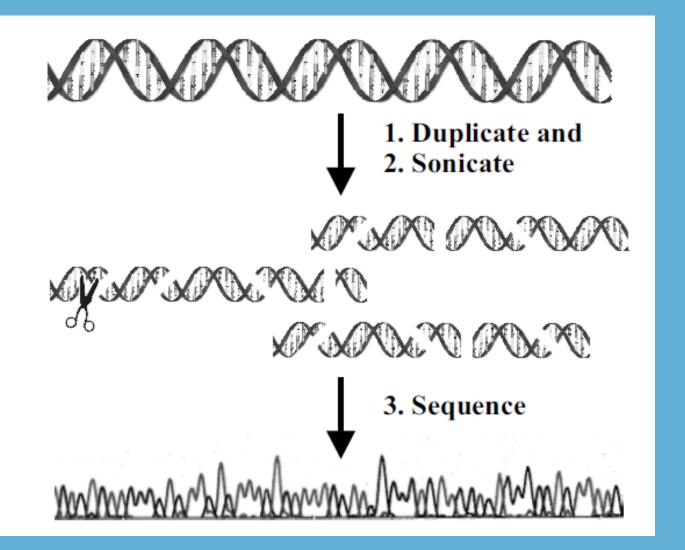


Sameer Ravi and Jesun Sahariar Firoz



### **DNA Sequencing Process**





### DNA Sequencing Process (cont.)



CCGTAGCCGGGATCCCGTCC
CCCGAACAGGCTCCCGCCGTAGCCG
AAGCTTTTCTCCCGAACAGGCTCCCG

## Output of DNA FAP

6. Call Consensus

AAGCTTTTCTCCCGAACAGGCTCCCGCCGTAGCCGGGATCCCGTCC

Figure. : Graphical representation of DNA sequencing and assembly



- Behavior of a Honey Bee Swarm:
  - ✓ Food Sources
  - ✓ Employed foragers
  - ✓ Unemployed foragers: scouts and onlookers



#### Algorithm 1 Generic ABC Algorithm

- Initialize potential food sources for employed bees.
- 2: while Requirements are not met do
- 3: Each employed bee goes to a food source in her memory and determines a neighbour source, then evaluates its nectar amount and dances in the hive
- 4: Each onlooker watches the dance of employed bees and chooses one of their sources depending on the dances, and then goes to that source. After choosing a neighbour around that, she evaluates its nectar amount.
- Abandoned food sources are determined and are replaced with the new food sources discovered by scouts.
- 6: The best food source found so far is registered
- 7: end while



#### • Initialization:

- √ Food Source => A permutation of fragments
- ✓ Generated randomly (No seeding technique involved)

#### Calculate Fitness:

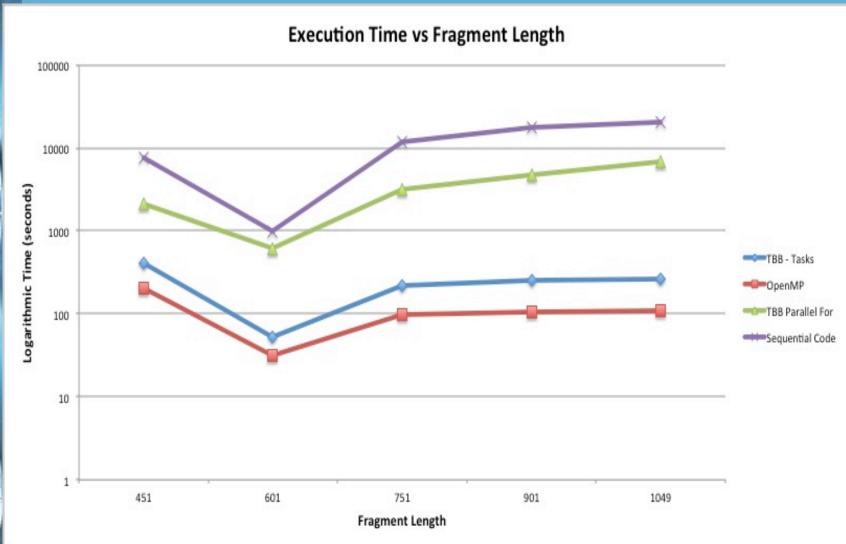
✓ By summing up the overlap amount of consecutive fragments.

$$F_{\mu} = \sum_{i=0}^{n-2} w(f[i], f[i+1]),$$



- Memorize the best solution.
- Send Employer bees
  - ✓ Estimate nectar amount => evaluate fitness
  - ✓ Modify the solution using Problem aware local search (PALS).

## **Experimental Results**





## Questions?