## **Clonorchis sinensis Genome**

Introduction to genomics (485 course)

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## <u>Outline</u>

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- The organism's : Taxonomy

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- Structural and Functional Annotation
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#### **The Selected Paper**

Wang et al. Genome Biology 2011, 12:R107 http://genomebiology.com/2011/12/10/R107



#### RESEARCH

**Open Access** 

## The draft genome of the carcinogenic human liver fluke *Clonorchis sinensis*

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#### Abstract

**Background:** Clonorchis sinensis is a carcinogenic human liver fluke that is widespread in Asian countries. Increasing infection rates of this neglected tropical disease are leading to negative economic and public health consequences in affected regions. Experimental and epidemiological studies have shown a strong association between the incidence of cholangiocarcinoma and the infection rate of *C. sinensis*. To aid research into this organism, we have sequenced its genome.

## **Clonorchis sinensis Taxonomy**



#### Origin of the name



#### **Geographical distribution**



**Trends in Parasitology** 

Named as **the Chinese OR the Oriental liver Fluke** because it orients to affect the Asian countries such as Korea , China, Taiwan and northern Vietnam

#### **Clonorchis sinensis physical Characteristics**

- Flat, leave shaped with slightly elongated and slender body
- length VS width = 15-20 mm VS 3-4 mm



## **Clonorchis sinensis physical Characteristics**

- Small, sharply curved with a clear lid-like structure named as operculum at the narrower end and a stem-like knob at the broader end.
- Miracium = larva =not free = trapped inside the fertilized egg



#### C.sinensis Life Cycle



#### Significance of studying C.sinensis 's genome

- Clonorchis sinensis is widespread and affect more than 35 million people in Asian countries.
- The Increasing infection rates of this neglected tropical disease are leading to negative economic and public health consequences in affected regions.
- Also, experimental and epidemiological studies have shown a strong association between the incidence of cholangiocarcinoma and the infection rate of C. sinensis.





#### Sample taken to sequence C.sinensis 's genome

 Adult C. sinensis flukes were isolated from cat livers (Henan Province, China) and then a single adult fluke was chosen for DNA genomic extraction.





### Sequencing method



#### First chemistry cycle: determine first base

To initiate the first sequencing cycle, add all four labeled reversible terminators, primers, and DNA polymerase enzyme to the flow cell.



#### Image of first chemistry cycle

After laser excitation, capture the image of emitted fluorescence from each cluster on the flow cell. Record the identity of the first base for each cluster.

#### Before initiating the next chemistry cycle

The blocked 3' terminus and the fluorophore from each incorporated base are removed.



#### Sequence read over multiple chemistry cycles

Repeat cycles of sequencing to determine the sequence of bases in a given fragment a single base at a time.

#### **Sequencing strategy** = Whole genome Shotgun



### <u>Genome Assembly</u>

#### Table 1 Summary of the C.sinensis genome assembly

	Total length (Mb)	Number	N50 <sup>a</sup> (bp)	N90 <sup>a</sup> (bp)	Longest (bp)
Contig <sup>b</sup>	515.56	60,796	14,708	4,079	137,874
Scaffold <sup>b</sup>	516.46	31,822	30,195	7,299	238,094
Super-scaffold <sup>b</sup>	516.47	26,446	42,632	8,441	400,764

- Genome coverage : 5000x
- Genome size : 516 Mb
- Number of chromosome : (2n=56) 20 pairs of small chromosomes

20 pairs of small chromosomes8 pairs of large chromosomes

- N50 contig :14,708 bp (15 kb)
- Total no. of contigs : 60,796 contigs
- N50 scaffolds: 30,195 bp (30kb)
- Total no of scaffolds: 31,822 scaffold
- MORE contigs = smaller in size / LESS Scaffolds = large in size

#### **Structural Annotation**



## <u>Functional</u> <u>Annotation</u>

 The major protein domains of
 C.sinensis are
 shared with other
 taxa and C. sinensis
 has the fewest
 unique domains.



## **Interesting information**

- Adult **liver flukes** can produce up to **4000 eggs** per day for at least six months.
- Worms can live up to **8 years** in humans.
- An infected person may pass viable eggs up to **30 years**.
- A human host with an average infection will have **two or three dozen** worms; heavily infected individuals have been found with as many as **20,000 worms**.

## **Interesting information**

#### How can I prevent *Clonorchis* infection?

1. Avoiding raw or undercooked freshwater fish or even lightly salted, smoked, or pickled fish.

2. The FDA recommendations :

#### Cooking

- Cook fish adequately (to an internal temperature of at least 145° F [~63° C]).
- •Freezing
  - At -4°F (-20°C) or below for at least 7 days (total time)
  - At -31°F (-35°C) or below until solid.

## <u>Questions</u>



• What is the common name of *Clonorchis sinensis*?

• What is the name of the sequencing method used ?

## THANK YOU FOR LISTENING TO MY PRESENTATION

# I HOPE YOU LIKED IT