Bioinformatics tutorials Documentation Release 1

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Contents:

CHAPTER 1

Welcome to 2014 MSU QB summer microbial genomics tutorial

Day	Schedule
Wed 8/13	 2-3pm: Get started with Amazon EC2 3-5pm: Read mapping and variant calling
Thu 8/14	• 2-5pm: Genome assembly

Sequencing technologes

- 454
- Illumina
- PacBio
- NanoPore

Resources

• Biostar

A high quality question & answer Web site.

• SEQanswers

A discussion and information site for next-generation sequencing.

• Software Carpentry lessons

A large number of open and reusable tutorials on the shell, programming, version control, etc.

Table Of Contents

Amazon Web Services instructions

Start up an EC2 instance

Here, we're going to startup an Amazon Web Services (AWS) Elastic Cloud Computing (EC2) "instance", or computer.

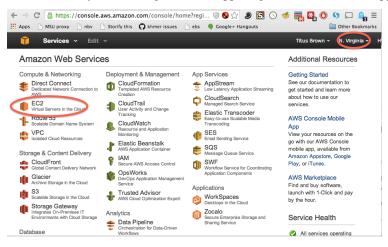
Go to 'https://aws.amazon.com' in a Web browser.

Select 'My Account/Console' menu option 'AWS Management Console."

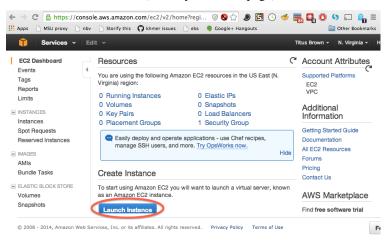
Log in with the following:

- username: qb2014msu@gmail.com
- password: TemporaryQB2014

Make sure it says North Virginia in the upper right, then select EC2 (upper left).



Select "Launch Instance" (midway down the page).



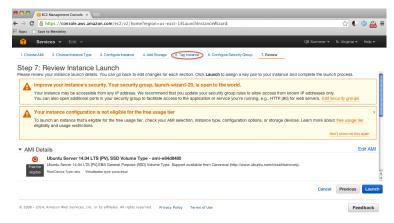
Next, scroll down the list of operating system types until you find Ubuntu 14.04 LTS (PV) – it should be at the very bottom. Click 'select'. (See *Starting up a custom operating system* if you want to start up a custom operating system instead of Ubuntu 14.04.)

1. Choose AMI 2. Choose Instance Ty	pe 3. Configure Instance	4. Add Storage 5. Tag Instance 6.	Configure Security Group
_	General Purpos preinstallei; Ag available. Root device type: Ubuntu Ubuntu Eros ser elgbe Ubuntu Server Type. Support i	er 14.04 LTS (PV), SSD Volume Ty 0 (64-bit) / ami-384d8450 (32-bit) 14.04 LTS (PV),EBS General Purpose (SS available from Canonical untu.com/cloud/services).	pe - Select

Scroll down the list of instance types until you find "m1.xlarge". Select the box to the left, and then click "Review and Launch."

Î	Services - Edit	¥				Titus Brown 👻 N. Vi	rginia ~
1. Cho	ose AMI 2. Choose Insta	nce Type 3. Col	nfigure Instance	4. Add Storage	5. Tag Instance 6. Co	nfigure Security Group	
Step	2: Choose an	Instance T	ype	50	2 × 00 (000)	160	
	General purpose	m1.small	1	1.7	1 x 160	-	
	General purpose	m1.medium	1	3.7	1 x 410		Ν
	General purpose	m1.large	2	7.5	2 x 420	Yes	Ν
	General purpose	m1.xlarge	4	15	4 x 420	Yes	
	Compute optimized	c3.large	2	3.75	2 x 16 (SSD)	-	N
	Compute optimized	c3.xlarge	4	7.5	2 x 40 (SSD)	Yes	N
			Cancel Pr	evious Review	w and Launch Next	: Configure Instance	Details

Before the launch, we need to name our intances, so we can tell ours from others'. Click on "5.Tag Instance" at the top of that page.



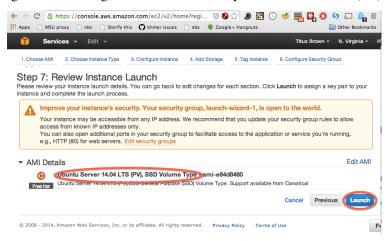
Add an unique value at the "Value" column, e.g. your name. Then click on "6.Configure Security Group".

C Attps://console.aws.amazon.com/ec2/v2/home?region=us-east-	1#LaunchInstanceWizard: 🏠 🕼 🖉 🤮
pps 🗋 Save to Mendeley	
🎁 Services 🗸 Edit 🗸	QB Summer 👻 N. Virginia 👻 Help 👻
1. Choose AMI 2. Choose Instance Type 3. Configure Instance 4. Add Storage 5.	Tag Instance 6. Configure Security Group 7. Review
ten C. Ten lastenen	
tep 5: Tag Instance	with key = Name and value = Webserver. Learn more about tagging your Amazon EC2 resources.
ag consists of a case-sensitive key-value pair. For example, you could deline a tag	with key = Name and value = webserver. Learn more about tagging your Amazon EC2 resources.
(127 characters maximum)	Value (255 characters maximum)
., (
	(your_name)
Name	

Choose "Select an existing security group", select "for_class" under column "Name" and click "Review and Launch":

1. Choose AMI			QB Summer + M	i. Virginia 👻 Help
	2. Choose Instance Type 3. Configure I	nstance 4. Add Storage 5. Tag Instance 6. Co	nfigure Security Group 7. Review	
	elow. Learn more about Amazon EC2 sec Assign a security group: O Creat	curity groups.	to the HTTP and HTTPS ports. You can create a new security gro	oup or select from a
Secur	ity Group ID	Name	Description	Actions
sg-6e4	4e060b	default	default VPC security group	Copy to new
sg-5e4	460d3b	for_class	allowing ssh with port 22 from any IP	Copy to new
	61e7a	launch-wizard-1	launch-wizard-1 created 2014-08-12T18:02:35.64	Copy to new
sg-1f5				0
sg-1f5		launch-wizard-10	launch-wizard-10 created 2014-08-13T14:09:46.0	Copy to new
sg-be4		-	launch-wizard-1 created 2014-08-12T18:02:35.64	Copy to n

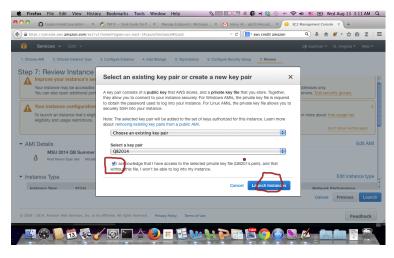
Ignore the warning, double check that it says "Ubuntu 14.04 LTS (PV)" at AMI Details, and cick "Launch".



Normally you will need to "create a new key pair" the first time through. However, for the purpose of easier management of our shared accout,

Select "Choose an existing key pair" and Select a key pair "QB2014".

Select "Launch Instance."



Click on the link of instance you just started.

	/2/home?region=us-east-1#LaunchInstanceWizard:		9 🖡 合 🖉・合 自 Z
Services 🗸 Edit 🗸			QB Summer 👻 N. Virginia 👻 Help
aunch Status			
Your instance is now launch	ing 💦		
 The following instance launch has 	been initiated I-5b4a9271 View launch log		
Get notified of estimated ch			
	arges il notification when estimated charges on your AWS bill e	waaad oo ooguust ugu dafino ifar oxoonala. If ugu ay	energ the free upper tied
orease ching assis to get ar ente	internet of the second of a ges of your parts on e	keese ar anount you deline (or example, if you ex	cood the nee dauge cont.
w to connect to your instance			
	a few minutes until it is in the running state, when it will b	be ready for you to use. Usage hours on your new in	stance will start immediately and continu
rue until you stop or terminate your inst	ance.		
k View Instances to monitor your insta	nce's status. Once your instance is in the running state, y	ou can connect to it from the Instances screen. Fit	nd out how to connect to your instance.
Here are some helpful resour	ces to get you started		
How to connect to your Linux instance	Amazon EC2: User Guide		
Learn about AWS Free Usage Tier	 Amazon EC2: Discussion Forum 		
	an also		
ile your instances are launching you			
ile your instances are launching you	and the second sec		
2008 - 2014, Amazon Web Services, Inc. er	its affiliates. All rights reserved. Privacy Policy Terms of U		Feedbac

Then you should see a "pending" line in the menu.

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🎁 Services 🗸 🗆	Edit 🗸				Titus Brown 🗸	N. Virginia	- Hel	р т
EC2 Dashboard Events	Launch Instanc	Connect	Actions *				0 ¢	> (
Tags	Filter: All insta	inces 👻 All ins	tance types 👻	Q i-00538	e2c	×		
Reports Limits					< < 1 to	1 of 1 Inst	ances	> >
INSTANCES	Name		 Instance Ty 	pe 👻 Availab	ility Zone 🕤 Instance St	ate - Stat	us Check	ks ·
Instances	0	i-00538e2c	m1.xlarge	us-east-	1e 🥥 pending	X	Initializir	ng
Spot Requests Reserved Instances	Instance: i-00	538e2c Public	DNS: -	000				
IMAGES	Description	Status Checks	Monitoring	Tags				
AMIs Bundle Tasks		Instance ID	i-00538e2c		Public DNS	-		
		Instance state	pending		Public IP			
ELASTIC BLOCK STORE		Instance type	m1.xlarge		Elastic IP	-		
Volumes Snapshots		Private DNS	-		Availability zone	us-east-1e		
		Private IPs	-		Security groups	launch-wiz	ard_1	

Wait until it turns green, then make a note of the "Public DNS" (we suggest copying and pasting it into a text notepad somewhere). This is your machine name, which you will need for logging in.

🧊 Services 🗸	Edit 🗸		Titus Brown 🗸	N. Virginia • Help •
EC2 Dashboard Events	Launch Instance Co	Actions *		ତ 🕈 (
Tags	Filter: All instances ¥	All instance types 👻 🔾	i-00538e2c	×
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INSTANCES	■ Name ♀ - In	stance ID 🔺 Instance Type	Availability Zone - Instance Sta	ate v Status Checks v
Instances	i-C	0538e2c m1.xlarge	us-east-1e 🥥 running	🖉 Initializing
Spot Requests Reserved Instances	Instance: i-00538e2c	Public DNS: ec2-54-205-96	-232.compute-1.amazonaws.com	
IMAGES AMIS	Description Status	Checks Monitoring Ta	gs	
Bundle Tasks	Inst	ance ID i-00538e2c	Public DNS	ec2-54-205-96- 232.compute- 1.amazonaws.com
ELASTIC BLOCK STORE	Instan	ce state running	Public IP	54.205.96.232
Snapshots	Instar	ice type m1.xlarge	Elastic IP	-
		to DNS in-10-235-34-	Availability zone	us-osst-1o

Then, go to Logging into your new instance "in the cloud" (Windows version) or Logging into your new instance "in the cloud" (Mac or Linux version)

You might also want to read about Terminating (shutting down) your EC2 instance.

Logging into your new instance "in the cloud" (Mac or Linux version)

OK, so you've created a running computer. How do you get to it?

The main thing you'll need is the network name of your new computer. To retrieve this, go to the instance view and click on the instance, and find the "Public DNS". This is the public name of your computer on the Internet.

Copy this name, and connect to that computer with ssh under the username 'ubuntu', as follows:

First we need a private key file to log in; it is the .pem file we already created during the setup of amazon instance. For convience of managing the shared account, we want you all to use the same .pem file, which can be downloaded from:

http://lyorn.idyll.org/~gjr/public/QB2014/QB2014.pem

If your browser is firefox with default settings, it should be downloaded into your "Downloads" folder. Move it onto "Desktop".

If your browser is chrome with default settings, the above link will be shown as a page with text. right click and select "Save As". In the pop up window, change the directory to "Desktop" and the file name is still "QB2014.pem"

Other browsers should be similar to either of the above two case.

Next, start Terminal:

- in Applications... Utilities... for mac
- use shortcut "CTRL+ATL+T" for linux

and then type:

chmod og-rwx ~/Desktop/QB2014.pem

to set the permissions on the private key file to "closed to all evildoers".

Then type:

ssh -i ~/Desktop/QB2014.pem ubuntu@ec2-???-???-???.compute-1.amazonaws.com

(but you have to replace the stuff after the '@' sign with the name of the host).

Here, you're logging in as user 'ubuntu' to the machine 'ec2-174-129-122-189.compute-1.amazonaws.com' using the authentication key located in 'QB2014.pem' on your Desktop.

You should now see a text line that starts with something like ubuntu@ip-10-235-34-223:~\$. You're in! Now type:

sudo bash cd /root

to switch into superuser mode (see: http://xkcd.com/149/) and go to your home directory.

This is where the rest of the tutorials will start!

You might also want to read about Terminating (shutting down) your EC2 instance.

To log out, type:

exit	
logout	

or just close the window.

Logging into your new instance "in the cloud" (Windows version)

Download .pem file

First we need a private key file to log in; it is the .pem file we already created during the setup of amazon instance. For convience of managing the shared account, we want you all to use the same .pem file, which can be downloaded from:

http://lyorn.idyll.org/~gjr/public/QB2014/QB2014.pem

If your browser is firefox with default settings, it should be downloaded into your "Downloads" folder. Move it onto "Desktop".

If your browser is chrome with default settings, the above link will be shown as a page with text. right click and select "Save As". In the pop up window, change the directory to "Desktop" and the file name is still "QB2014.pem"

Other browsers should be similar to either of the above two case.

Download Putty and Puttygen from here: http://www.chiark.greenend.org.uk/~sgtatham/putty/download.html

Generate a ppk file from your pem file

(You only need to do this once!)

Open puttygen; select "Load".

Ŀ	PuTTY Key Generator	×
	File Key Conversions Help	
	Key	
L	No key.	
	Actions	
	Generate a public/private key pair	Generate
	Load an existing private key file	Load
	Save the generated key	Save public key Save private key
	Parameters	
	Type of key to generate:	SSH-2 DSA
	 SSH-1 (RSA) SSH-2 RSA Number of bits in a generated key: 	1024
	Number of bits in a generated key.	1024

Find and load your '.pem' file; it's probably in your Downloads folder. Note, you have to select 'All files' on the bottom.

	😴 Load private key:	×
File Key Conversions Help Key No key.	Organize * New folder	• 4 Search aws
Actions Generates a public fundas for gore Last an executing public largy Spane for generated largy Spane for generated largy Formations	Kontek Kontek	AWS-independent AWS-independent stdat PHO
Type of key to generate: SSH-1 (RSA) ® SSH-2 RSA © SSH-2 DSA Number of bits in a generated key: 2048	File narte amezon pem	All Files (*.*) Open Cancel

Load it.



Now, "save private key". Put it somewhere easy to find.

PuTTY Key Generator			×		
File Key Conversion	s Help				
Key Public key for pasting in	to OpenSSH authorized	d_keys file:			
ssh-rsa AAAAB3NzaC1yc2EAAAADAQABAAABAQCjb0udZ3CpTWZ4auWYxDJsS57mR7k4p/ waH72t+3xwZc+/0pYxaUBzjdHy314gEQpc/bgJy0RNfblY+hnhcbOj +61zo7niom4/GkM32GLySSga3p +fqRlYW79k1LWJt0n3cwiuyPoA2pyg]6y47V6vxmlmVJEf1AXMBr50ngke1h/i3SJCDZz +					
Key fingerprint:	ssh-rsa 2048 c7:cd:91fd:e7:d3:c6fe:c7:c7f2:32:0f:87:08:41				
Key comment:	imported-openssh-key				
Key passphrase:					
Confirm passphrase:					
Actions					
Generate a public/private key pair Generate					
Load an existing private key file					
Save the generated key	,	Save public key	Save private key		
Parameters					
Type of key to generate SSH-1 (RSA)	e:	© SSI	H-2 DSA		
Number of bits in a gene	arated key:		1024		

Logging into your EC2 instance with Putty

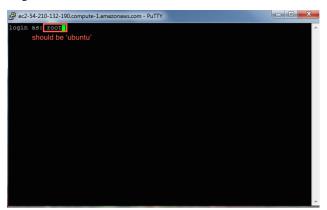
Open up putty, and enter your hostname into the Host Name box.

Reputty Configuration	×
Category:	
🖃 Session 🔺	Basic options for your PuTTY session
Logging	Specify the destination you want to connect to
En Terminal	Host Name (or IP address) Port
Bell	-72-179-190.compute-1.amazonaws.com
Features	Connection type:
🖶 Window	🔘 Raw 🔘 Telnet 🔘 Rlogin 💿 SSH 🔘 Serial
Appearance	Load, save or delete a stored session
Behaviour	Saved Sessions
Selection	
Colours	Default Settings
Data Proxy	Save
Telnet	Delete
Rlogin	
⊟- SSH	
Kex	Close window on exit:
	Always Never Only on clean exit
About	Open Cancel
About	Open Cancel
2 nuthrann	6/2/2012 10/41 DM Application

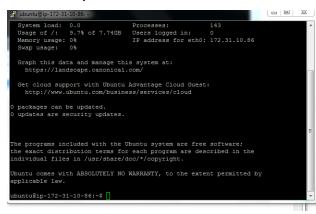
Now, go find the 'SSH' section and enter your ppk file (generated above by puttygen). Then select 'Open'.

PuTTY Configuratio	n	X
Category: - Terminal - Keyboard - Bell - Features - Window - Appearance - Behaviour - Translation - Selection - Colours - Connection - Data - Proxy - Teinet - Riogin - SSH - Key - Auth - X11 - Tunnels - Bug	A III	Options controlling SSH authentication Bypass authentication entirely (SSH-2 only) Display pre-authentication banner (SSH-2 only) Authentication methods Authentication methods Attempt authentication using Pageant Attempt TIS or CryptoCard auth (SSH-1) Attempt TiS or CryptoCard auth (SSH-2) Authentication parameters Allow agent forwarding Allow attempted changes of usemame in SSH-2 Private key file for course record. C:\Users\Aswathyseb\Desktop\aws\put
About		Open Cancel

Log in as "ubuntu".



Declare victory!



Here, you're logging in as user 'ubuntu' to the machine 'ec2-174-129-122-189.compute-1.amazonaws.com' using the authentication key located in 'amazon.pem' on your Desktop.

You should now see a text line that starts with something like ubuntu@ip-10-235-34-223:~\$. You're in! Now type:

sudo bash cd /root

to switch into superuser mode (see: http://xkcd.com/149/) and go to your home directory.

This is where the rest of the tutorials will start!

You might also want to read about Terminating (shutting down) your EC2 instance.

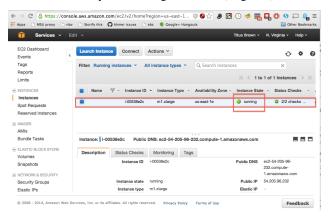
To log out, type:

exit	
logout	

or just close the window.

Terminating (shutting down) your EC2 instance

While your instance is running, Amazon will happily charge you on a per-hour basis – check out the pricing for more information. In general, you will want to shut down your instance when you're done with it; to do that, go to your EC2 console and find your running instances (in green).



Then, select one or all of them, and go to the 'Actions...' menu, and then select 'Terminate'. Agree.

After a minute or two, the console should show the instance as "terminated".

	e.aws.amazon.com/ec2/v2/home?region=us-east-1 💿 🔇 🏠 🤌 🖺 🗅 Storify this 🖸 khmer issues 🗋 ebs 🎈 Google+ Hangouts	. () 🥌 🜉 🛃 () () 💭 🔒 Ξ ☐ Other Bookmarks
🎁 Services 🗸 Ed	it ~	Titus Brown 👻 N. Virginia 👻 Help 👻
EC2 Dashboard Events Tags	Launch Instance Connect Actions * Filter: All Instances * All Instance types * Q. Search Instances	ତ କ ଡ ×
Reports Limits		<pre> < < 1 to 2 of 2 Instances > > </pre>
INSTANCES	Name 🖓 - Instance ID 🔺 Instance Type - Availability Zone -	- Instance State - Status Checks - A
Instances Spot Requests	i-00538e2c m1.xlarge us-east-1e	terminated
Reserved Instances	i-983decb4 m1.xlarge us-east-1e	terminated /
IMAGES AMIs Bundle Tasks	Instance: I-00538e2c Public DNS: -	
ELASTIC BLOCK STORE	Description Status Checks Monitoring Tags	
Volumes Snapshots	Instance ID i-00538e2c Publ Instance state terminated Pu	Ic DNS - Iblic IP Istic IP -
Security Groups Elastic IPs	Private DNS - Availabilit Duivate IDa Comunities	
© 2008 - 2014, Amazon Web Se	rvices, Inc. or its affiliates. All rights reserved. Privacy Policy Terms of Use	Feedback

Starting up a custom operating system

The instructions in *Start up an EC2 instance* tell you how to start up a machine with Ubuntu Linux version 14.04 on it, but that machine comes with very little software installed. For anything where you are executing actual analyses, you're going to want to have a bunch of basic software installed.

Therefore, we make custom versions of Ubuntu available as well, that come with some software pre-installed.

To boot these, go to EC2/Launch and select "Community AMIs" instead of the default "Quick Start";

Then type in the AMI number or name in the search box. We need to use:

ami-d25283ba

for this whole workship. Or you can just type "MSU" in the search box and hit ENTER. "MSU 2014 QB Summer Course" is the one needed.

Then proceed with the rest of *Start up an EC2 instance*.

Firefox File Edit View	History Bookmarks Tools Window		🔸 🍪 🕙 🚸 🛜 🐠 💻 💽 🕅 14msu@ × 👔 EC2 Management Conse	
https://console.aws.amazon.co	m/ec2/v2/home?region=us-east-1#LaunchInstanc	eWizard: 🗸 🦉 🖉 🛪 av	vs credit amazon Q 🖡 🏫	#・☆ 🖻 Z 😑
🎁 Services 🗸 Edit 🗸			QB Summer 🕶	N. Virginia 👻 Help 👻
1. Choose AMI 2. Choose Instance	e Type 3. Configure Instance 4. Add Storage	5. Tag Instance 6. Configure Security Group	7. Review	
An AMI is a template that contains	mazon Machine Image (AMI the software configuration (operating system, ap tplace; or you can select one of your own AMIs.		unch your instance. You can select an A	Cancel and Exit MI provided by AWS, our
Quick Start	Q MSU X		< ≤ 1 to	6 of 6 AMIs > >
My AMIs		J" on AWS Marketplace		
AWS Marketplace		pre-configured to run on AWS		
Community AMIs	ci-days-ipython-nb-demo	- ami-c5de82ac		Select
Operating system Amazon Linux	IPython Notebook server demo Root device type: ebs Virtualizati	for MSU CI Days on type: paravirtual		64-bit
🗆 Cent OS 🛛 🎆	MSU 2014 QB Summer Co	ourse - ami-d25283ba		Select
 Debian Pedora 	-	on type: paravirtual		64-bit
Gentoo	msu-assembly-2013.04.07	' - ami-d6e581bf		Salaat
© 2008 - 2014, Amazon Web Services	, Inc. or its affiliates. All rights reserved. Privacy P	olicy Terms of Use		Feedback
) = E W % P		

Variant calling

The goal of this tutorial is to show you the basics of variant calling using Samtools.

We'll be using data from one of Rich Lenski's LTEE papers, the one on the evolution of citrate consumption in LTEE.

Booting an Amazon AMI

Start up an Amazon computer (m1.large or m1.xlarge) using AMI ami-d25283ba (MSU 2014 QB Summer Course) (see *Start up an EC2 instance* and *Starting up a custom operating system*).

Log in the cloud machine from our local computer with Windows or from Mac OS X.

Logging in

Log in and type:

sudo bash

to change into superuser mode.

Software

Softwares are already installed in the ami loaded. Tools used in this tutorial are:

- bwa (mapping reads to reference genome)
- samtools (a fast and versatile tool for processing DNA alignment in SAM/BAM format.)

Download data

Data are already download in "/mnt" directory:

ls /mnt

to check the data files:

- assembly (directory of data for tomorrow's workshop)
- REL606.fa (reference genome)
- SRR098038.fastq.gz (reads from illumina sequencing)

Do the mapping

Now let's map all of the reads to the reference. Start by indexing the reference genome:

```
cd /mnt
```

bwa index REL606.fa

Now, do the mapping of the raw reads to the reference genome:

bwa aln REL606.fa SRR098038.fastq.gz > SRR098038.sai

Make a SAM file (this would be done with 'sampe' if these were paired-end reads):

bwa samse REL606.fa SRR098038.sai SRR098038.fastq.gz > SRR098038.sam

This file contains all of the information about where each read hits on the reference.

Next, index the reference genome with samtools:

samtools faidx REL606.fa

Convert the SAM into a BAM file:

samtools import REL606.fa.fai SRR098038.sam SRR098038.bam

Sort the BAM file:

samtools sort SRR098038.bam SRR098038.sorted

And index the sorted BAM file:

samtools index SRR098038.sorted.bam

Visualizing alignments

'samtools tview' is a text interface that you use from the command line; run it like so:

```
samtools tview SRR098038.sorted.bam REL606.fa
```

The '.'s are places where the reads align perfectly in the forward direction, and the ','s are places where the reads align perfectly in the reverse direction. Mismatches are indicated as A, T, C, G, etc.

You can scroll around using left and right arrows; to go to a specific coordinate, use 'g' and then type in the contig name and the position. For example, type 'g' and then 'rel606:553093<ENTER>' to go to position 553093 in the BAM file.

Use 'q' to quit.

Counting alignments

This command:

samtools view -c -f 4 SRR098038.bam

will count how many reads DID NOT align to the reference (214518).

This command:

samtools view -c -F 4 SRR098038.bam

will count how many reads DID align to the reference (6832113).

And this command:

gunzip -c SRR098038.fastq.gz | wc

will tell you how many lines there are in the FASTQ file (28186524). Reminder: there are four lines for each sequence.

Calling SNPs

You can use samtools to call SNPs like so:

```
samtools mpileup -uD -f REL606.fa SRR098038.sorted.bam | bcftools view -bvcg - >_

→SRR098038.raw.bcf
```

(See the 'mpileup' docs here.)

Now convert the BCF into VCF:

bcftools view SRR098038.raw.bcf > SRR098038.vcf

You can check out the VCF file by using 'tail' to look at the bottom:

tail *.vcf

Each variant call line consists of the chromosome name (for E. coli REL606, there's only one chromosome - rel606); the position within the reference; an ID (here always '.'); the reference call; the variant call; and a bunch of additional information.

Again, you can use 'samtools tview' and then type (for example) 'g' 'rel606:4616538' to go visit one of the positions. The format for the address to go to with 'g' is 'chr:position'.

You can read more about the VCF file format here.

Questions/discussion items

Why so many steps?

Assembling E. coli sequences with Velvet

The goal of this tutorial is to show you the basics of assembly using the Velvet assembler.

We'll be using data from Efficient de novo assembly of single-cell bacterial genomes from short-read data sets, Chitsaz et al., 2011.

Booting an Amazon AMI

Start up an Amazon computer (m1.large or m1.xlarge) using AMI ami-d25283ba (see *Start up an EC2 instance* and *Starting up a custom operating system*).

Log in with Windows or from Mac OS X.

Logging in

Log in and type:

sudo bash

to change into superuser mode.

Packages to install

Softwares and packages are already installed:

- velvet (An assmbler)
- khmer (A package for preprocess and assemble large data)
- quast (A tools for evaluating assembly with known genomes)

Data

Data is already downloaded in /mnt/assembly:

ls /mnt/assembly

to see data in "/mnt/assembly":

- ecoli_ref-5m-trim.fastq.gz: quality trimmed pair end data sets from E. coli genome sequence data.
- ecoli-reads-5m-dn-paired.fa.gz: data set is a specially processed data set using digital normalization that will assemble quickly.

Running an assembly

Go inside the data directory:

cd /mnt/assembly

Now... assemble the small, fast data sets, using the Velvet assembler. Here we will set the required parameter k=21:

```
velveth ecoli.21 21 -shortPaired -fasta.gz ecoli-reads-5m-dn-paired.fa.gz
velvetg ecoli.21 -exp_cov auto
```

Check out the stats for the assembled contigs for a cutoff of 1000:

```
python /usr/local/share/khmer/sandbox/assemstats3.py 1000 ecoli.*/contigs.fa
```

Also try assembling with k=23 and k=25:

```
velveth ecoli.23 23 -shortPaired -fasta.gz ecoli-reads-5m-dn-paired.fa.gz
velvetg ecoli.23 -exp_cov auto
velveth ecoli.25 25 -shortPaired -fasta.gz ecoli-reads-5m-dn-paired.fa.gz
velvetg ecoli.25 -exp_cov auto
```

Now check out the stats for the assembled contigs for a cutoff of 1000:

python /usr/local/share/khmer/sandbox/assemstats3.py 1000 ecoli.*/contigs.fa

(Also read: What does k control in de Bruijn graph assemblers?.)

Comparing and evaluating assemblies - QUAST

Run QUAST to mapping contigs (ecoli.23/contigs.fa, and ecoli.25/contigs.fa) reference genome (ecoliMG1655.fa):

gunzip ecoliMG1655.fa.gz
/root/quast-2.3/quast.py -R ecoliMG1655.fa ecoli.*/contigs.fa

Note that here we're looking at *all* the assemblies we've generated.

Now look at the results:

more quast_results/latest/report.txt

The first bits to look at are Genome fraction (%) and # misassembled contigs, I think.

Searching assemblies – BLAST

Build BLAST databases for the assemblies you've done:

```
cd /mnt/assembly
for i in 21 23 25
do
    extract-long-sequences.py -o ecoli-$i.fa -l 500 ecoli.$i/contigs.fa
    formatdb -i ecoli-$i.fa -o T -p F
done
```

BLAST is already installed. Let's search for a specific gene (CRP, a transcription regulator).

The commandline for BLAST is:

blastall -i crp.fa -d ecoli-21.fa -p tblastn -b 1 -v 1

Questions and Discussion Points

Why do we use a lower cutoff of 1kb for the assemstats3 links, above? Why not 0?

CHAPTER 2

Indices and tables

- genindex
- modindex
- search