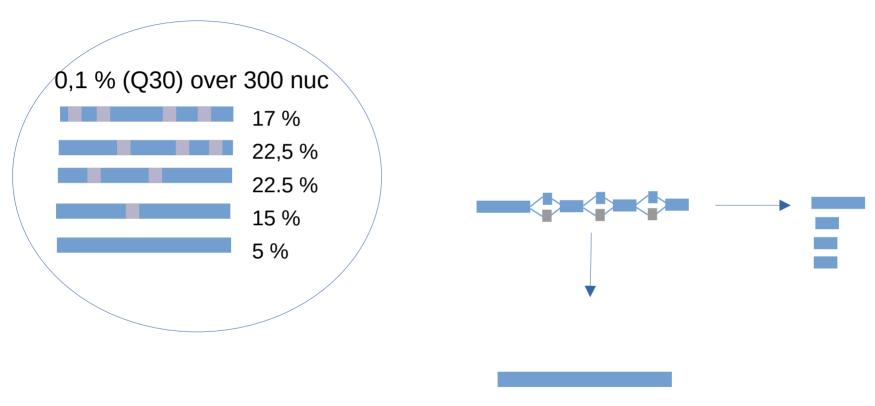
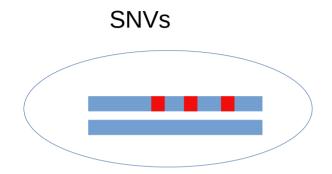
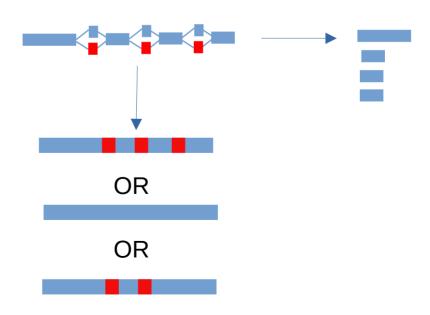


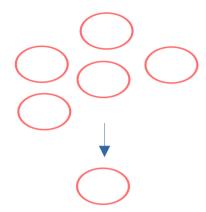
Sequencing errors



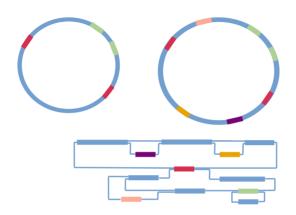




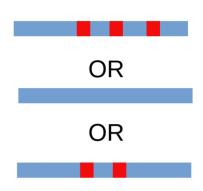
### **Summary**



Dereplication of sequence at the scale of the read length

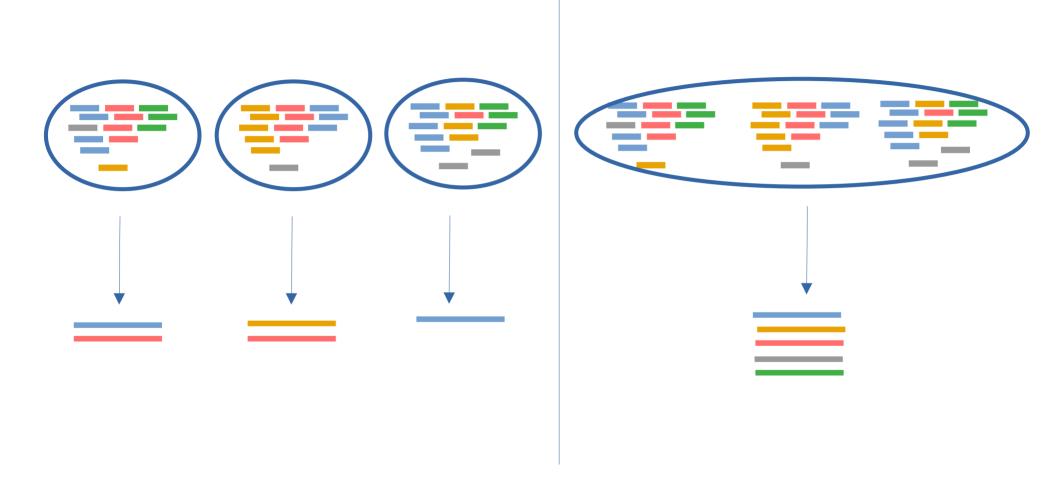


Strain diversity increase complexity → smaller contigs



Assembly does not conserve strain level informations

# **Single Sample Assembly VS Assembly**



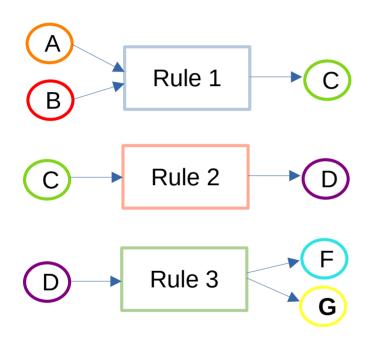
# **Single Sample Assembly VS Assembly** Strain Strain Strain2 Strain2

Median = 40 N50 ~ 80

Total length = 340 Half = 170

# Input

# Output

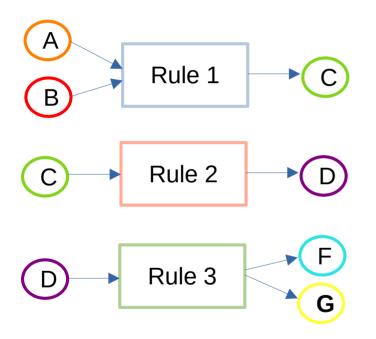


## Request: generate G

```
→ G exists ?
```

# Input

# Output



### **Request:** generate G

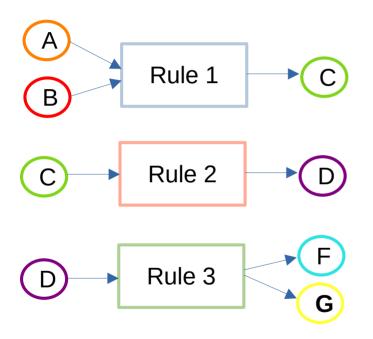
→ G exists? Can it be generated? No, Rule3 can

#### Schedule:

Rule 1

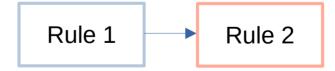
# Input

#### **Output**



#### Request: generate G

- → G exists ? Can it be generated ? No, Rule3 can
- → Rule 3 needs D, exists, generated ? Doesn't exist, Rule 2 can



# Input **Output** Rule 1 Rule 2 Rule 3

G

#### Request: generate G

- → G exists ? Can it be generated ? No, Rule3 can
- → Rule 3 needs D, exists, generated ? Doesn't exist, Rule 2 can
- → Rule 2 needs C, exists, generated ? Doesn't exist, Rule 1 can



# Input **Output** Rule 1 Rule 2 Rule 3 G

#### Request: generate G

- → G exists ? Can it be generated ? No, Rule3 can
- → Rule 3 needs D, exists, generated ?

  Doesn't exist, Rule 2 can
- → Rule 2 needs C, exists, generated ?

  Doesn't exist, Rule 1 can
- → Rule 1 needs A,B, exist, generated ? YES!



#### rule Hello\_world:

input: "/home/ubuntu/requirement.txt"

output: "{path}/snakemake.txt"

shell: "echo HELLO WORLD > {output}"

#### rule Hello\_world:

input: "{path}/requirement.txt"
output: "{path}/snakemake.txt"

shell: "echo HELLO WORLD > {output}"

#### rule Hello world:

input: "/home/ubuntu/requirement.txt"

output: "{macha\_green\_tea}/snakemake.txt"

shell: "echo HELLO WORLD > {output}"

