



COMPASS 软件介绍

数据分析

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PROTEIN SIMPLE -- 创新蛋白质分析技术专家



Jess/Abby/Wes

全自动 Western



FluorChem

多功能成像



Ella

微流控全自动 ELISA



Milo

单细胞 Western



MFI

微流成像颗粒计数分类仪



Maurice

CE-SDS + iIEF 双功能 CE

MEET SIMPLE WESTERN

JESS ABBY WES



GEL-RUNNING AUTO
TRANSFER-FREE
BLOT-FREE
HANDS-FREE

超微量样品+自动化+定量

1 Simple Western 工作原理

2 Simple Western 优势及应用

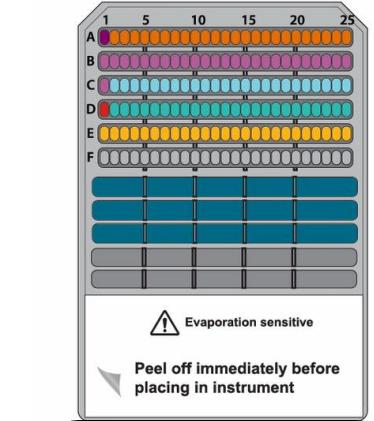
3 Simple Western 实验操作简介

您需要做的工作



1. 配制样品和试剂

- 加样
- 混匀
- 加热



2. 加样

- 加marker
- 加样品
- 加封闭液
- 加一抗
- 加二抗
- 加发光液
- 加洗液



3. 运行

- 放毛细管卡盒
- 放加样板
- 运行程序

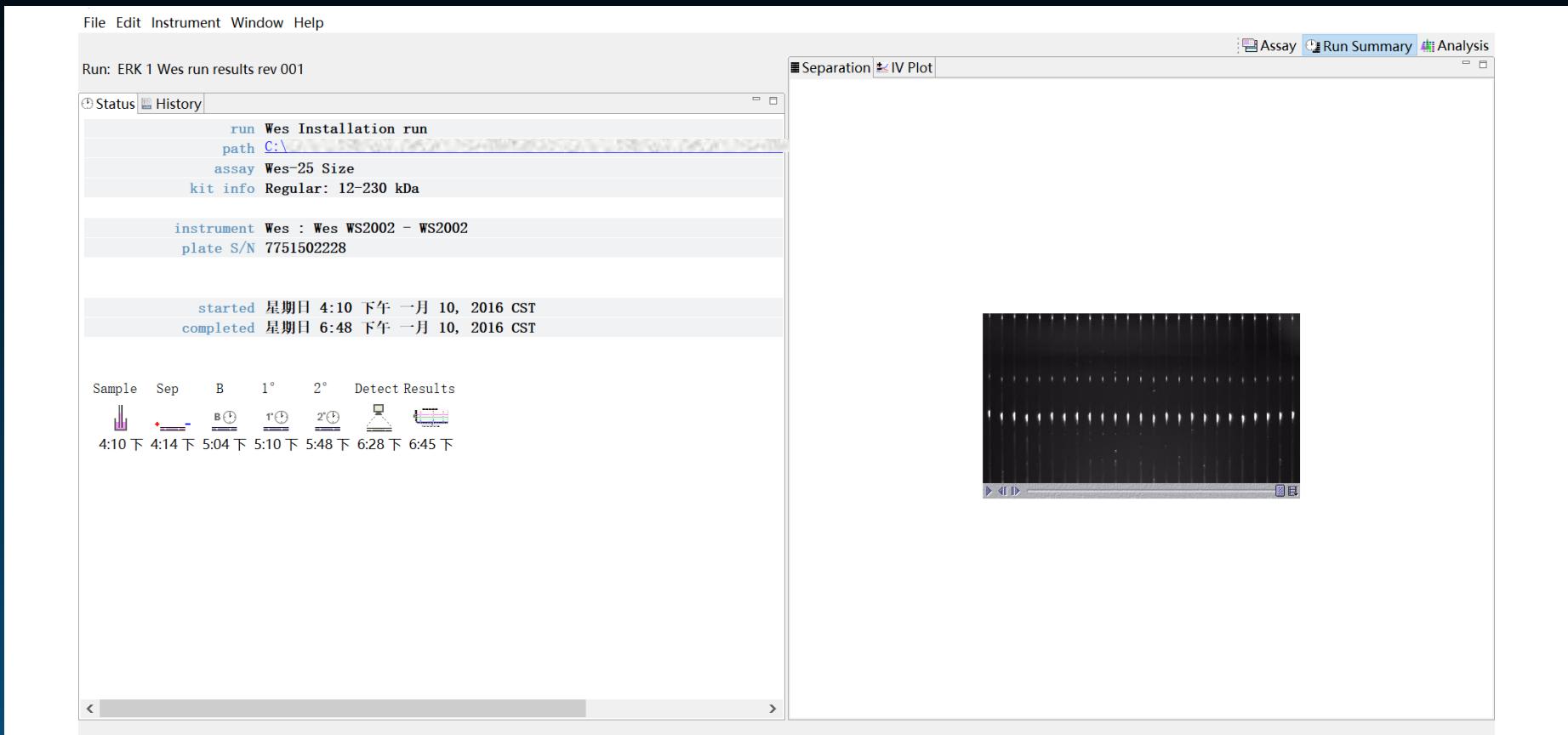


Compass for SW

*样品不能加 Loading Buffer

One Pair of
Pants Later...

运行结束后



01

Compass 介绍

02

数据分析前检查

03

数据分析

01

Compass 介绍

02

数据分析前检查

03

数据分析

COMPASS 软件

COMPASS 有三个主要界面

- Assay – 实验前的设置
- Run summary – 运行过程中的实时情况
- Analysis – 分析界面



ASSAY 界面

File Edit Instrument Window Help

Assay: Wes Size

Protocol History Notes

Assay Run Summary Analysis

Layout

12-230 kDa

Value

Separation Matrix

Stacking Matrix

Sample

Separation Time (m) 25.0

Separation Voltage 375

Antibody Diluent Ti 5.0

Primary Antibody Ti 30.0

Secondary Antibody 30.0

Detection

Well Row E1

Detection Profile HDR

Template

Edit

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25
A	Bio...																								
B																									
C	Ant...																								
D	Str...																								
E																									

ASSAY 界面

File Edit Instrument Window Help

Assay: Wes Size

Protocol History Notes

Assay Run Summary Analysis

Layout

12-230 kDa

Value

- Separation Matrix
- Stacking Matrix
- Sample
- Separation Time (m) 25.0
- Separation Voltage 375
- Antibody Diluent Ti 5.0
- Primary Antibody Ti 30.0
- Secondary Antibody 30.0
- Detection
- Well Row E1
- Detection Profile HDR

Template

Edit

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25
A	Bio...														Sample										
B																Antibody Diluent									
C	Ant...															Primary Antibody									
D	Str...															Secondary HRP Conjugate									
E																Luminol/Peroxide									

10 ASSAY 界面

File Edit Instrument Window Help

Assay: Wes Size

Protocol History Notes

Assay Run Summary Analysis

Layout Add Remove

12-230 kDa

Value

Separation Matrix

Stacking Matrix

Sample

Separation Time (m) 25.0

Separation Voltage 375

Antibody Diluent Ti 5.0

Primary Antibody Ti 30.0

Secondary Antibody 30.0

Detection

Well Row E1

Detection Profile HDR

Well Content

Name: Sample

Attribute:

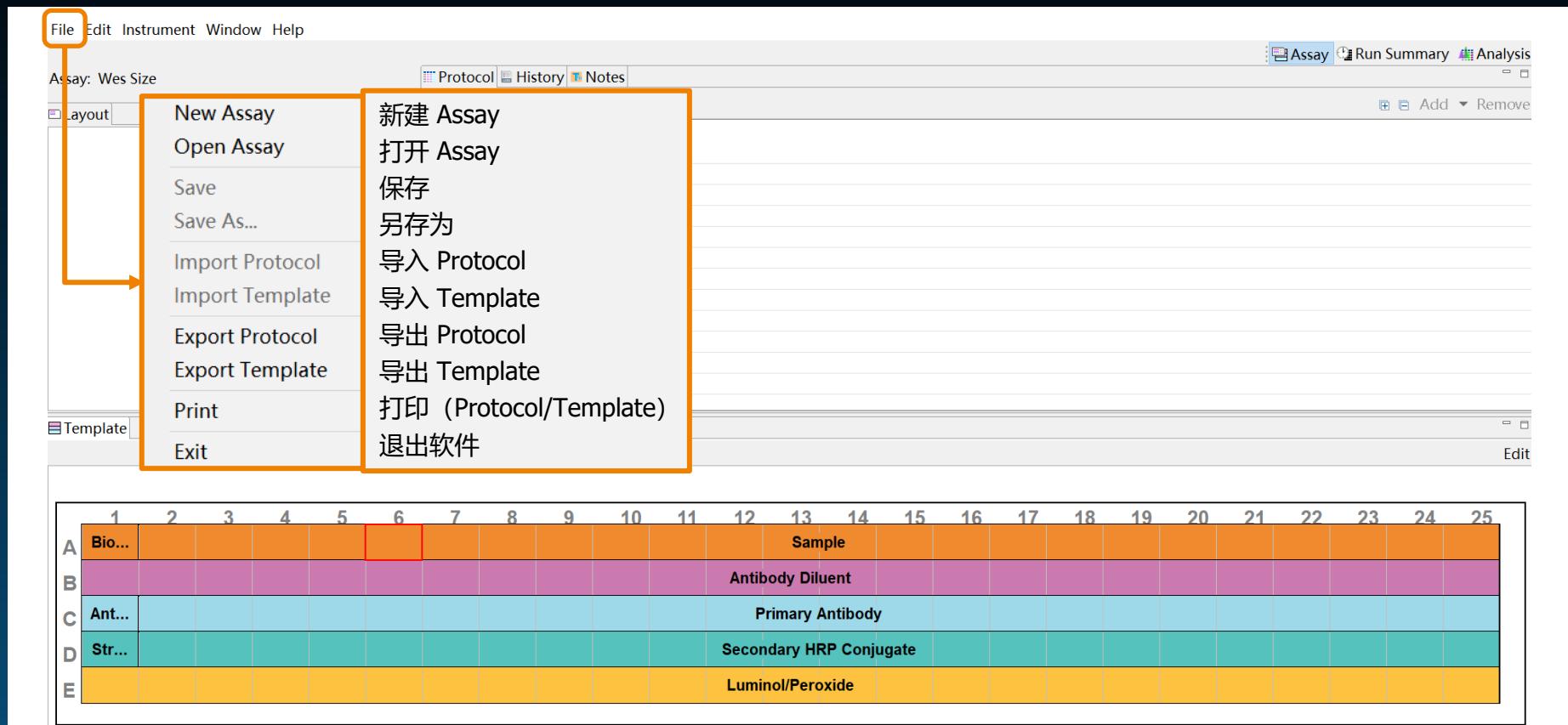
OK Cancel

Template Edit

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25

A	Bio...														Sample										
B															Antibody Diluent										
C	Ant...														Primary Antibody										
D	Str...														Secondary HRP Conjugate										
E															Luminol/Peroxide										

ASSAY 界面



ASSAY 界面

File **Edit** Instrument Window Help

Assay: Wes Size

Protocol History Notes

Assay Run Summary Analysis

Layout

Cut

Copy **Ctrl+C**

Paste **Ctrl+V**

Default Analysis...

Default Analysis View...

Preferences

剪切

拷贝

粘贴

默认分析设置

默认分析预览

参数设置

Well Row E1

Detection Profile HDR

Template

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25
A	Bio...																								
B																									
C	Ant...																								
D	Str...																								
E																									

10 ASSAY 界面

The screenshot displays the Bio-Techne Assay software interface. The main window shows a 10x25 well layout for a Western blot analysis. The wells are color-coded by sample: A (Bio...), B (Antibody Diluent), C (Primary Antibody), D (Secondary HRP Conjugate), and E (Luminol/Peroxide). The 6th well in the first row is highlighted in red. The software interface includes a top menu bar (File, Edit, Instrument, Window, Help), a protocol tab, a history tab, and a notes tab. A context menu is open over the 6th well, showing options: Cut, Copy (Ctrl+C), Paste (Ctrl+V), Default Analysis..., Default Analysis View..., and Preferences. A second context menu is open over the text '剪切' (Cut), showing options: 剪切 (Cut), 拷贝 (Copy), and 粘贴 (Paste). A 'Standards' dialog box is open, titled 'Default Analysis: Wes-25 Size.assay'. It contains a table of fluorescent peaks with their MW (kDa) and Position, and a 'Fit' column with checkboxes. The table data is as follows:

MW (kDa)	Position	Fit
1	170	<input type="checkbox"/>
29	350	<input checked="" type="checkbox"/>
230	650	<input checked="" type="checkbox"/>

Buttons for Import..., Export..., OK, Cancel, and Apply are at the bottom of the dialog box. The bottom of the image shows the well layout for lanes 21 to 25.

10 ASSAY 界面

The screenshot shows the Bio-Techne Assay software interface. The main window displays a 12-well plate layout with rows A-E and columns 1-12. The plate is partially filled with samples: Row A (Bio...), Row B (Antibody Diluent), Row C (Primary Antibody), Row D (Secondary HRP Conjugate), and Row E (Luminol/Peroxide). Well 6 is highlighted in orange. The 'Edit' menu is open, showing options: Cut, Copy (Ctrl+C), Paste (Ctrl+V), Default Analysis..., Default Analysis View..., and Preferences. A context menu is also open over Well 6, showing: 剪切 (Cut), 复制 (Copy), 复贴 (Paste), 默认分析设置 (Default Analysis Settings), 默认分析预览 (Default Analysis Preview), and 参数设置 (Parameter Settings). A 'Graph View Options' dialog box is open, titled 'Default Analysis View Wes-25 Size.assay'. It contains two tabs: 'Graph View Options' and 'Lane View Options'. Under 'Graph View Options', checkboxes are checked for Matching Peak Names, Peak Names, and Peak Values. Under 'Plot Label', checkboxes are checked for Sample, Capillary, Attribute, Primary, Secondary, and Exposure. Other checkboxes are unchecked for Fitted Peaks, Baseline Fit, Noise Region, All Exposures, and Grid Lines. Buttons at the bottom of the dialog are OK, Cancel, and Apply.

File Edit Instrument Window Help

Assay: Wes Size

Protocol History Notes

Cut

Copy Ctrl+C

Paste Ctrl+V

Default Analysis...

Default Analysis View...

Preferences

剪切

拷贝

粘贴

默认分析设置

默认分析预览

参数设置

Default Analysis View Wes-25 Size.assay

Graph View Options

Lane View Options

Matching Peak Names

Peak Names

Peak Values

Fitted Peaks

Baseline Fit

Noise Region

All Exposures

Grid Lines

Plot Label

Sample

Attribute

Primary

Attribute

Secondary

Attribute

Capillary

Exposure

OK Cancel Apply

Assay Run Summary Analysis

Add Remove

	1	2	3	4	5	6	7	8	9	10	11	12		1	22	23	24	25
A	Bio...												Sample					
B													Antibody Diluent					
C	Ant...												Primary Antibody					
D	Str...												Secondary HRP Conjugate					
E													Luminol/Peroxide					

ASSAY 界面

The screenshot displays the ASSAY software interface with several windows and toolbars.

Top Menu Bar: File, Edit, Instrument, Window, Help.

Protocol Tab: Shows tabs for Protocol, History, and Notes.

Layout Tab: Shows a sample plate layout with columns 1-13 and rows A-E. Well 6 is highlighted in red, and E1 is highlighted in yellow. Detection settings: Well Row E1, Detection Profile HDR.

Edit Context Menu (highlighted by an orange box):

- Cut
- Copy **Ctrl+C**
- Paste **Ctrl+V**
- Default Analysis...
- Default Analysis View...
- Preferences

Protocol Context Menu (highlighted by an orange box):

- 剪切 (Cut)
- 拷贝 (Copy)
- 粘贴 (Paste)
- 默认分析设置 (Default Analysis Settings)
- 默认分析预览 (Default Analysis Preview)
- 参数设置 (Parameter Settings)

Preferences Dialog (highlighted by an orange box):

Access Control

- Enable
- Server: 127.0.0.1
- Auto Lock Inactivity (min): 20

These settings apply to all users.

Buttons: Restore Defaults, Apply, OK, Cancel, Edit.

Sample Plate Layout:

	1	2	3	4	5	6	7	8	9	10	11	12	13	1
A	Bio...													
B														Antibody Diluent
C	Ant...													Primary Antibody
D	Str...													Secondary HRP Conjugate
E														Luminol/Peroxide

Number 25 is located in the bottom right corner of the sample plate layout.

10 ASSAY 界面

File Edit Instrument **Window** Help

Assay: Wes Size

Layout Assay Run Summary Analysis

Notes

Value

25.0

375

Antibody Diluent Ti 5.0

Primary Antibody Ti 30.0

Secondary Antibody 30.0

Detection

Well Row E1

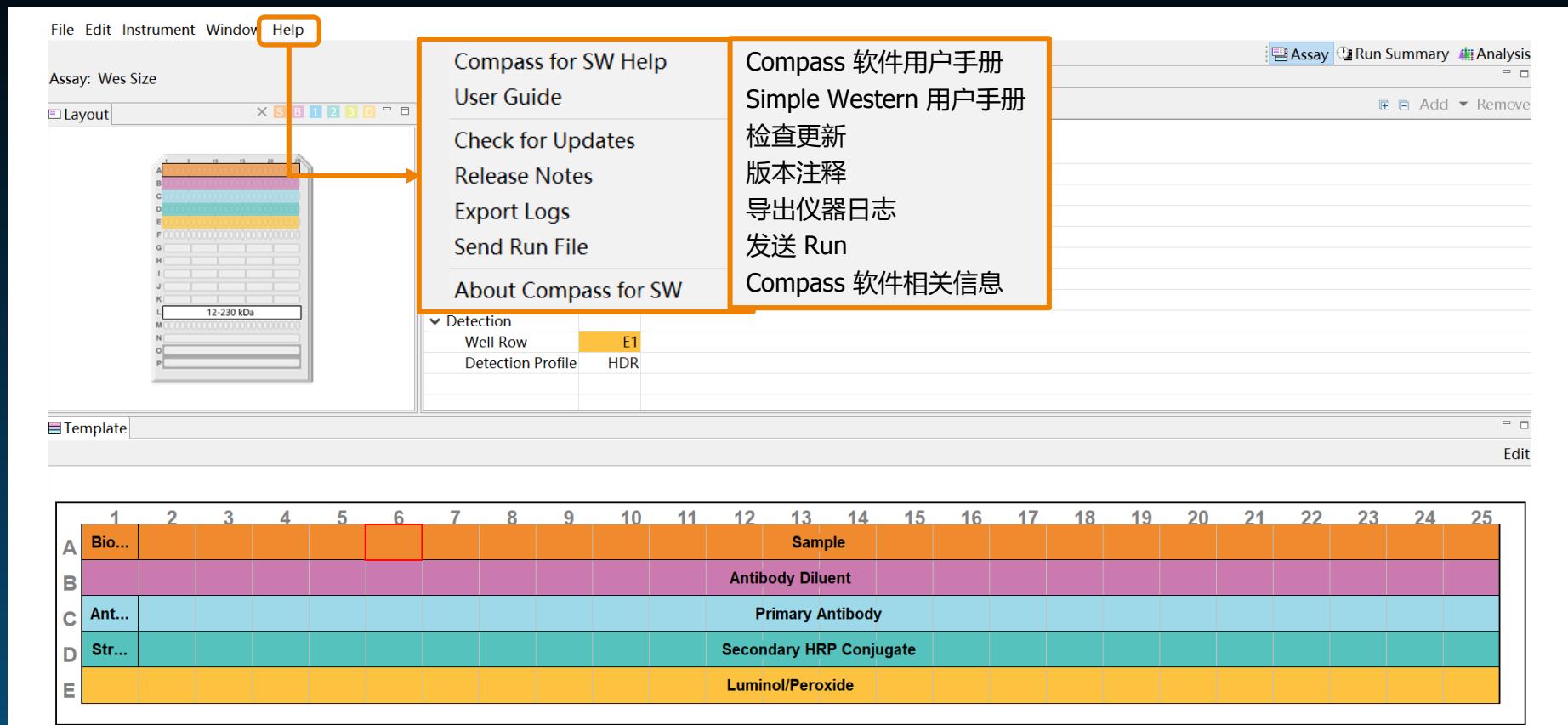
Detection Profile HDR

Template Edit

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25
A	Bio...																								
B																									
C	Ant...																								
D	Str...																								
E																									

ASSAY 界面



RUN SUMMARY 界面

File Edit Instrument Window Help

Run: ERK 1 Wes run results rev 001

Separation  Assay  Analysis

Status 

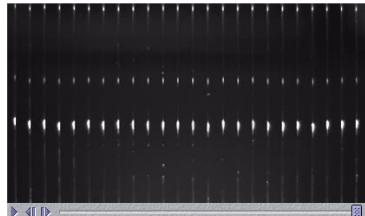
run Wes Installation run
path C:\
assay Wes-25 Size
kit info Regular: 12-230 kDa

instrument Wes : Wes WS2002 - WS2002
plate S/N 7751502228

started 星期日 4:10 下午 一月 10, 2016 CST
completed 星期日 6:48 下午 一月 10, 2016 CST

Sample Sep B 1° 2° Detect Results

4:10 下 4:14 下 5:04 下 5:10 下 5:48 下 6:28 下 6:45 下



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RUN SUMMARY 界面

File Edit Instrument Window Help

Run: ERK 1 Wes run results rev 001

Status History

run Wes Installation run
path C:\
assay Wes-25 Size
kit info Regular: 12-230 kDa

instrument Wes : Wes WS2002 - WS2002
plate S/N 7751502228

started 星期日 4:10 下午 一月 10, 2016 CST
completed 星期日 6:48 下午 一月 10, 2016 CST

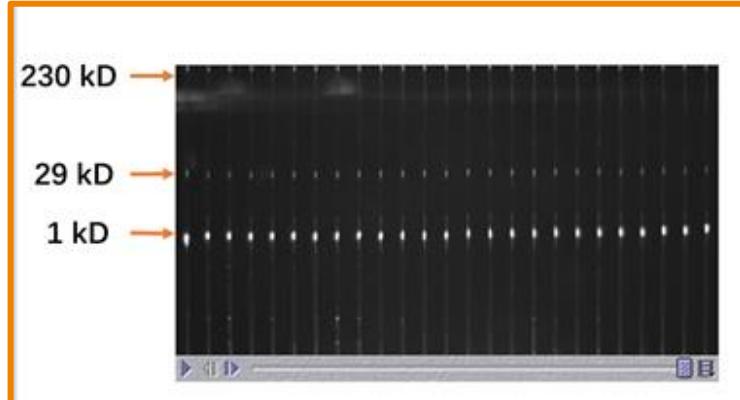
Sample Sep B 1° 2° Detect Results

4:10 下 4:14 下 5:04 下 5:10 下 5:48 下 6:28 下 6:45 下

- ▶ 12-230 kit: 1 kDa, 29 kDa , 230 kDa
- ▶ 66-440 kit: 57 kDa, 280 kDa
- ▶ 2-40 kit: 1 kDa, 26 kDa

Separation IV Plot

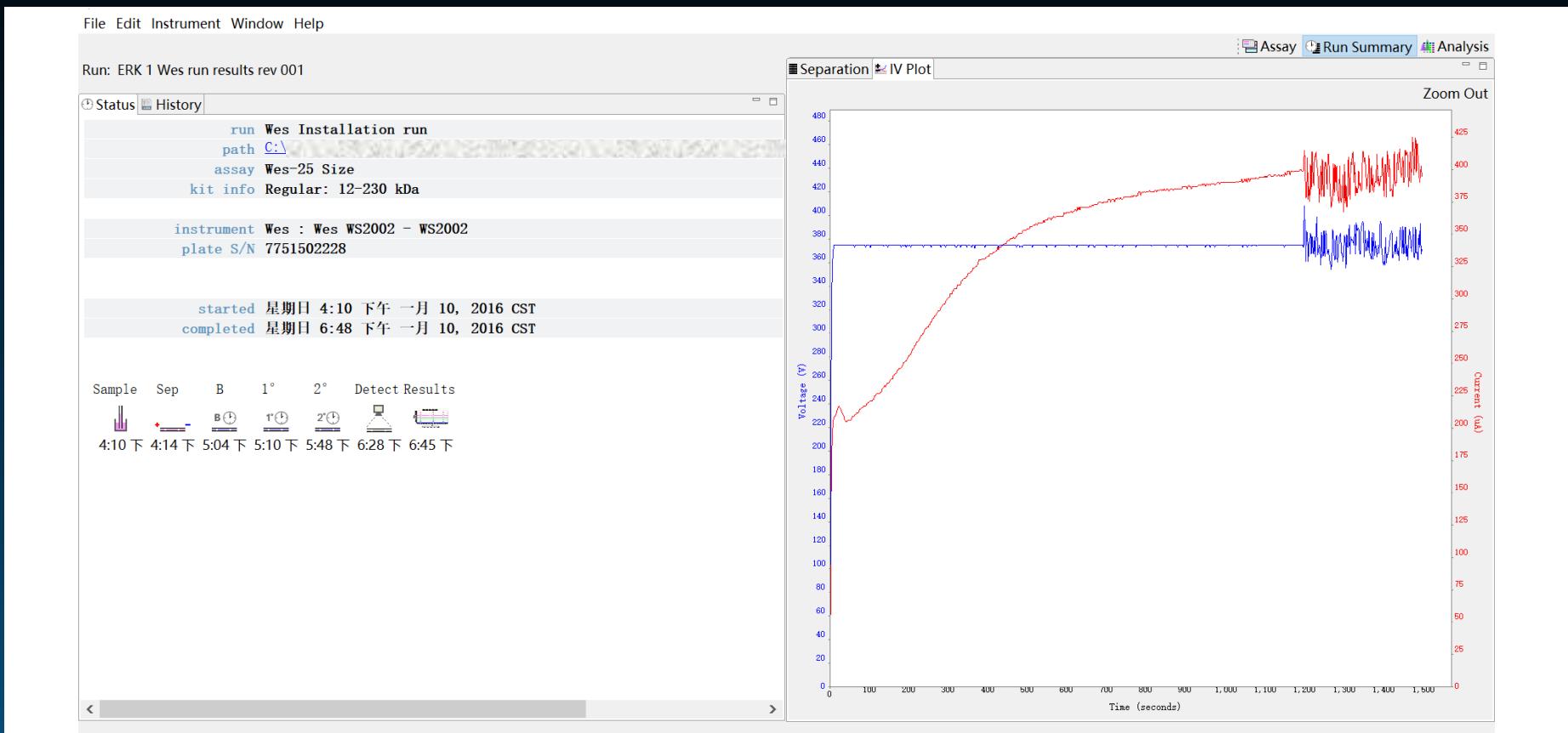
Assay Run Summary Analysis



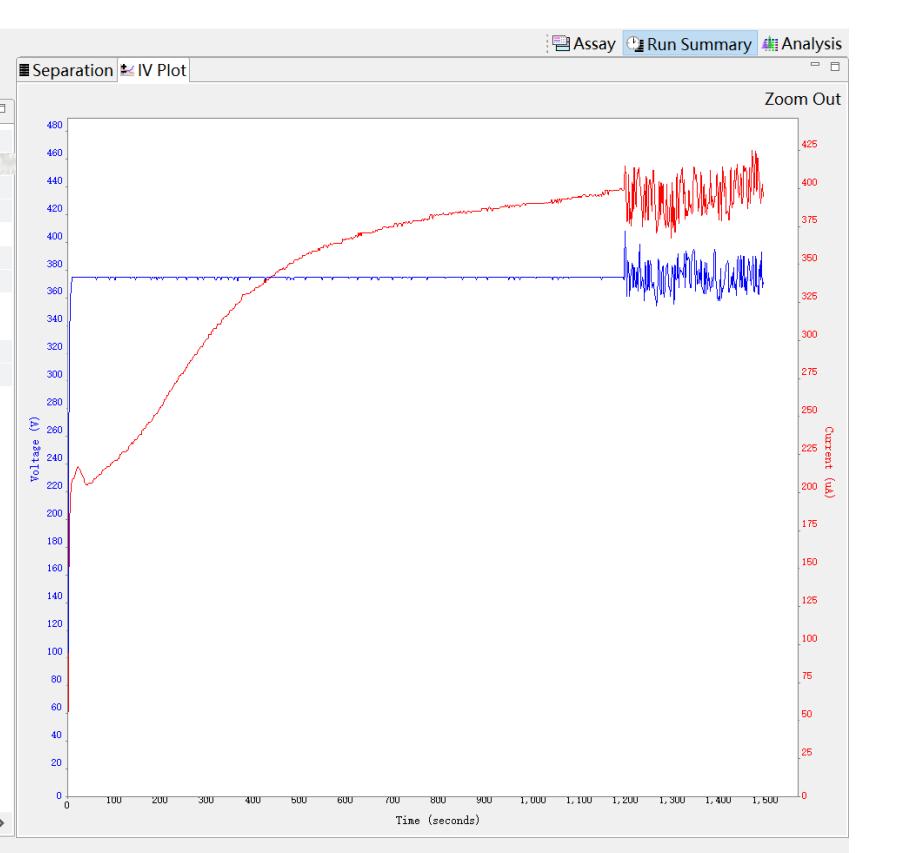
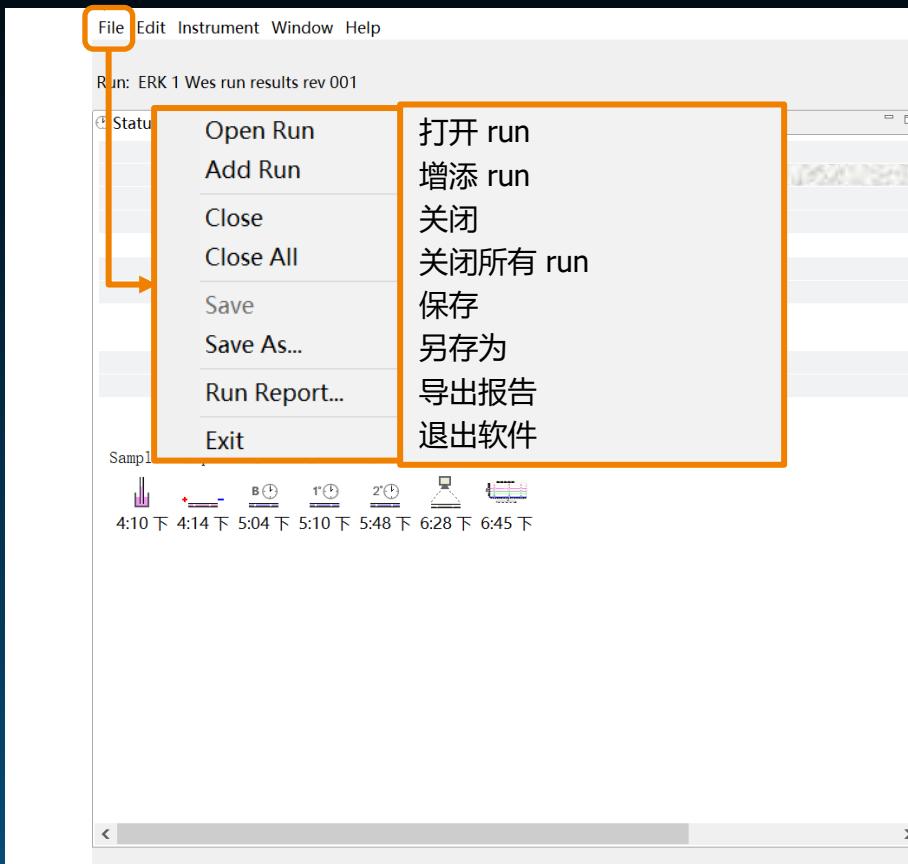
230 kDa
29 kDa
1 kDa

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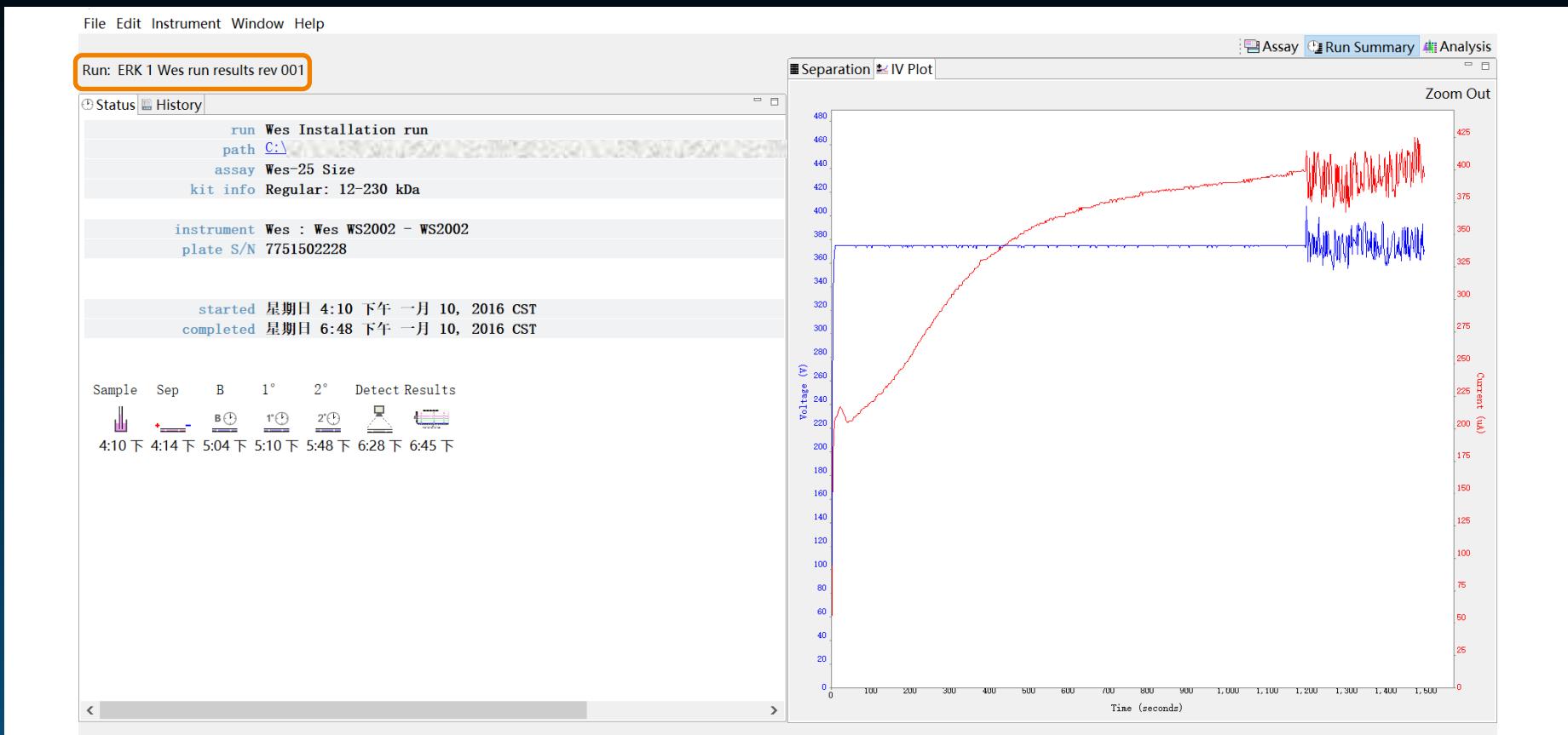
RUN SUMMARY 界面



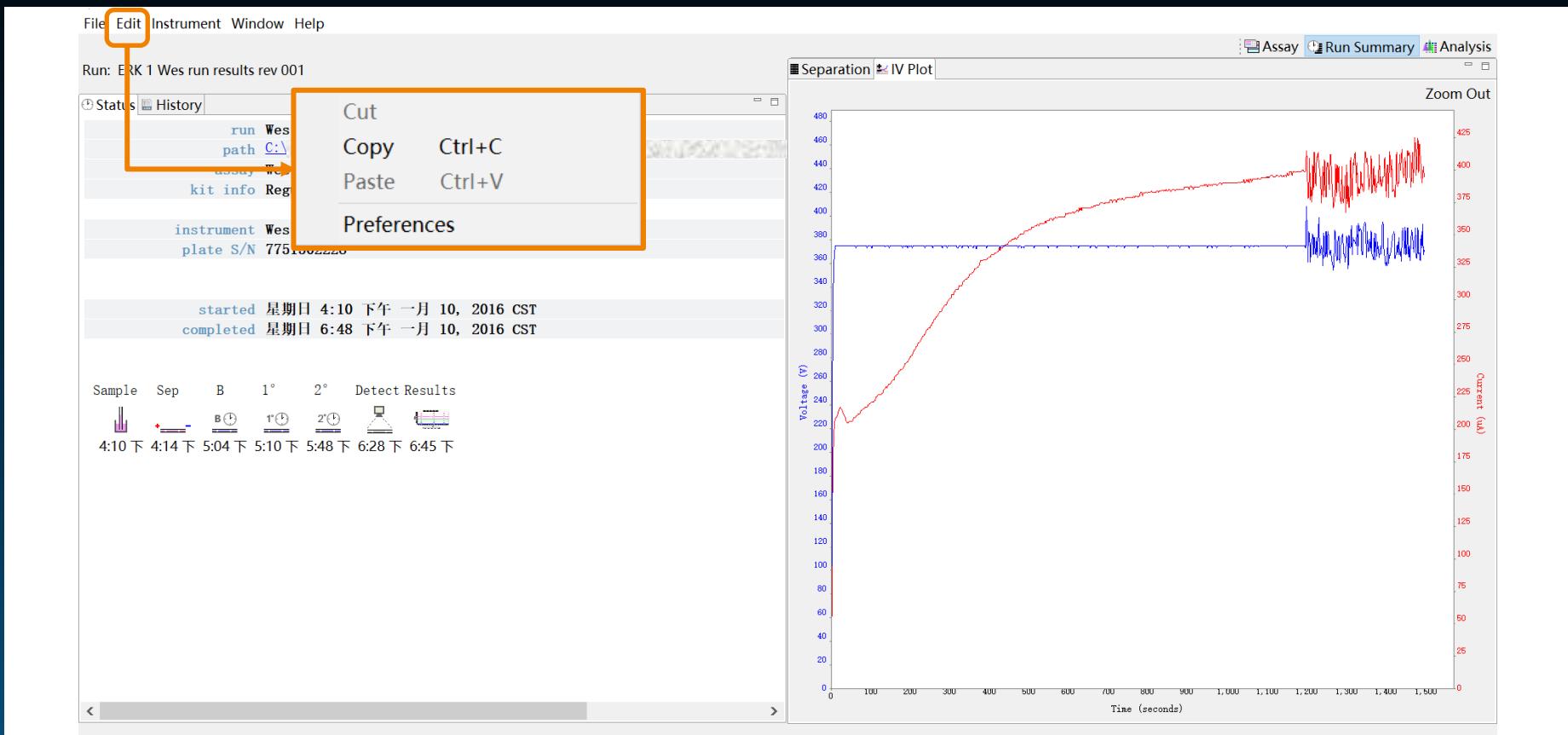
RUN SUMMARY 界面



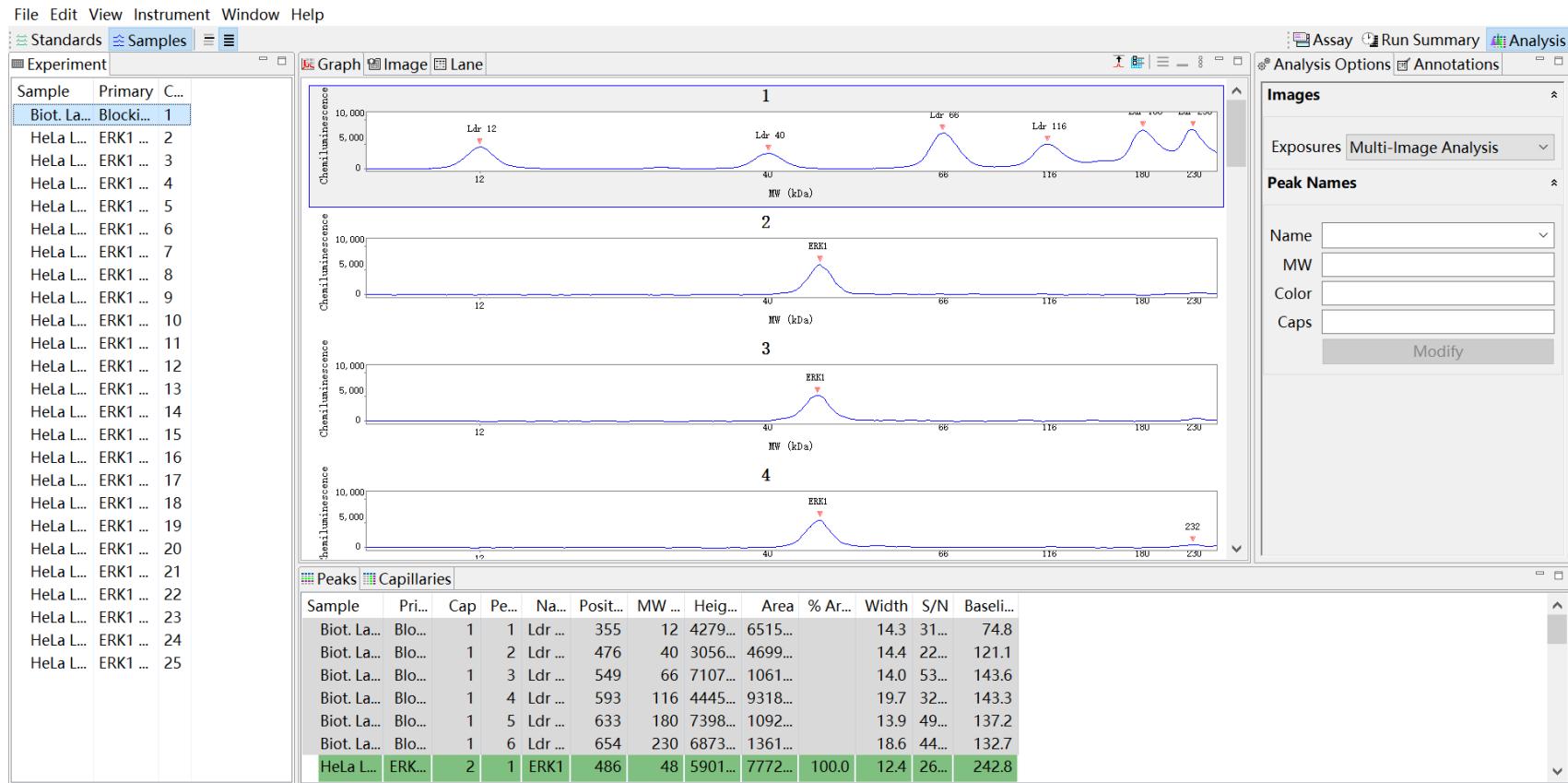
RUN SUMMARY 界面



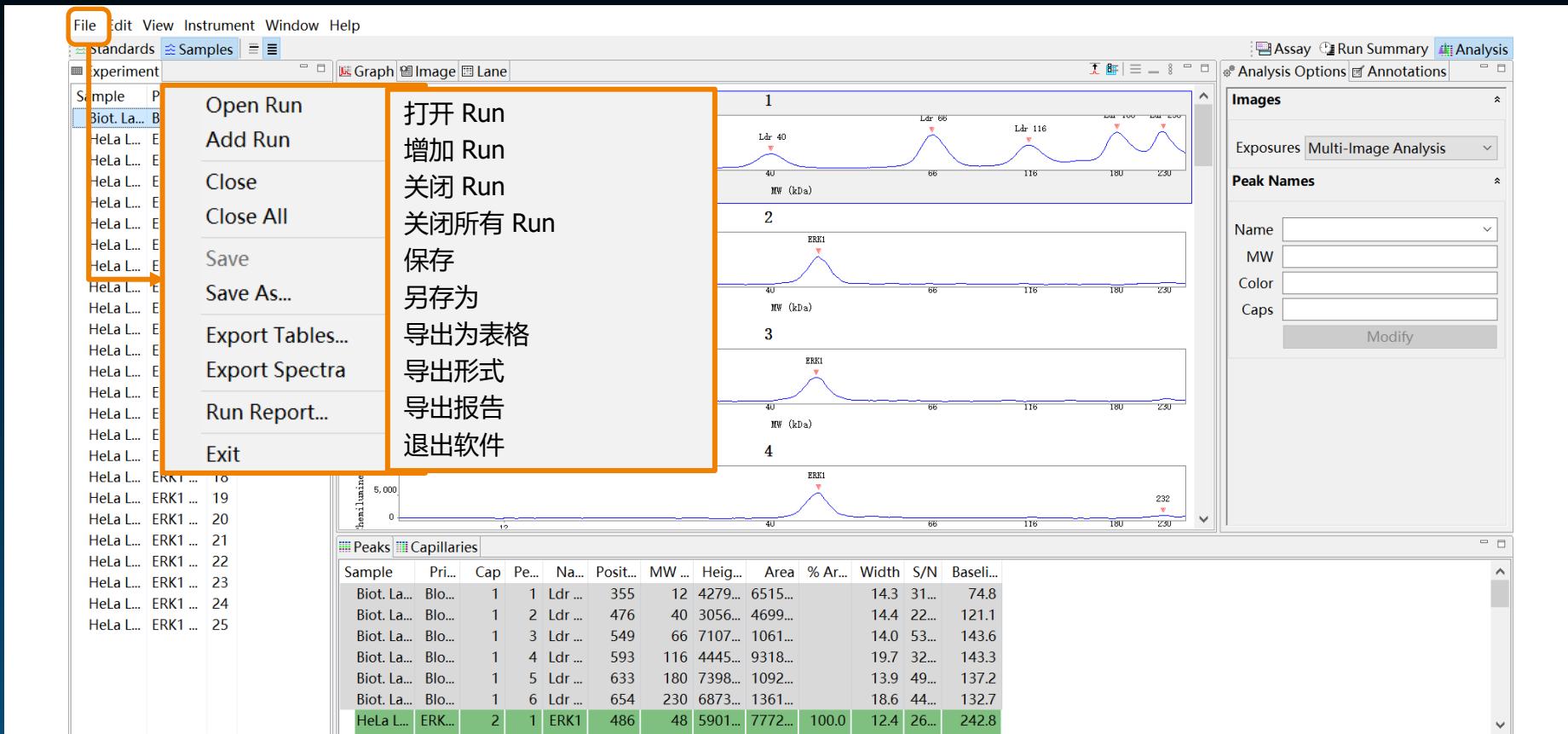
RUN SUMMARY 界面



ANALYSIS 界面

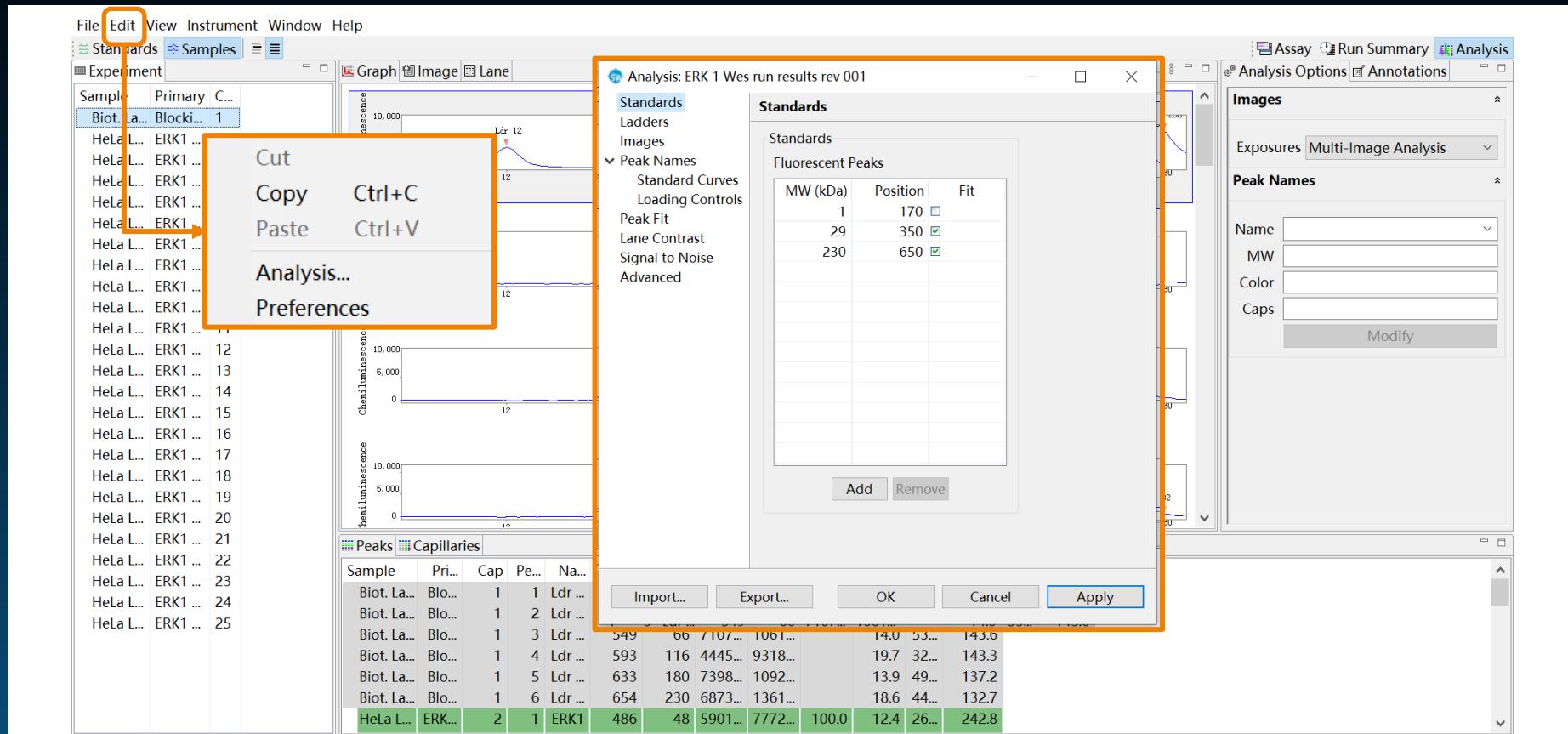


ANALYSIS 界面

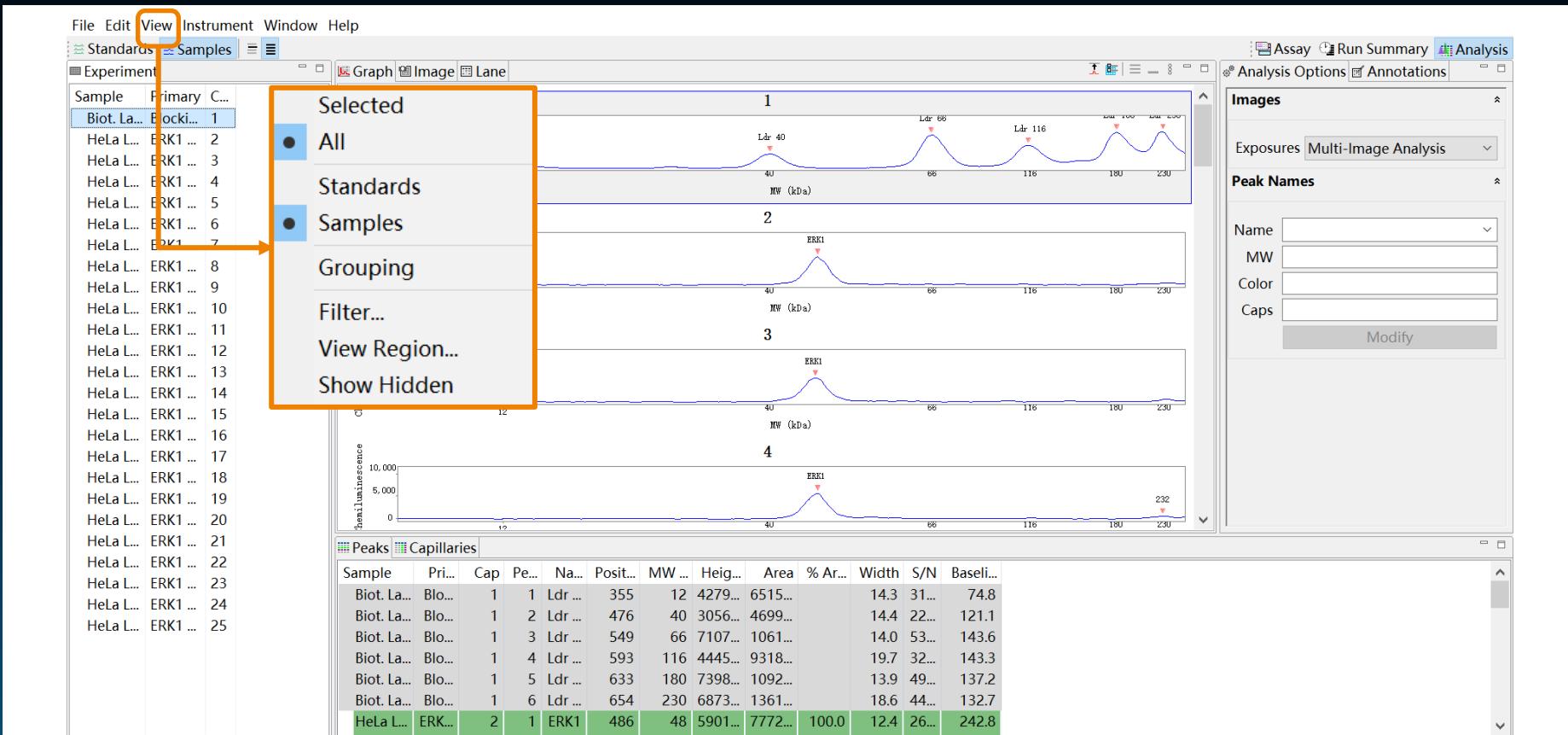


ANALYSIS 界面

ANALYSIS 界面



ANALYSIS 界面



ANALYSIS 界面

File Edit View Instrument Window Help

Standards Samples

Experiment

Graph Image Lane

Selected

All

Standards

Samples

Grouping

Filter...

View Region...

Show Hidden

Capillaries

1

2

3

4

5

6

7

8

9

10

11

Show named peaks only

OK Cancel

Sample Pri... Cap Pe... Na... Posit... MW ... Heig

Biot. La... Blo... 1 1 Ldr ... 355 12 4279

Biot. La... Blo... 1 2 Ldr ... 476 40 3056

Biot. La... Blo... 1 3 Ldr ... 549 66 7107... 1061... 14.0 53... 143.6

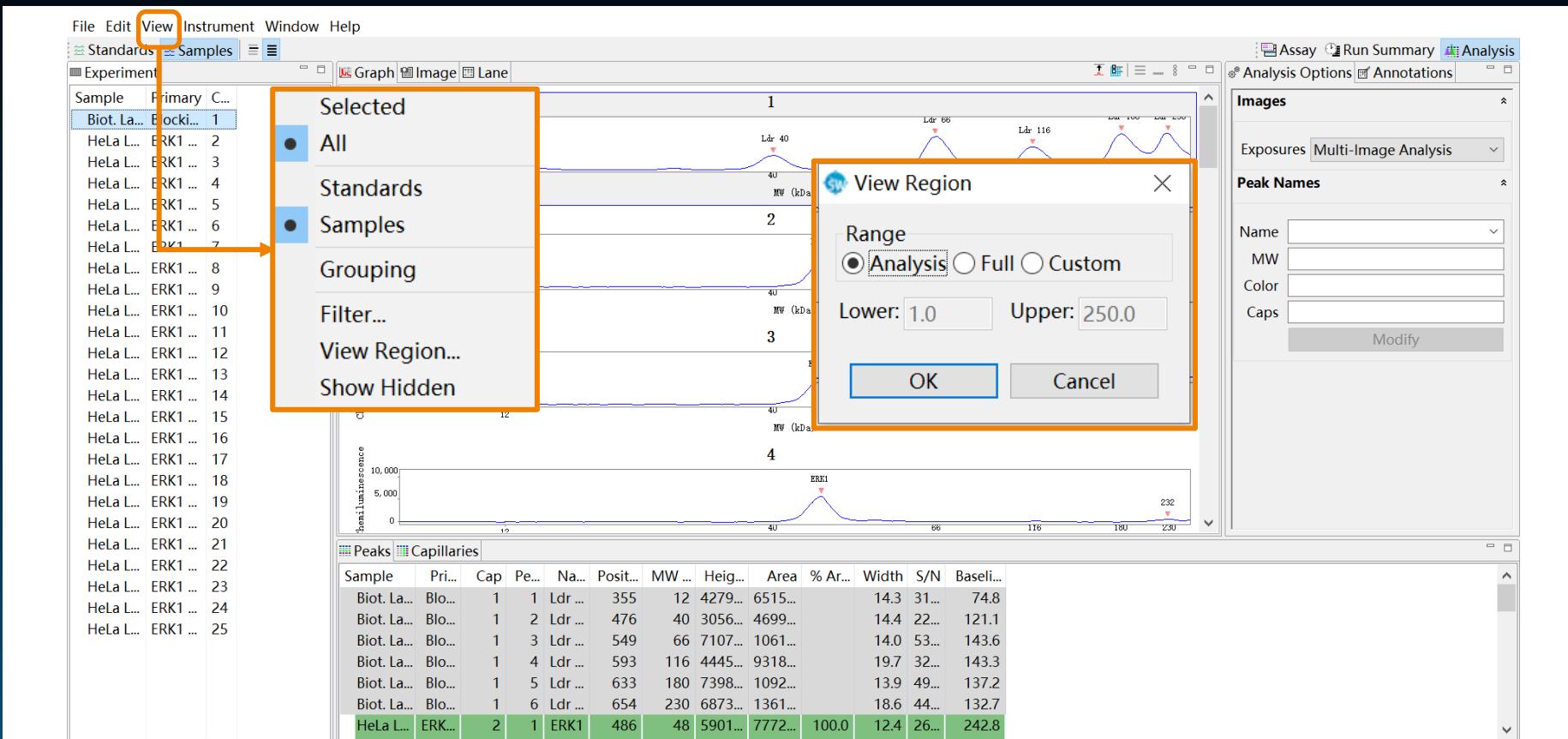
Biot. La... Blo... 1 4 Ldr ... 593 116 4445... 9318... 19.7 32... 143.3

Biot. La... Blo... 1 5 Ldr ... 633 180 7398... 1092... 13.9 49... 137.2

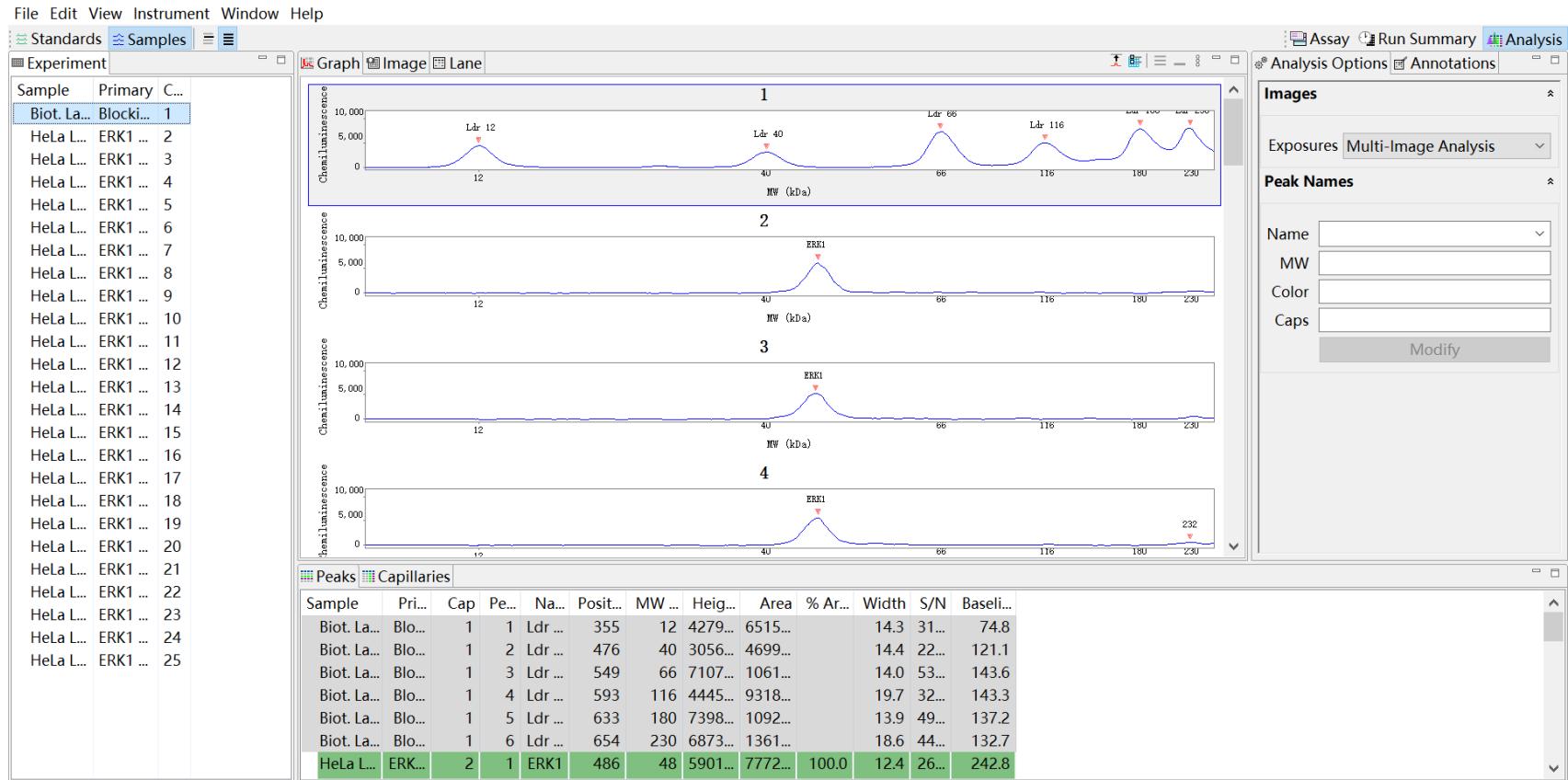
Biot. La... Blo... 1 6 Ldr ... 654 230 6873... 1361... 18.6 44... 132.7

HeLa L... ERK... 2 1 ERK1 486 48 5901... 7772... 100.0 12.4 26... 242.8

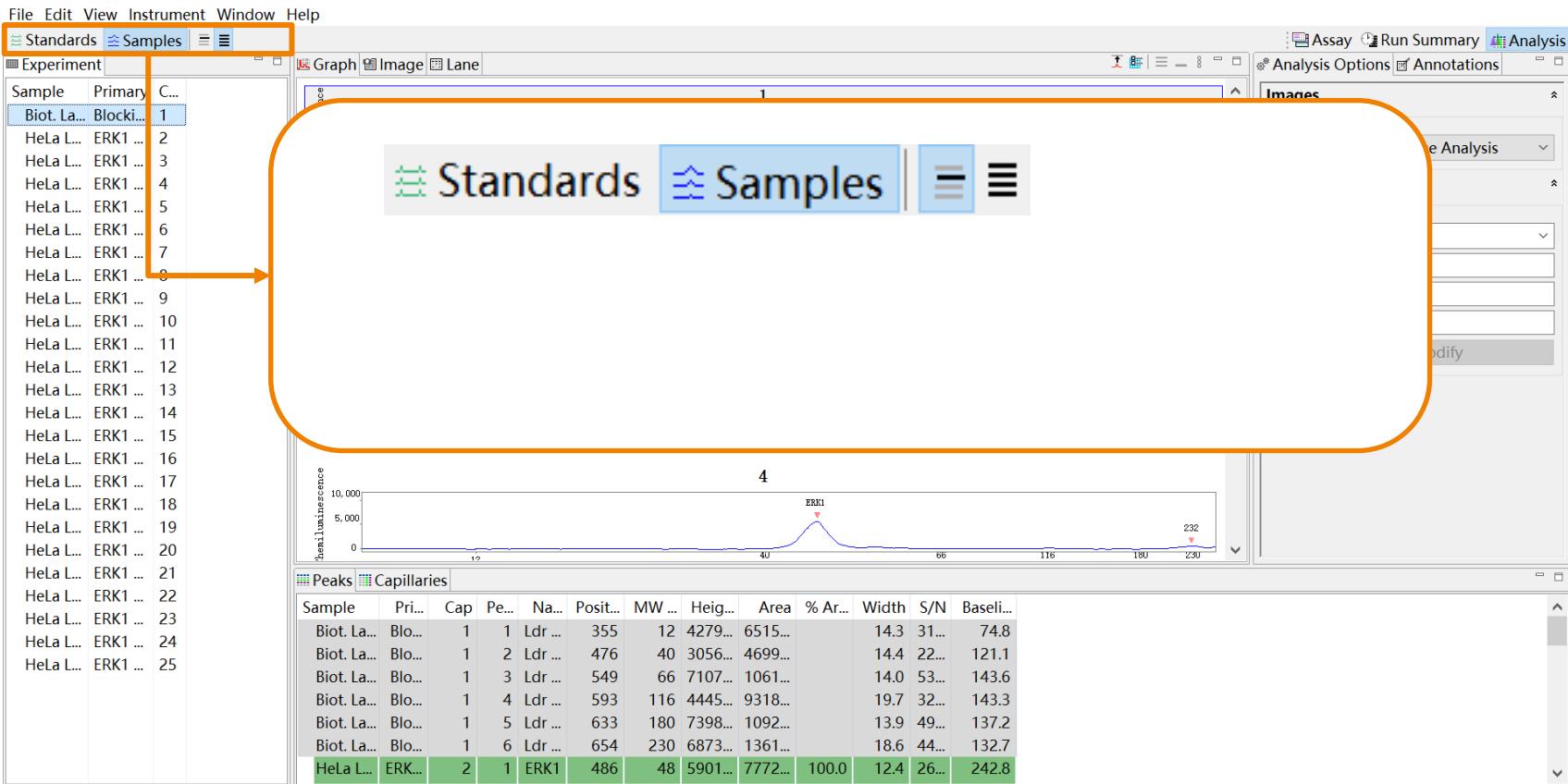
ANALYSIS 界面



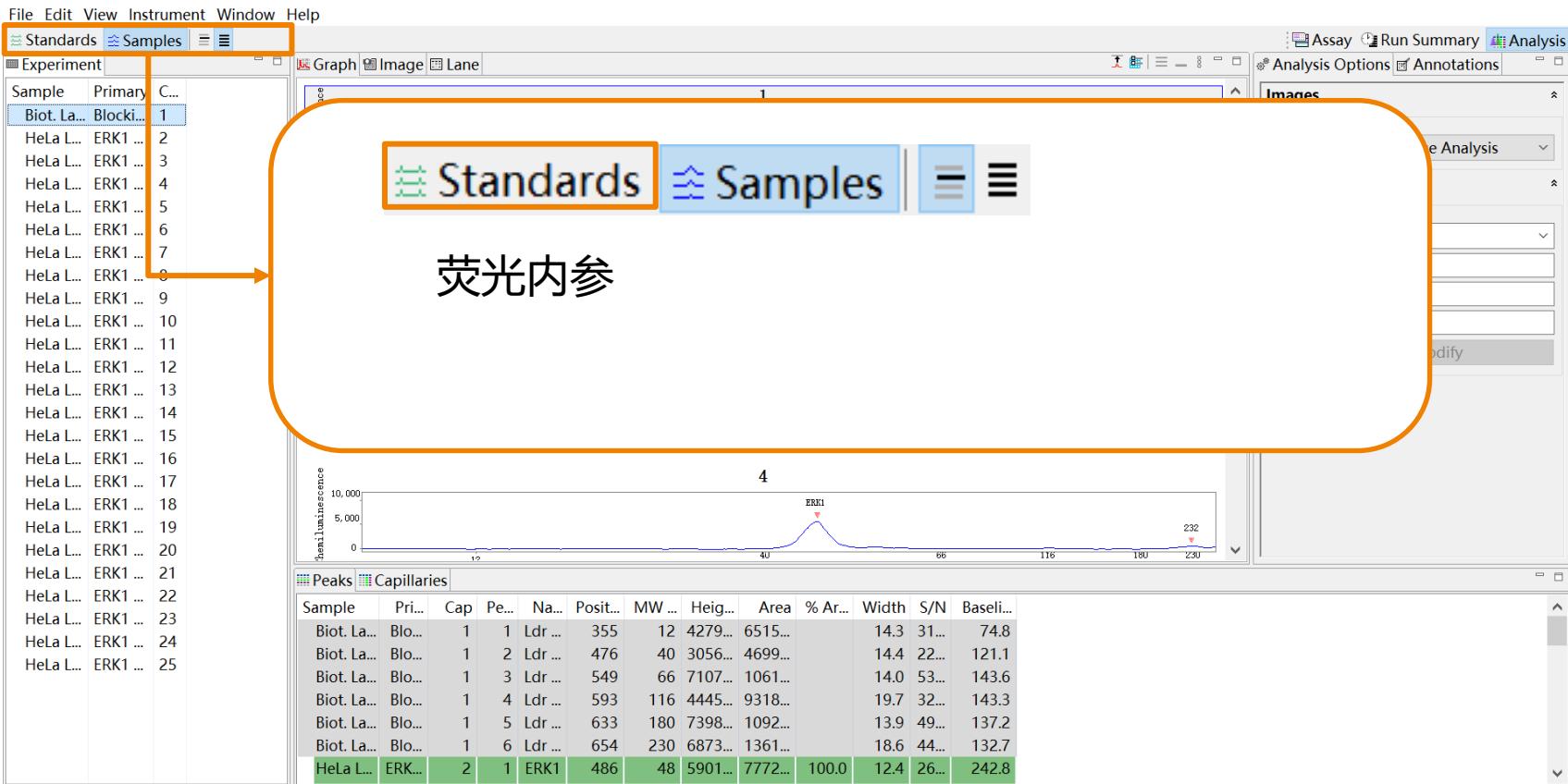
ANALYSIS 界面



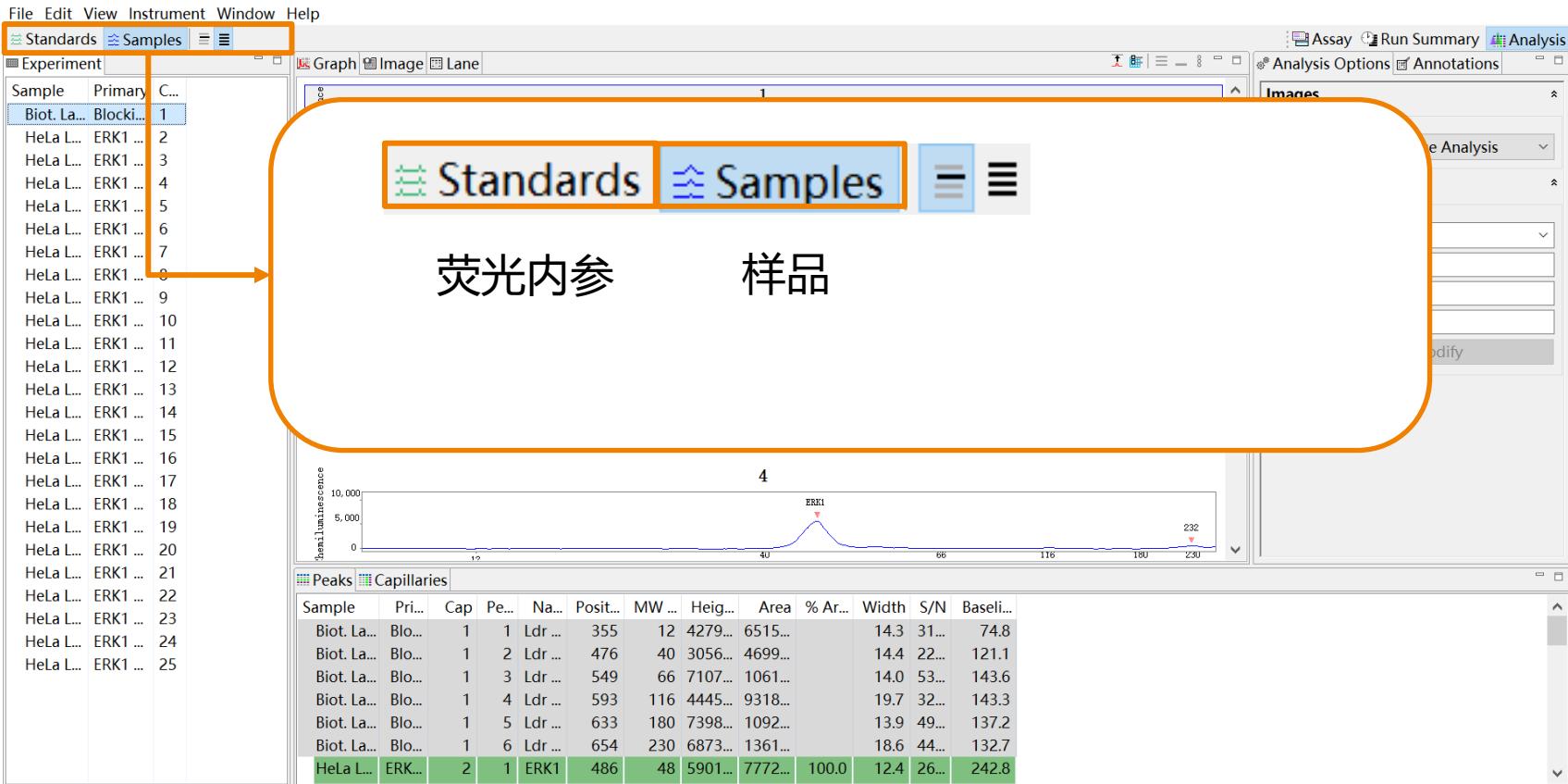
13 ANALYSIS 界面



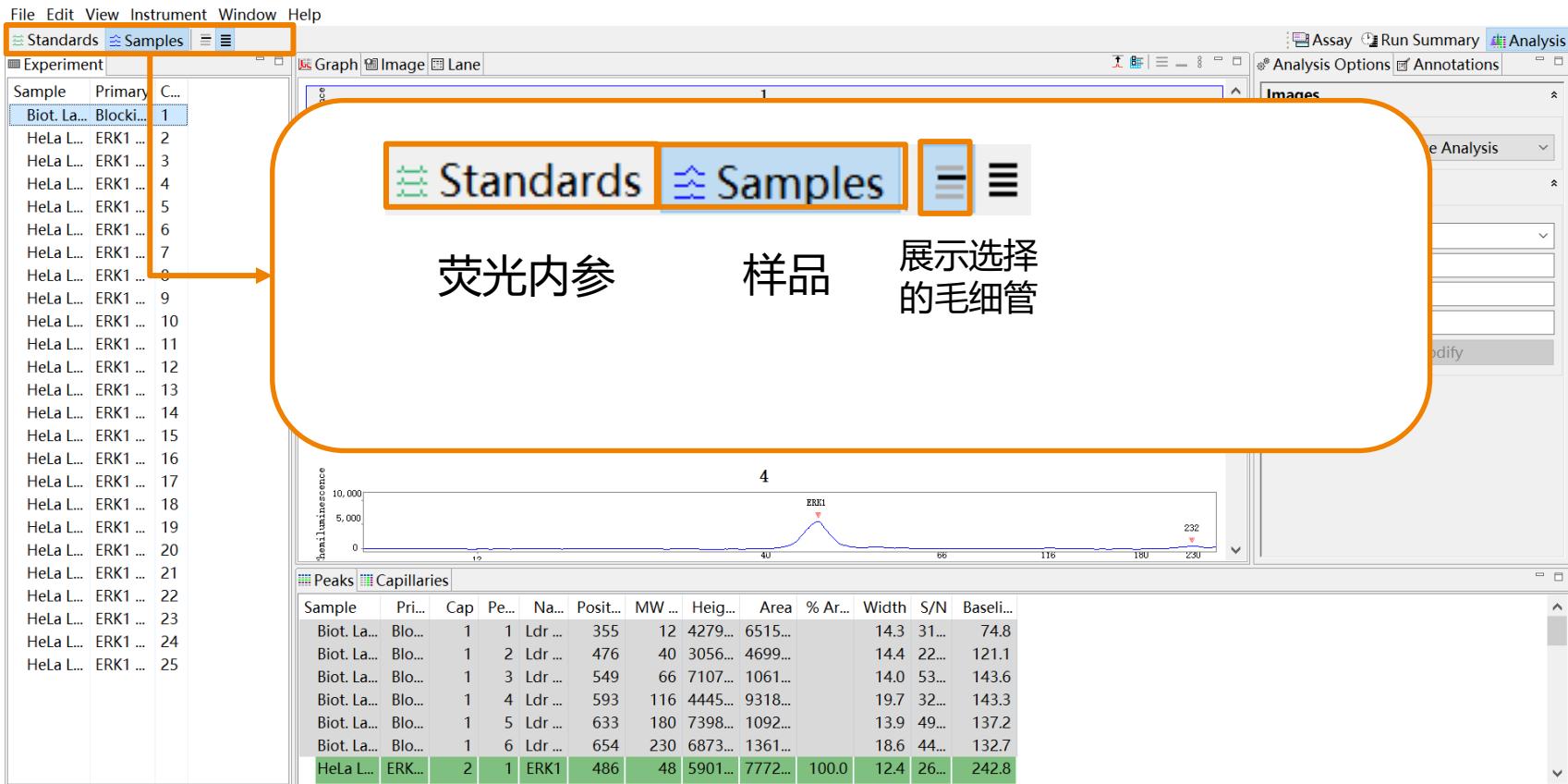
13 ANALYSIS 界面



13 ANALYSIS 界面



13 ANALYSIS 界面



13 ANALYSIS 界面

File Edit View Instrument Window Help

Standards Samples

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

Graph Image Lane

1

Standards Samples

荧光内参 样品 展示选择的毛细管

展示所有毛细管

4

Chemiluminescence

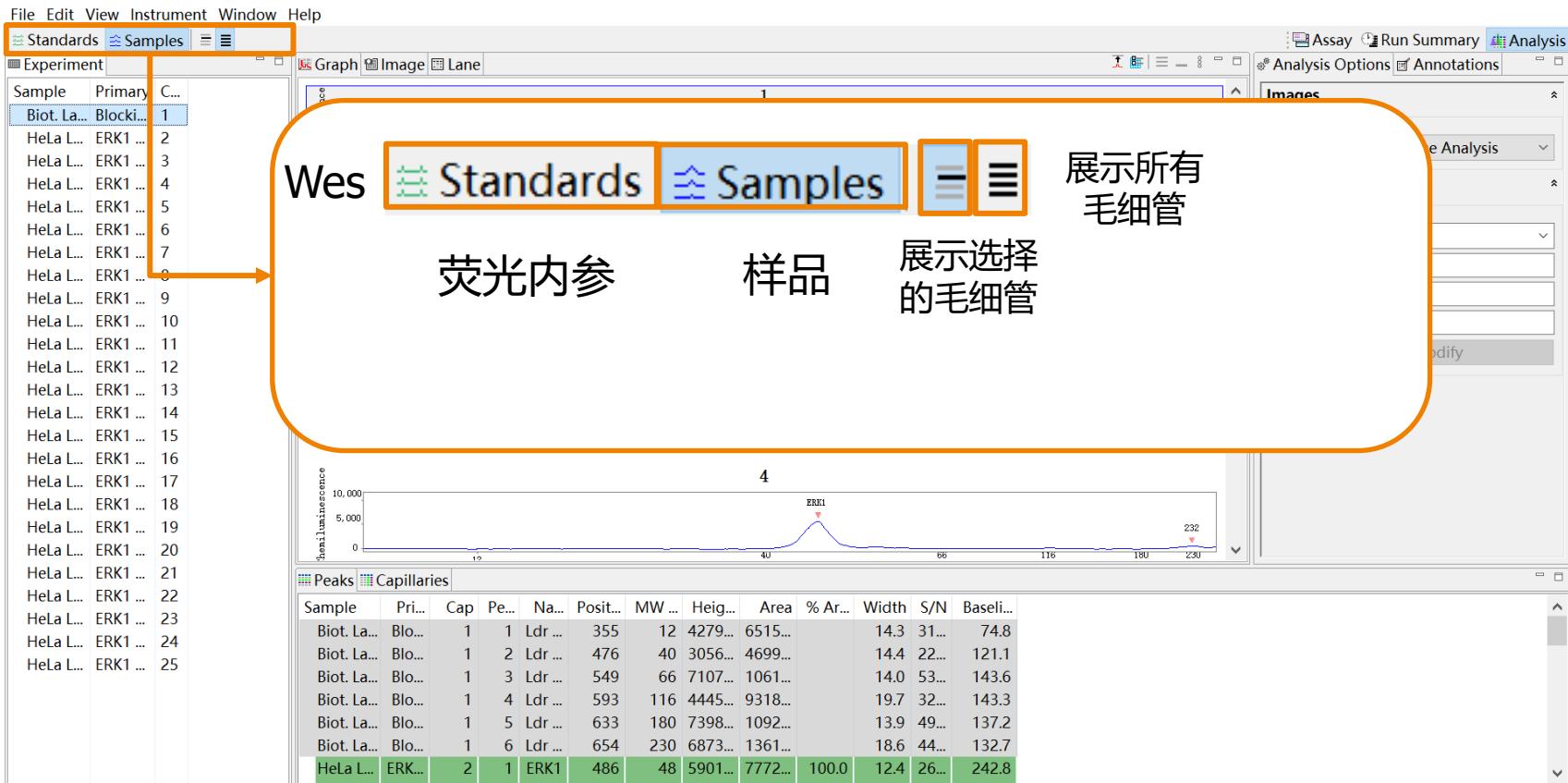
ERK1

232

Peaks Capillaries

Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

13 ANALYSIS 界面



13 ANALYSIS 界面

File Edit View Instrument Window Help

Standards Samples

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

Graph Image Lane

Wes Standards Samples

荧光内参 样品 展示选择的毛细管

Jess

1

4

ERK1

232

Chemiluminescence

10,000

5,000

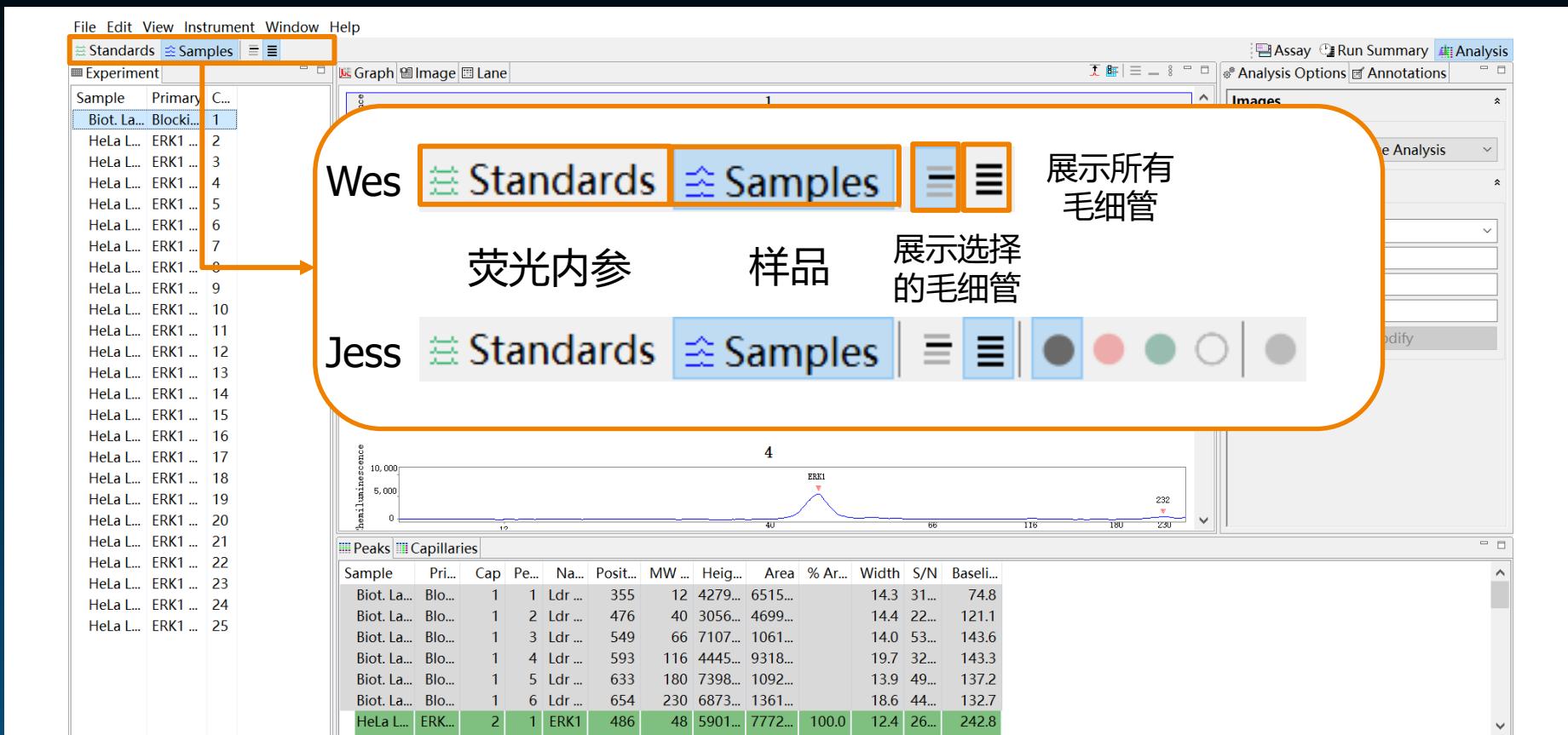
0

40 66 116 180 230

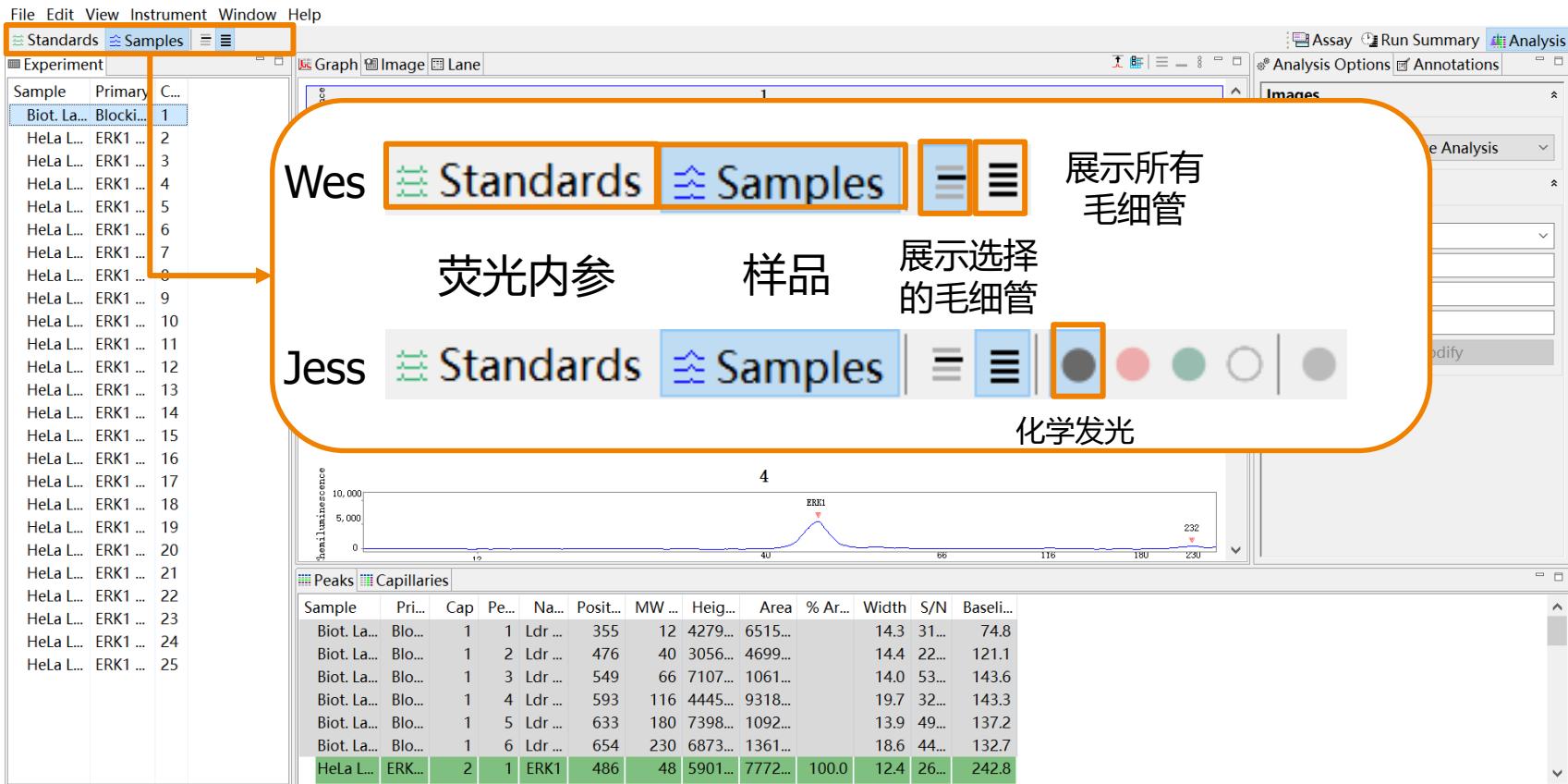
Peaks Capillaries

Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

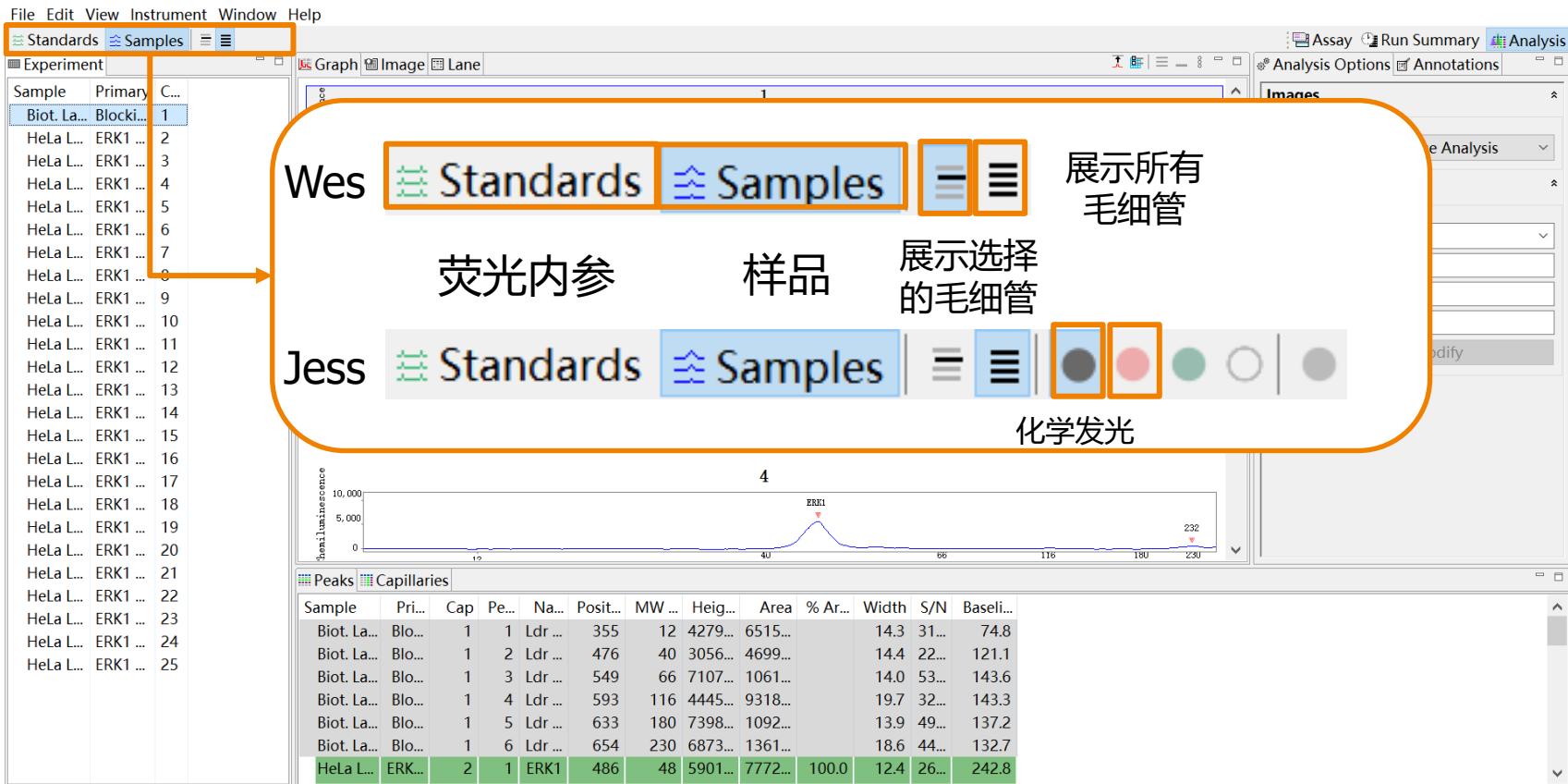
13 ANALYSIS 界面



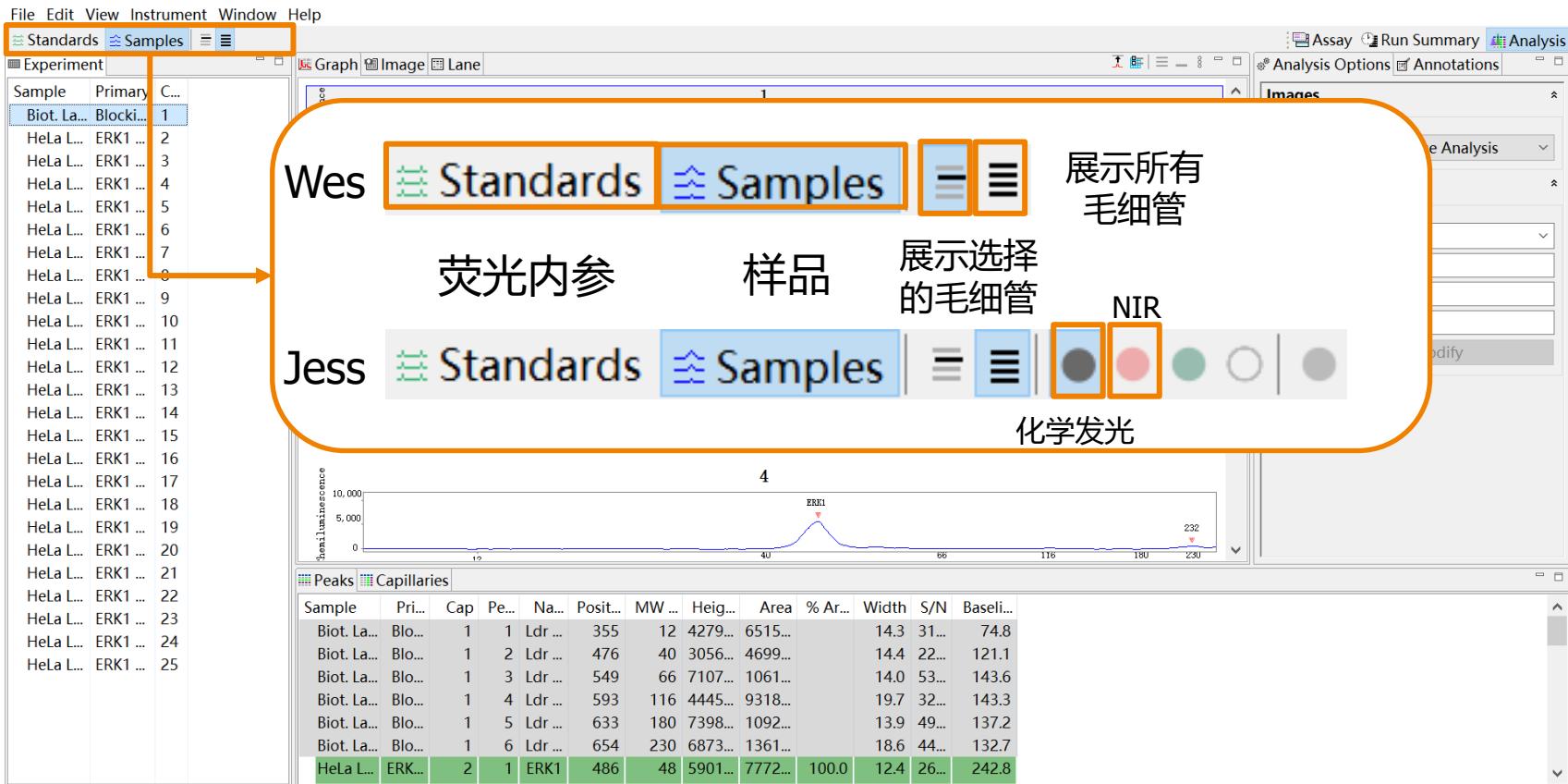
13 ANALYSIS 界面



13 ANALYSIS 界面



13 ANALYSIS 界面



13 ANALYSIS 界面

File Edit View Instrument Window Help

Standards Samples  

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

Graph Image Lane

Wes Standards Samples  

荧光内参 样品 展示选择的毛细管

Jess Standards Samples  

NIR 

化学发光

4

Chemiluminescence

10,000
5,000
0

ERK1

232

Peaks Capillaries

Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

Assay Run Summary Analysis

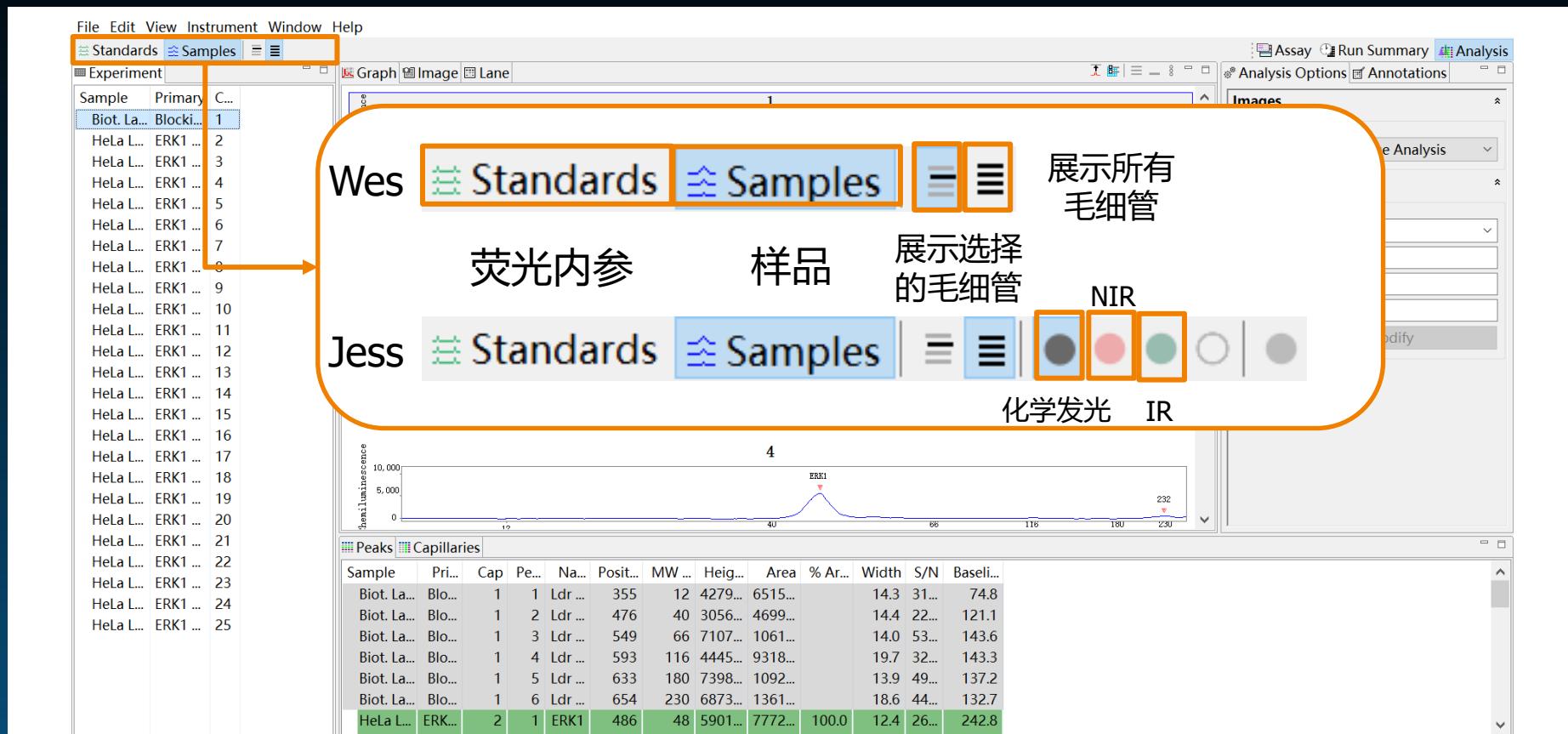
Analysis Options Annotations

Images

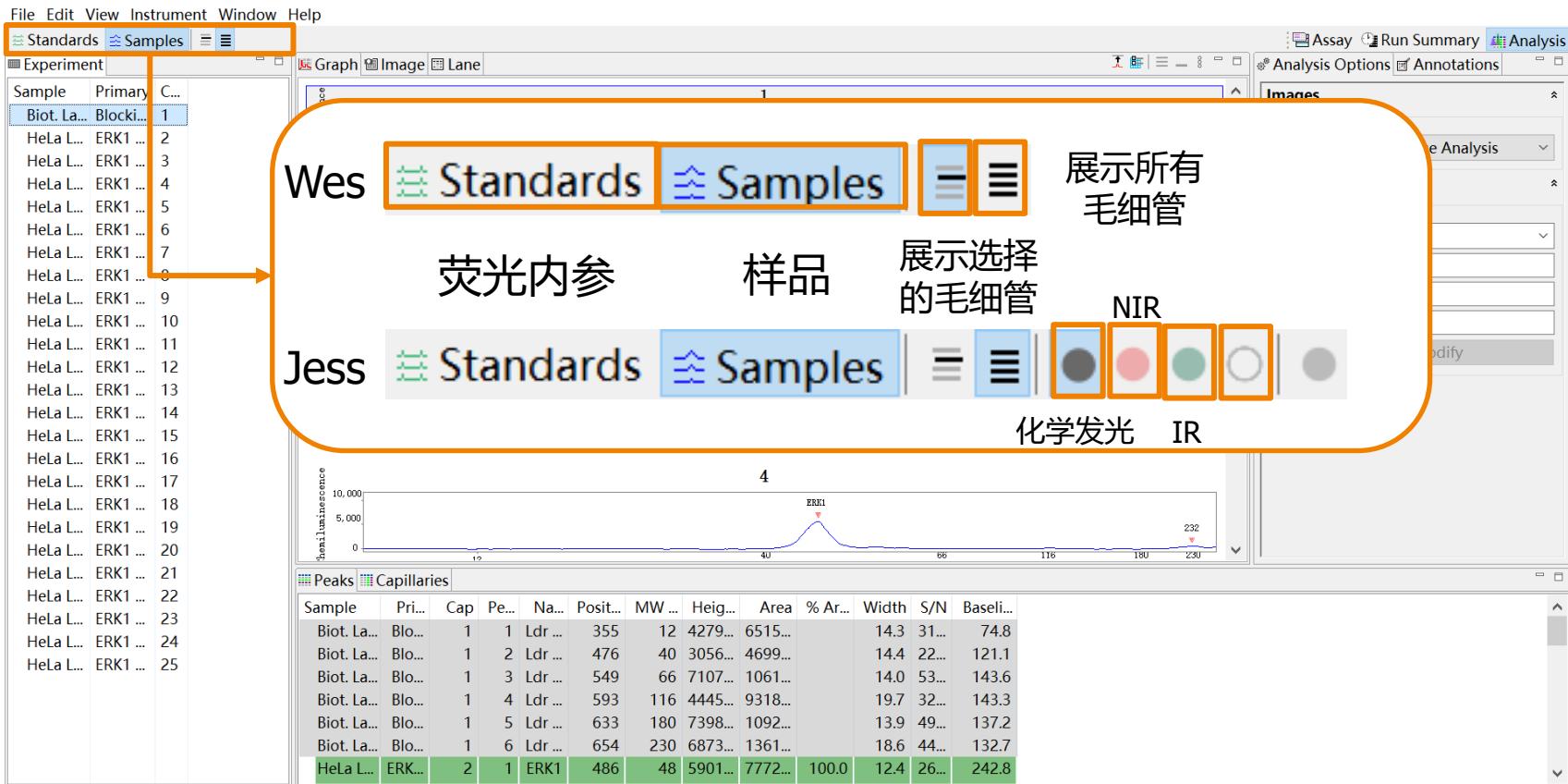
bio-analysis

Modify

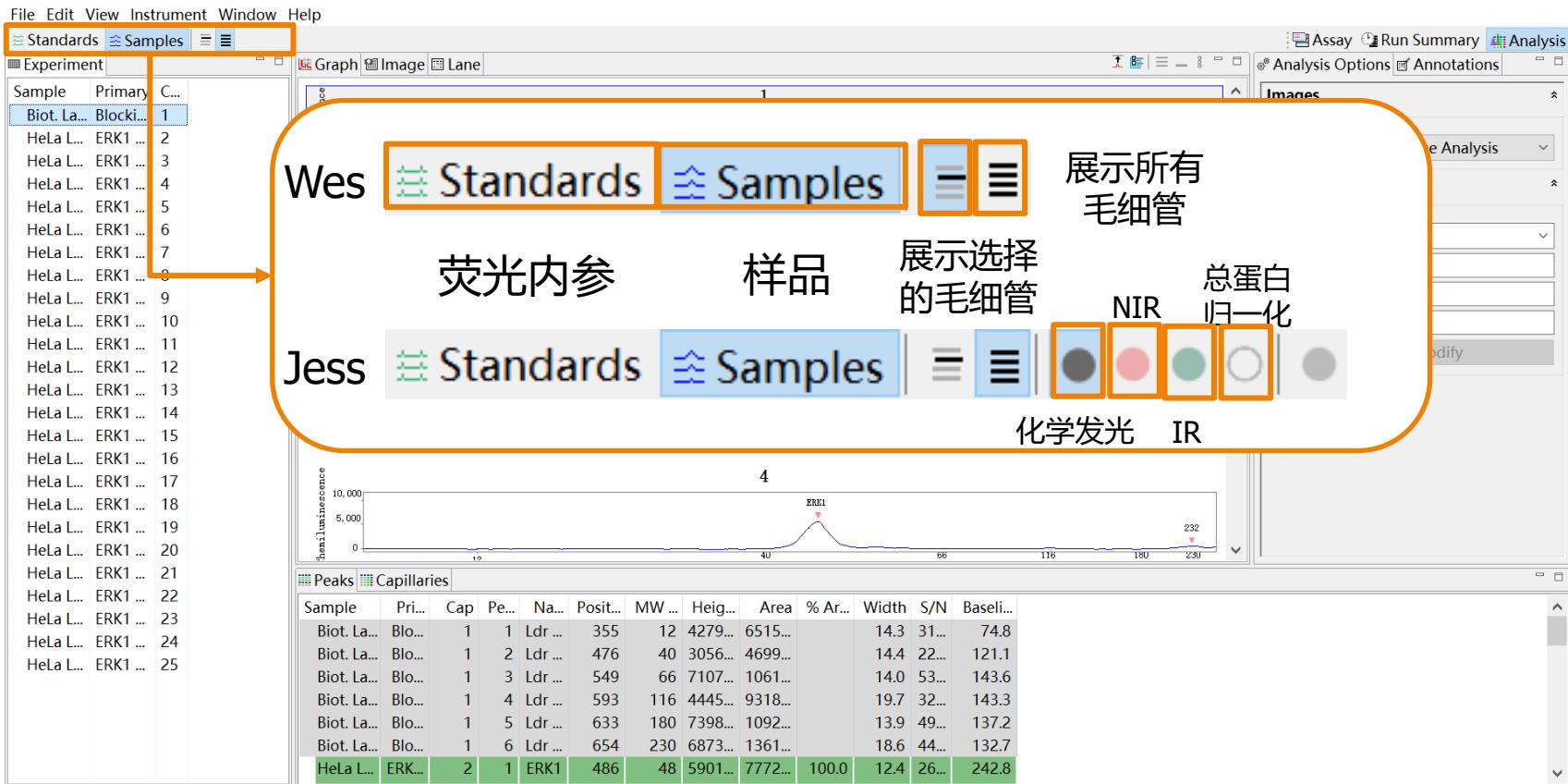
13 ANALYSIS 界面



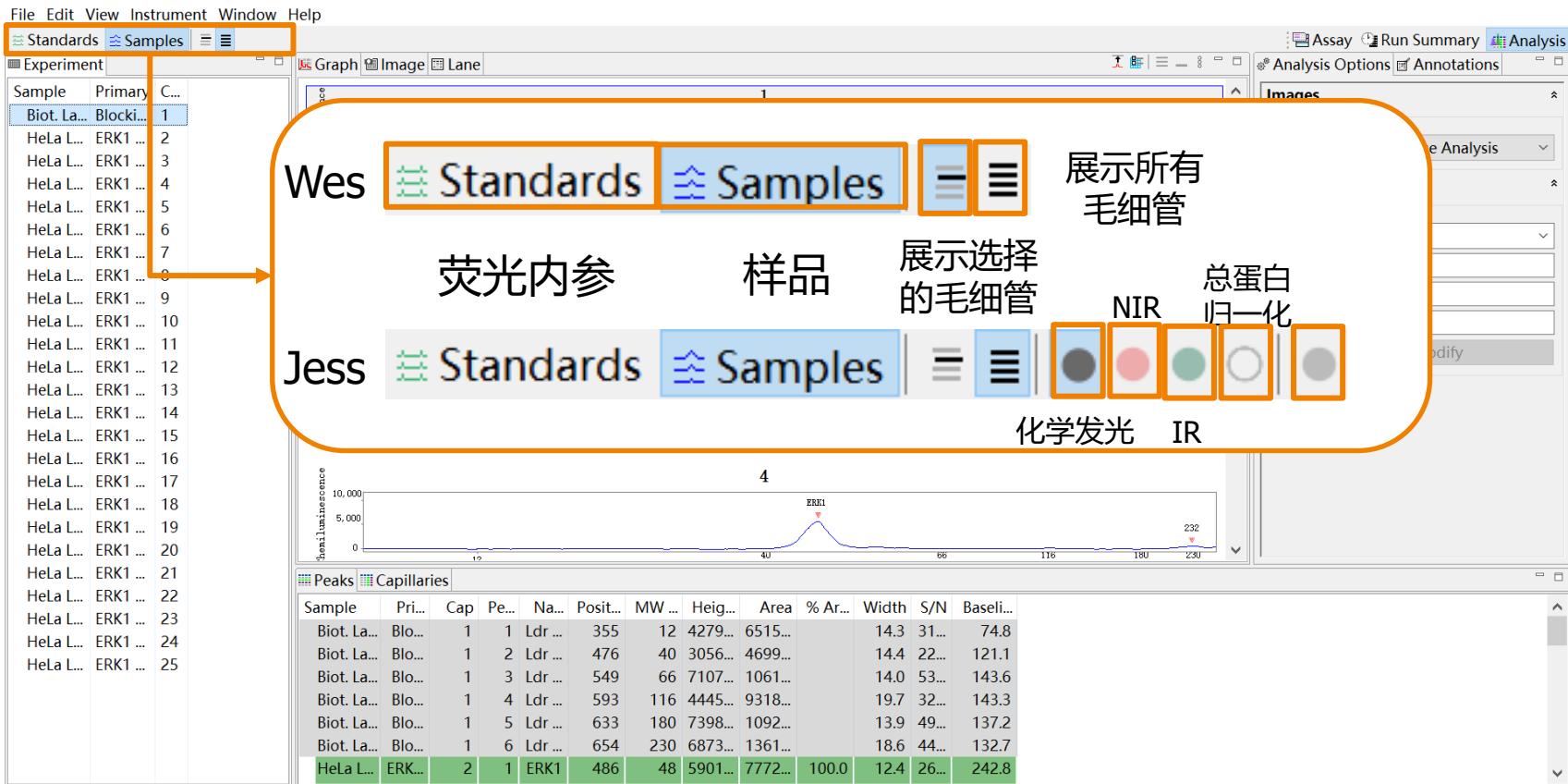
13 ANALYSIS 界面



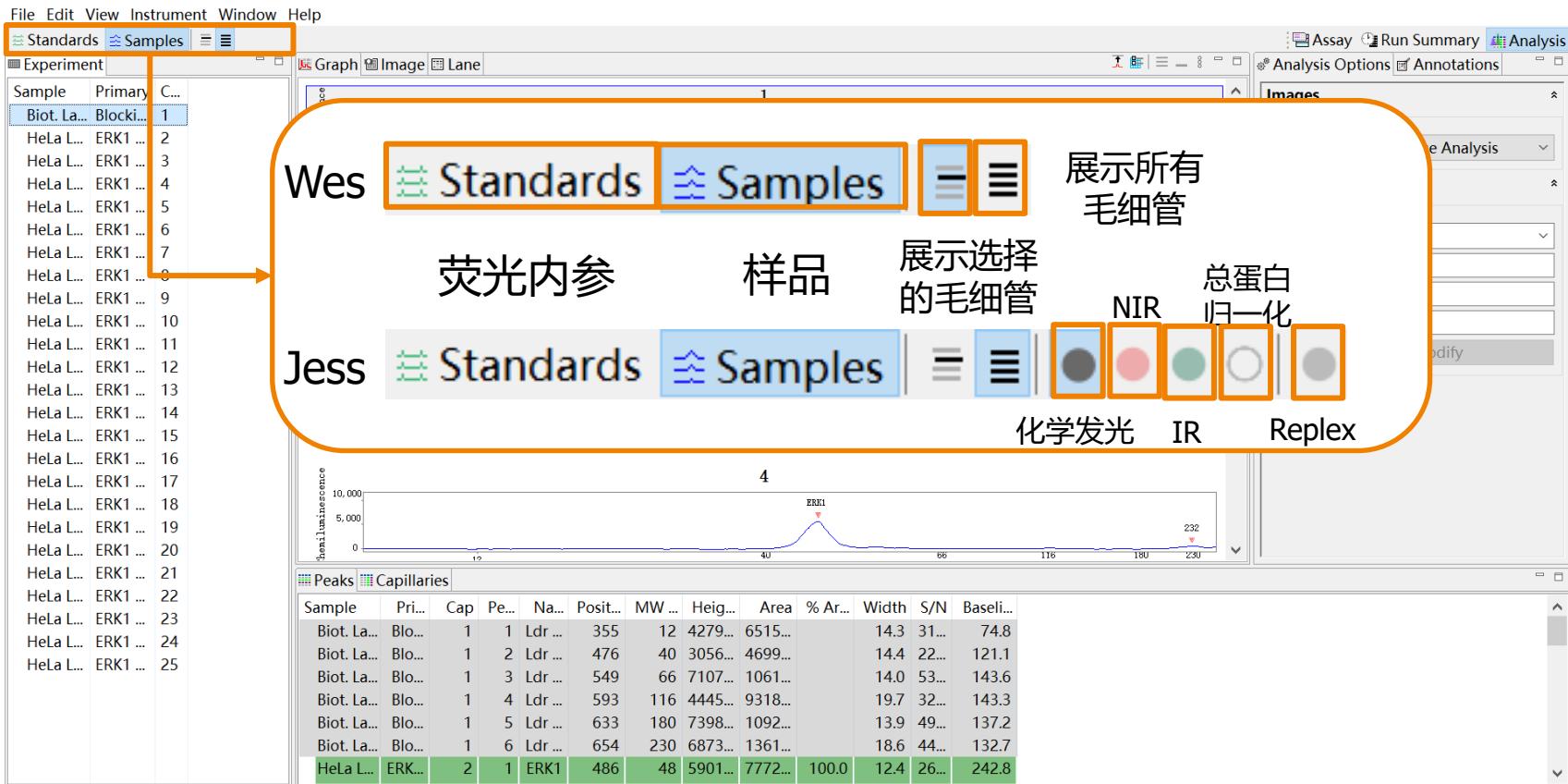
13 ANALYSIS 界面



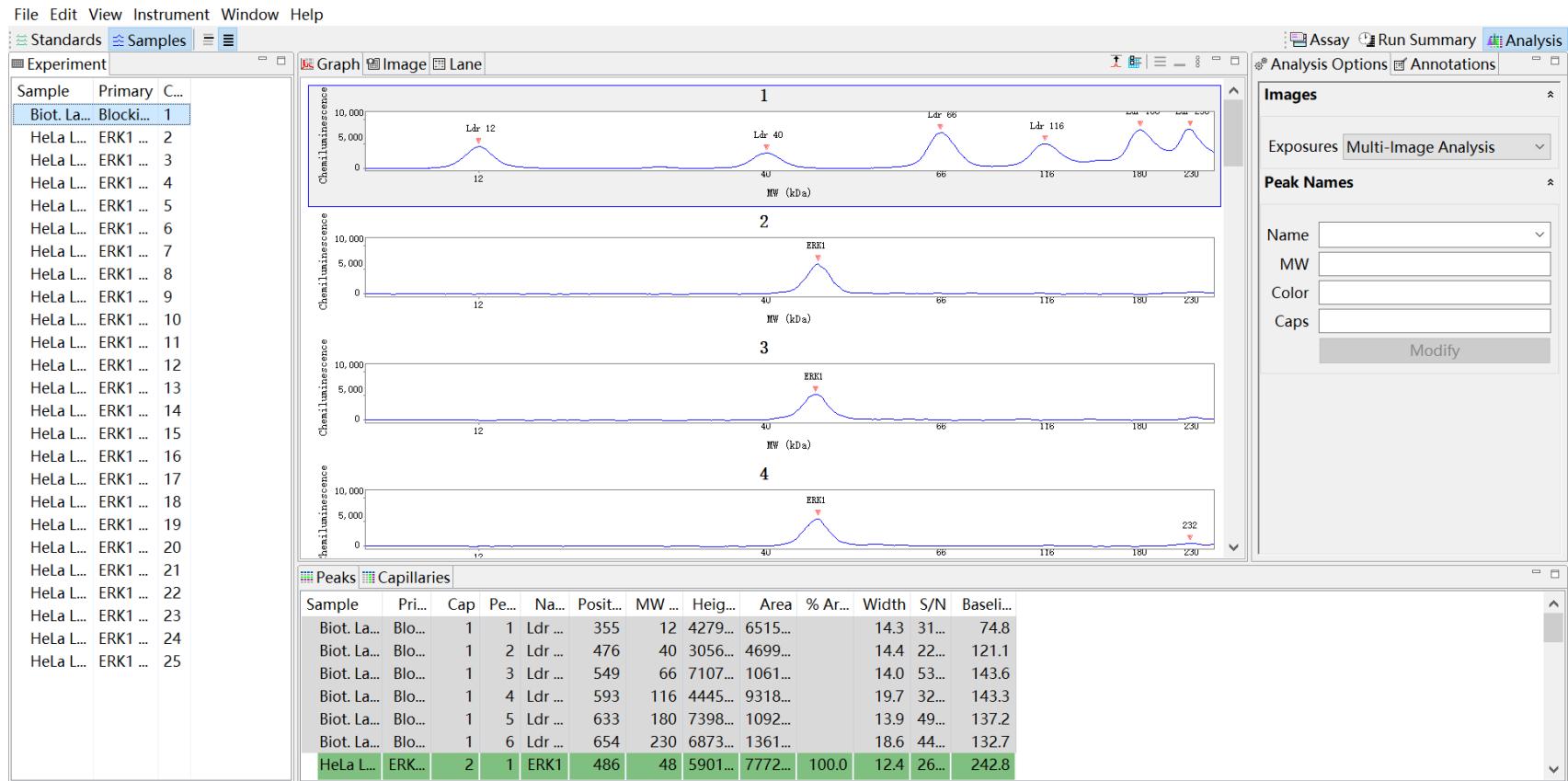
13 ANALYSIS 界面



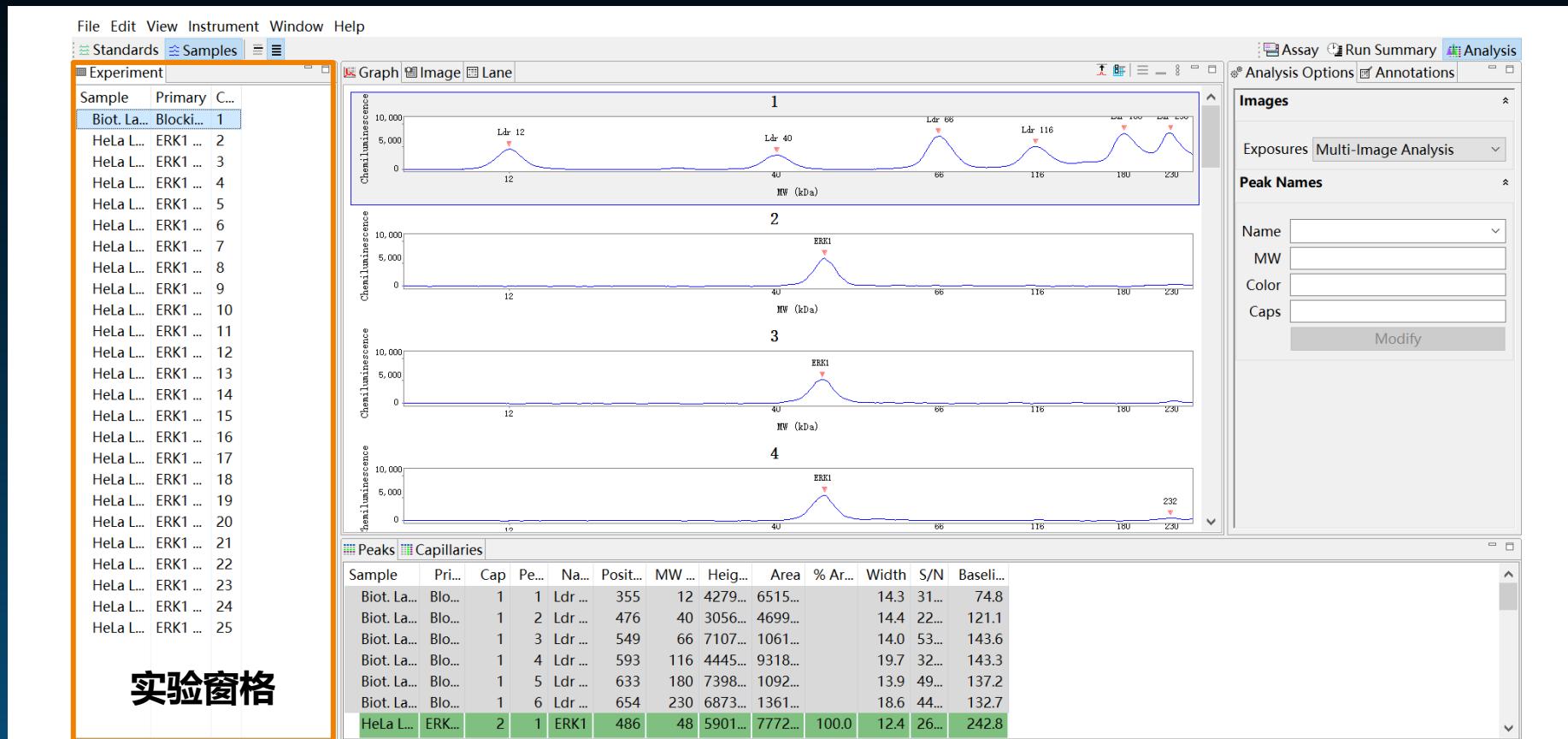
13 ANALYSIS 界面



ANALYSIS 界面



13 ANALYSIS 界面



13 ANALYSIS 界面

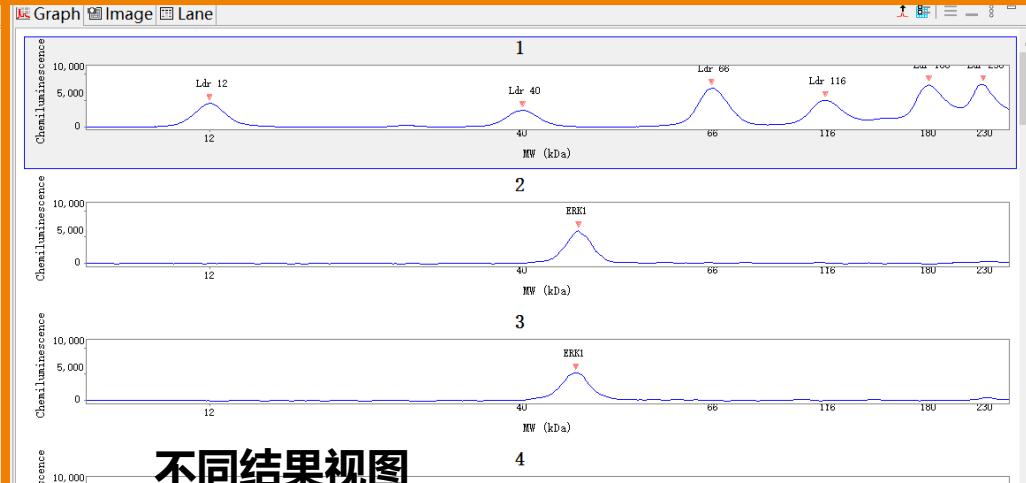
File Edit View Instrument Window Help

Standards Samples

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

实验窗格



Peaks Capillaries

Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

Assay Run Summary Analysis

Analysis Options Annotations

Images

Exposures Multi-Image Analysis

Peak Names

Name:

MW:

Color:

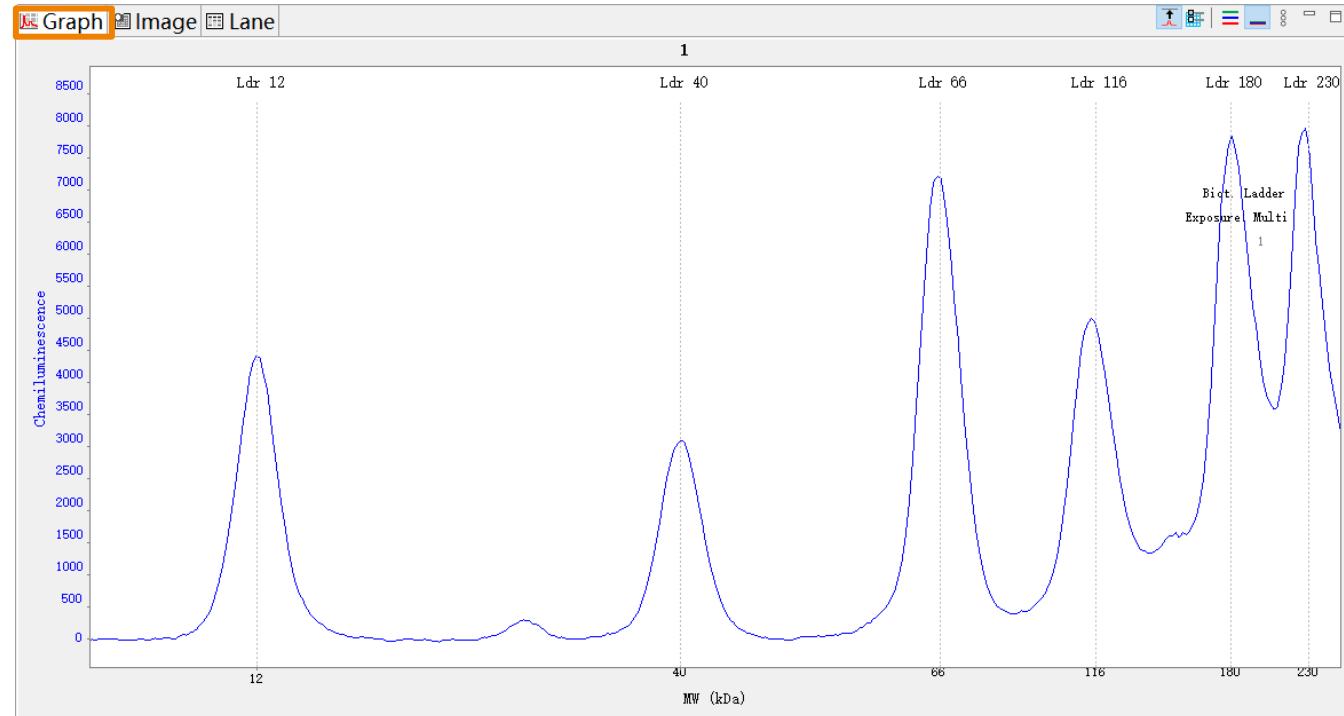
Caps:

Modify

14 ANALYSIS 界面

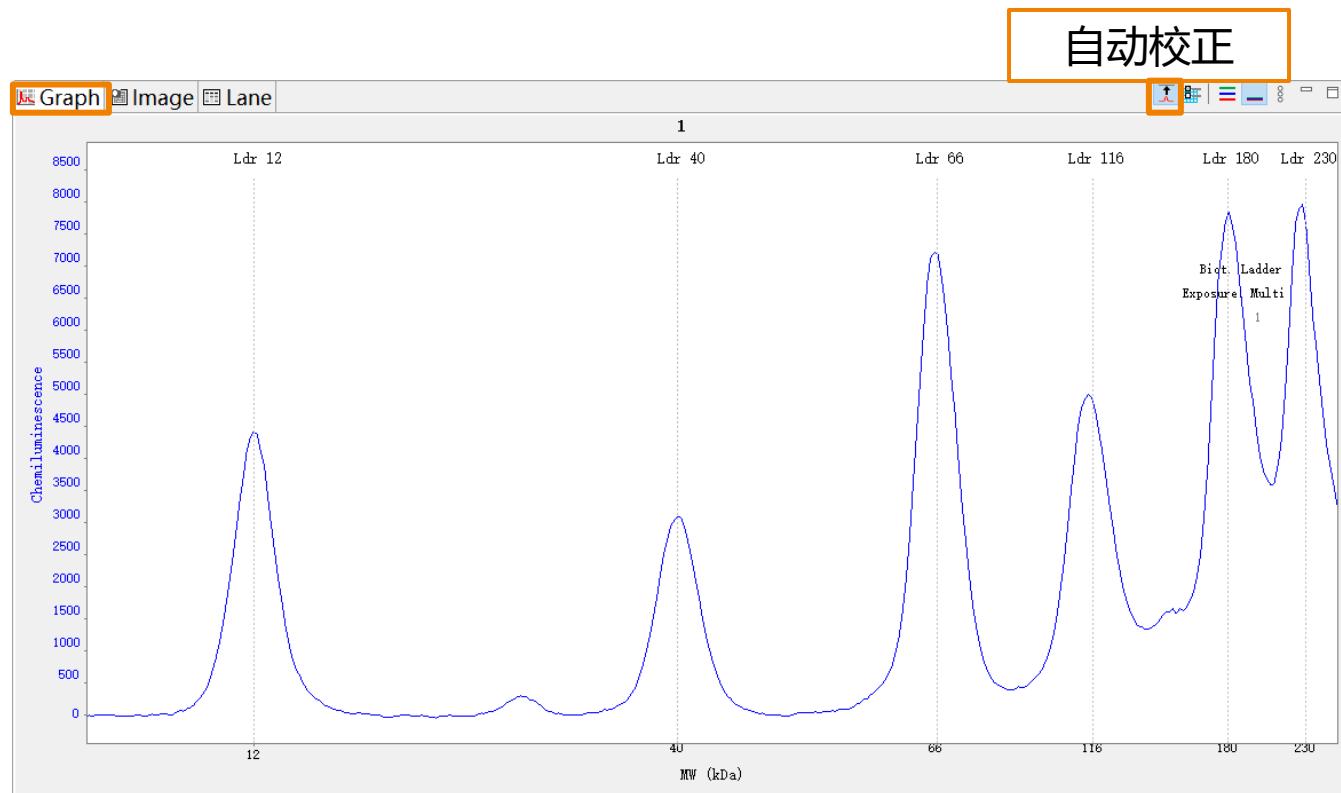
峰型图

- 峰形图信息量最大，数据质量分析推荐用峰形图
- 选择不同的通道，将有不止一个Y轴
- 点击 Auto Scale 将自动调整所有毛细管的量程



峰型图

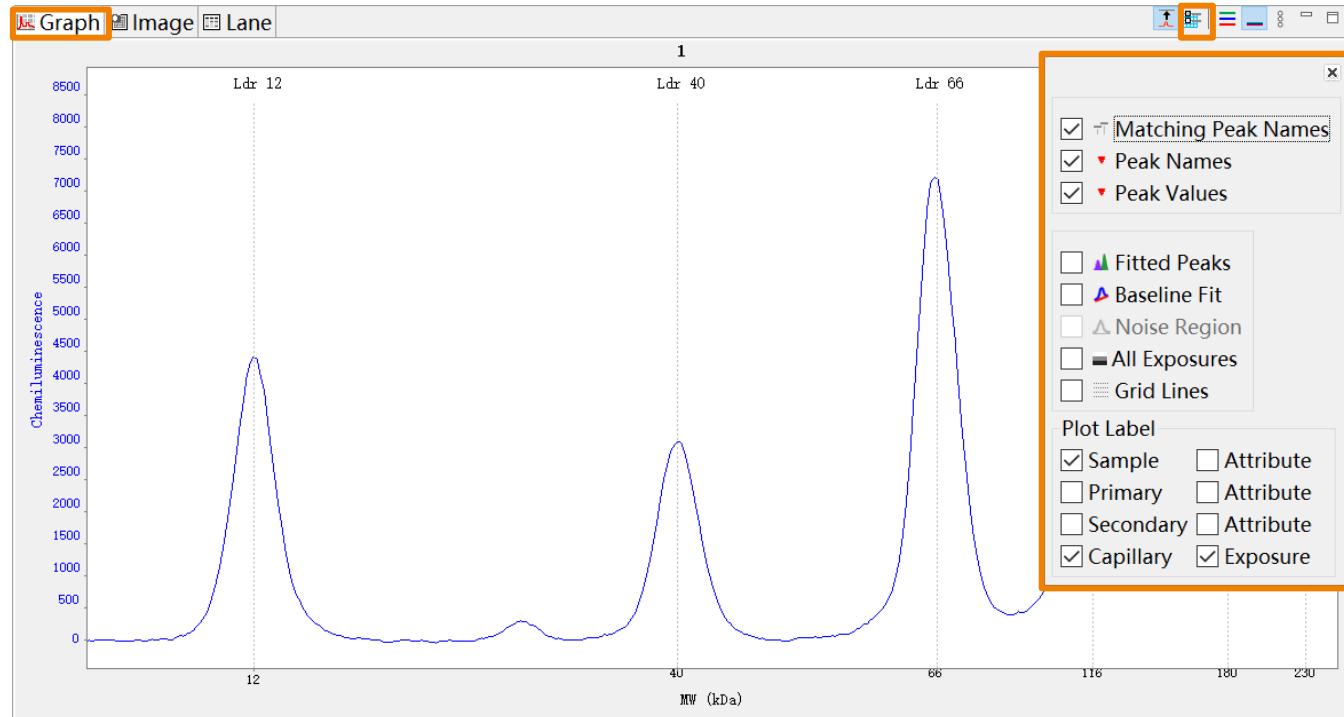
- 峰形图信息量最大，数据质量分析推荐用峰形图
 - 选择不同的通道，将有不止一个Y轴
 - 点击 Auto Scale 将自动调整所有毛细管的量程



14 ANALYSIS 界面

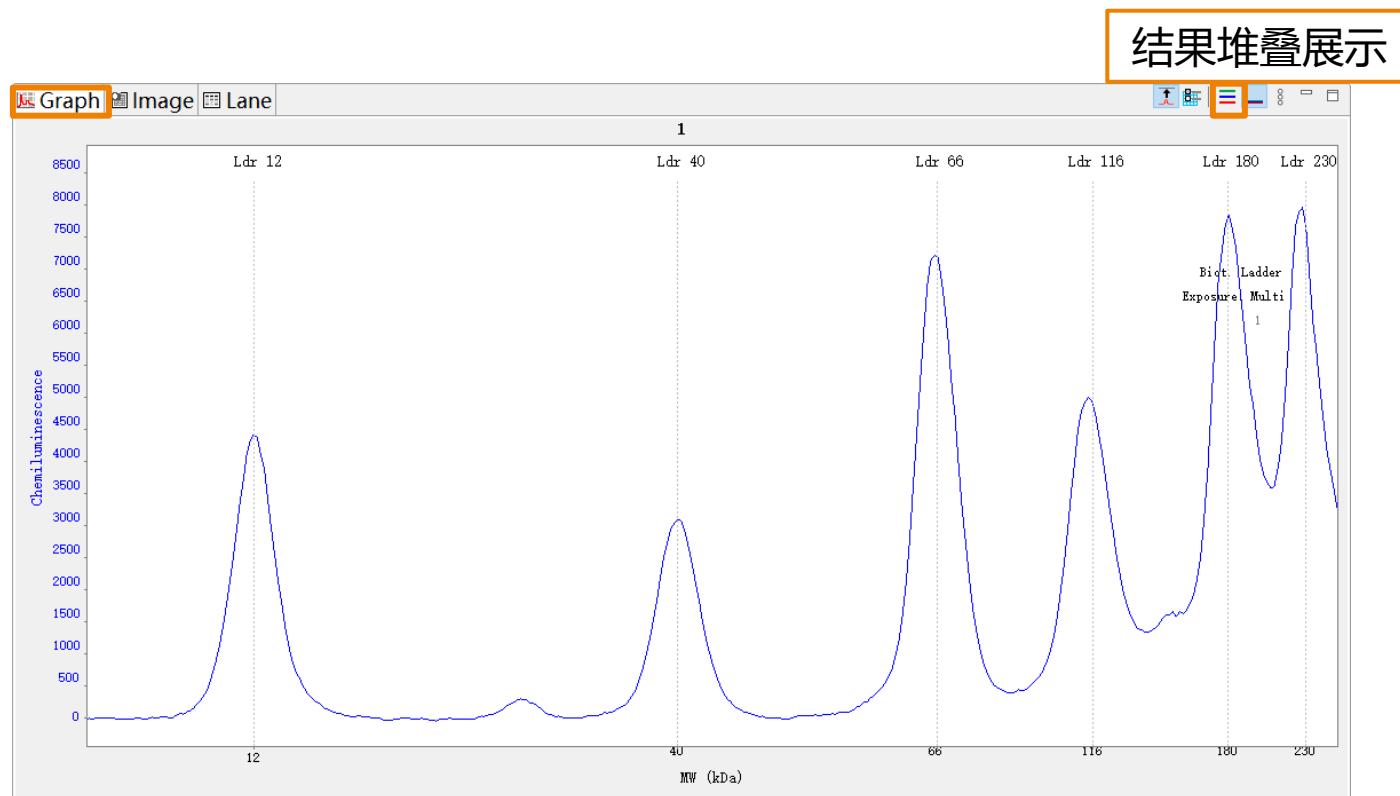
峰型图

- 峰形图信息量最大，数据质量分析推荐用峰形图
- 选择不同的通道，将有不止一个Y轴
- 点击 Auto Scale 将自动调整所有毛细管的量程



峰型图

- 峰形图信息量最大，数据质量分析推荐用峰形图
 - 选择不同的通道，将有不止一个Y轴
 - 点击 Auto Scale 将自动调整所有毛细管的量程

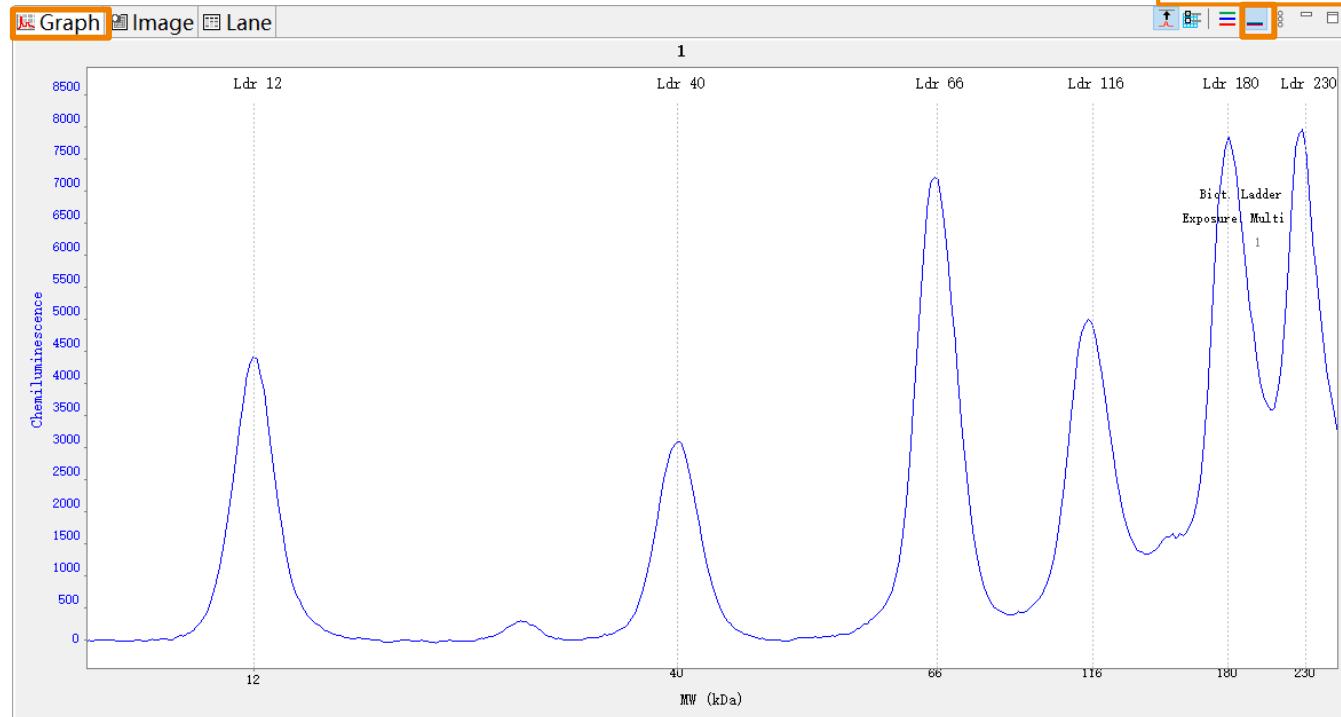


14 ANALYSIS 界面

峰型图

- 峰形图信息量最大，数据质量分析推荐用峰形图
- 选择不同的通道，将有不止一个Y轴
- 点击 Auto Scale 将自动调整所有毛细管的量程

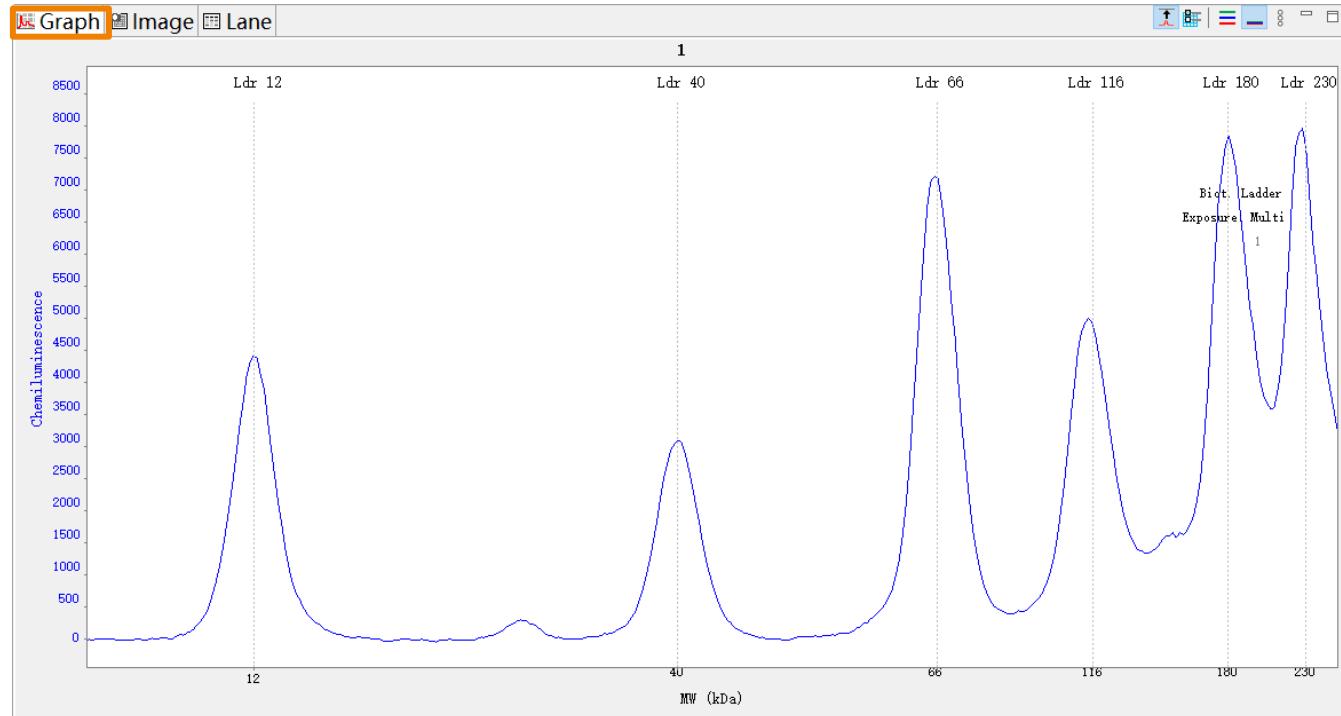
结果合并展示



14 ANALYSIS 界面

峰型图

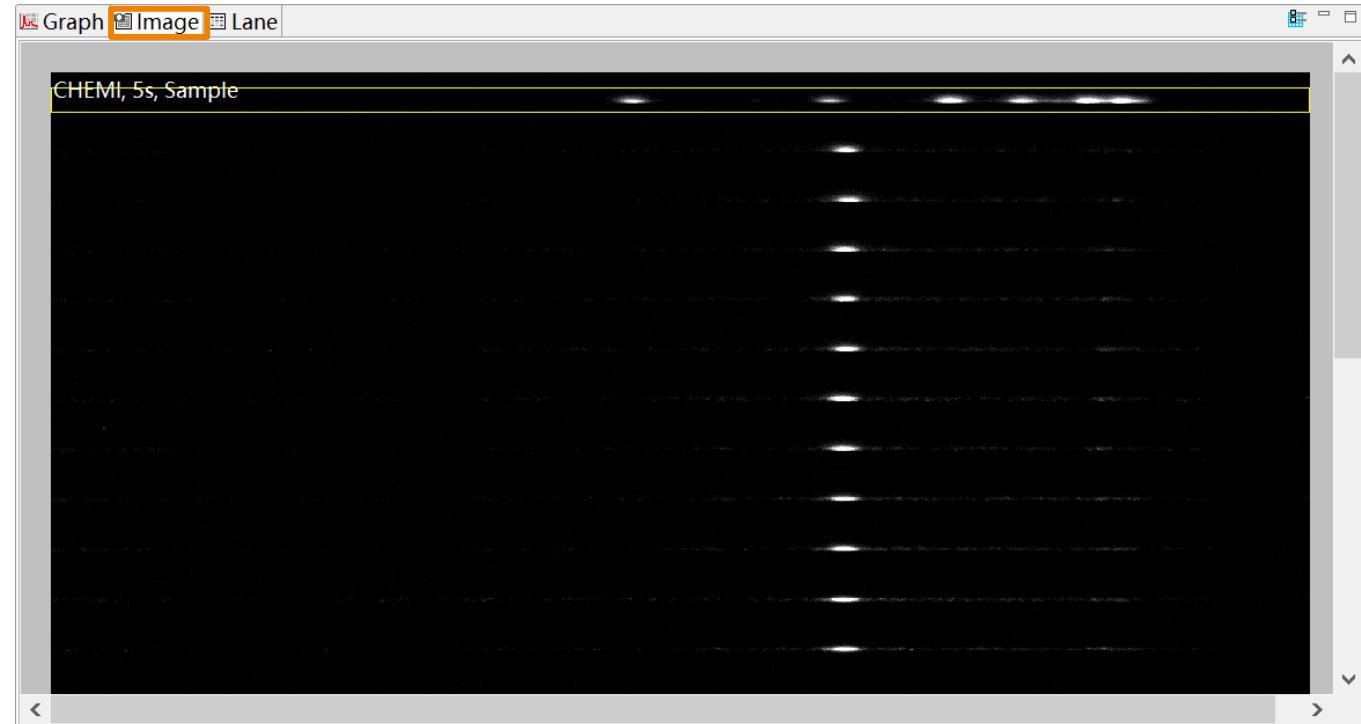
- 峰形图信息量最大，数据质量分析推荐用峰形图
- 选择不同的通道，将有不止一个Y轴
- 点击 Auto Scale 将自动调整所有毛细管的量程



ANALYSIS 界面

成像图

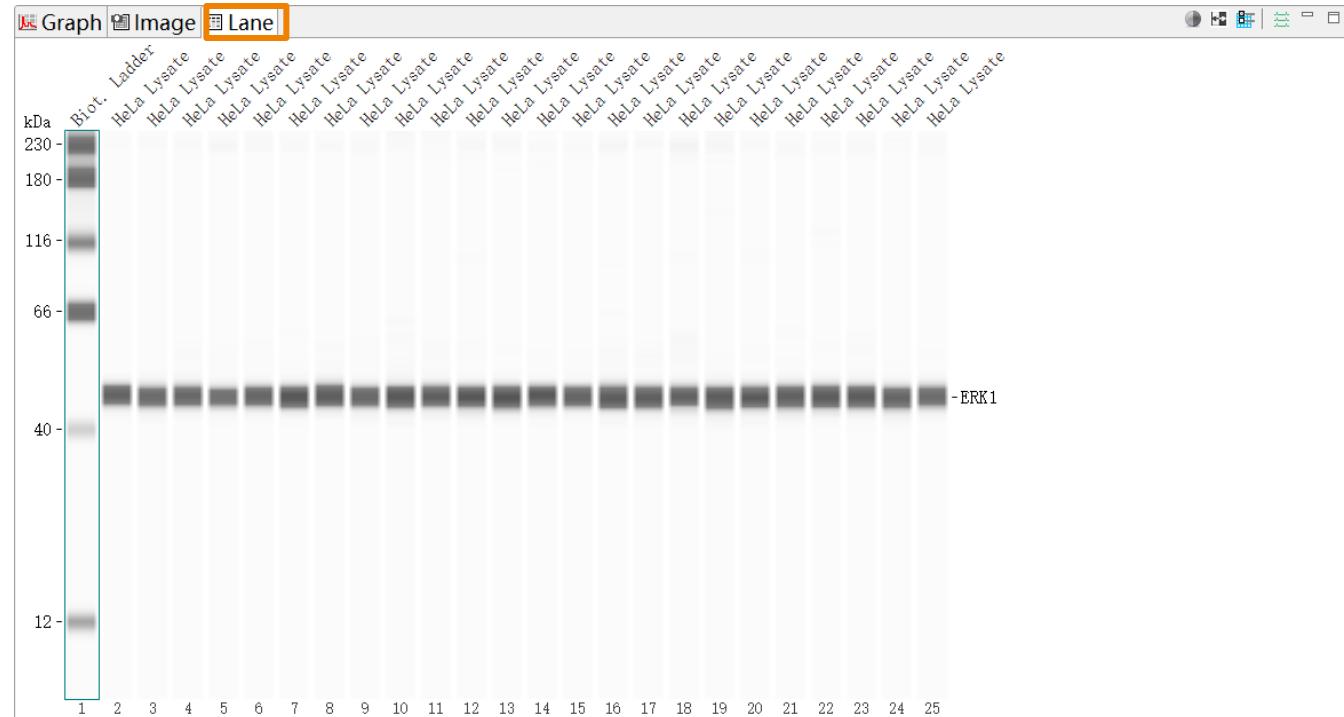
毛细管实际曝光
结果



ANALYSIS 界面

泳道图

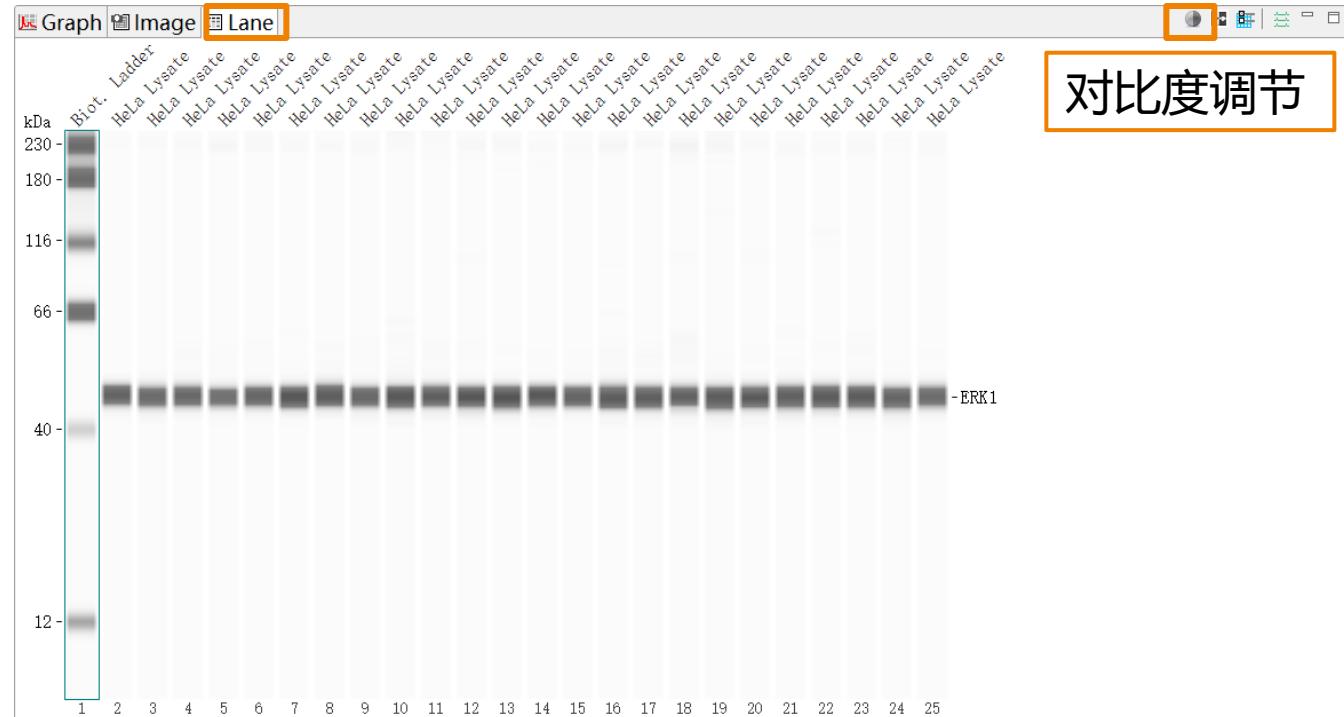
- 大部分报告的结果形式
- 可以任意显示或横向移动泳道
- 可以单独调整每个通道的对比度
 - 对比度调整仅改变图片中条带呈现效果, 峰定量结果不变



ANALYSIS 界面

泳道图

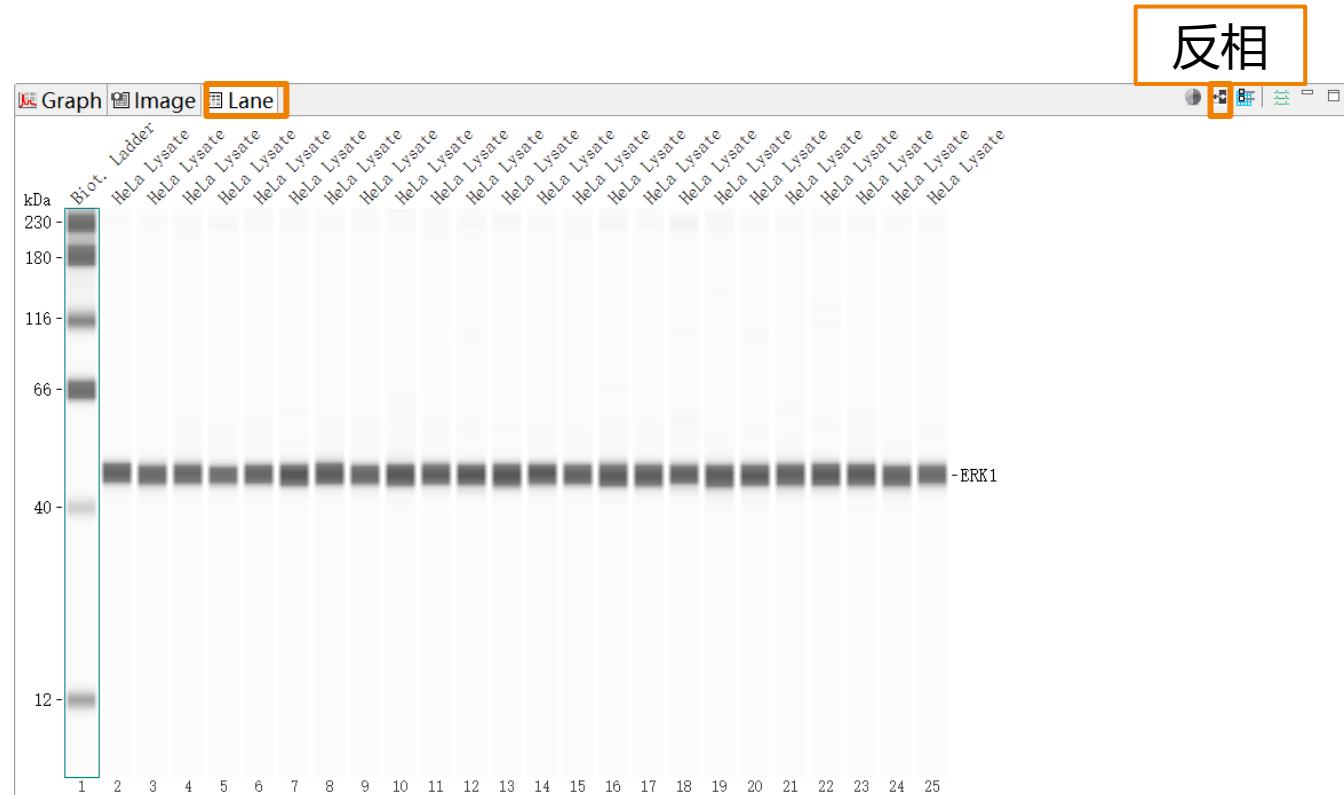
- 大部分报告的结果形式
- 可以任意显示或横向移动泳道
- 可以单独调整每个通道的对比度
 - 对比度调整仅改变图片中条带呈现效果, 峰定量结果不变



ANALYSIS 界面

泳道图

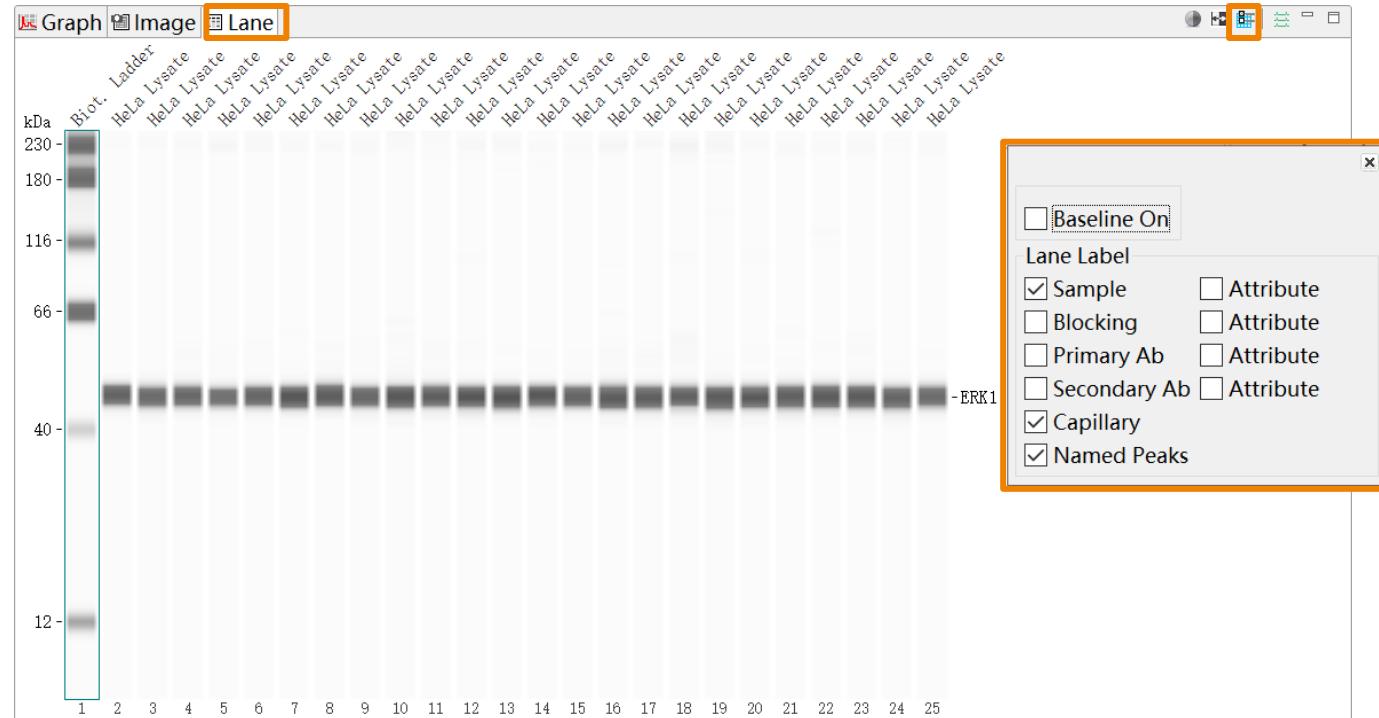
- 大部分报告的结果形式
- 可以任意显示或横向移动泳道
- 可以单独调整每个通道的对比度
 - 对比度调整仅改变图片中条带呈现效果, 峰定量结果不变



ANALYSIS 界面

泳道图

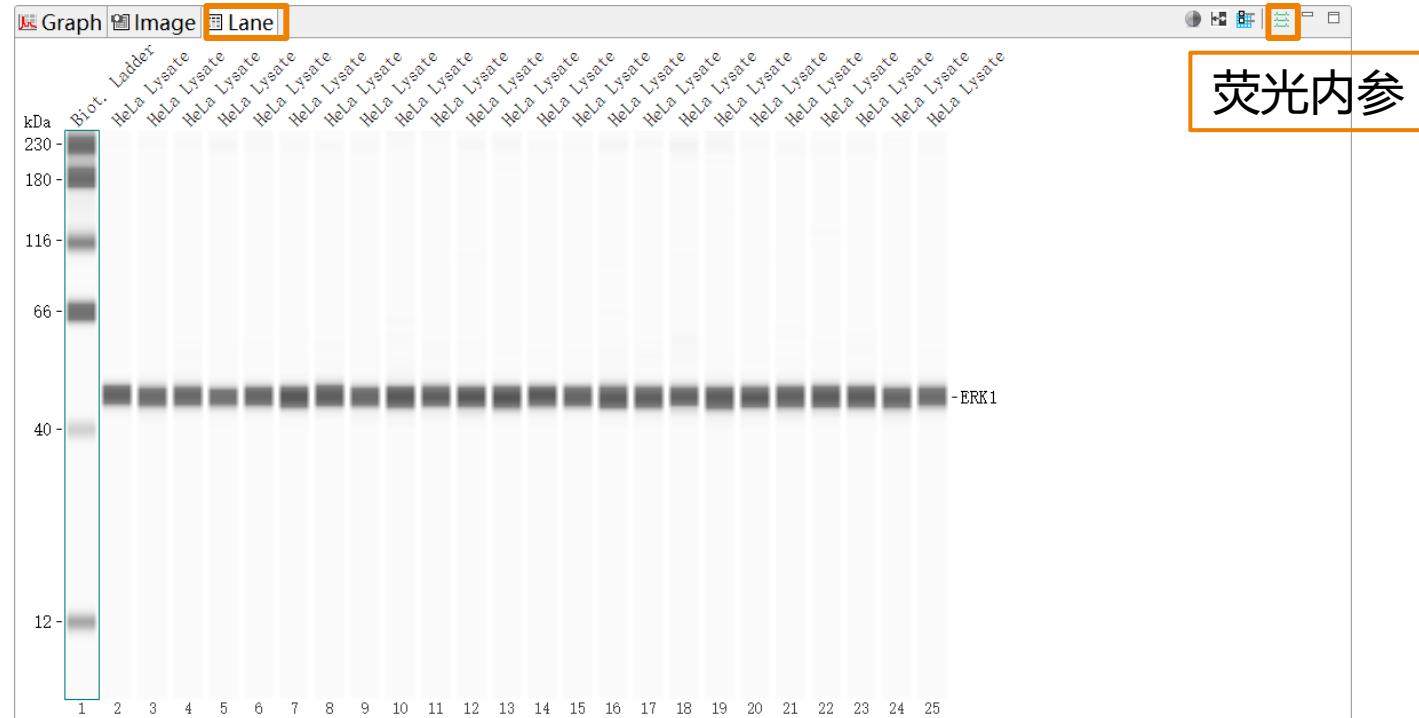
- 大部分报告的结果形式
- 可以任意显示或横向移动泳道
- 可以单独调整每个通道的对比度
 - 对比度调整仅改变图片中条带呈现效果, 峰定量结果不变



ANALYSIS 界面

泳道图

- 大部分报告的结果形式
- 可以任意显示或横向移动泳道
- 可以单独调整每个通道的对比度
 - 对比度调整仅改变图片中条带呈现效果, 峰定量结果不变



ANALYSIS 界面

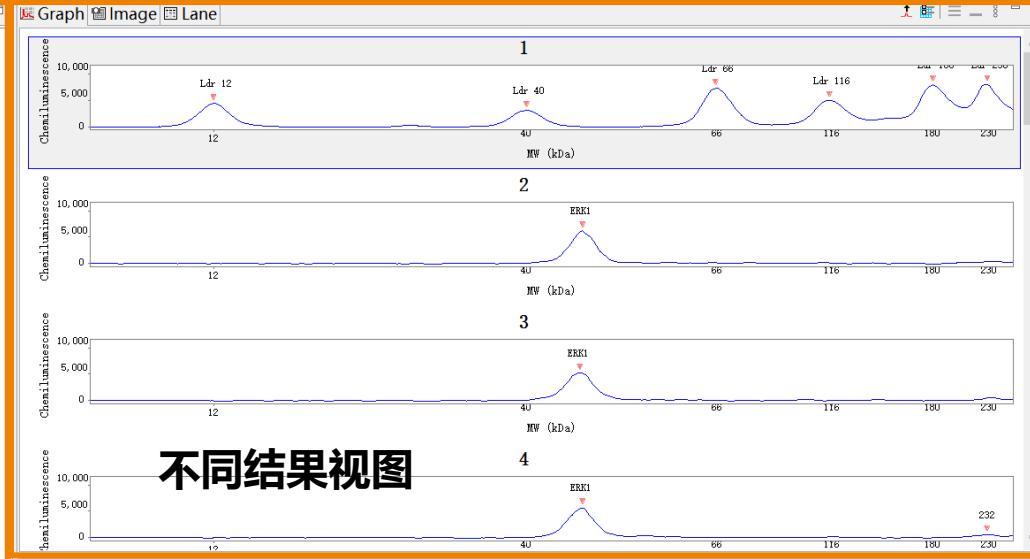
File Edit View Instrument Window Help

Standards Samples

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

实验窗格



Peaks Capillaries

Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

Assay Run Summary Analysis

Analysis Options Annotations

Images

Exposures Multi-Image Analysis

Peak Names

Name	<input type="text"/>
MW	<input type="text"/>
Color	<input type="text"/>
Caps	<input type="text"/>
Modify	

Modify

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ANALYSIS 界面

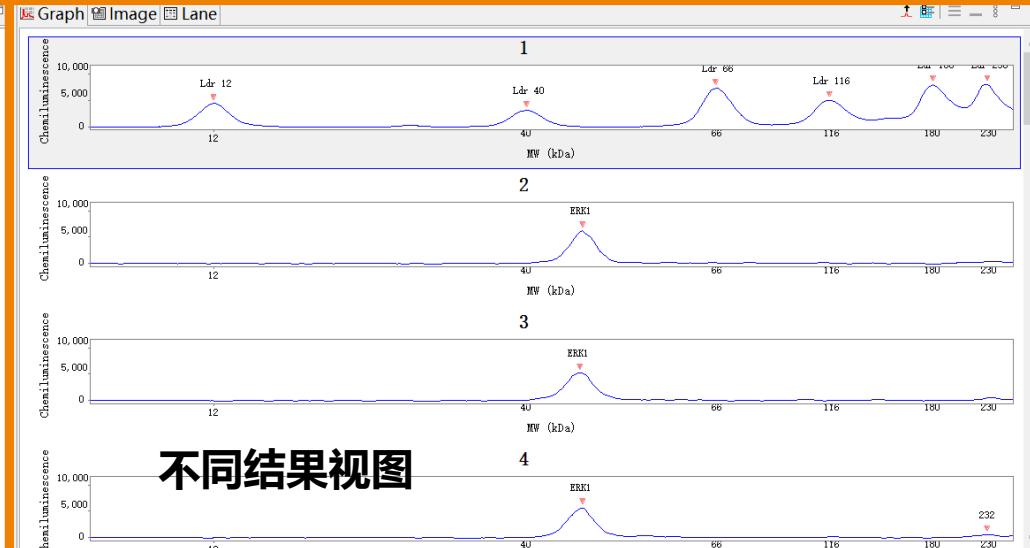
File Edit View Instrument Window Help

Standards Samples

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

实验窗格



Assay Run Summary Analysis

Analysis Options Annotations

Images

Exposures Multi-Image Analysis

Peak Names

Name	<input type="text"/>
MW	<input type="text"/>
Color	<input type="text"/>
Caps	<input type="text"/>
Modify	

峰统计数据表

Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

ANALYSIS 界面

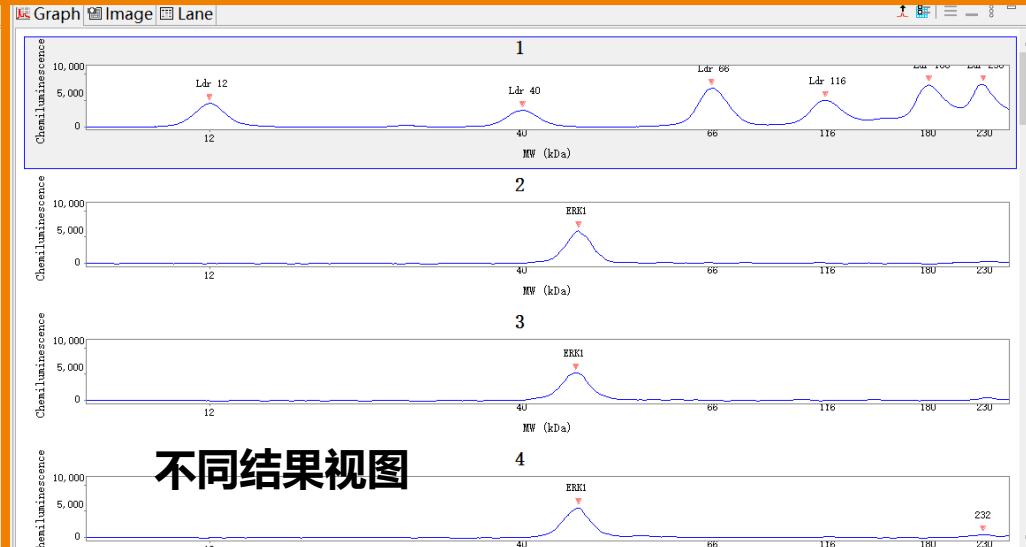
File Edit View Instrument Window Help

Standards Samples

Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25

实验窗格



Sample	Pri...	Cap	Pe...	Na...	Posit...	MW ...	Heig...	Area	% Ar...	Width	S/N	Baseli...
Biot. La...	Blo...	1	1	Ldr ...	355	12	4279...	6515...	14.3	31...	74.8	
Biot. La...	Blo...	1	2	Ldr ...	476	40	3056...	4699...	14.4	22...	121.1	
Biot. La...	Blo...	1	3	Ldr ...	549	66	7107...	1061...	14.0	53...	143.6	
Biot. La...	Blo...	1	4	Ldr ...	593	116	4445...	9318...	19.7	32...	143.3	
Biot. La...	Blo...	1	5	Ldr ...	633	180	7398...	1092...	13.9	49...	137.2	
Biot. La...	Blo...	1	6	Ldr ...	654	230	6873...	1361...	18.6	44...	132.7	
HeLa L...	ERK...	2	1	ERK1	486	48	5901...	7772...	100.0	12.4	26...	242.8

峰统计数据表

Assay Run Summary Analysis

Analysis Options Annotations

Images

Exposures Multi-Image Analysis

Peak Names

Name	<input type="text"/>
MW	<input type="text"/>
Color	<input type="text"/>
Caps	<input type="text"/>
Modify	

分析选项窗格

01

Compass 介绍

02

数据分析前检查

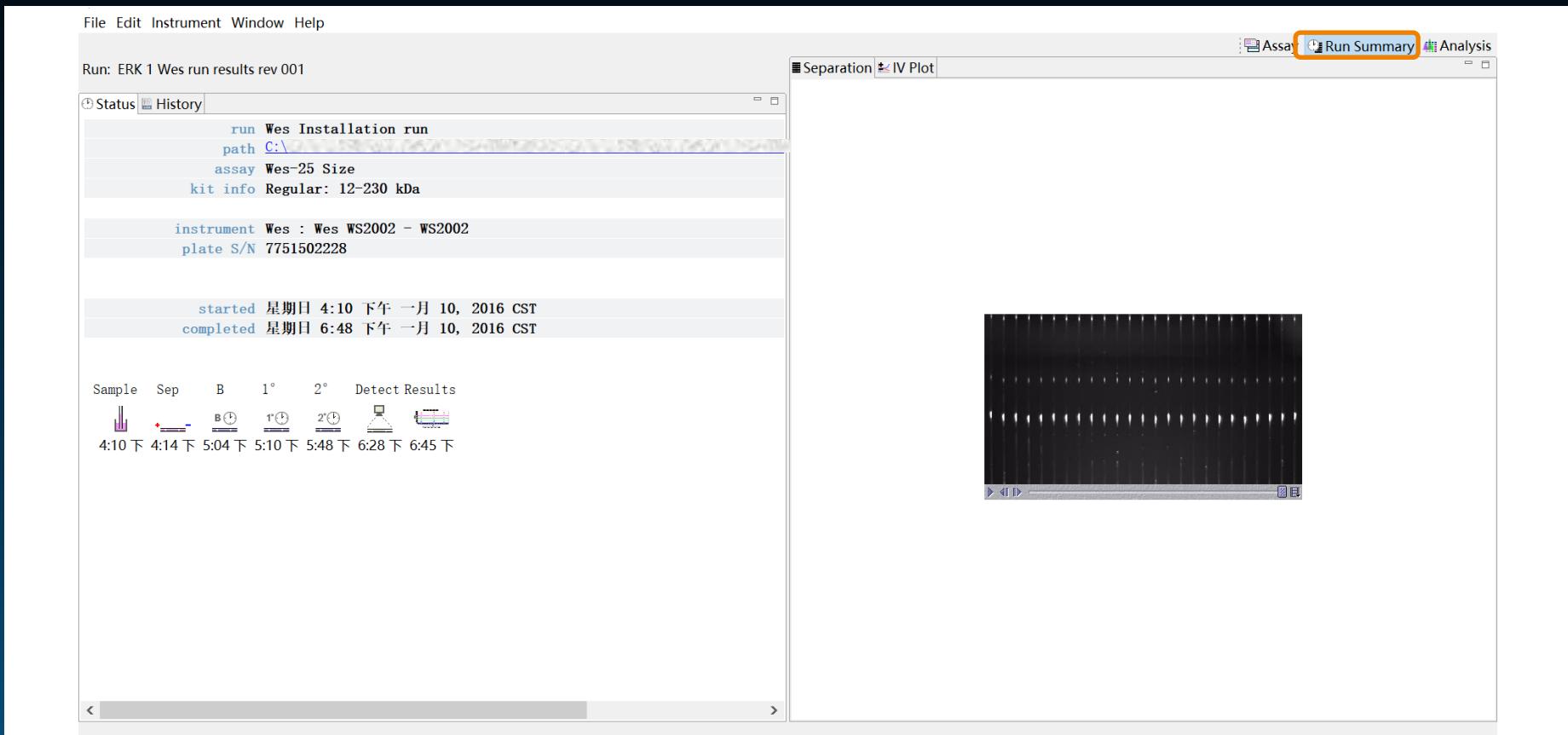
03

数据分析

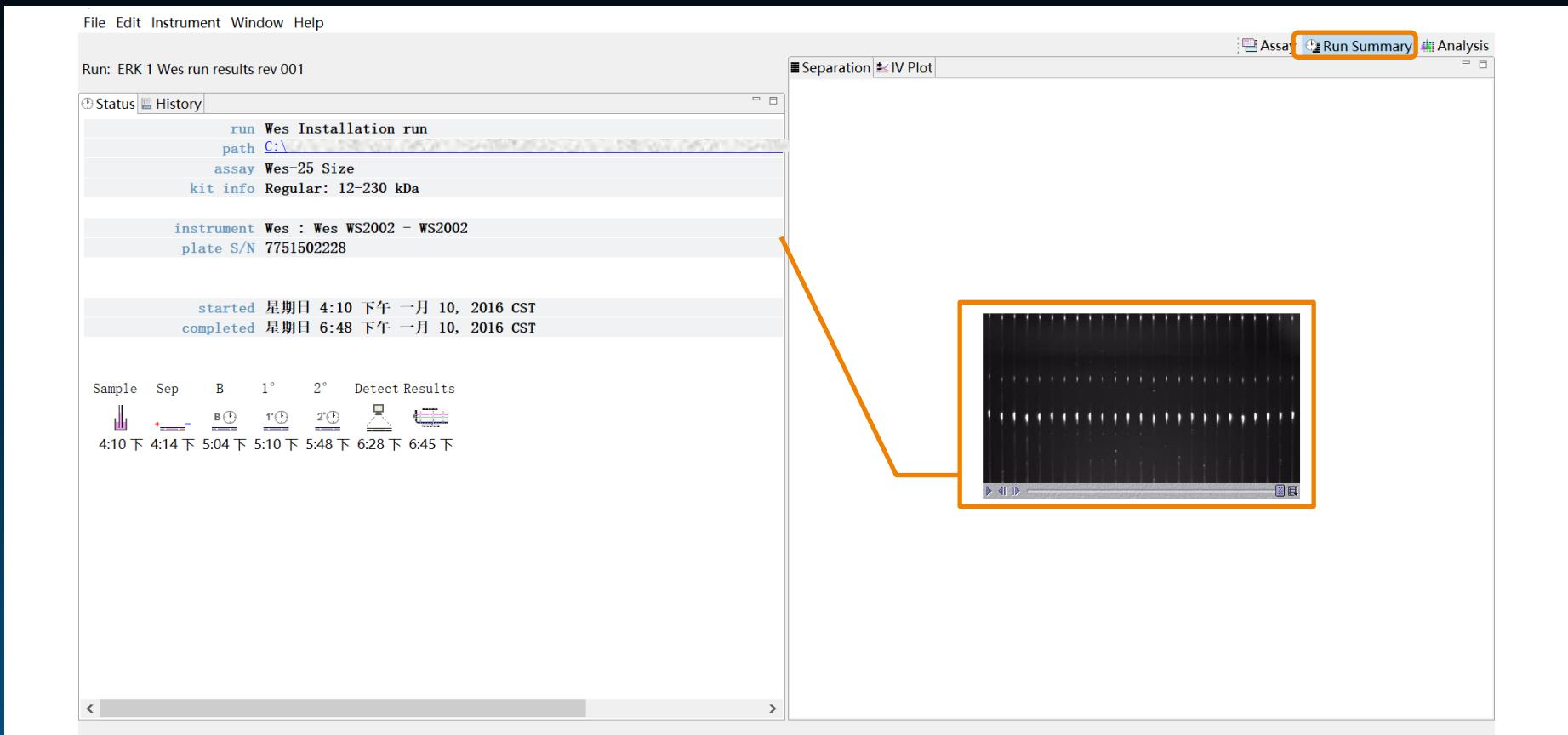
数据分析前检查

- 请确保在查看数据前，先通过几个部分的检查确认实验没有问题，包括荧光内参确认、Ladder确认、数据质量确认
 - 荧光内参整齐并识别良好
 - Ladder 识别良好
 - 化学发光信号无过曝
 - 荧光通道相机无饱和 (Jess)
 - 背景无过高
 - 峰面积正常

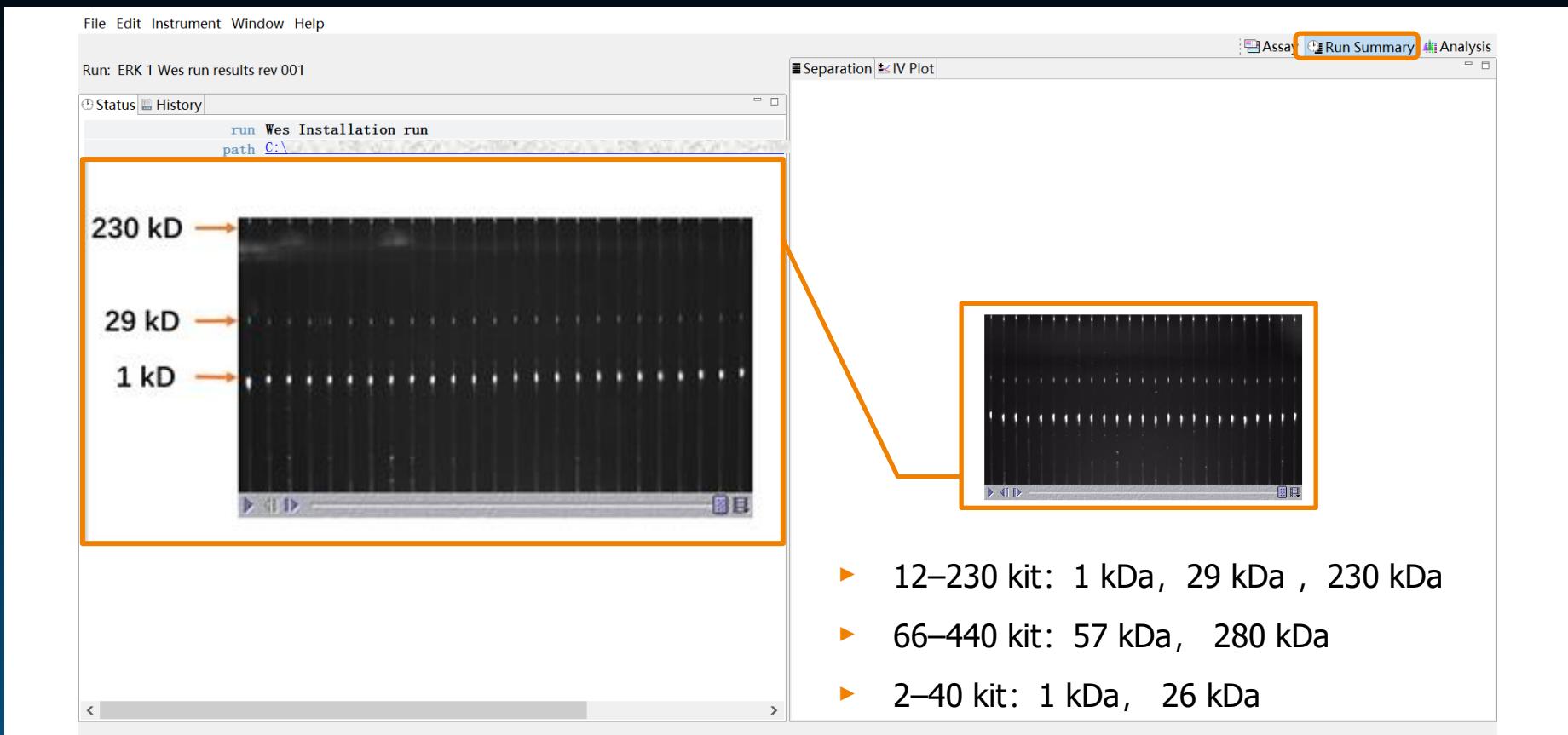
20 1. 荧光内参整齐



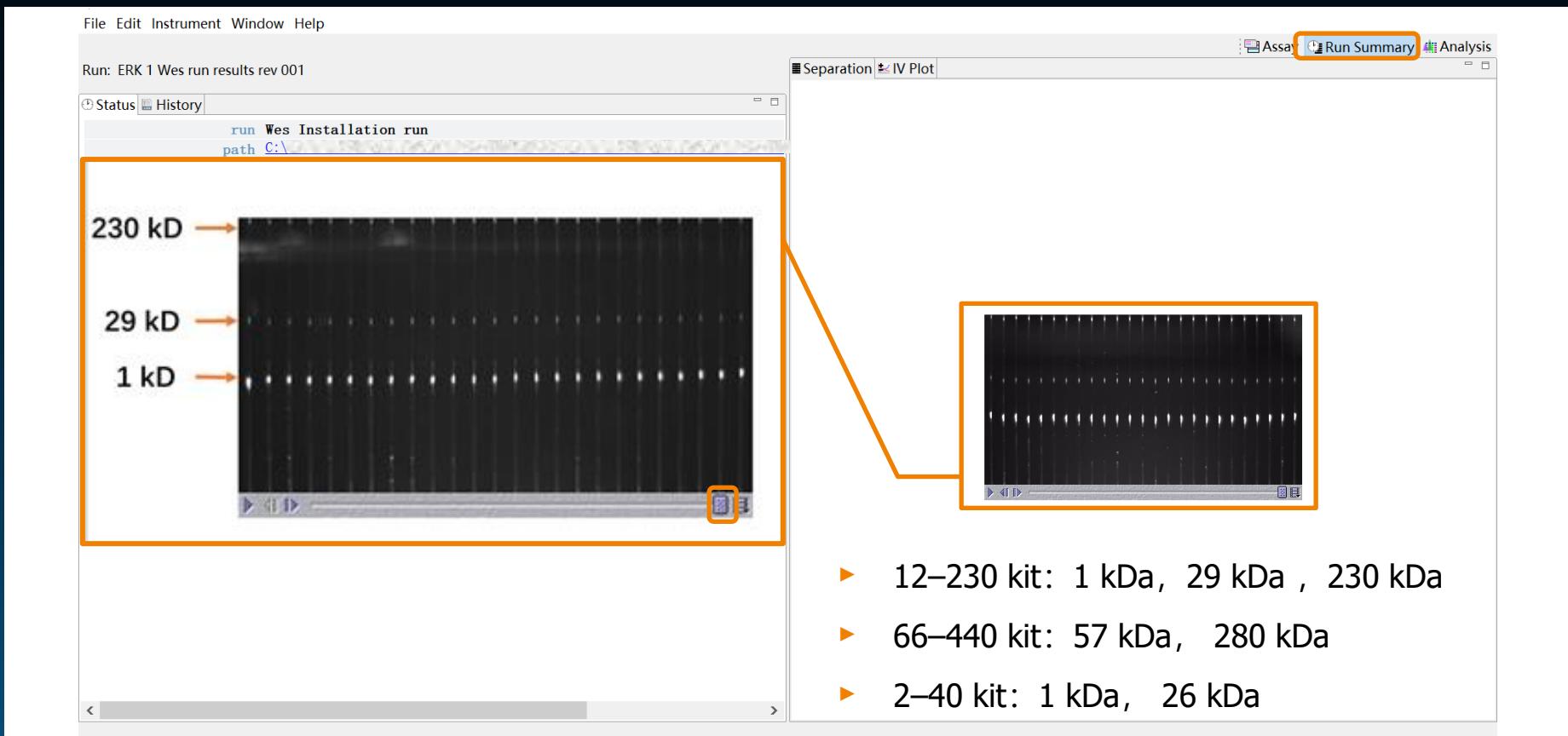
20 1. 荧光内参整齐



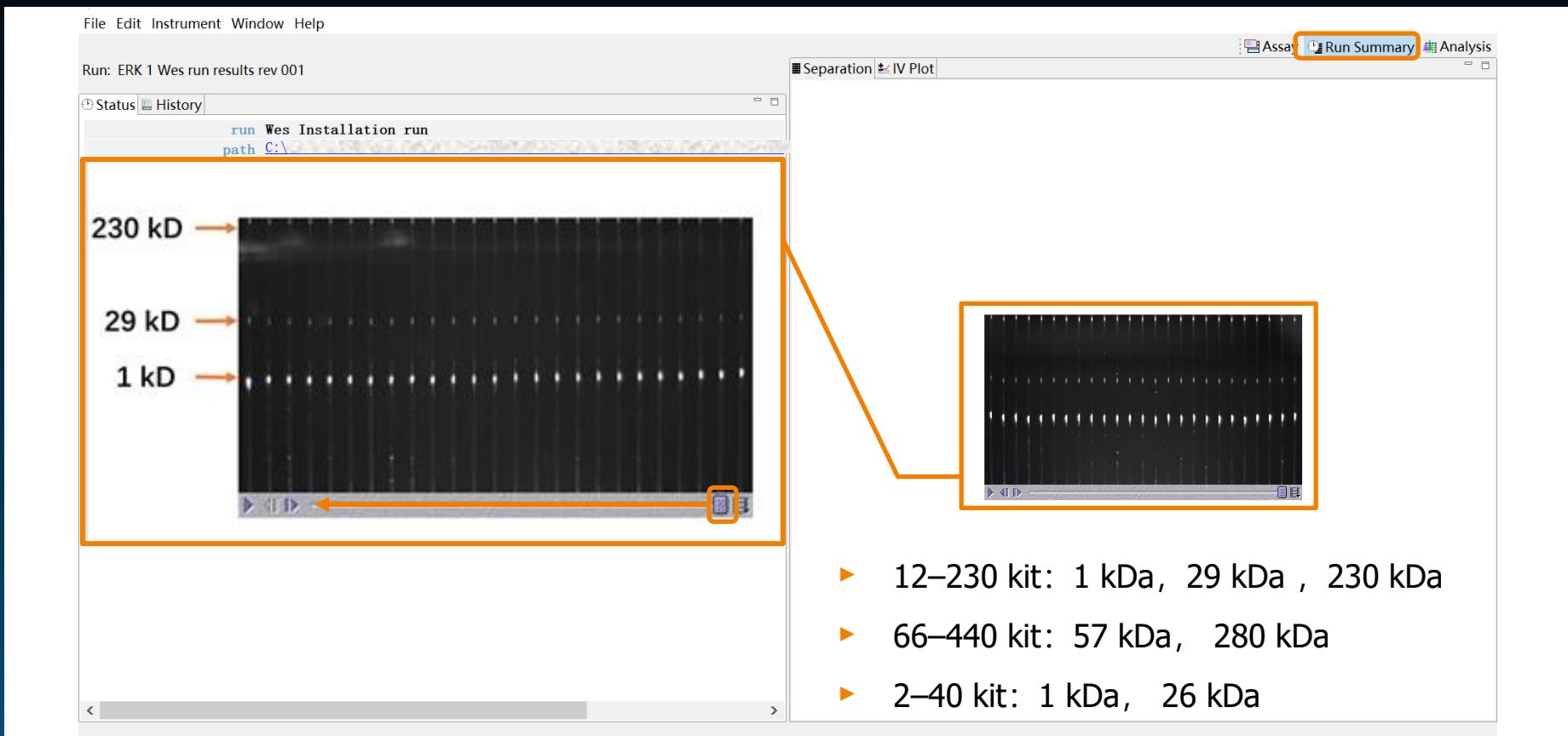
1. 荧光内参整齐



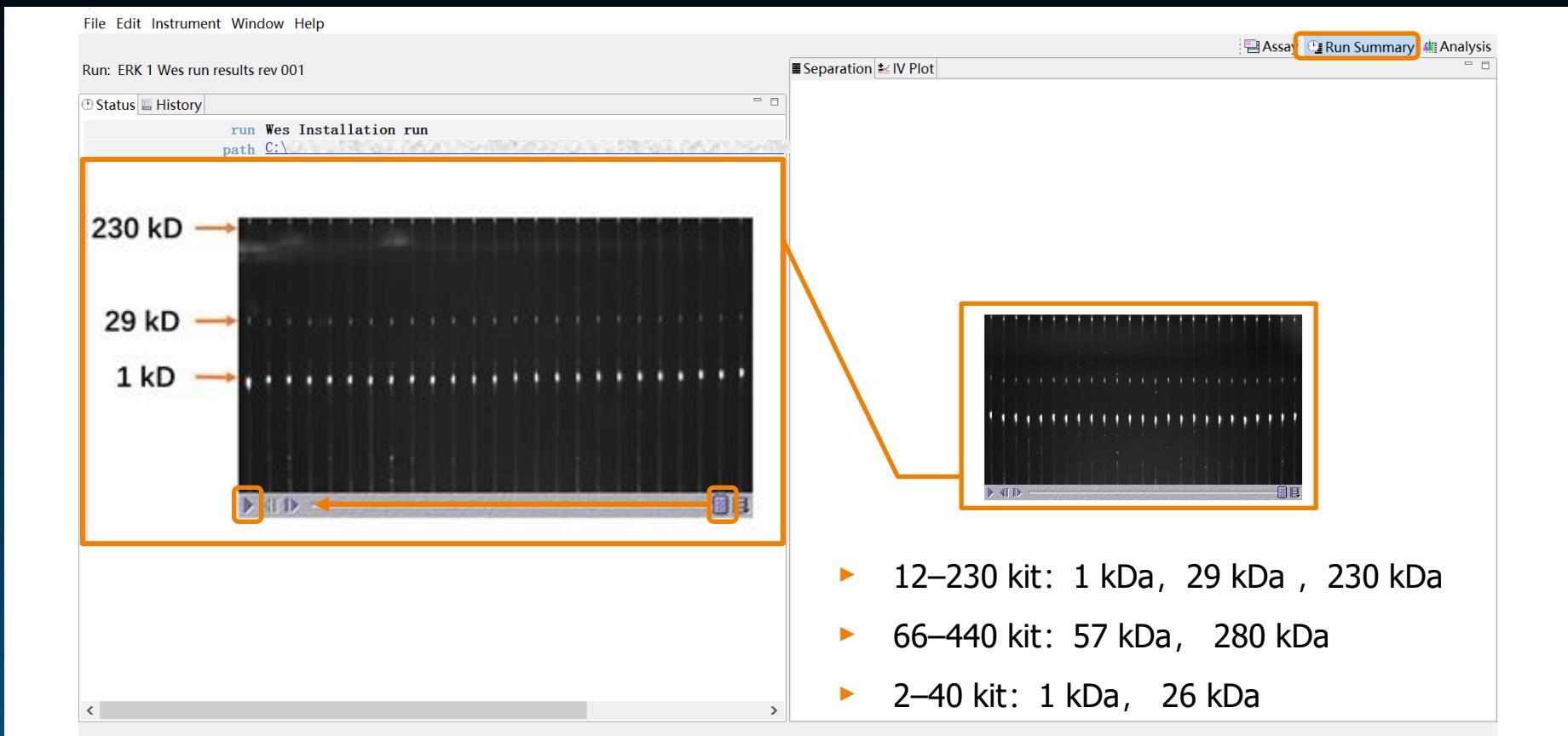
1. 荧光内参整齐



1. 荧光内参整齐



1. 荧光内参整齐



21 1. 荧光内参整齐

1. 荧光内参整齐



1. 荧光内参整齐

File Edit View Instrument Window Help

Standards Samples

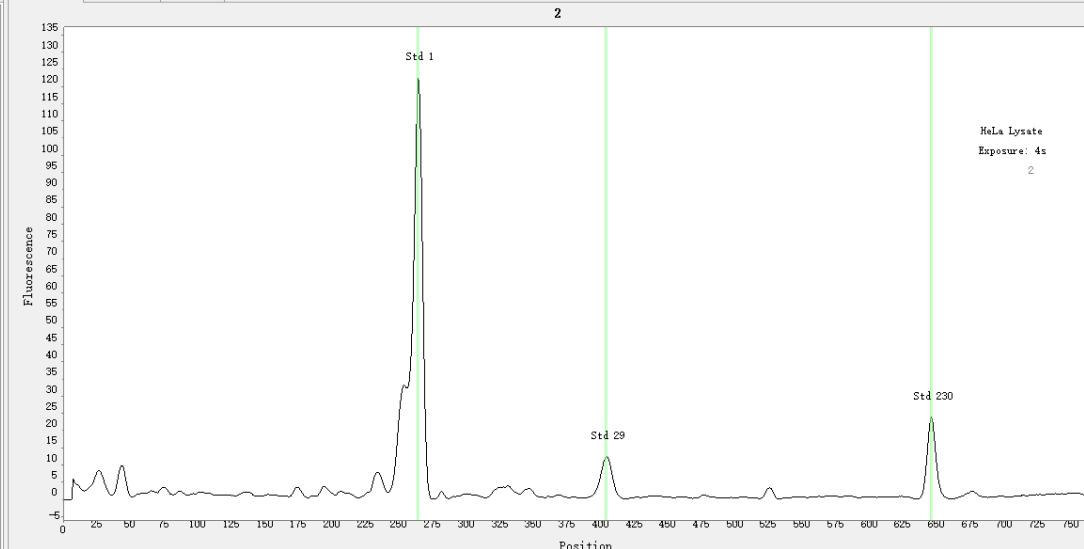
Experiment

Sample	Primary	C...
Biot. La...	Blocki...	1
HeLa L...	ERK1 ...	2
HeLa L...	ERK1 ...	3
HeLa L...	ERK1 ...	4
HeLa L...	ERK1 ...	5
HeLa L...	ERK1 ...	6
HeLa L...	ERK1 ...	7
HeLa L...	ERK1 ...	8
HeLa L...	ERK1 ...	9
HeLa L...	ERK1 ...	10
HeLa L...	ERK1 ...	11
HeLa L...	ERK1 ...	12
HeLa L...	ERK1 ...	13
HeLa L...	ERK1 ...	14
HeLa L...	ERK1 ...	15
HeLa L...	ERK1 ...	16
HeLa L...	ERK1 ...	17
HeLa L...	ERK1 ...	18
HeLa L...	ERK1 ...	19
HeLa L...	ERK1 ...	20
HeLa L...	ERK1 ...	21
HeLa L...	ERK1 ...	22
HeLa L...	ERK1 ...	23
HeLa L...	ERK1 ...	24
HeLa L...	ERK1 ...	25



Graph

Image Lane



Peaks Capillaries

Sample	Pri...	Cap	Pe...	Posit...	Heig...
HeLa L...	ERK...	2	1	262.6	122.6
HeLa L...	ERK...	2	2	403.9	12.4
HeLa L...	ERK...	2	3	646.0	24.0

Assay Run Summary

Analysis

Analysis Options Annotations

Images

Exposures Multi-Image Analysis

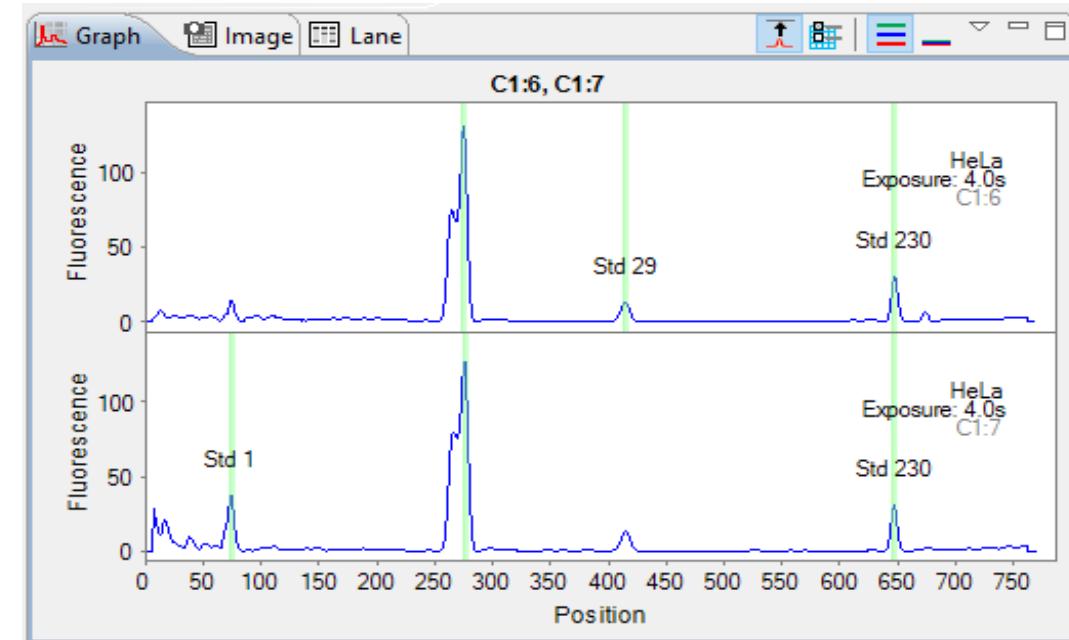
Peak Names

Name	<input type="text"/>
MW	<input type="text"/>
Color	<input type="text"/>
Caps	<input type="text"/>
<input type="button" value="Modify"/>	

1. 荧光内参整齐

Compass 软件一般能自动识别校正荧光内参，但是偶尔也会识别不正确

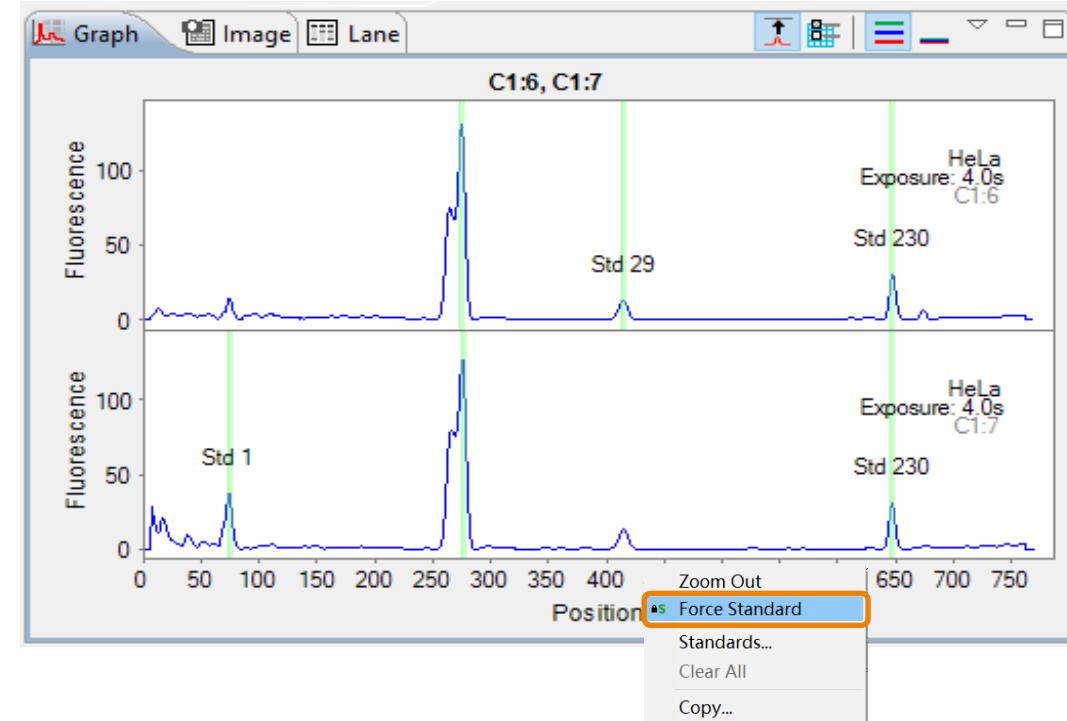
荧光内参识别不正确，将导致**分子量计算错误**



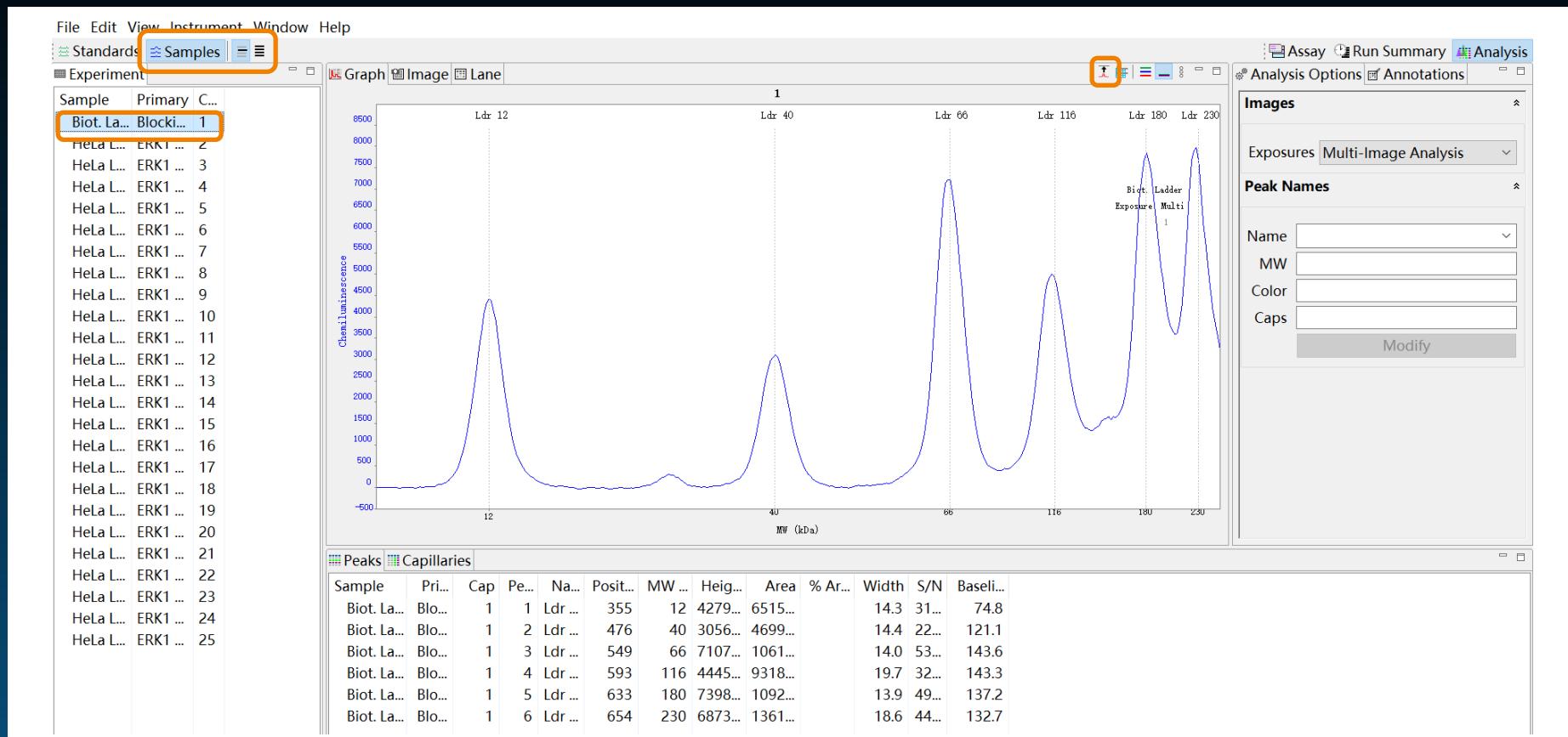
1. 荧光内参整齐

Compass 软件一般能自动识别校正荧光内参，但是偶尔也会识别不正确

荧光内参识别不正确，将导致**分子量计算错误**



2. LADDER 识别良好



2. LADDER 识别良好

1. 如果 ladder 峰识别错误, 点击右键进行校正

- 注意您的 ladder 通道要选择正确

2. 如果有一个 ladder 峰不可见, 可能调整下视野

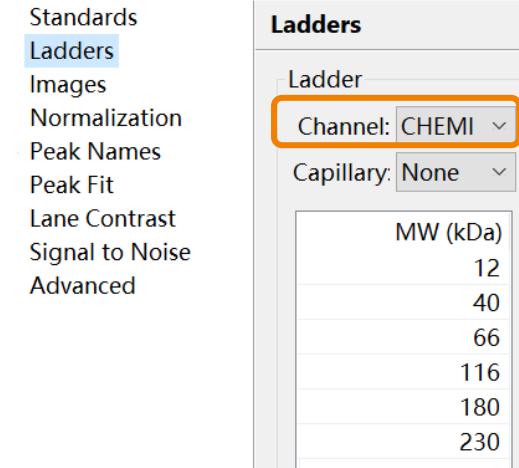
- 点击 View → View Region , 选择 *Full Range*, 可查看整个毛细管的视野数据

3. 如果不能通过识别来校正某个 *ladder* 峰

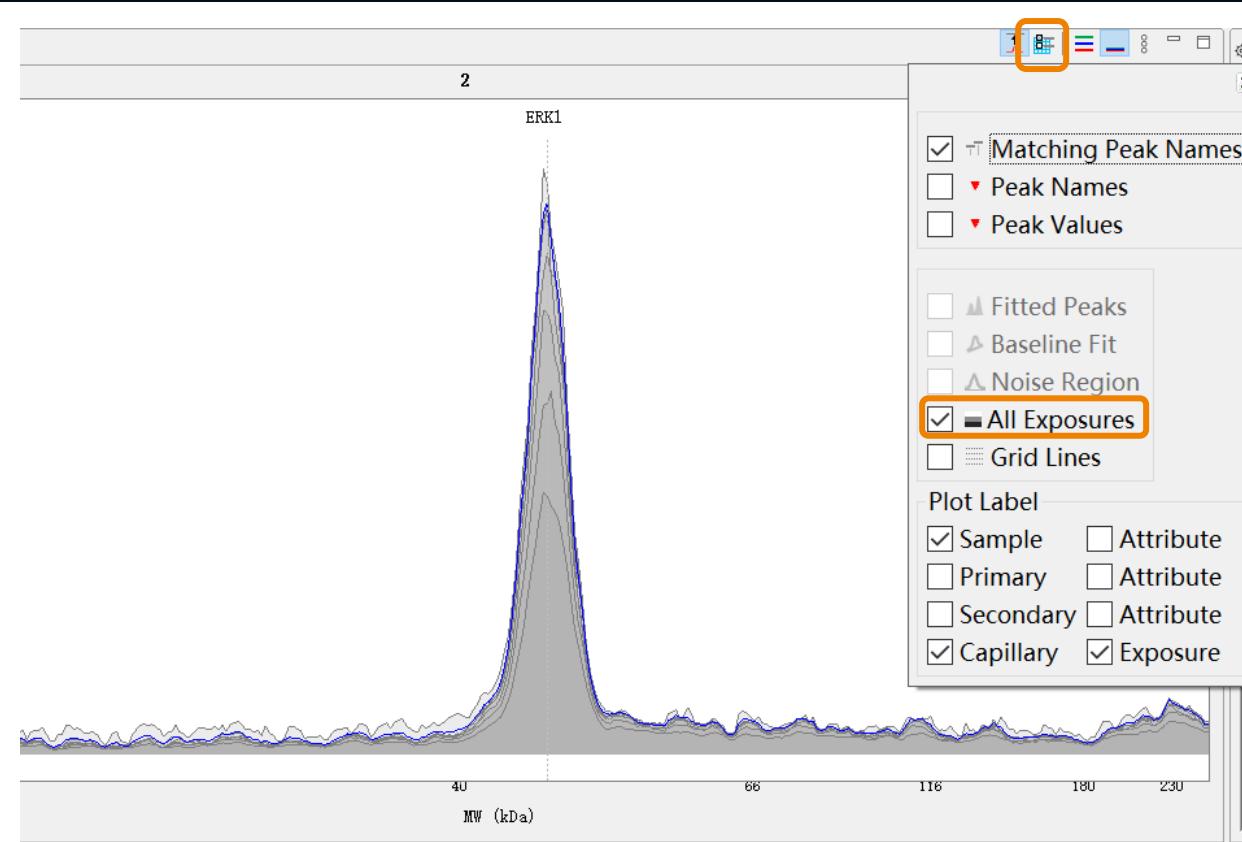
点击 Edit → analysis → ladders.

- 如有必要, 移除某个Ladder; 或选择 “none” 关闭 ladder 选项

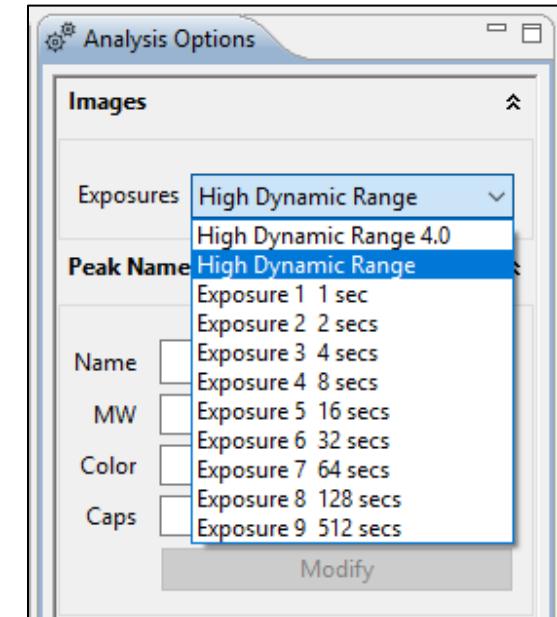
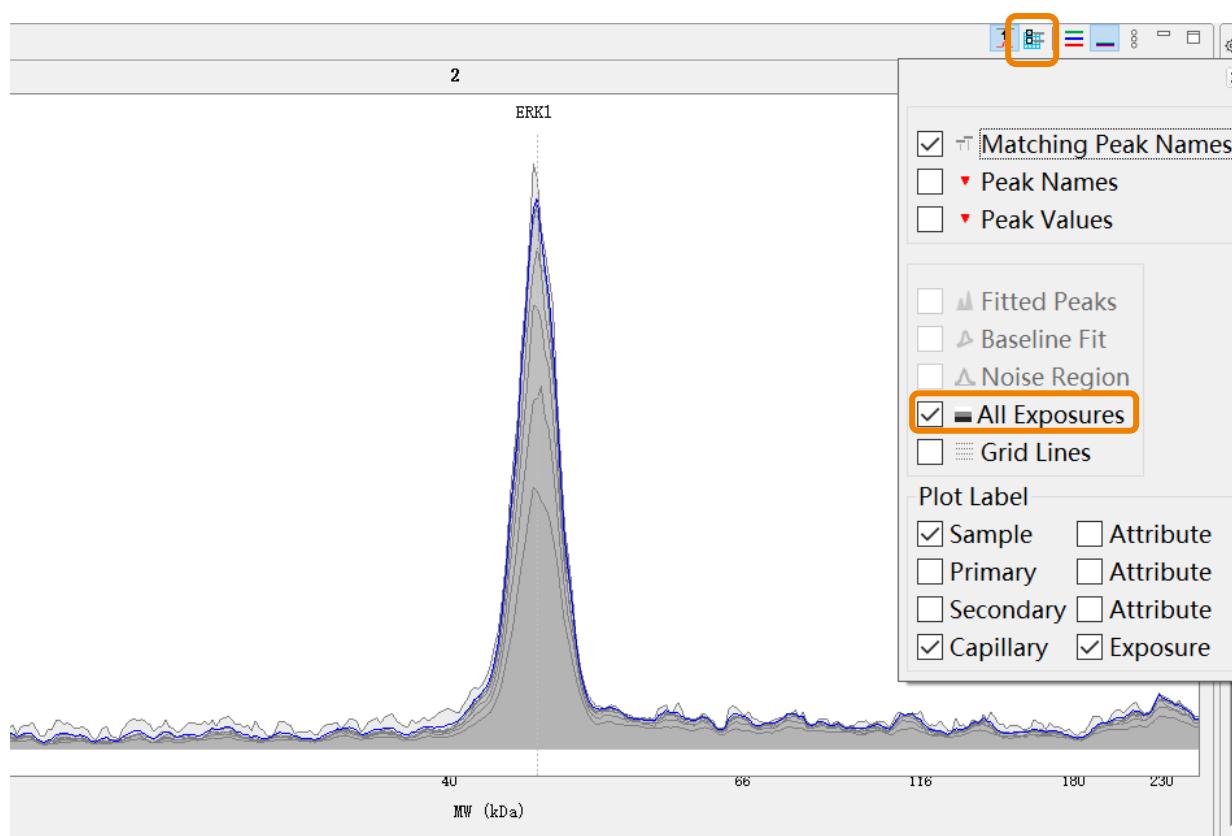
* 在没有 ladder 识别的情况下, Compass 软件依赖于每个毛细管内的荧光内参进行分子量计算



3. 化学发光信号无过曝

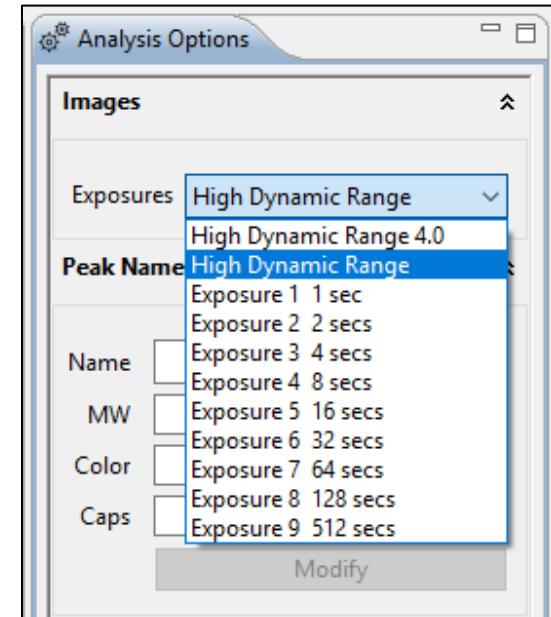
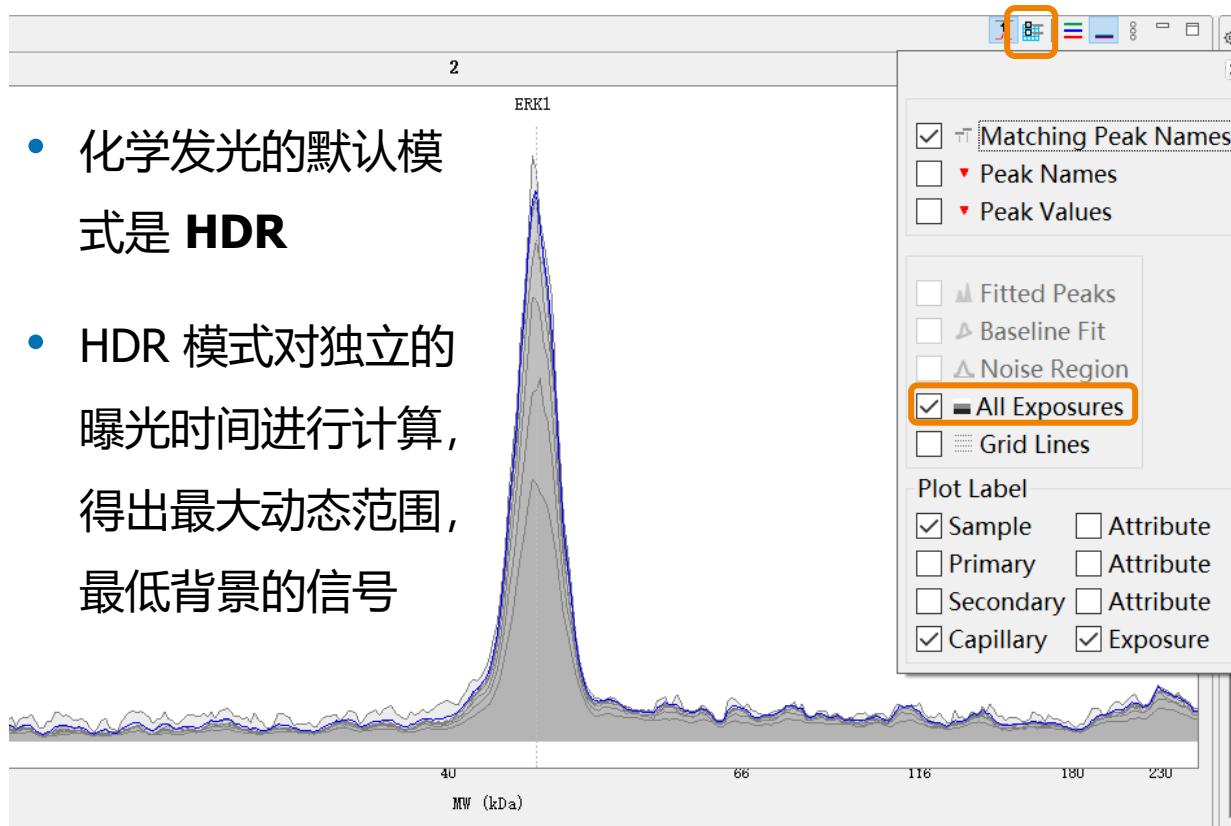


3. 化学发光信号无过曝



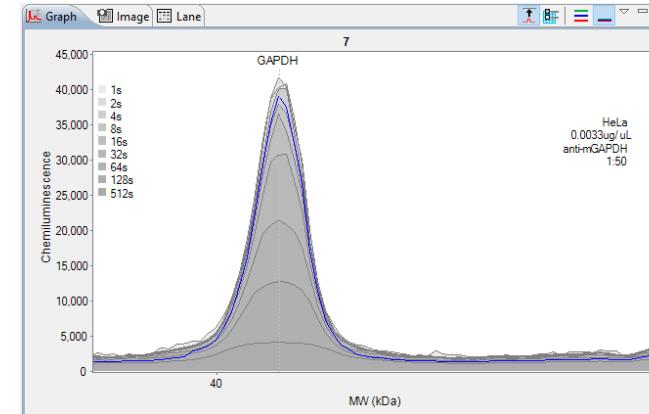
3. 化学发光信号无过曝

- 化学发光的默认模式是 **HDR**
- HDR 模式对独立的曝光时间进行计算，得出最大动态范围，最低背景的信号

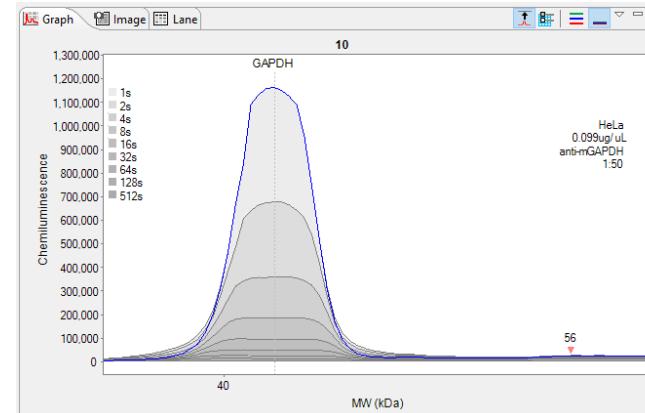


3. 化学发光信号无过曝

- 化学发光信号实际上是信号/时间的比值.
 - 在长曝光时间中，信号衰减，所以峰高下降
- 如果您看到在前几个曝光时间中，某个峰的信号明显下降，说明：
 - 化学发光底物消耗完
 - 该信号超出了动态范围
- 保持化学发光的峰高小于 **300,000**



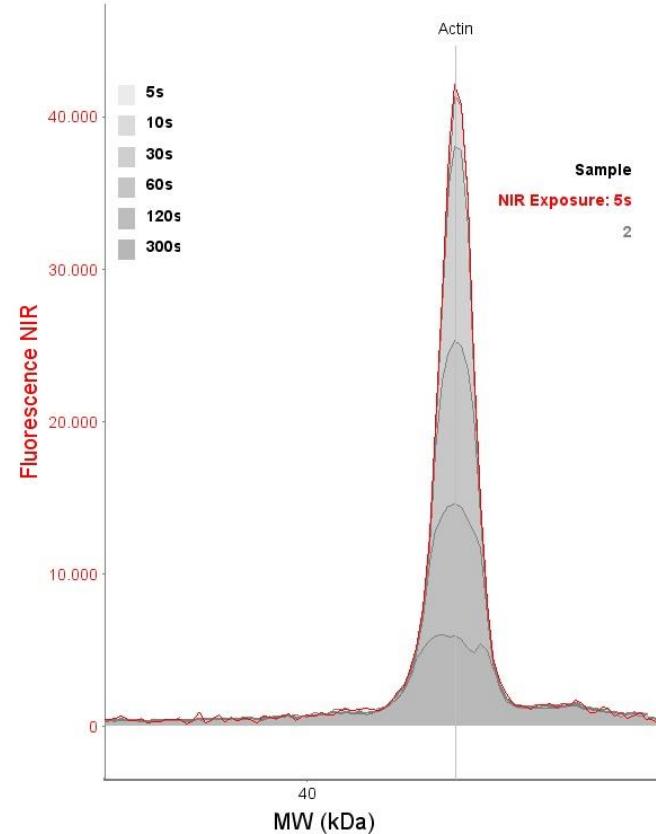
正常信号



过曝信号

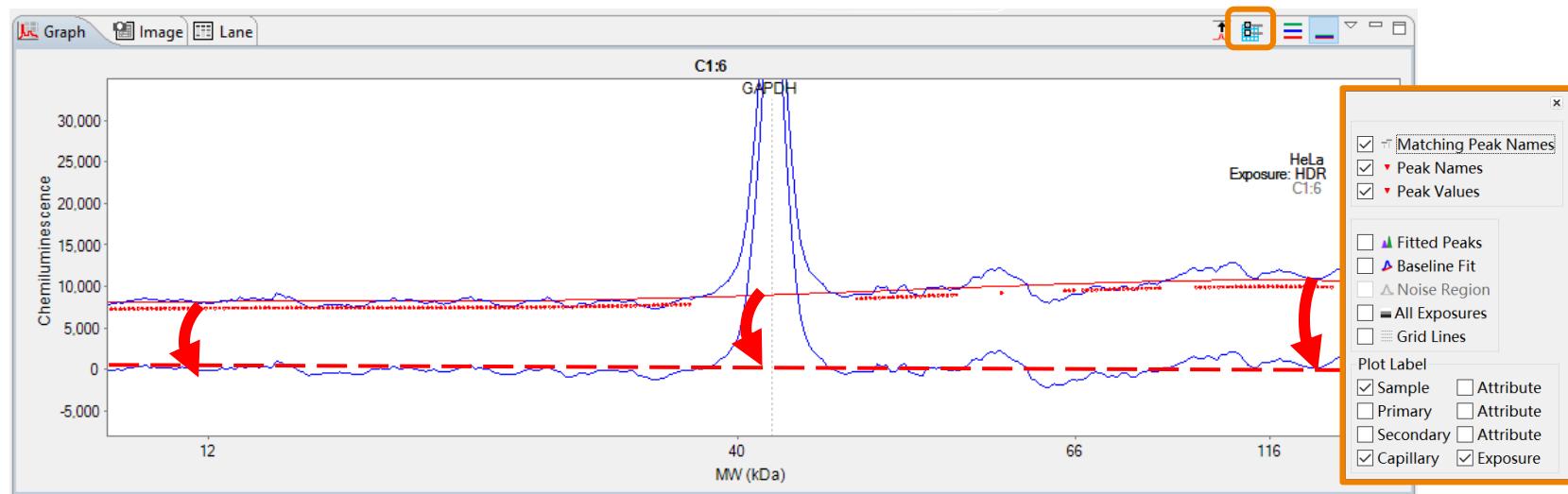
4. 荧光通道相机无饱和 (JESS)

- 在荧光通道中，信号太强有可能导致相机像素饱和
- 一旦观察到信号开始下降，说明相机饱和
- 分析数据时，选取信号稳定的曝光时间
- 因此，最小的两个曝光时间峰应该重叠，否则说明相机较快饱和
- 保证荧光检测的峰高低于 **60,000**



29. 5. 背景无过高——基线拟合

- 软件默认显示扣除背景后的信号曲线 (背景设为0)
- 要观察基线, 需要勾选 “Baseline fit”; 基线参数设置在 Edit→Analysis...中
- 软件根据基线下面的点拟合出基线, 可添加或删除基线点



5. 背景无过高——基线拟合

- 为什么高背景是个问题?
 - 信号饱和风险
 - 显著减弱动态范围
 - 降低检测灵敏度

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
Biot. Ladder	Blocking	1	6	Ldr 230	654	230	6921.9	137776.2		18.7	2454.0	41.8
HeLa Lysate	ERK1 RTU	2	1	ERK1	486	48	5940.1	80205.3	100.0	12.7	412.8	183.9

5. 背景无过高——基线拟合

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 - 降低检测灵敏度

基线拟合

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
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Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
Biot. Ladder	Blocking	1	6	Ldr 230	654	230	6921.9	137776.2		18.7	2454.0	41.8
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样品

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
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Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
Biot. Ladder	Blocking	1	6	Ldr 230	654	230	6921.9	137776.2		18.7	2454.0	41.8
HeLa Lysate	ERK1 RTU	2	1	ERK1	486	48	5940.1	80205.3	100.0	12.7	412.8	183.9

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 - 降低检测灵敏度

一抗

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
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Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
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毛细管号

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
Biot. Ladder	Blocking	1	6	Ldr 230	654	230	6921.9	137776.2		18.7	2454.0	41.8
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峰序号

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
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Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
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峰名

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
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Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
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迁移位置

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
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分子量

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
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Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
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 - 降低检测灵敏度

峰高

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
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峰面积

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
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相对峰面积

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
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峰宽

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
Biot. Ladder	Blocking	1	6	Ldr 230	654	230	6921.9	137776.2		18.7	2454.0	41.8
HeLa Lysate	ERK1 RTU	2	1	ERK1	486	48	5940.1	80205.3	100.0	12.7	412.8	183.9

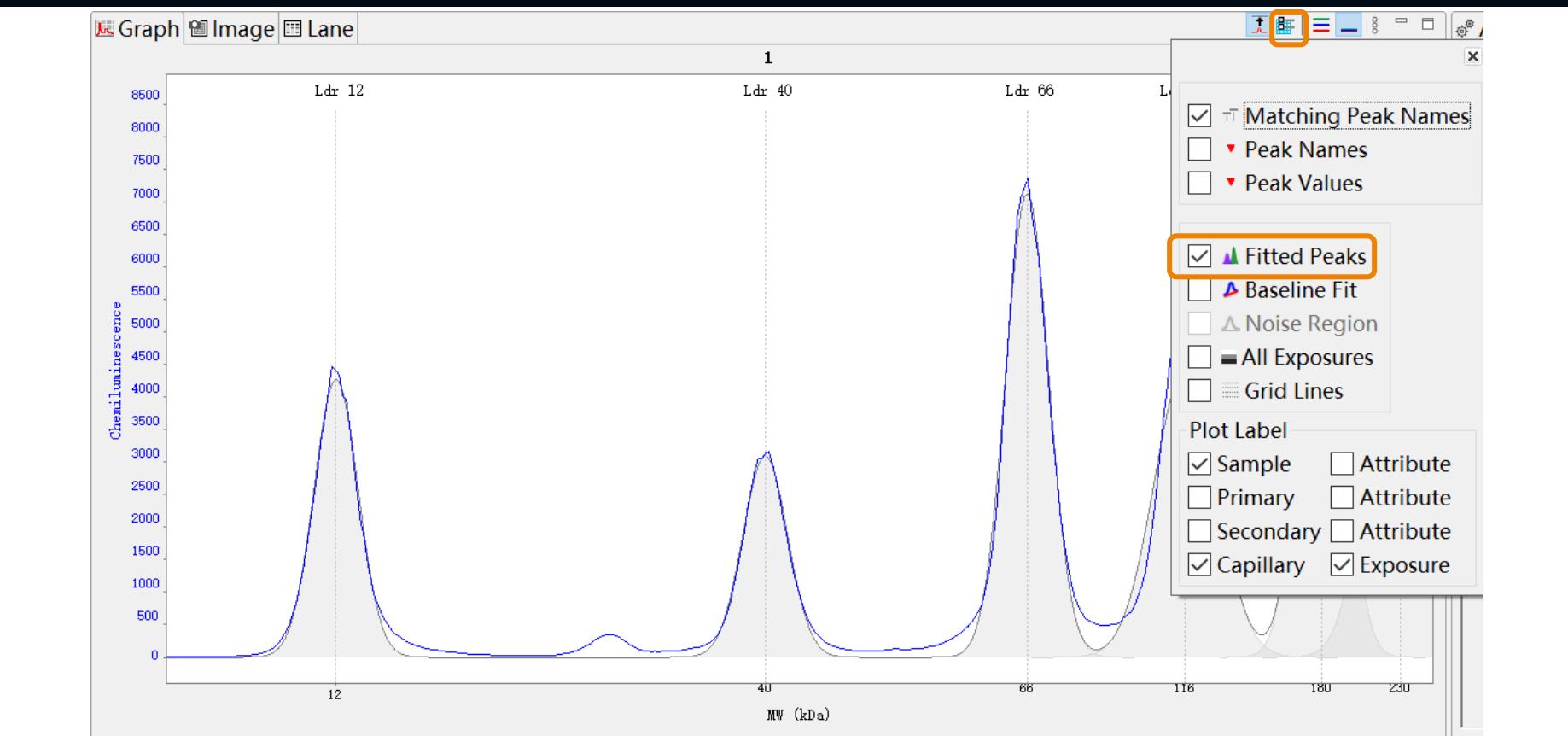
5. 背景无过高——基线拟合

- 为什么高背景是个问题?
 - 信号饱和风险
 - 显著减弱动态范围
 - 降低检测灵敏度

信噪比

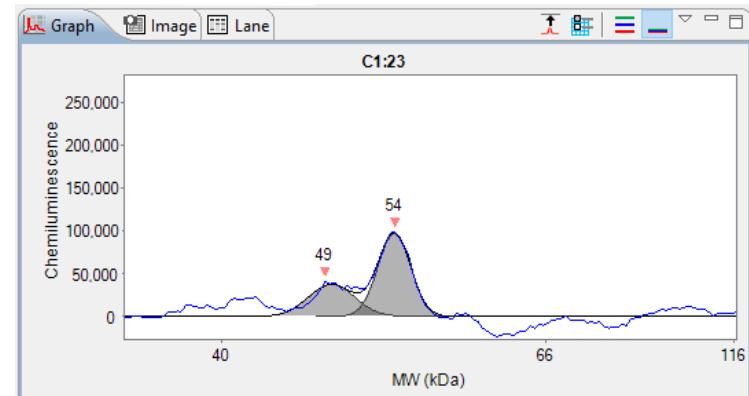
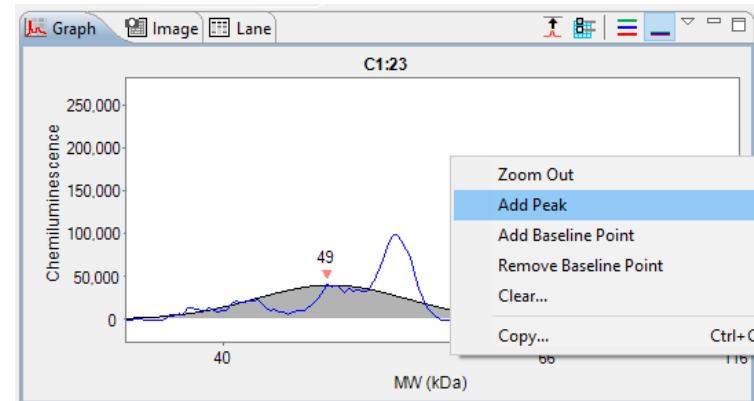
Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline
Biot. Ladder	Blocking	1	1	Ldr 12	355	12	4279.4	66444.9		14.6	1667.0	22.5
Biot. Ladder	Blocking	1	2	Ldr 40	476	40	3095.9	49008.2		14.9	1180.9	35.6
Biot. Ladder	Blocking	1	3	Ldr 66	549	66	7139.7	109106.0		14.4	2831.7	40.0
Biot. Ladder	Blocking	1	4	Ldr 116	593	116	4454.0	97617.1		20.6	1743.1	41.4
Biot. Ladder	Blocking	1	5	Ldr 180	633	180	7471.2	111393.5		14.0	2652.4	41.8
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HeLa Lysate	ERK1 RTU	2	1	ERK1	486	48	5940.1	80205.3	100.0	12.7	412.8	183.9

5. 峰面积正常



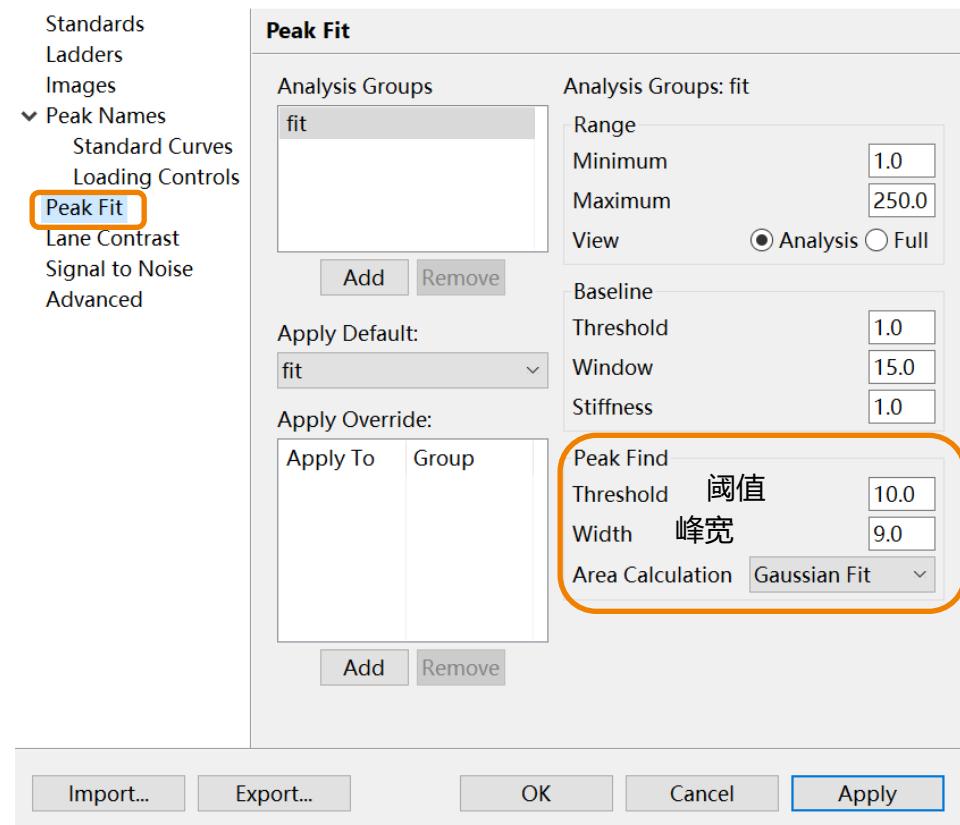
5. 峰面积正常

- 灰色部分的面积即峰面积.
- 需确认拟合峰跟实际峰形拟合良好
- 右键单击可通过“add peak” 或 “remove peak” 增加或去除峰
- 在 analysis 菜单中, 可选择拟合方式, 包括 gaussian fit (高斯拟合) vs dropped lines (垂直拟合, 推荐总蛋白定量时使用)



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数据分析前检查

- 确认完成了以下检查
 - ✓ 荧光内参整齐并识别良好
 - ✓ Ladder 识别良好
 - ✓ 化学发光信号无过曝
 - ✓ 荧光通道相机无饱和 (Jess)
 - ✓ 背景无过高
 - ✓ 峰面积正常

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可以开始分析数据了

01

Compass 介绍

02

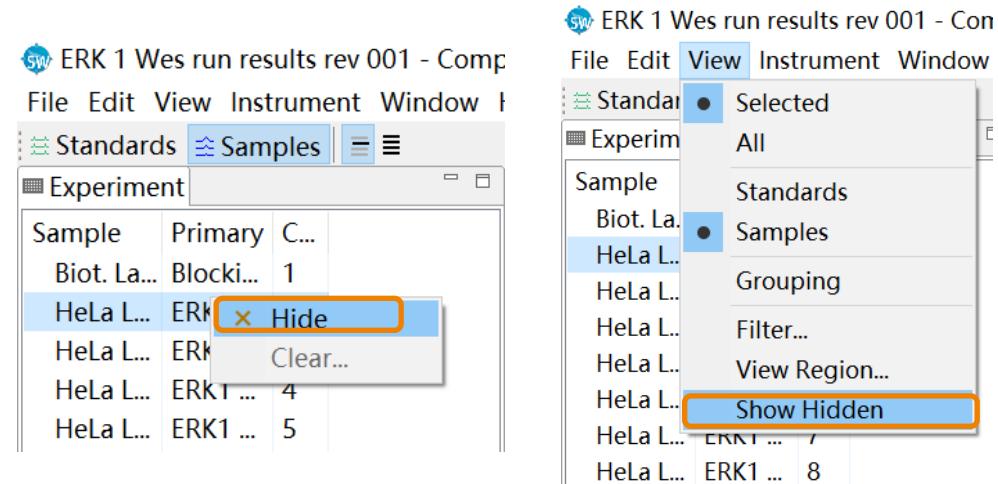
数据分析前检查

03

数据分析

编辑实验信息

- 样品及抗体信息等在 Assay 界面输入
 - Assay template 里输入的信息可以随时更改（运行前/后）
- 右键点击可隐藏泳道
- 如果要显示隐藏的泳道，点击：
 - *View→Show Hidden*
 - 右键点击 unhide 恢复



选择检测通道

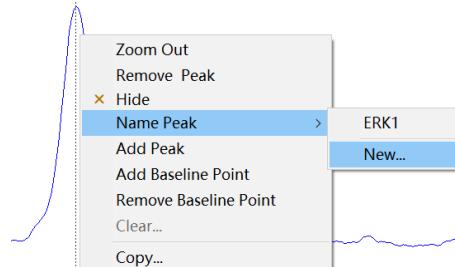
- 毛细管/泳道可选择性浏览或者全局浏览
- 检测通道可进行开/关
- 选择不同的通道，将有不止一个Y轴



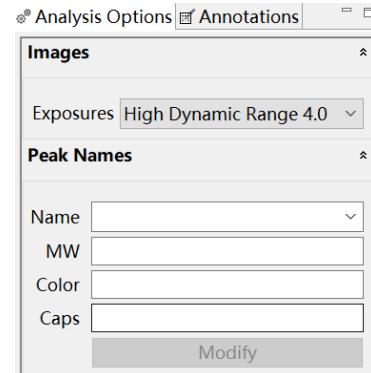
峰命名

当您查看数据时，标记您的峰值很有帮助

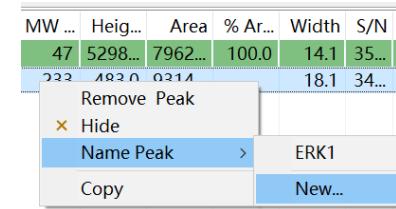
- 选项 1: 右键点击峰形图中的峰, 选择 Name Peak → New... 进行命名



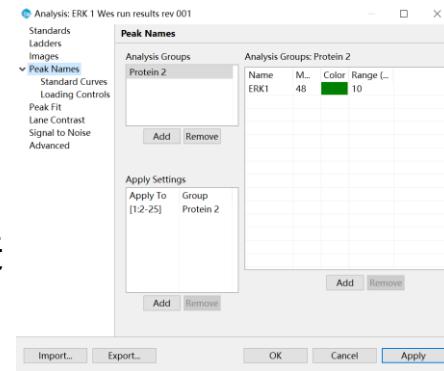
- 选项 2: Analysis Options 窗格里
 - 在 Name 中: 下拉选中 [New]
 - 输入峰名称等
 - 单击 Create



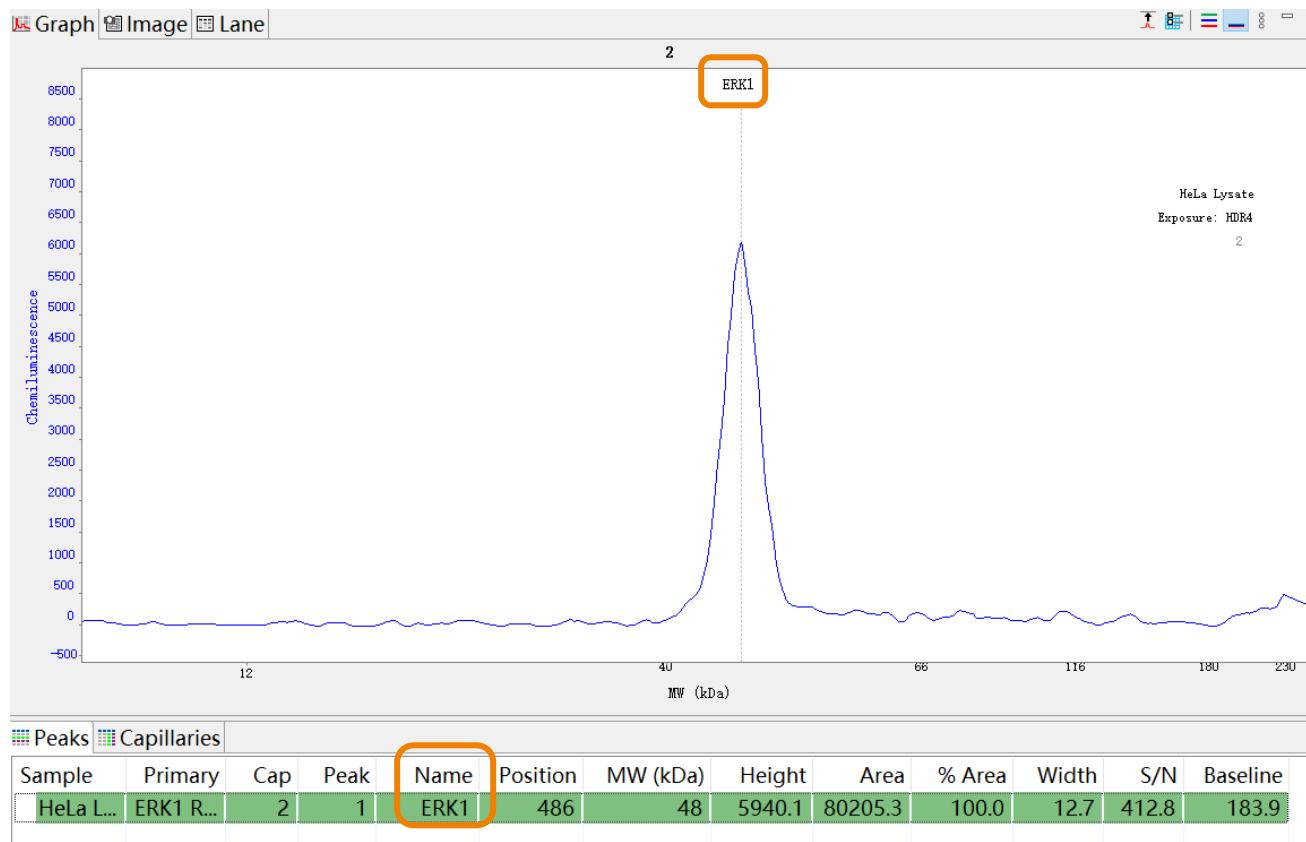
- 选项 3: 右键点击峰统计表中的峰, 选择 Name Peak → New... 进行命名



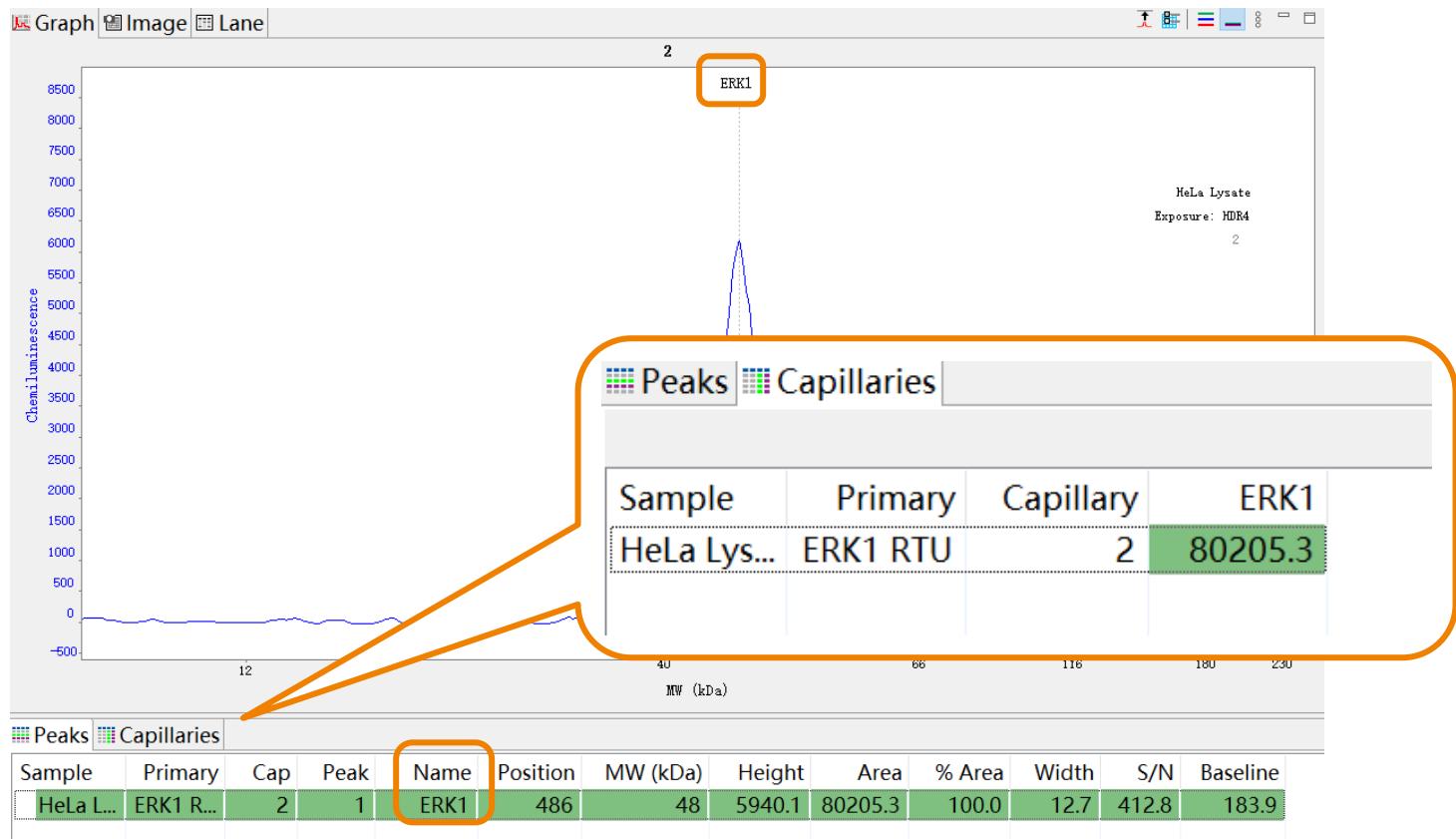
- 选项 4: 在 Edit>Analysis 菜单栏中
 - 在 Peak Names 栏中进行峰命名



峰命名



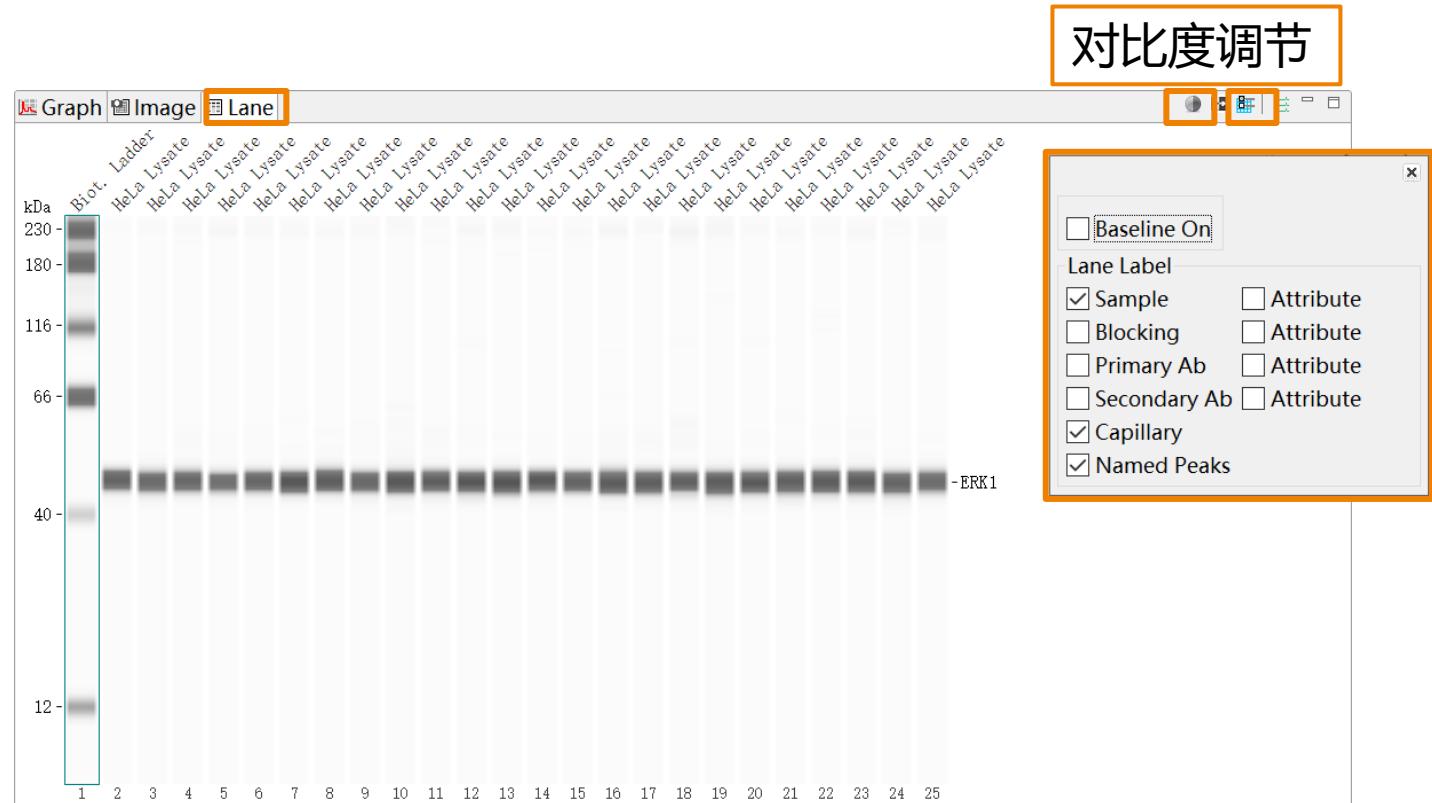
峰命名



ANALYSIS 界面

泳道图

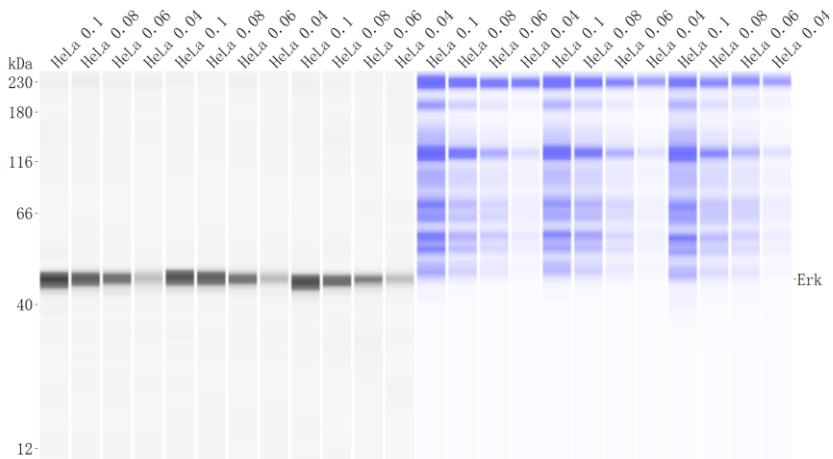
- 大部分报告的结果形式
- 可以任意显示或横向移动泳道
- 可以单独调整每个通道的对比度
 - 对比度调整仅改变图片中条带呈现效果, 峰定量结果不变



导出数据

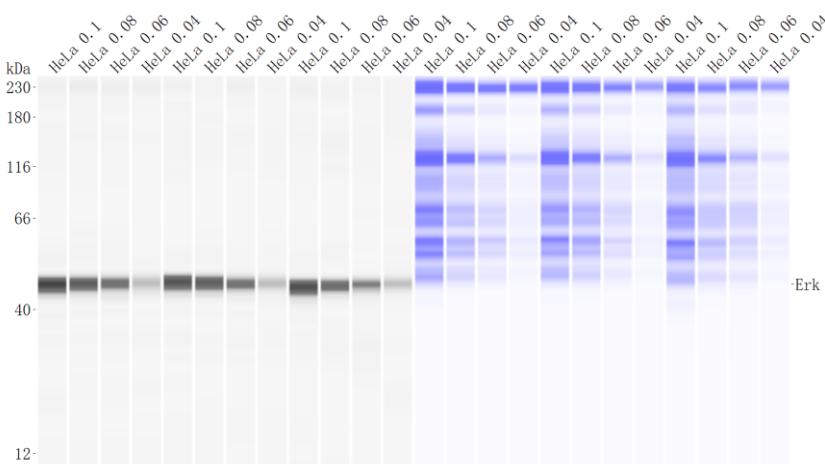
- 图像拷贝
 - 在峰形图或者泳道图中，右键点击Copy 复制或保存图像
- 选中峰统计表中的行（一行或多行），右键点击 Copy 复制数据
- 点击 File→export tables
 - 软件将建立一个文件夹导出各类统计数据
- 点击 File→run report
 - 软件将建立一个 PDF 文档，详细报告图和表

能否直接注释结果？

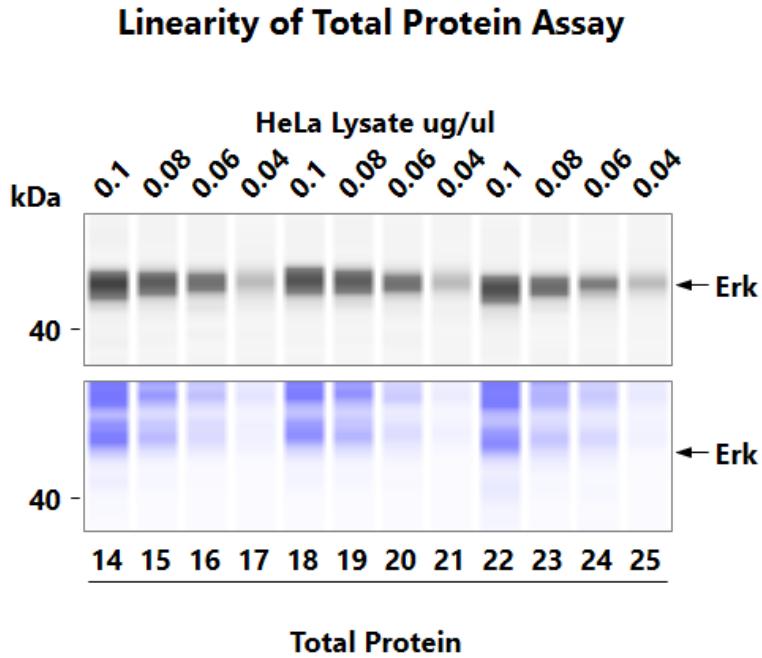


1;141;151;161;171;181;191;201;211;221;231;241;252;142;152;162;172;182;192;202;212;222;232;242;25

能否直接注释结果？

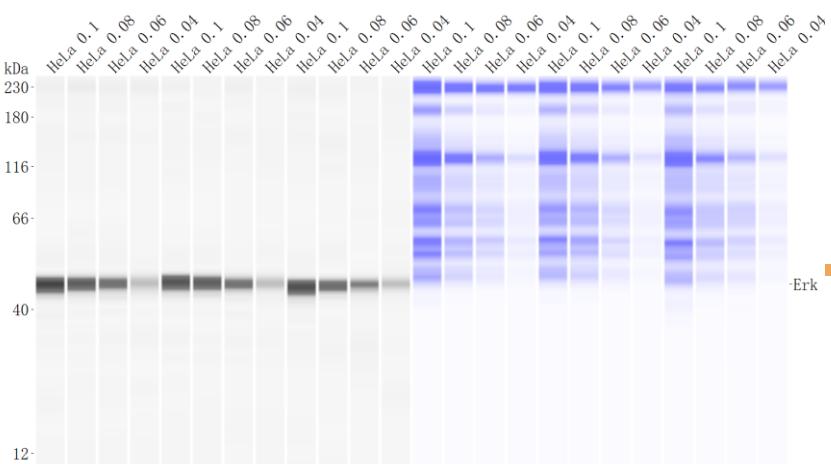


1;141;151;161;171;181;191;201;211;221;231;241;252;142;152;162;172;182;192;202;212;222;232;242;25

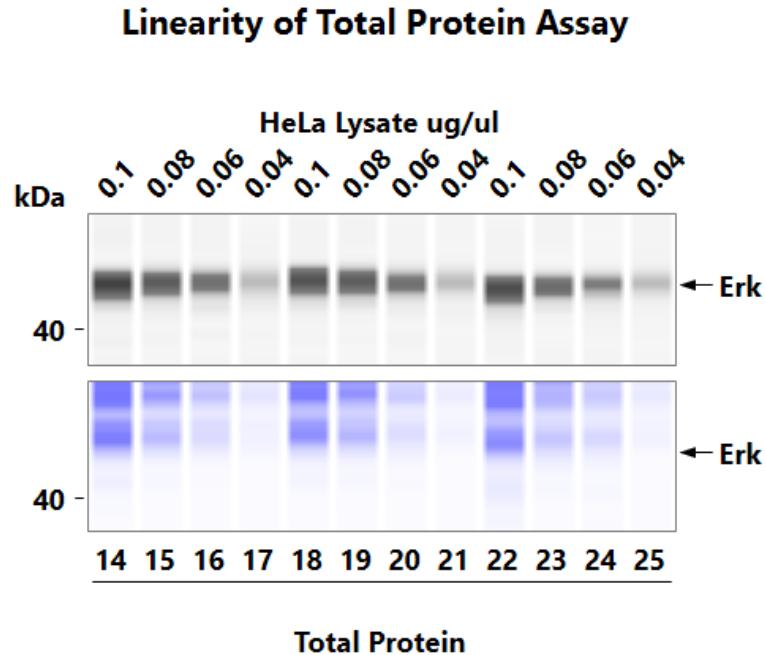


HeLa/Erk with RePlex and TP Lane_annotation_training_file.cbz

能否直接注释结果？

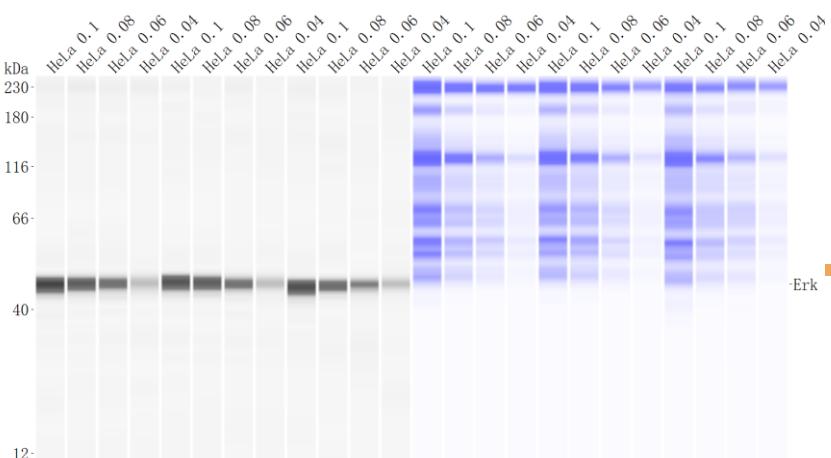


1;141;151;161;171;181;191;201;211;221;231;241;252;142;152;162;172;182;192;202;212;222;232;242;25



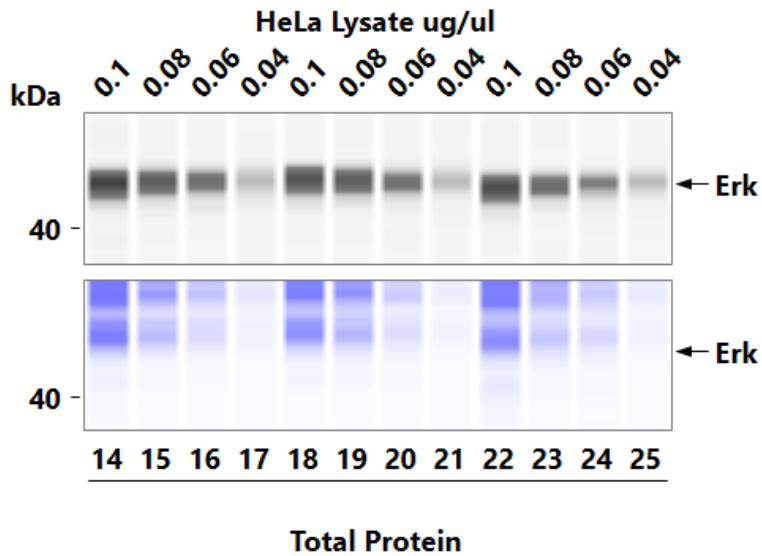
HeLa/Erk with RePlex and TP Lane_annotation_training_file.cbz

能否直接注释结果?



1:141:151:161:171:181:191:201:211:221:231:241:252:142:152:162:172:182:192:202:212:222:232:242:25

Linearity of Total Protein Assay



HeLa/Erk with RePlex and TP
Lane_annotation_training_file.cbz

公众号报修流程



09:40 ProteinSimple 领域的应用技术研讨会

活动回顾 | 蛋白质分析技术在转化医学领域的应用技术研讨会 · 成都站

2021年6月25日, Bio-Techne集团 (NASDAQ:TECH)旗下ProteinSimple与转化医学网携手,在成都锦江宾馆成功举办了蛋白质分析技术在转化医学领域的应用技术研讨会, 50多位专家和学者, 共聚蓉城参与此次盛会!

2021年6月29日 17:00 2021太仓国际生物医药创新和产业化峰会暨2021中国蛋白药物质量与技术创新研讨会

2021年7月2日-4日 太仓市宝龙禧来登酒店

展位号 23

邀请函 | ProteinSimple 邀您共赴 2021 太仓国际生物医药创新和产业化峰会

2021年7月2日-4日, ProteinSimple与您相约, 来登酒店, 共赴 2021 太仓国际生物医药创新和产业化峰会。23号展位, 不见不散!

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protein simple

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*产品型号 请选择
*产品序列号 请手工输入

咨询人联系方式

*姓名 王婉婷
*电话 18516606926
*电子邮件 xianting.wang@bio-techne.com
*单位名称 ProteinSimple
*省份 请选择
*城市 请选择
区级 请选择
*详细地址 街道、楼牌号等

技术问题描述

可添加至多3张咨询图片

技术问题细节描述

请描述技术问题细节...

添加图片

提交

09:41

电子邮件 xianting.wang@bio-techne.com
单位名称 ProteinSimple
省份 请选择
城市 请选择
区级 请选择
详细地址 街道、楼牌号等

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- **Simple ELISA**



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- **Simple Sc-Western**



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- **Simple Particle Analysis**



Maurice

- **Simple iCIEF + CE-SDS**