

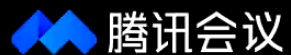
BGI华大

文献阅读与科研展示

第三组-王硕 吕志远 赵万东 孙士瑶 吕丰源 苏光烨

2021.10.09

1. 背景介绍
2. 文献讲解-Roche公司的454技术（合成法）
3. 文献讲解-ABI公司的SOLiD技术（连接法）



vuan的快速会议

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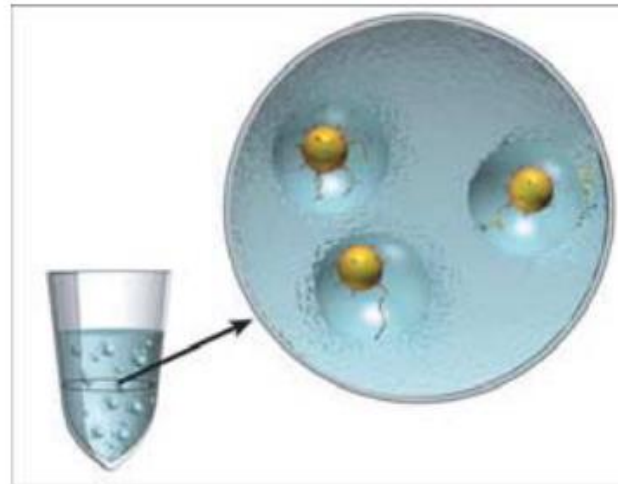
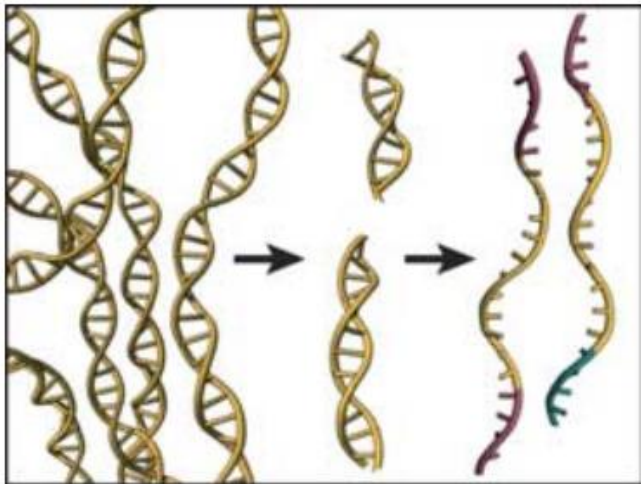
ARTICLES

Genome sequencing in microfabricated high-density picolitre reactors

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Sample preparation

top left

Genomic DNA is isolated, fragmented, ligated to adapters and separated into single strands

top right

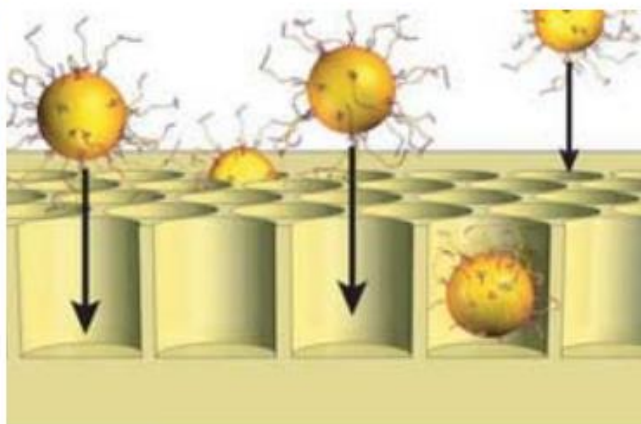
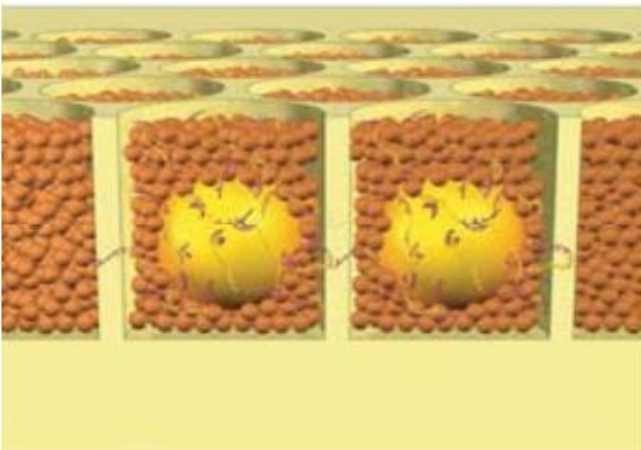
Emulsion-based PCR

bottom right

The emulsion is broken, the DNA strands are denatured, and beads carrying single-stranded DNA clones are deposited into wells of a fibre-optic slide

bottom left

Smaller beads carrying immobilized enzymes required for pyrophosphate sequencing are deposited into each well

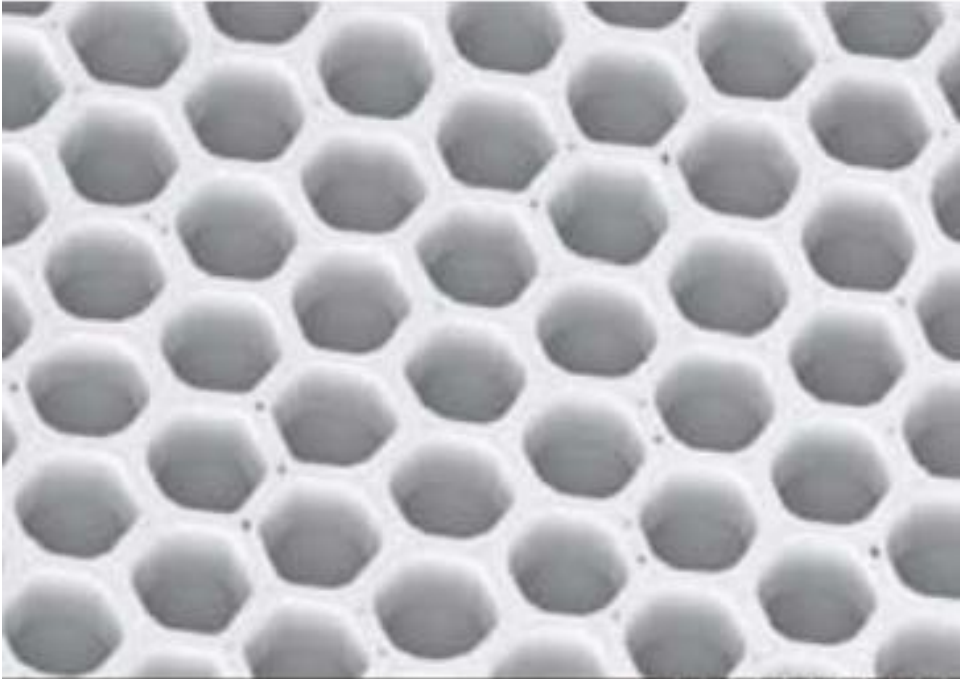




Microscope photograph of emulsion showing droplets containing a bead and empty droplets. The thin arrow points to a 28- μ m bead; the thick arrow points to an approximately 100- μ m droplet.



Scanning electron micrograph of a portion of a fibre-optic slide, showing fibre-optic cladding and wells before bead deposition.



Fibre-optic core diameter : 44 μm

Fibre-optic core cladding : 2–3

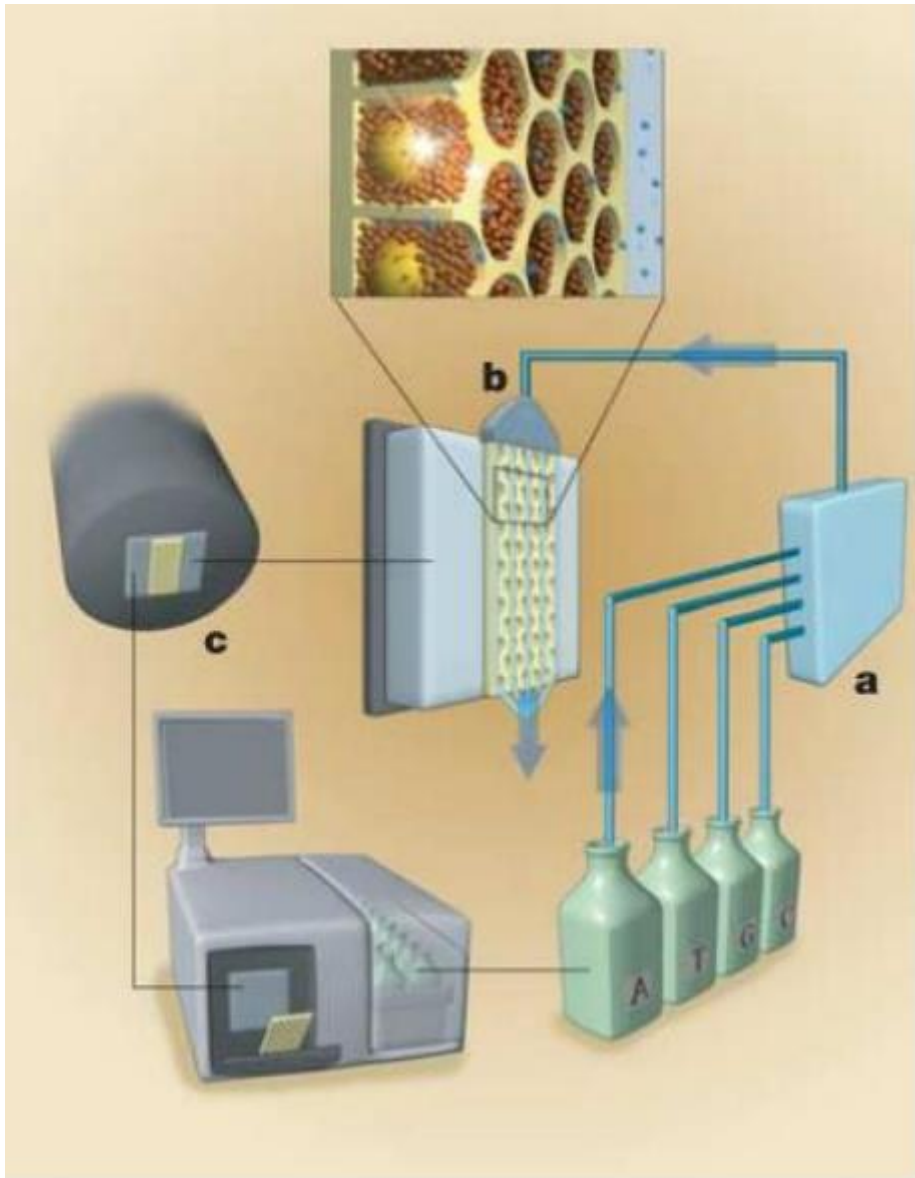
Fibre-optic core depth : 55 μm

Fibre-optic core centre-to-centre distance : 50 μm

The well density 480 wells μm^{-2}

The slide contain 1.6 million wells

The flow chamber designed to create a 300- μm high channel



Sequencing instrument.

(a) 流体组件

(b) 一个流动室，包括含有孔的光纤载玻片

(c) A CCD 电荷耦合器件传感器 camera-based imaging assembly



Accurate Multiplex Polony Sequencing of an Evolved Bacterial Genome

Jay Shendure *et al.*

Science **309**, 1728 (2005);

DOI: 10.1126/science.1117389

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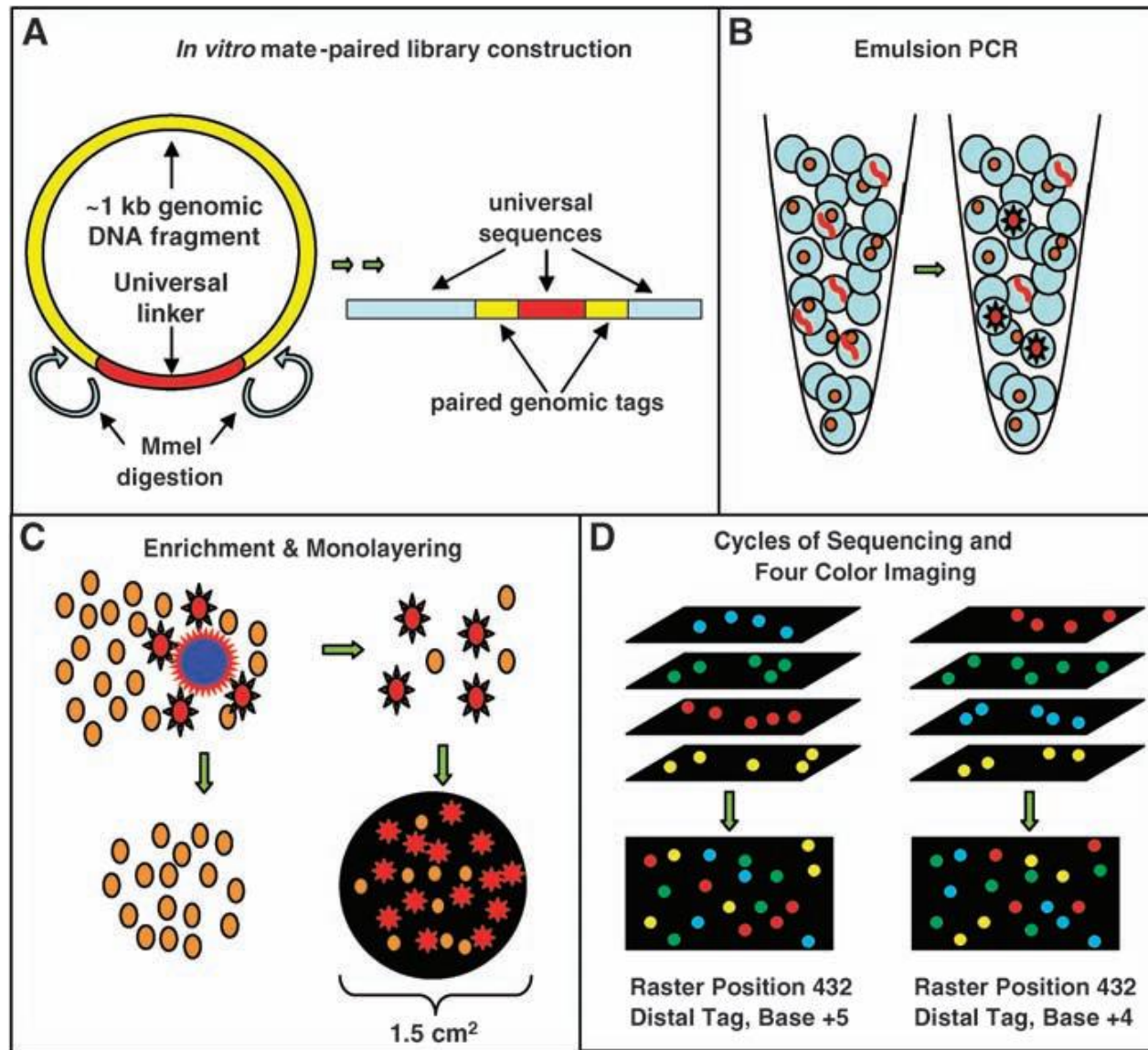
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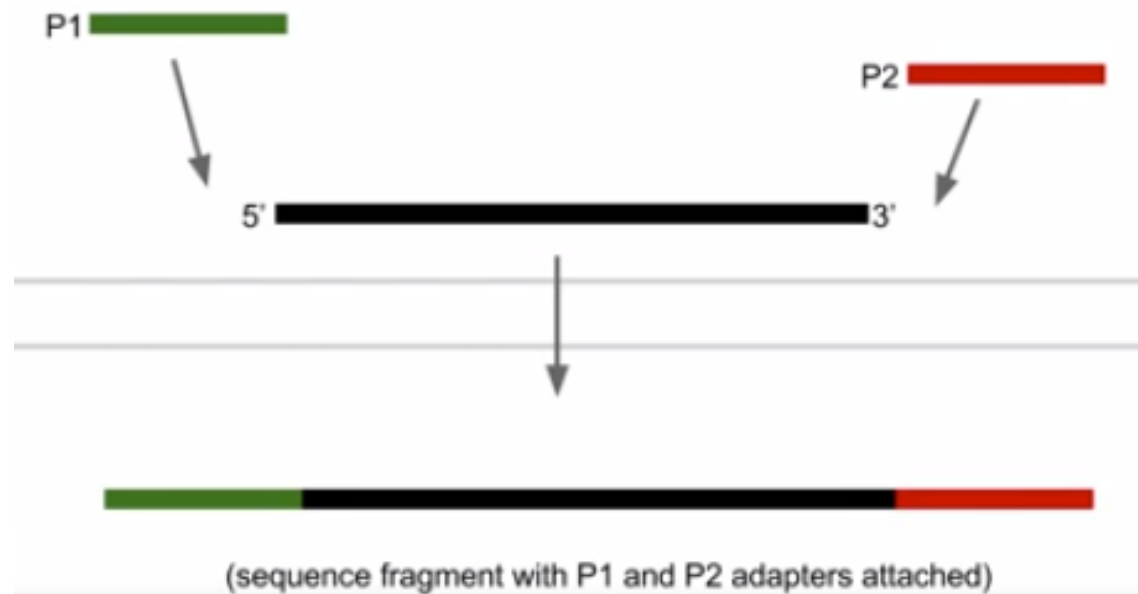
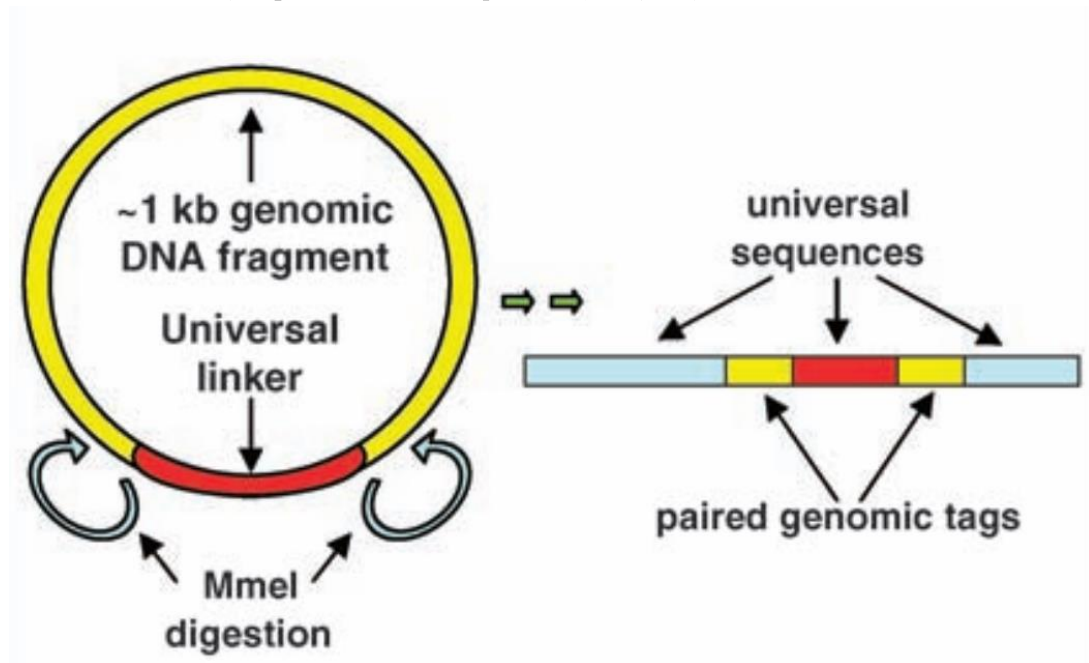
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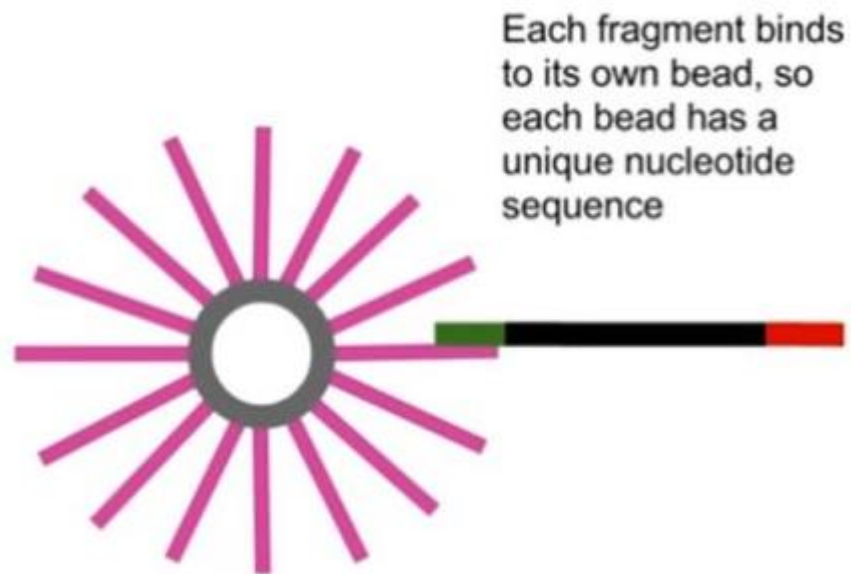
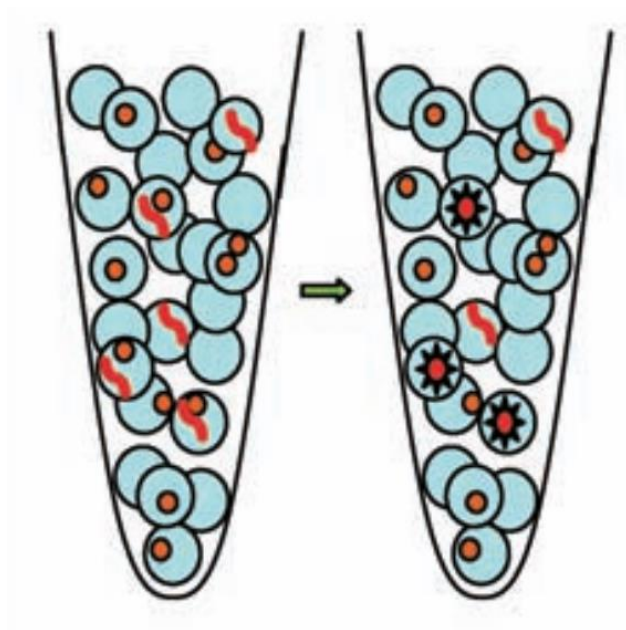
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SOLiD流程1 打断，构建文库

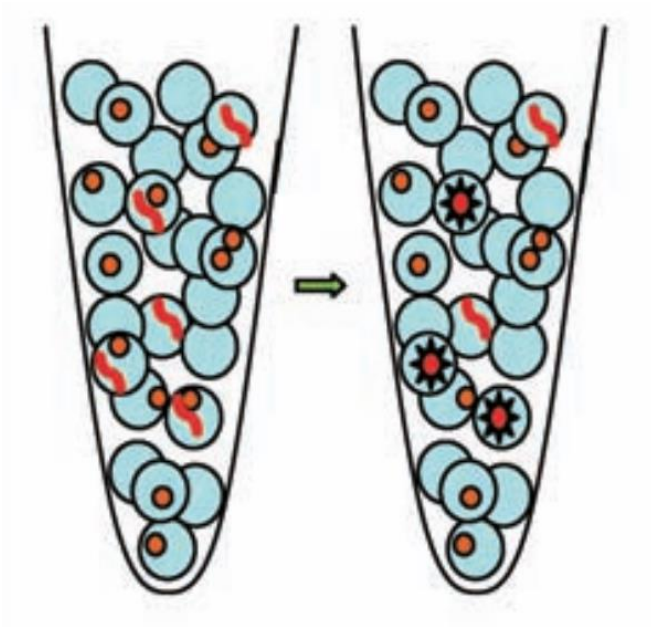


SOLiD流程



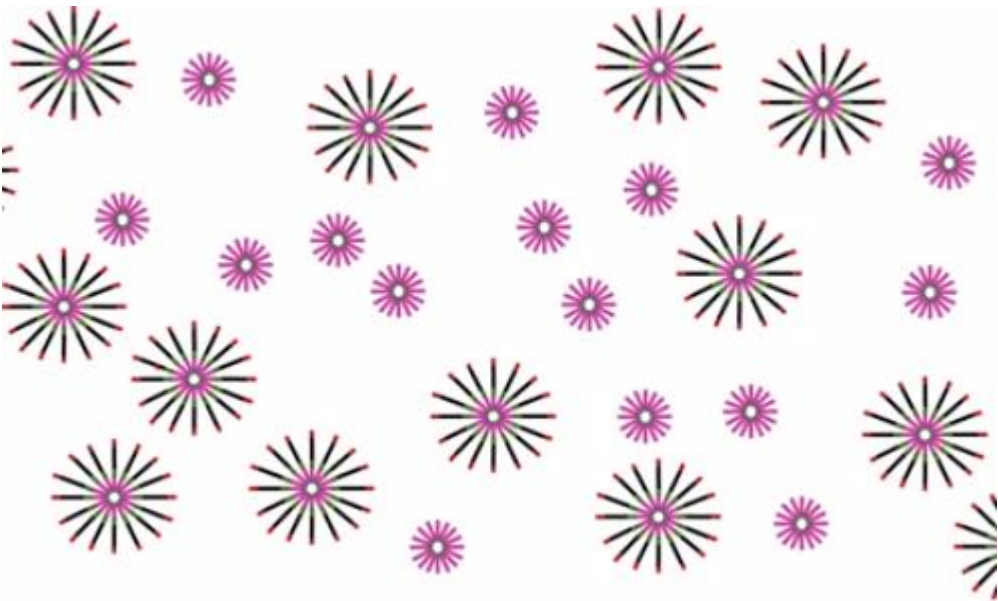
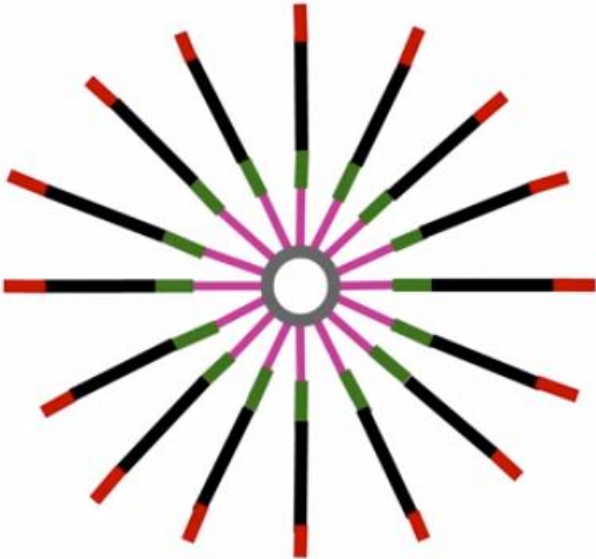
- fragment可以结合特定bead，不是所有bead都有结合的fragment
- 结合有fragment的bead，通过离心筛选出来，然后结合到glass slide

SOLiD流程

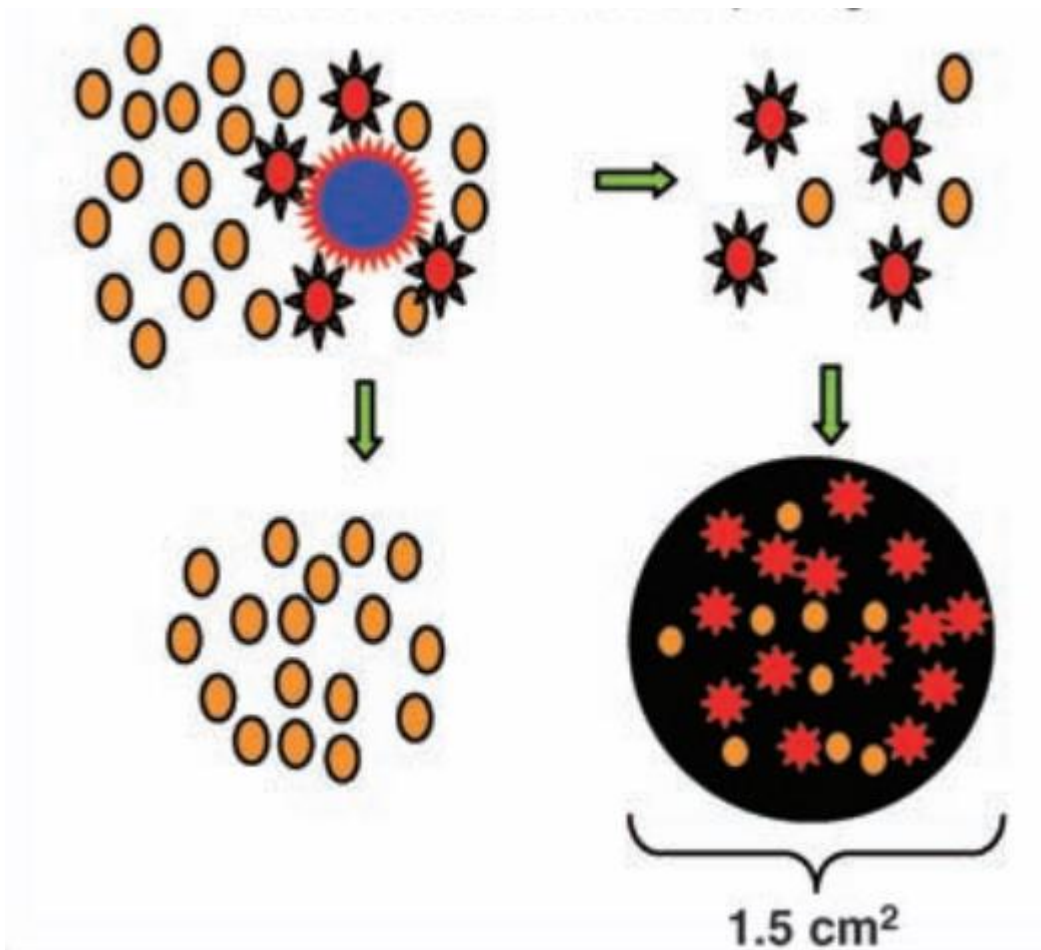


PCR

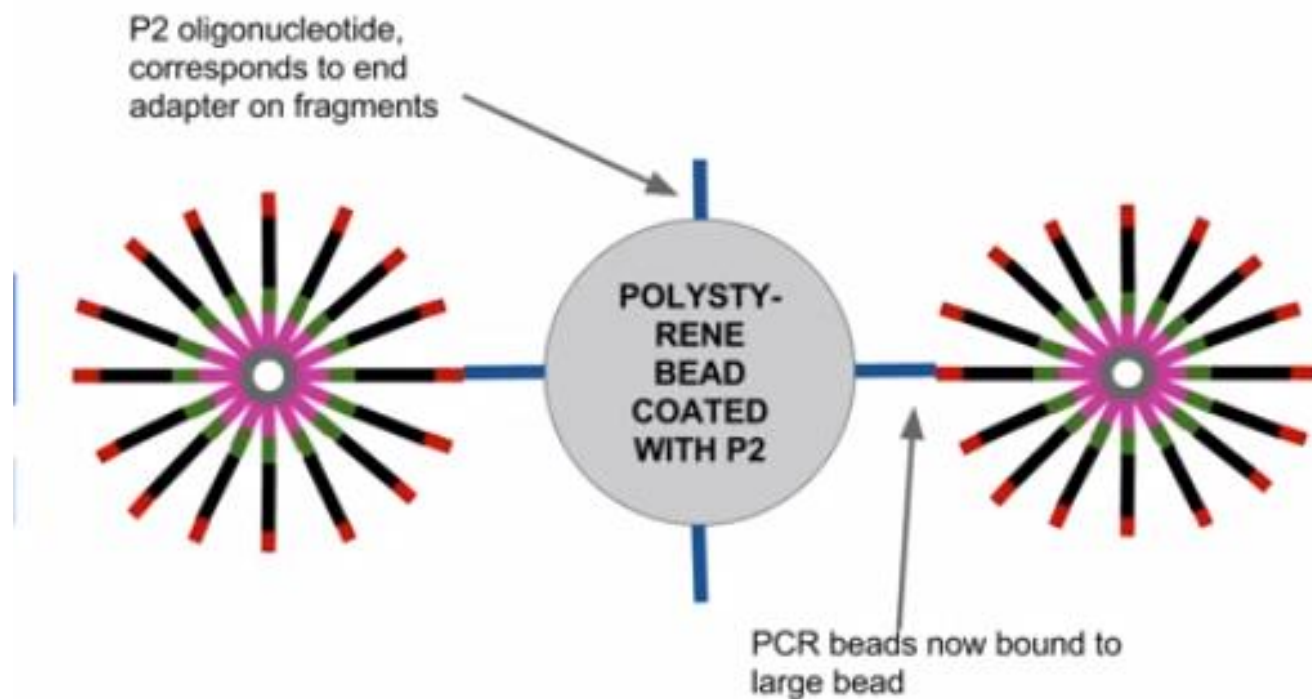
Millions of copies of template strand on each bead



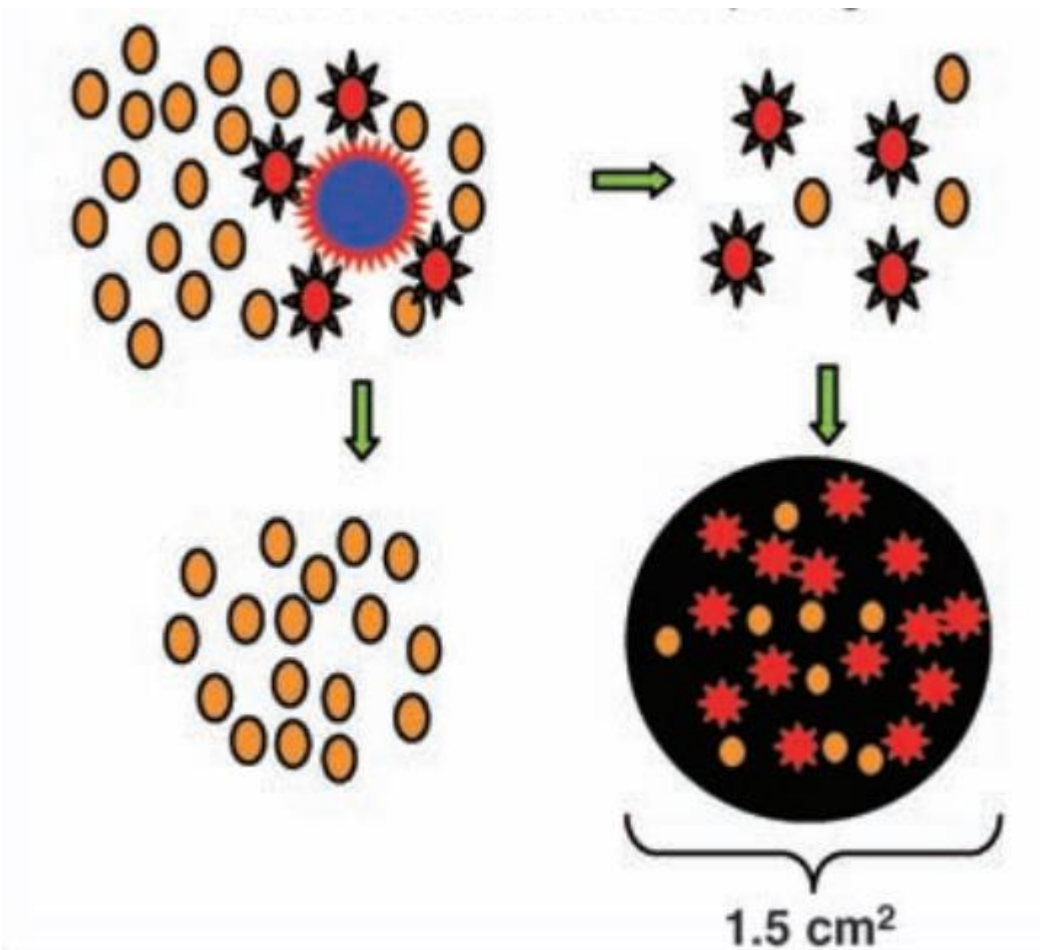
SOLiD流程



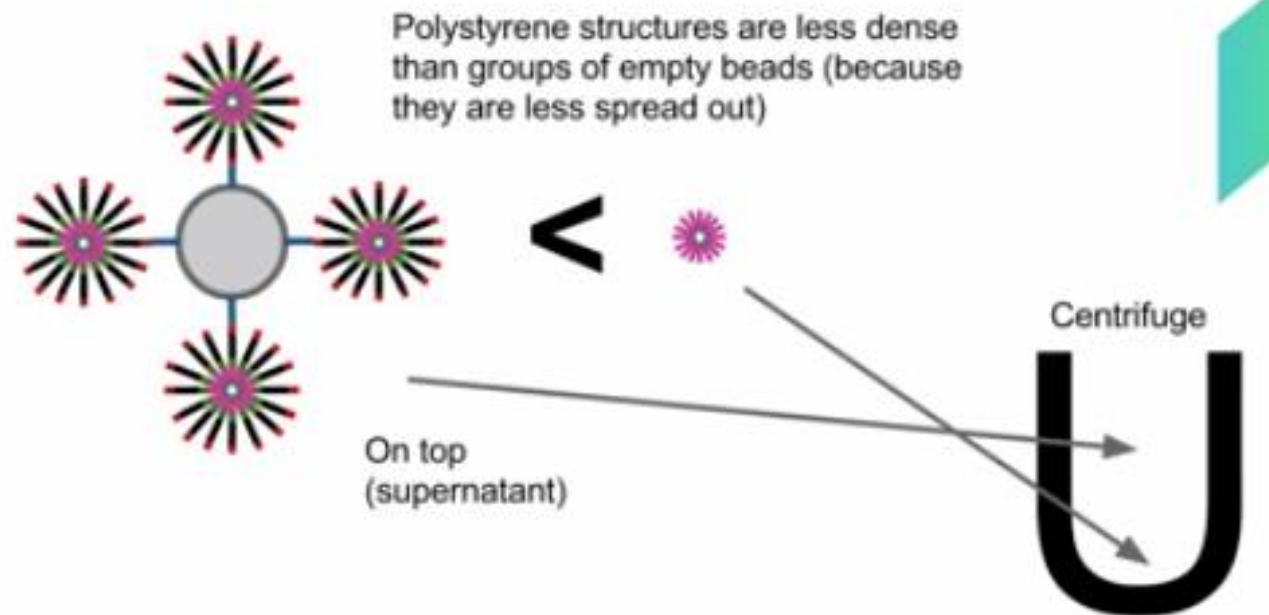
First, gather all the good beads...



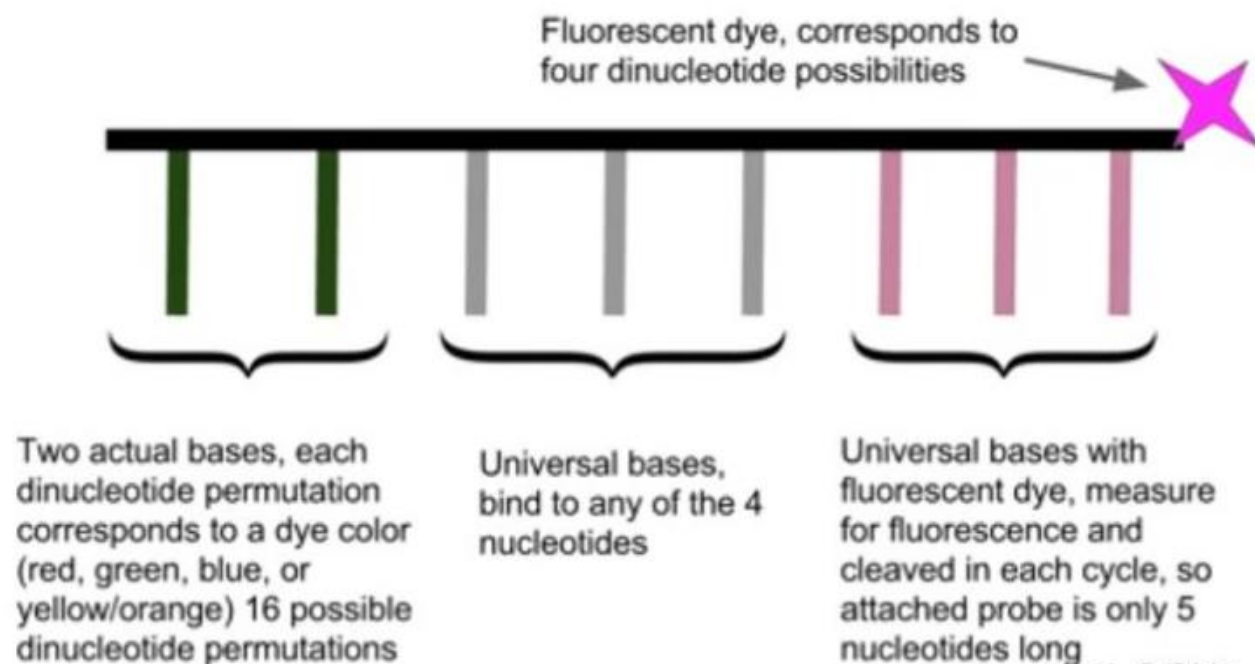
SOLiD流程



Then centrifuge out the bad ones!

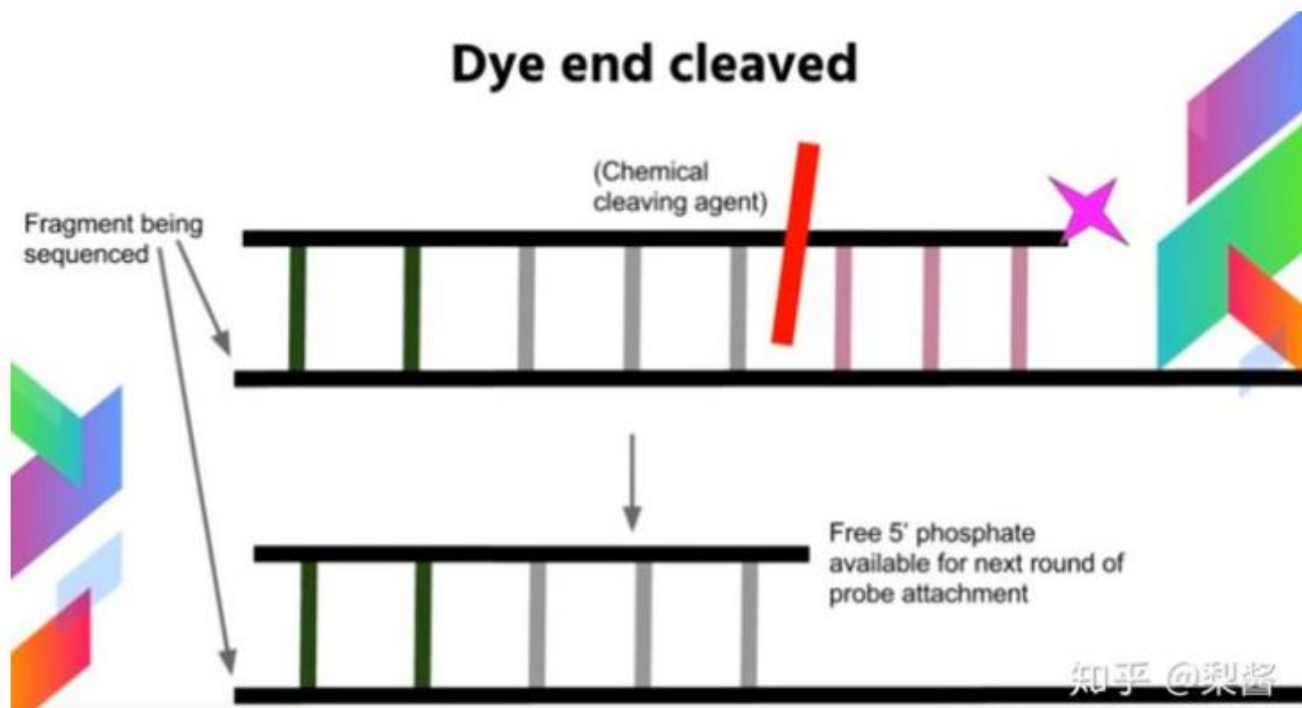


Probe Anatomy



带有不同荧光标记的8碱基探针（different fluorescently dyed oligonucleotide 8-mer bases）

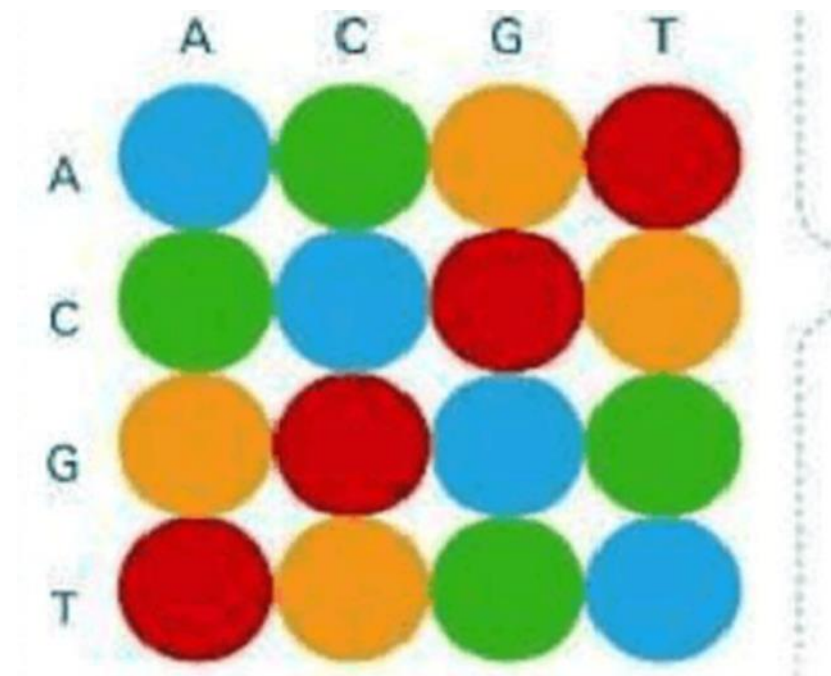
- 前2个碱基是真实的碱基。组合起来可以有16种可能，AA\AT\AG\AC, GG\GA\GT\GC.....每种排列对应一种荧光颜色（红、绿、蓝、黄）。**注意！**也就是说每种颜色可以对应4种排列。
- 中间3个通用碱基，
- 后3个有荧光标记的碱基



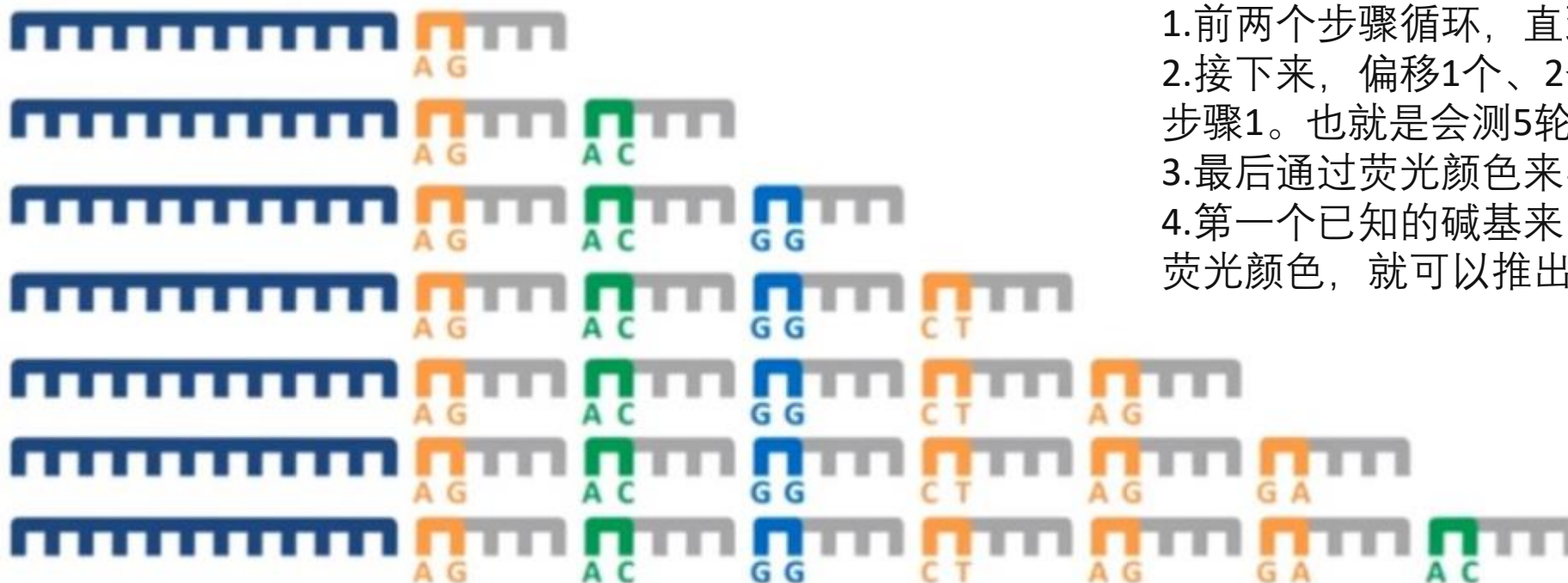
- 1.加上8碱基探针后：
- 2.用激光激发荧光。
- 3.用化学方法剪切掉后面3个荧光标记的碱基，只留下前5个碱基和游离的磷酸基团，以便加上第二个探针。

Result of one round → incomplete data

1. 加上8碱基探针后：
2. 用激光激发荧光。
3. 用化学方法剪切掉后面3个荧光标记的碱基，只留下前5个碱基和游离的磷酸基团，以便加上第二个探针。



Complete more cycles!



- 1.前两个步骤循环，直到合成整条链。
- 2.接下来，偏移1个、2个、3个、4个碱基，重复步骤1。也就是会测5轮
- 3.最后通过荧光颜色来判断
- 4.第一个已知的碱基来自5'或3'的adapter，又已知荧光颜色，就可以推出另外一个碱基

But we only have fluorescence measurements for every 5th base....

The entire process is repeated four times, each time with the primer offset by 1 base



偏移1个碱基，再测4轮

- 1.接下来，偏移1个、2个、3个、4个碱基，重复步骤1。也就是会测5轮
- 2.最后通过荧光颜色来判断
- 3.第一个已知的碱基来自5'或3'的adapter，又已知荧光颜色，就可以推出另外一个碱基



优缺点

a) 优点：

- 目前二代测序技术中准确度最高
- 除了测序和重测序之外，还能进行全基因组表达图谱分析、SNP、microRNA、甲基化等分析。

b) 缺点：

- 读长短，拼接复杂
- 双碱基对应一个荧光信号，如果发生读码错误，将发生连锁读码反应。

二者区别

- ePCR的过程中，SOLiD采用的磁珠直径更小，仅有1微米；而454使用两种磁珠，小磁珠上有固定酶，大磁珠上有reads
- SOLiD的独特之处在于没有使用常用的DNA聚合酶，而是使用DNA连接酶

相同之处

- 均使用ePCR技术
- 均使用磁珠进行扩增

THANKS

OMICS FOR ALL

基因科技造福人类