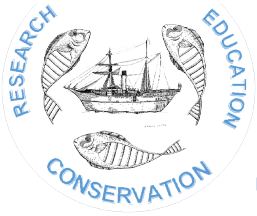
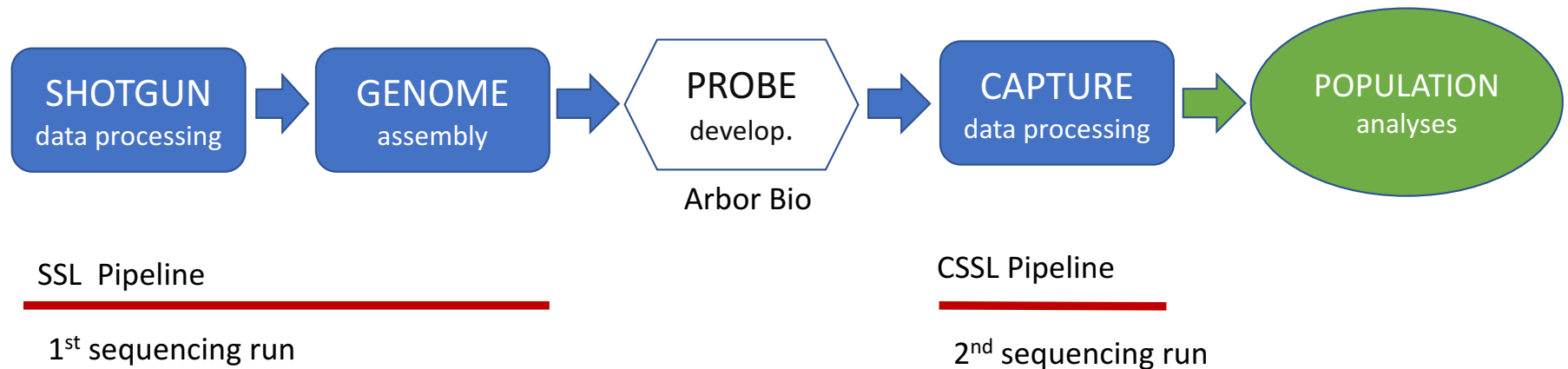


The Philippines PIRE Project



# The Philippines PIRE PIPELINE

by Dr. Eric Garcia



# Organism's Genome



## Shearing/Digestion

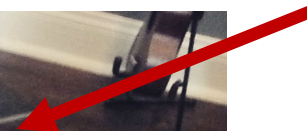




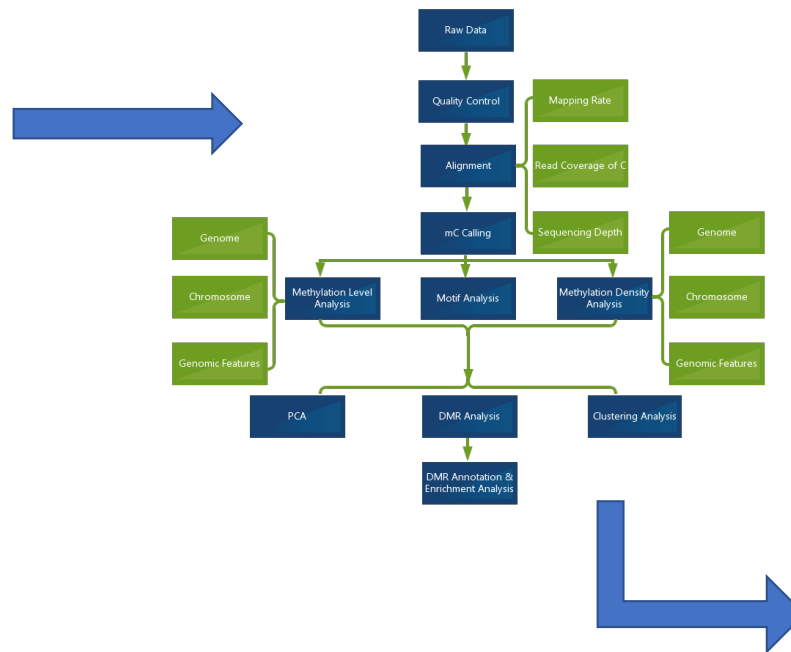
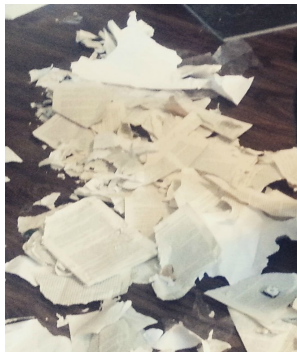
**Shearing/Digestion**



**Shotgun  
boy**



# PIPELINES: making sense of genomic data



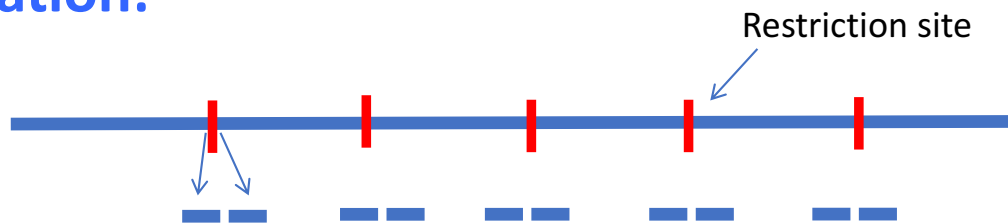
# PPP Pipeline

---

## RAD library preparation:

DNA Prep

Sbf1 genome  
digestion



# PPP Pipeline - SSL

---

## Shotgun library preparation:

DNA Prep

Shotgun



# PPP Pipeline - SSL

## Shotgun library preparation:

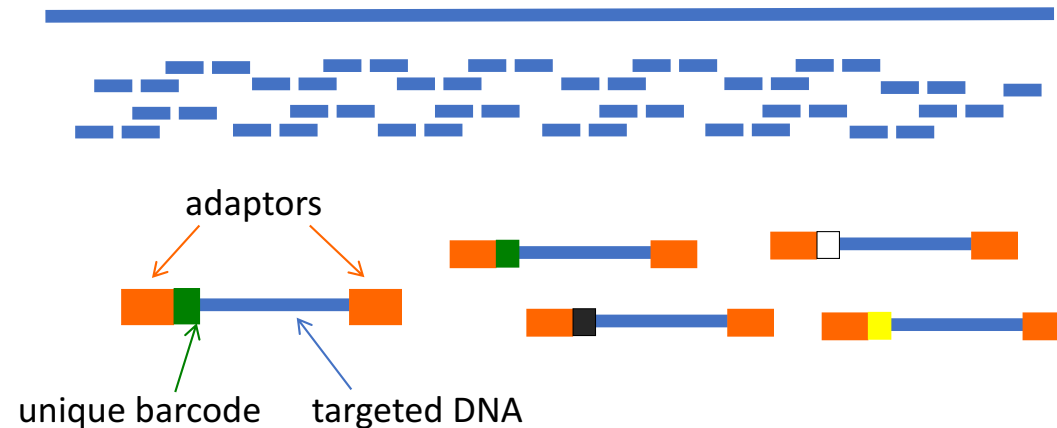
DNA Prep

Shotgun

Lib. Prep

Ligation of Adaptors  
and unique barcode

1<sup>st</sup> Sequencing Run



Next Generation, high throughput *illumina* sequencing

r1 - forward

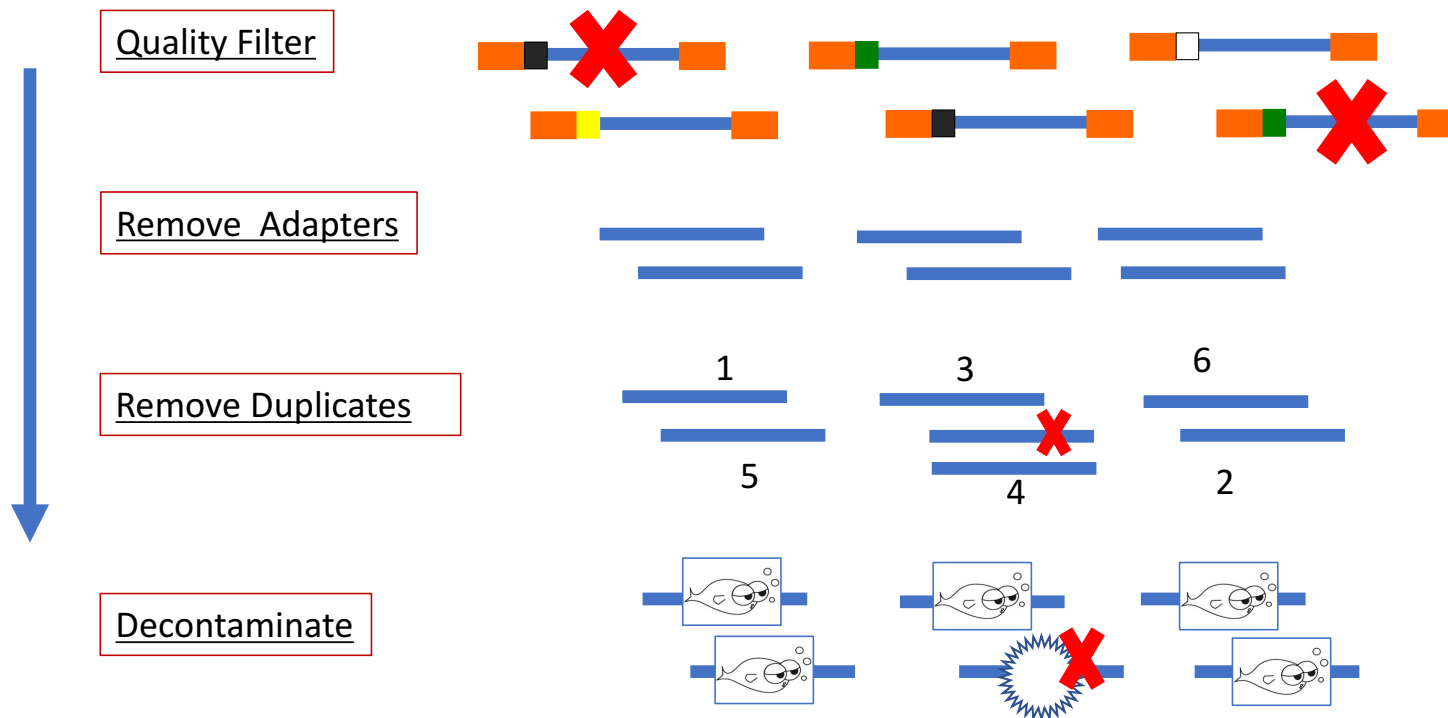


Pair-end Sequencing



# PPP Pipeline - SSL

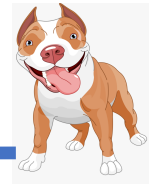
## Bioinformatics: fq pre-processing



# PPP Pipeline - SSL

---

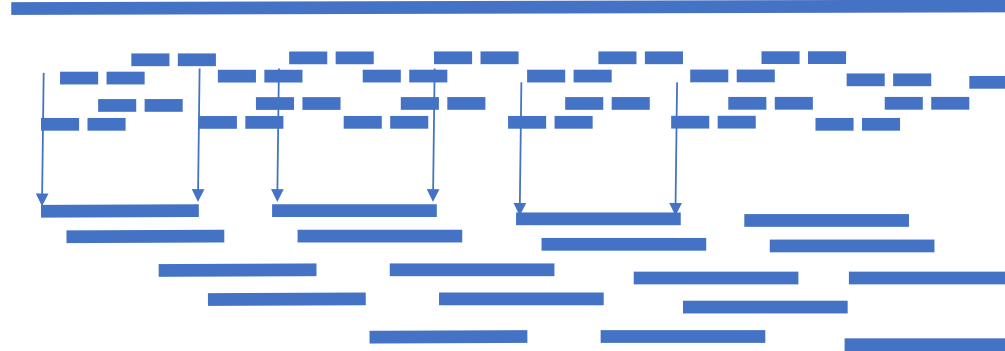
Genome



reads

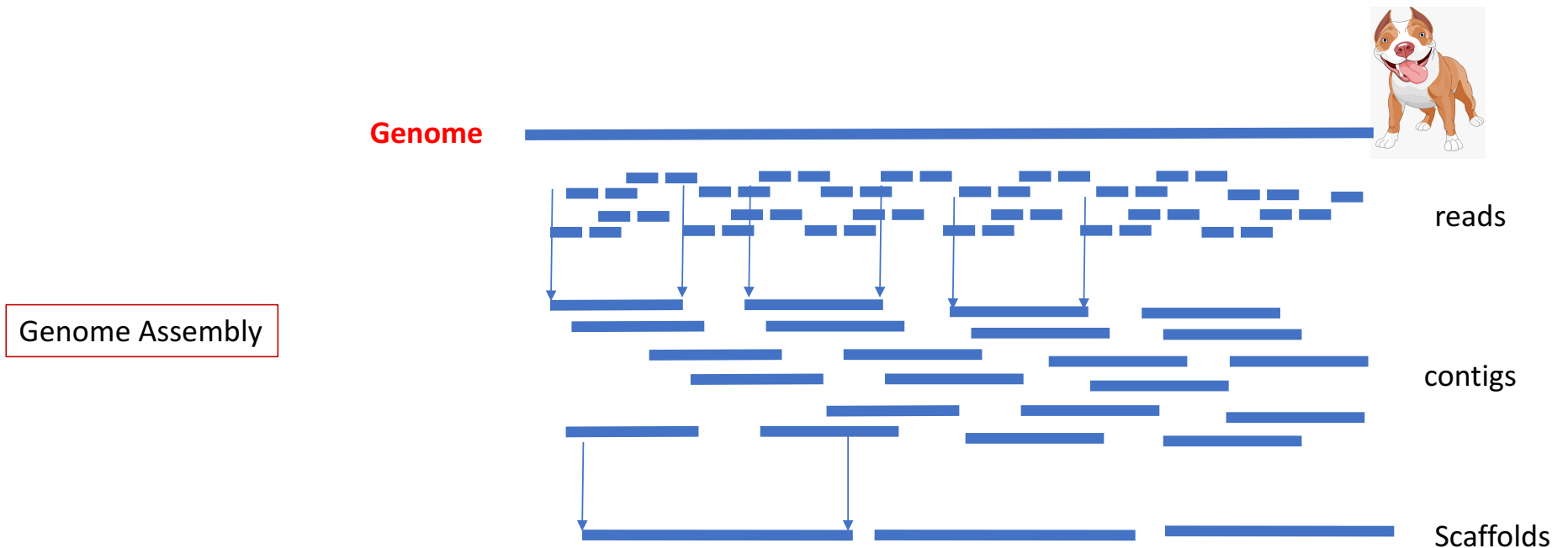
Genome Assembly

contigs



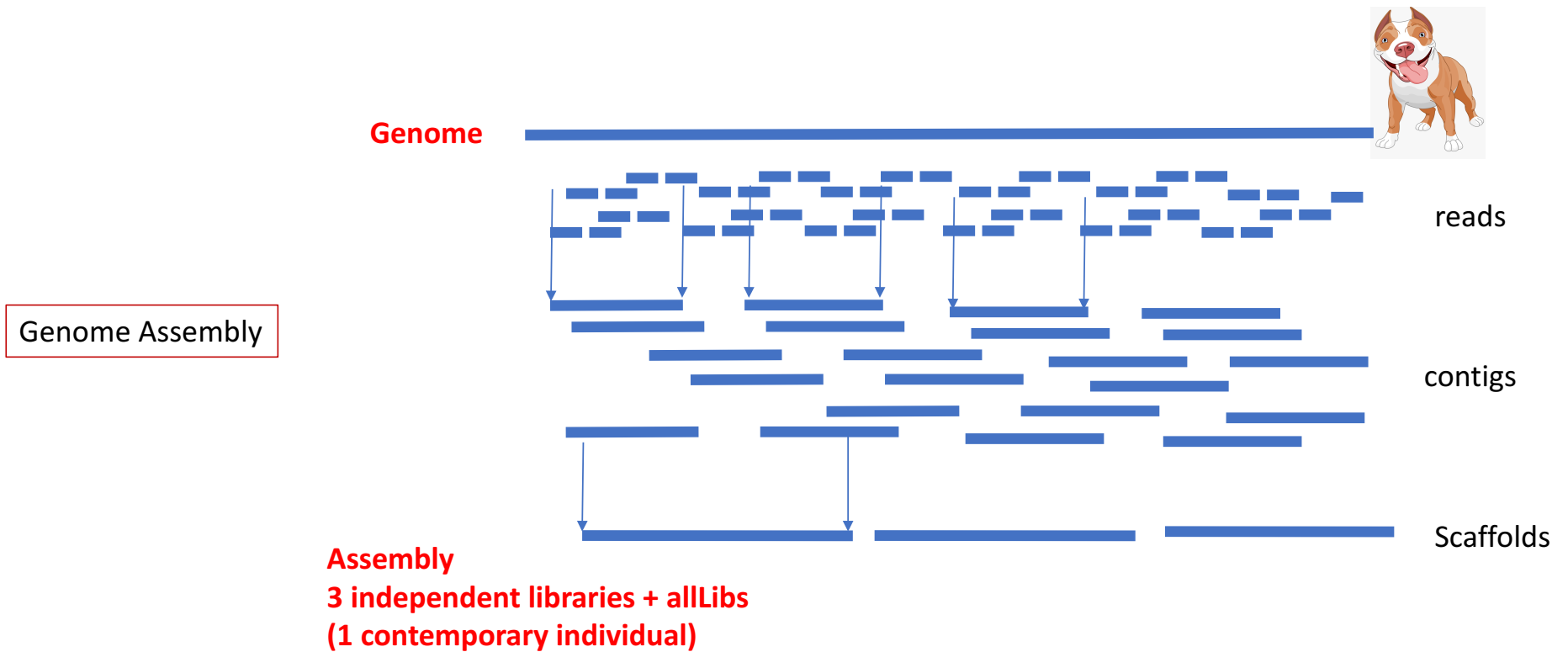
# PPP Pipeline - SSL

---



# PPP Pipeline - SSL

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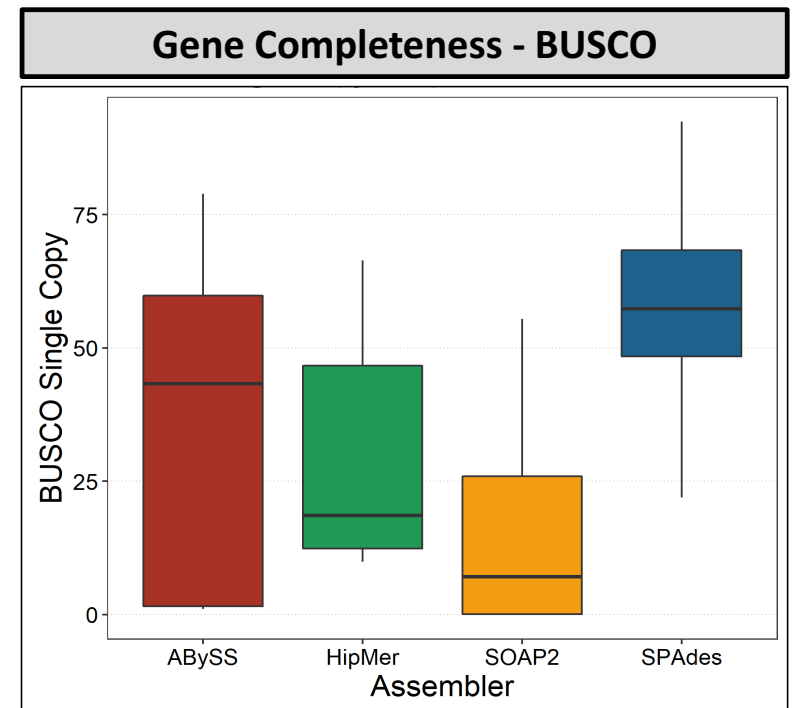
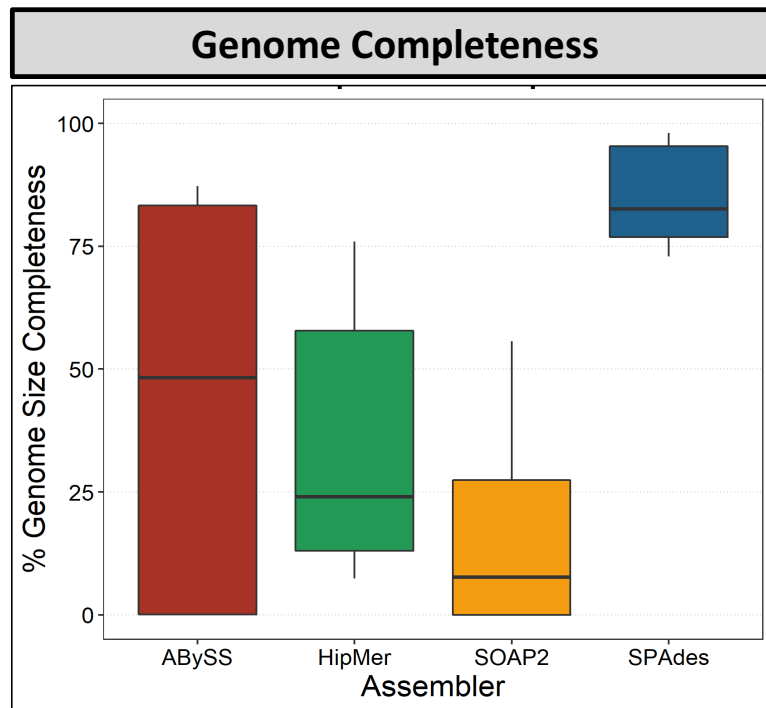
# PPP Pipeline - SSL

## Bioinformatics | assembler comparison

### QUALITY

Assembler

- ABYSS
- HipMer
- SOAP2
- SPAdes



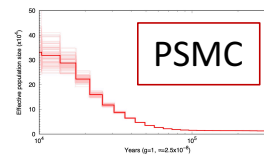


# PPP Pipeline - SSL

Best assembly

QUAST

BUSCO



mtDNA

Probe Development



# PPP Pipeline - SSL

Best assembly

QUAST

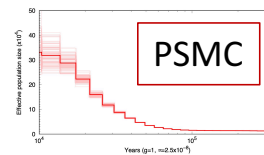
BUSCO

Probe Development

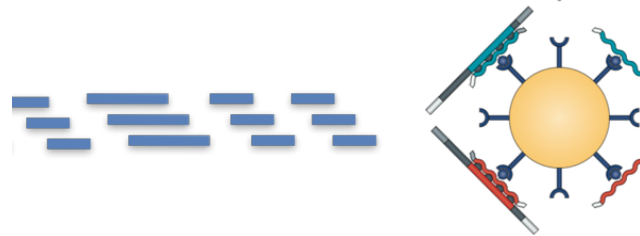
Capture

all individuals  
historical and contemporary

2<sup>nd</sup> illumina Sequencing Run



mtDNA



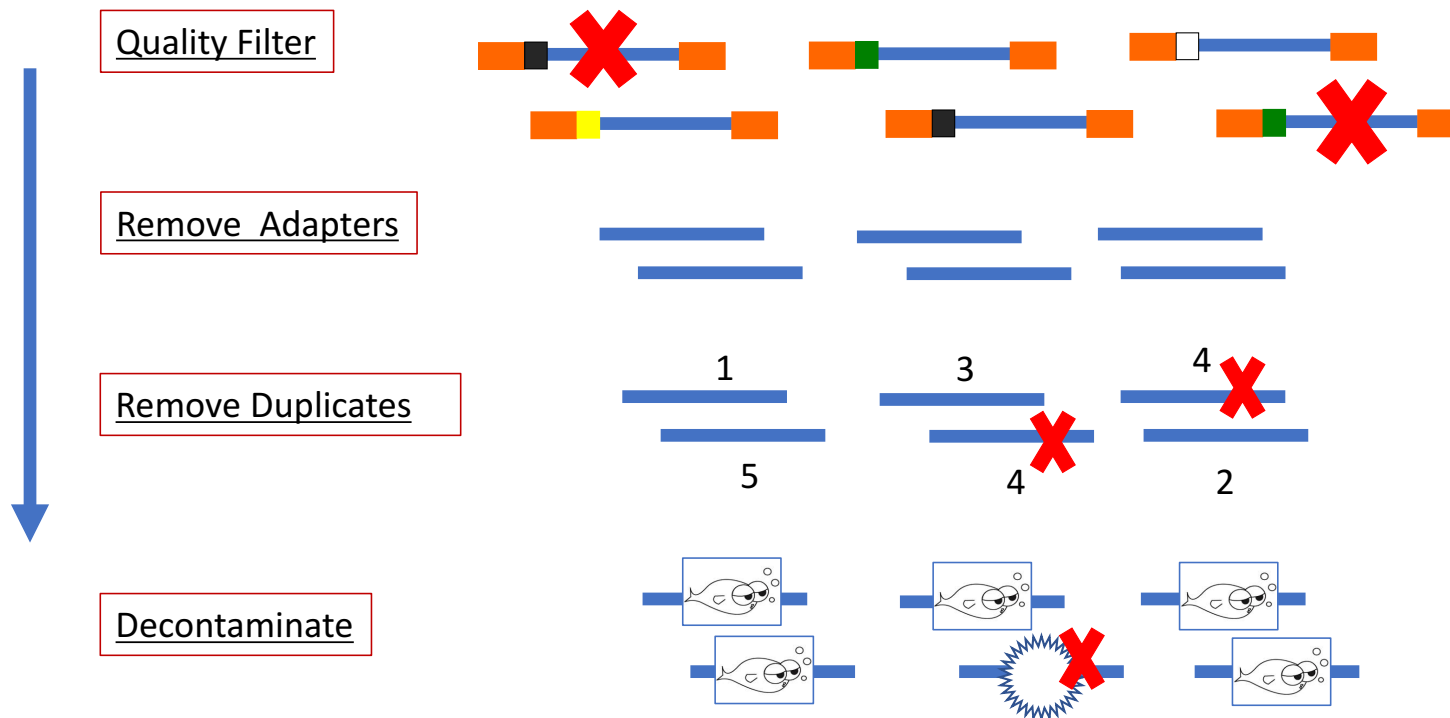
r1 - forward

r2 - reverse

Pair-end Sequencing

# PPP Pipeline - CSSL

## Bioinformatics: fq pre-processing



# PPP Pipeline - CSSL

## Bioinformatics: fq pre-processing

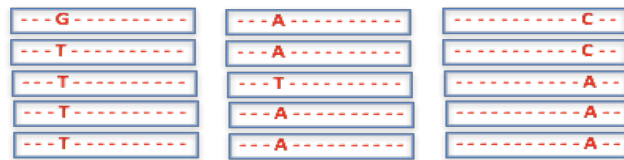
Quality Filter

Remove Adapters

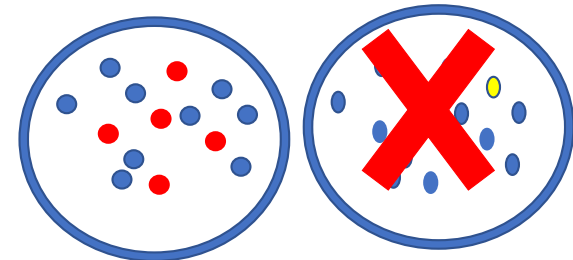
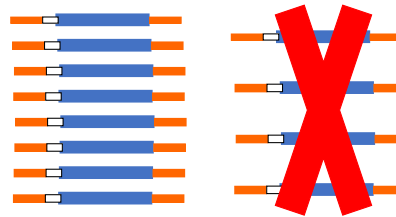
Remove Duplicates

Decontaminate

Genotyping



Depth Filters  
Missing data Filters  
Individual Filters  
Population Filters



## Population analyses

