Viewer	
32 ~ ```	😂 New Folder 🛛 🝳 Upload 🝳 Delete 👍	Rename 🛛 🎡 More = 🤇
34 If any of the preceding lines return 'FALSE', double check	> / > home > FestivalWorkshopSC > BrainA	
1:1 Dismantling the bulk: examining neuronal heterogeneity using single-cell te	A Name Size	Modified
Console /home/FestivalWorkshopSC/  IS a COLLAPORATIVE project with many contributors.  (ype 'contributors()' for more information and	CPIL CLASSIFICATION, CSV 11/20 ND	Aug 26, 2016, 2:13 PM
citation()' on how to cite R or R packages in publications.		Aug 26, 2016, 2:13 PM
<pre>ype 'demo()' for some demos, 'help()' for on-line help, or help.start()' for an HTML browser interface to help.</pre>	Cell metadata.csv 187.7 KB	Aug 26, 2016, 2:13 PM
ype 'q()' to quit R.		Aug 26, 2016, 2:13 PM
		Aug 26, 2016, 2:13 PM

# End Result

MINISTRATION AND AND AND AND AND AND AND AND AND AN	Charles and the second s			
	0 +	iii kdkorthauer.github.io	C	0 1
	RStudio write.csv(exp.airr(exp.airr) , quote = PALST)	Dismantling the bulk	examining neuronal heterogenetics.cov , rov.names = rect	eneity using single-cell t +
	To get more insight into the models fit I visualize the results for a particular gen neuronal subtypes for the Differentially	by SCDE, we can use the related sode.test.expression. For example, we can view the cell-specific posterior dis Expressed gene Gad7:	on.difference function to stributions for the two different	
	<pre># visualize the results for exp.diff(rownames(exp.diff))</pre>			
	## 1b mle ## Gad1 0.5899266 0.9075793 ## GPVal ## Gad1 7.806413e-11	ub ce E cE 1.179853 0.5899266 7.153782 6.504284 0.4419	Pval D38e-13	
	<pre>scde.test.gene.expression.di</pre>	ifference("Gad1", models = err.mod, counts =	ets, prior = prior.mod)	
	a construction of the cons	Excitatory	0.00 0.04 0.08 joint posterior	
	01- 10	MLE: -10.03 95% CI: -10.48 : -9.60 Z = -7.16 aZ = -7.16 10 log2 expression ratio	2	
	88.	Inhibitory	-	

https://kdkorthauer.github.io/FestivalWorkshopVignettes/