

REVEALING SMRT™ BIOLOGY

On-site System Training Workflow

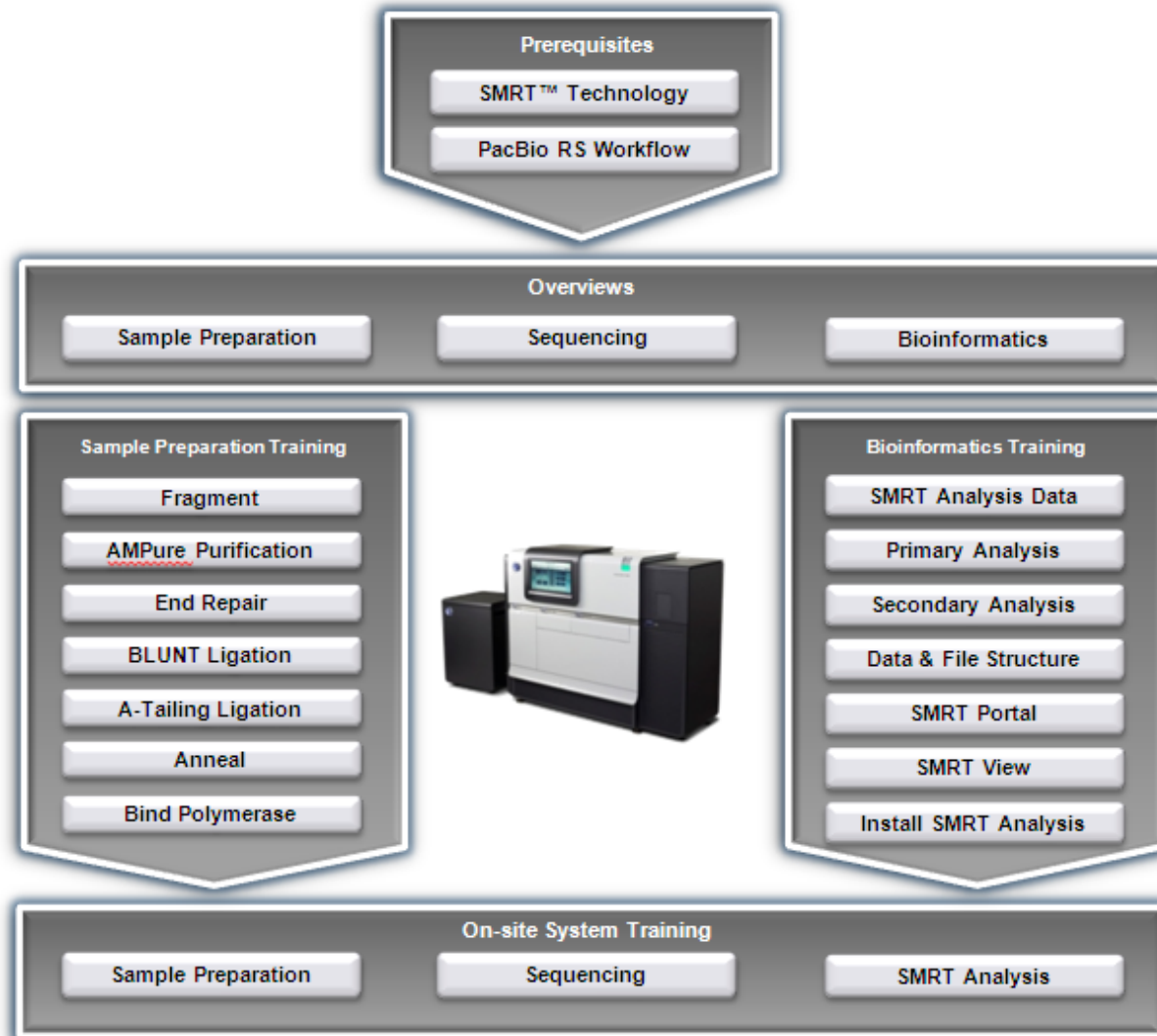
Pacbio RSII 系统标准客户培训流程

FU Wei

Senior Application Specialist

Login to the [PacBio Portal](http://www.pacificbiosciences.com/support/pubmap/training.html) and perform following tasks

<http://www.pacificbiosciences.com/support/pubmap/training.html>



Download documents



Template Preparation and Sequencing Guide



Procedure & Checklist - 2 kb Template Preparation and Sequencing

Before You Begin

To perform this procedure, you must have the Pacific Biosciences®:

- DNA Template Prep Kit (verify you have the correct kit for your insert size)
- DNA/Polymerase Binding Kit
- DNA Sequencing Kit
- DNA Control Complex
- SMRT® Cells for standard sequencing

This procedure can be used to prepare 1500 bp to 3 kb libraries from 500 ng up to 750 ng of sheared and concentrated DNA. If preparing larger amounts of DNA, scale all the reaction volumes proportionally (e.g., if the input amount of DNA is double the amount set forth in this procedure, then double all the reaction volumes listed in the tables).

| Insert Size Target | Insert Size Range | Sheared and Concentrated DNA Amount | Ligation | DNA Damage Repair |
|--------------------|-------------------|-------------------------------------|----------|-------------------|
| 2 kb | ~ 1500 bp to 3 kb | | | |

For DNA sequencing, it is recommended

Fragment and Concentrate DNA

Use a Covaris® System (with Adaptive Focused Acoustic) to concentrate your DNA sample. Note that some loss is to be expected as a result of inefficient amounts of starting DNA in order to (25 ng/μL) for the End-Repair reaction.

| Target Shear Size | Tube | Shearing Volume | Temp. Chills |
|-------------------|-----------------|-----------------|--------------|
| 2.0 kb | Mini-Tube Clear | 200.0 | 4°C |



Procedure & Checklist - Low-Input 10 kb Library Preparation and Sequencing (MagBead Station)

Before You Begin

To perform this procedure, you must have the PacBio®:

- DNA Template Prep Kit (verify you have the correct kit for your insert size)
- DNA/Polymerase Binding Kit
- MagBead Kit
- DNA Sequencing Kit
- DNA Control Complex
- SMRT® Cells for standard sequencing.

This procedure can be used to prepare 10 kb libraries from 1 μg up to 5 μg of sheared and concentrated DNA. If preparing libraries with DNA input amounts greater than 5 μg, scale all the reaction volumes proportionally (e.g., if the input amount of DNA is 10 μg, which is double the maximum input amount set forth in this procedure, then double all the reaction volumes listed in the tables).

| Insert Size Target | Insert Size Range | Sheared and Concentrated DNA Amount | Ligation | DNA Damage Repair |
|--------------------|-------------------|-------------------------------------|----------|-------------------|
| 10 kb | 8 kb to 12 kb | 1 to 5 μg | Blunt | Required |

For DNA sequencing, it is recommended to perform titrations to ensure optimal loading.

Fragment and Concentrate DNA

Use a Covaris® g-TUBE™ to shear your DNA sample. The most up-to-date guidance on how to use the g-TUBE, along with recommended centrifuges and centrifugation speeds, can be found in the g-TUBE user manual available for download from the Covaris website. Depending upon the quality of your sample, approximately 20% to 50% sample loss is to be expected as a result of the shearing and concentration process. Therefore, be sure to have sufficient amounts of starting DNA in order to have at least 1 μg of sheared and concentrated DNA (30 ng/μL) for the Damage Repair reaction.

Note that a Hydroshear® Shearing Device can also be used to shear your DNA sample. However, because Hydrodynamic shearing performance can vary with each shearing assembly, we recommend optimizing the shearing whenever a new shearing assembly is used.



Procedure & Checklist - 10 kb Template Preparation and Sequencing

Before You Begin

To perform this procedure, you must have the Pacific Biosciences®:

- DNA Template Prep Kit (verify you have the correct kit for your insert size)
- DNA/Polymerase Binding Kit
- DNA Sequencing Kit
- DNA Control Complex
- SMRT® Cells for standard sequencing.

This procedure can be used to prepare 10 kb libraries from 5 μg of sheared and concentrated DNA. For sheared DNA libraries and PCR products greater than 4 kb, any DNA damage (generated during DNA extraction and PCR amplifications) must be repaired using the DNA Damage Repair reagents provided by Pacific Biosciences and according to the instructions as set forth in this Procedure. Common types of damage may include abasic sites, cytosine deamination, and oxidation.

all the reaction volumes proportionally (e.g., if the input amount of procedure, then double all the reaction volumes listed in the tables).

| ge | Sheared and Concentrated DNA Amount | Ligation | DNA Damage Repair |
|----|-------------------------------------|----------|-------------------|
| | 5 μg | Blunt | Required |

to perform titrations to ensure optimal loading.

A

your DNA sample using the following conditions. Note that approximately 20% to 50% sample loss is to be expected as a result of the shearing and concentration process. Therefore, be sure to have sufficient amounts of starting DNA in order to have at least 1 μg of sheared and concentrated DNA (140 ng/μL) for the Damage Repair reaction.

since can vary with each shearing assembly, we recommend optimizing assembly is used.

| Speed Code | Cycles | Shearing Volume |
|------------|--------|-----------------|
| 0C8 - 0C10 | 20 | 200 μL |



Lab equipment and supplies

- Covaris
or Hydroshear



or



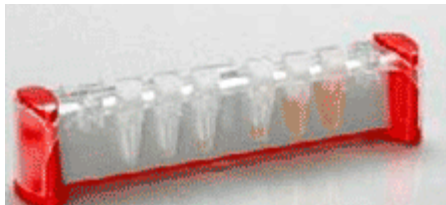
- Agilent 2100 Bioanalyzer



- Qubit™ Quantitation Platform
- Fluorometer



- Agencourt® AMPureXP® Beads



- Low bind tubes
- PCR System
- Centrifuge (Plate)
- Vortex Genie



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Gene Company Limited

基因有限公司

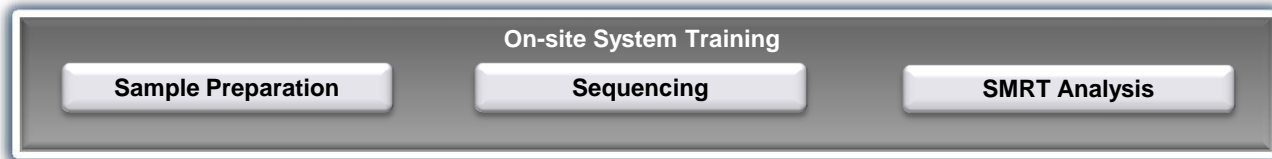
A Gene Group Company

P6/C4 reagents&Consumables

| 厂家 | 类型 | 货号 | 描述 |
|-----|----|-------------|---|
| PCB | 耗材 | 000-448-888 | Sample Plate (50 pack) |
| | | 000-882-901 | Sample Plate Septa (50 pack) |
| | | 000-947-365 | Mixing Plate (160 pack) |
| | | 100-243-500 | Mixing Plate (40 pack) |
| | | 100-237-900 | Sequencing plate septa |
| | | 001-292-541 | Tube septa |
| | | 100-154-200 | Pipet Tips (10 racks) |
| | | 001-252-068 | Pipet Tips (50 racks) |
| | 试剂 | 100-171-800 | SMRT Cell 8Pac V3 (8 Cells) |
| | | 100-209-300 | SMRT Cell Oil |
| | | 100-133-600 | MagBead Kit |
| | | 100-265-900 | AMPure PB (5ml) |
| | | 100-259-100 | SMRTbell Template Prep Kit 1.0 |
| | | 100-286-300 | SMRTbell HT Template Prep Kit 1.0 |
| | | 100-356-200 | DNA Sequencing Reagent 4.0 |
| | | 100-356-400 | DNA Sequencing Reagent 4.0 (10 package) |
| | | 100-372-700 | DNA/Polymerase Binding Kit 6.0 V2 |
| | | 100-356-500 | DNA Internal Control Complex (P6) |
| | 其他 | 100-514-900 | SMRTbell™ Barcoded Adapter Complete Prep Kit - 96 |
| | | 100-465-900 | SMRTbell™ Damage Repair Kit |
| | | 100-466-100 | Barcoded Universal F/R Primers Plate - 96 |

On-site System Training

- Duration: 5 days
- Covers the end-to-end workflow:



- Primary Users
 - Scientists
 - Research Associates
 - Bioinformaticians
- Standard Training
 - Lambda only
 - 10 kb
 - 5 days total
 - Standard Training Kit
 - Help customers with their samples once training is complete

Training Agenda:

| Time | Day 1 (Monday) | Day 2 (Tuesday) | Day 3 (Wednesday) | Day 4 (Thursday) |
|----------|--|-------------------------------------|-----------------------------|----------------------------|
| | Training Overview | 10 kb Library Construction (Contd.) | Annealing/Bindinf | Data Analysis |
| 9:00 AM | Technology Overview | Heat Kill Ligase | Data Analysis Overview | •SMRT Portal •SMRT View |
| 9:30 AM | Workflow Overview | Exo Treatment | | |
| 10:00 AM | | Double Ampure Purification | Calculator Overview | |
| 10:30 AM | 10 kb Lambda Library Shearing | Quantitation, Bioanalyzer, Break | | |
| 11:00 AM | | Quantitation, Bioanalyzer, Break | Hands On Pactice Calculator | Metrics Review |
| 11:30 AM | Purification of sheared DNA | MBS Binding | Lunch Break | |
| 12:00 PM | Lunch Break | | | |
| 1:00 PM | Quantitation (Qubit, nanodrop). BioAnalyzer | Training Run RS Remote Set Up | Troubleshooting Overview | Instrument RS Overview |
| 1:30 PM | | | 10Kb Library prep | RS remote Overview |
| 2:00 PM | PreCR | | | |
| 2:30 PM | End Repair | Instrument Run | Hands On Pactice RS remote | |
| 3:00 PM | Purify | Additional Training | | |
| 3:30 PM | Quantitation/Quality | | | |
| 4:00 PM | Blunt Ligation (safe stopping point), Store at 4 C overnight | | | |
| 4:30 PM | | | | |
| 5:00 PM | Review: Day 1 | Review: Day 2 | Review: Day 3 | Review: Day 4 |

Sequencing Workflow

Template
Preparation

Run
Design

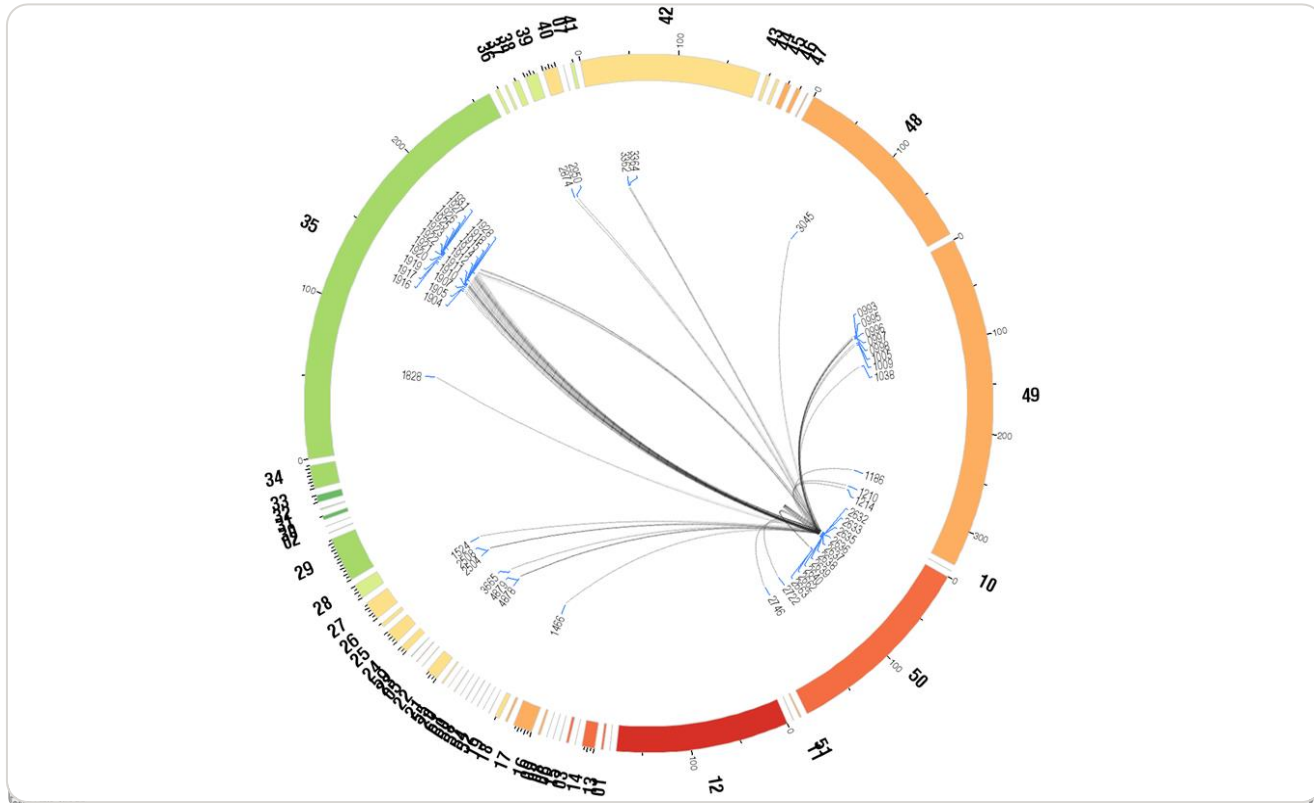
Polymerase
Binding

Instrument
Run

Primary
Analysis

Secondary
Analysis

Tertiary
Analysis



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SMRT Portals View

After Training

- Sign-off training completion form
- Complete the quiz
- Follow up on bioinformatics by PacBio Tech Support
 - Technical bioinformatics issues
 - Q&A
- Contact PacBio Tech support
 - Toll Free Telephone +1 (877) 920-7222
 - Email techsupport@pacificbiosciences.com
 - Logging a case on PacBio Portal



DISTRIBUTOR PORTAL

Case Edit

Search

Search All

search

[Advanced Search...](#)

Create New...

Recent Items

2011-05-23 Office Hour
- Low Yield
Troubleshooting, Error
Messages, PM

2011-05-02 Office Hour
- Web Based Calculator

2011-04-18 Office Hour
- Release v1.2.1 Field
Webinar

2011-03-28 Office Hour
- SMRT Cell Loading

Repair stage pi...

2011-03-16 Office Hour
- Annealing and Binding
Calculator

Cholera Webinar by Eric
Schadt

Recorded Webinar -
Release v1.2 End-User
Training

Case Edit

Submit

Save & Close

Submit & Add Attachment

Save & New

Cancel

Submit & Add Attachment

Case Information

= Required Information

Case Owner Yuanjie Zhou

Account Name Click lookup icon...

Contact Name

Parent Case

Case Manager

--None--

Visible in Customer Portal

Status

Submitted

Solution Release

--None--

Description Information

Type Complaint

Case Reason --None--

Case Reason Detail --None--

Subject

Description

Internal Comments

Other Information



SMRT® Cell

Thank you for attention