

# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### Default Table and Plot Settings

#### TABLE SETTINGS – ANALYSIS VIEW: SAMPLES SETTINGS

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The screenshot shows the 'Table Setting Editor' dialog box with the 'Samples' tab selected. The 'Column Settings' table is as follows:

	Show	Column	Filtering	Content
1	<input checked="" type="checkbox"/>	Status	Show All Records	N/A
2	<input type="checkbox"/>	Sample File	Show All Records	
3	<input checked="" type="checkbox"/>	Sample Name	Show All Records	
4	<input type="checkbox"/>	Sample ID	Show All Records	
5	<input type="checkbox"/>	Comments	Show All Records	
6	<input checked="" type="checkbox"/>	Sample Type	Show All Records	N/A
7	<input type="checkbox"/>	Specimen Category	Show All Records	N/A
8	<input checked="" type="checkbox"/>	Analysis Method	Show All Records	
9	<input checked="" type="checkbox"/>	Panel	Show All Records	
10	<input checked="" type="checkbox"/>	Size Standard	Show All Records	
11	<input type="checkbox"/>	Matrix	Show All Records	
12	<input checked="" type="checkbox"/>	Run Name	Show All Records	
13	<input type="checkbox"/>	Instrument Type	Show All Records	
14	<input type="checkbox"/>	Instrument ID	Show All Records	
15	<input type="checkbox"/>	Run Date & Time	Show All Records	
16	<input type="checkbox"/>	Reference Data	Show All Records	N/A
17	<input checked="" type="checkbox"/>	Sizing Quality Overridden	Show All Records	N/A
18	<input checked="" type="checkbox"/>	Sample File Not Found	Show All Records	N/A
19	<input type="checkbox"/>	Matrix Not Found	Show All Records	N/A
20	<input checked="" type="checkbox"/>	Size Standard Not Found	Show All Records	N/A
21	<input checked="" type="checkbox"/>	Off-scale	Show All Records	N/A
22	<input checked="" type="checkbox"/>	Sizing Quality	Show All Records	N/A
23	<input checked="" type="checkbox"/>	User Defined Column 1	Show All Records	
24	<input checked="" type="checkbox"/>	User Defined Column 2	Show All Records	
25	<input checked="" type="checkbox"/>	User Defined Column 3	Show All Records	

Font Settings:  
Font: Arial  
Size: 11

Buttons: Show, Hide, OK, Cancel

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### TABLE SETTINGS – ANALYSIS VIEW: GENOTYPES SETTINGS

Table Setting Editor

General Samples **Genotypes**

Genotypes Table Settings:

Column Settings:

	Show	Column	Filtering	Content
1	<input type="checkbox"/>	Sample File	Show All Records	
2	<input checked="" type="checkbox"/>	Sample Name	Show All Records	
3	<input type="checkbox"/>	Sample ID	Show All Records	
4	<input type="checkbox"/>	Run Name	Show All Records	
5	<input type="checkbox"/>	Panel	Show All Records	
6	<input checked="" type="checkbox"/>	Marker	Show All Records	
7	<input checked="" type="checkbox"/>	Dye	Show All Records	N/A
8	<input checked="" type="checkbox"/>	Allele	Show All Records	
9	<input type="checkbox"/>	Size	Show All Records	
10	<input type="checkbox"/>	Height	Show All Records	
11	<input type="checkbox"/>	Peak Area	Show All Records	
12	<input type="checkbox"/>	Data Point	Show All Records	
13	<input type="checkbox"/>	Mutation	Show All Records	
14	<input type="checkbox"/>	AE Comment	Show All Records	
15	<input type="checkbox"/>	Integration Comments	Show All Records	
16	<input checked="" type="checkbox"/>	Allele Display Overflow	Show All Records	N/A
17	<input checked="" type="checkbox"/>	Allele Edit	Show All Records	N/A
18	<input type="checkbox"/>	Omit From Clustering (SNPle:	Show All Records	N/A
19	<input checked="" type="checkbox"/>	Off-scale	Show All Records	N/A

Show Hide

Font Settings:

Font: Arial

Size: 11

Allele Settings

Number of Alleles   Keep Allele, Size, Height, Area, Data Point, Mutation and Comment together

OK Cancel

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### TABLE SETTINGS – CASEWORK VIEW: SAMPLES SETTINGS

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Table Setting Editor

General Samples Genotypes

Samples Table Settings:

Column Settings:

	Show	Column	Filtering	Content
1	<input checked="" type="checkbox"/>	Status	Show All Records	N/A
2	<input checked="" type="checkbox"/>	Sample File	Show All Records	
3	<input checked="" type="checkbox"/>	Sample Name	Show All Records	
4	<input type="checkbox"/>	Sample ID	Show All Records	
5	<input type="checkbox"/>	Comments	Show All Records	
6	<input checked="" type="checkbox"/>	Sample Type	Show All Records	N/A
7	<input checked="" type="checkbox"/>	Specimen Category	Show All Records	N/A
8	<input checked="" type="checkbox"/>	Analysis Method	Show All Records	
9	<input checked="" type="checkbox"/>	Panel	Show All Records	
10	<input checked="" type="checkbox"/>	Size Standard	Show All Records	
11	<input type="checkbox"/>	Matrix	Show All Records	
12	<input checked="" type="checkbox"/>	Run Name	Show All Records	
13	<input type="checkbox"/>	Instrument Type	Show All Records	
14	<input type="checkbox"/>	Instrument ID	Show All Records	
15	<input type="checkbox"/>	Run Date & Time	Show All Records	
16	<input type="checkbox"/>	Reference Data	Show All Records	N/A
17	<input checked="" type="checkbox"/>	Sizing Quality Overridden	Show All Records	N/A
18	<input type="checkbox"/>	Sample File Not Found	Show All Records	N/A
19	<input type="checkbox"/>	Matrix Not Found	Show All Records	N/A
20	<input checked="" type="checkbox"/>	Size Standard Not Found	Show All Records	N/A
21	<input type="checkbox"/>	Off-scale	Show All Records	N/A
22	<input type="checkbox"/>	Sizing Quality	Show All Records	N/A
23	<input checked="" type="checkbox"/>	User Defined Column 1	Show All Records	
24	<input checked="" type="checkbox"/>	User Defined Column 2	Show All Records	
25	<input checked="" type="checkbox"/>	User Defined Column 3	Show All Records	

Font Settings:

Font: Arial

Size: 11

Show Hide

OK Cancel

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### TABLE SETTINGS – CASEWORK VIEW: GENOTYPES SETTINGS

Table Setting Editor

General Samples **Genotypes**

Genotypes Table Settings:

Column Settings:

Show	Column	Filtering	Content
<input type="checkbox"/>	Sample File	Show All Records	
<input type="checkbox"/>	Sample Name	Show All Records	
<input type="checkbox"/>	Sample ID	Show All Records	
<input type="checkbox"/>	Run Name	Show All Records	
<input type="checkbox"/>	Panel	Show All Records	
<input checked="" type="checkbox"/>	Marker	Show All Records	
<input checked="" type="checkbox"/>	Dye	Show All Records	N/A
<input checked="" type="checkbox"/>	Allele	Show All Records	
<input checked="" type="checkbox"/>	Size	Show All Records	
<input type="checkbox"/>	Height	Show All Records	
<input type="checkbox"/>	Peak Area	Show All Records	
<input type="checkbox"/>	Data Point	Show All Records	
<input type="checkbox"/>	Mutation	Show All Records	
<input checked="" type="checkbox"/>	AE Comment	Show All Records	
<input type="checkbox"/>	Integration Comments	Show All Records	
<input type="checkbox"/>	Allele Display Overflow	Show All Records	N/A
<input checked="" type="checkbox"/>	Allele Edit	Show All Records	N/A
<input type="checkbox"/>	Omit From Clustering (SNPle:	Show All Records	N/A
<input type="checkbox"/>	Off-scale	Show All Records	N/A

Show Hide

Font Settings:

Font: Arial

Size: 11

Allele Settings

Number of Alleles: 15  Keep Allele, Size, Height, Area, Data Point, Mutation and Comment together

OK Cancel

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### TABLE SETTINGS – CASEWORK VIEW: GENOTYPES SETTINGS (continued)

The screenshot shows the 'Table Setting Editor' window with the 'Genotypes' tab selected. The 'Genotypes Table Settings' section contains a table with columns for 'Show', 'Column', 'Filtering', and 'Content'. The 'Font Settings' section shows 'Font: Arial' and 'Size: 11'. The 'Allele Settings' section shows 'Number of Alleles: 15' and a checkbox for 'Keep Allele, Size, Height, Area, Data Point, Mutation and Comment together'.

Show	Column	Filtering	Content
<input type="checkbox"/>	Sharp Peak (M)	Show All Records	N/A
<input type="checkbox"/>	One Basepair Allele (M)	Show All Records	N/A
<input type="checkbox"/>	Single Peak Artifact (M)	Show All Records	N/A
<input type="checkbox"/>	Split Peak (M)	Show All Records	N/A
<input type="checkbox"/>	Out of Bin Allele	Show All Records	N/A
<input type="checkbox"/>	Peak Height Ratio	Show All Records	N/A
<input type="checkbox"/>	Low Peak Height	Show All Records	N/A
<input type="checkbox"/>	Spectral Pull-up	Show All Records	N/A
<input type="checkbox"/>	Allele Number	Show All Records	N/A
<input type="checkbox"/>	Broad Peak	Show All Records	N/A
<input type="checkbox"/>	Double Peak (SNP)	Show All Records	N/A
<input type="checkbox"/>	Narrow Bin (SNP)	Show All Records	N/A
<input type="checkbox"/>	Control Concordance	Show All Records	N/A
<input type="checkbox"/>	Overlap (HID)	Show All Records	N/A
<input type="checkbox"/>	Cross Talk	Show All Records	N/A
<input type="checkbox"/>	Genotype Quality	Show All Records	N/A
<input type="checkbox"/>	User Defined Column 1	Show All Records	
<input type="checkbox"/>	User Defined Column 2	Show All Records	
<input type="checkbox"/>	User Defined Column 3	Show All Records	

Font Settings:  
Font: Arial  
Size: 11

Allele Settings:  
Number of Alleles: 15  
 Keep Allele, Size, Height, Area, Data Point, Mutation and Comment together

Buttons: Show, Hide, OK, Cancel

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### PLOT SETTINGS: ANALYSIS VIEW

#### Analysis View: Sample Header

Plot Settings Editor

General | **Sample Header** | Genotype Header | Sizing Table | Labels | Display Settings

Sample Header Settings:

	Show	Column
1	<input checked="" type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input checked="" type="checkbox"/>	Sizing Quality Overridden
5	<input checked="" type="checkbox"/>	Off-scale
6	<input checked="" type="checkbox"/>	Sizing Quality

Show Hide

OK Cancel

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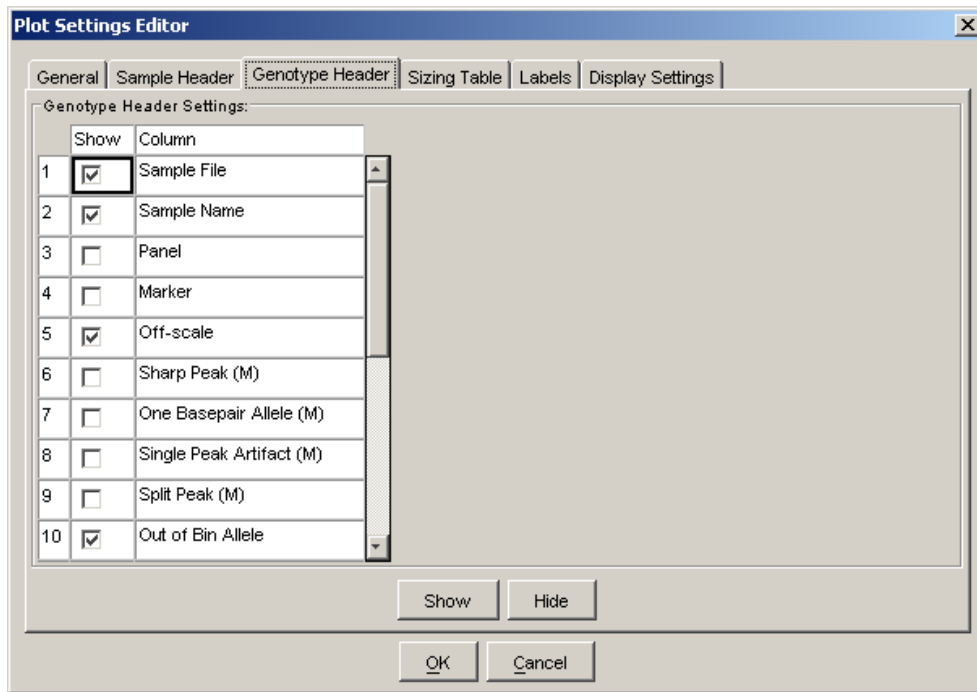
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### Analysis View: Genotype Header



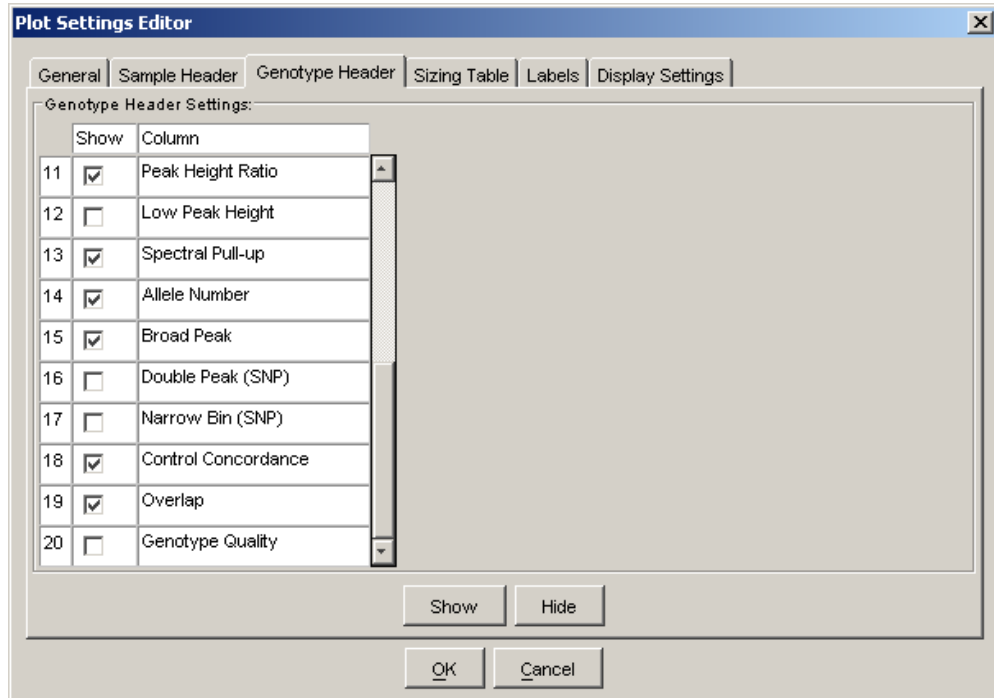
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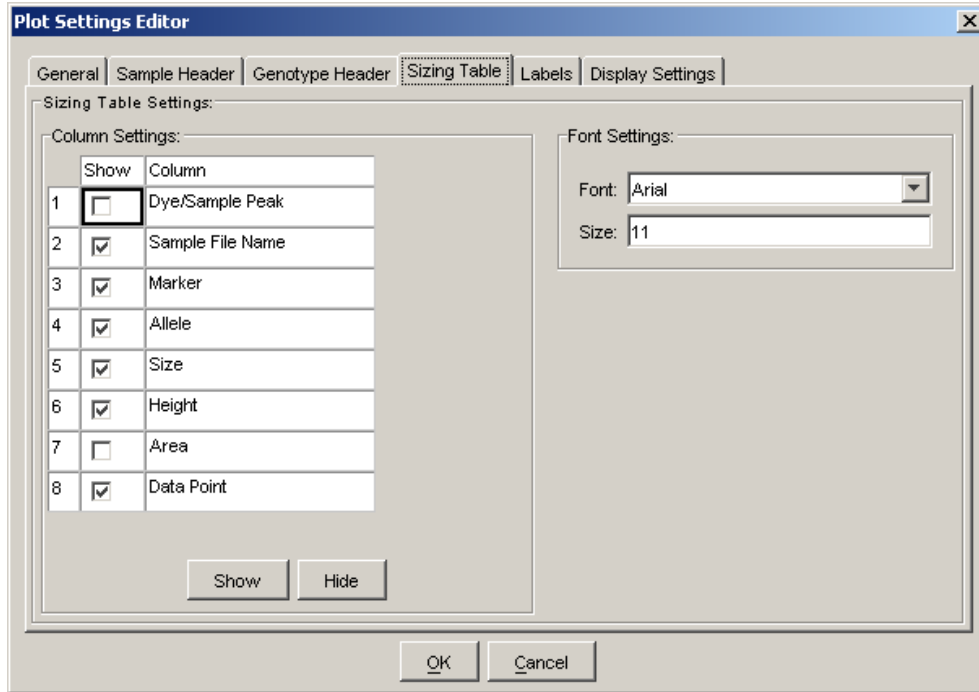
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### Analysis View: Sizing Table



### Analysis View: Labels

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**Plot Settings Editor**

General | Sample Header | Genotype Header | Sizing Table | **Labels** | Display Settings

Show Labels:

Label 1: Allele Call

Label 2: Size

Label 3: Height

Label 4: AE Comment

Font: Times New Roman

Size: 9

When opening the Plot Window:

Show data type prefixes

Show type of edit

Invert mutant labels

Label Color: Dye Color-Border

OK Cancel

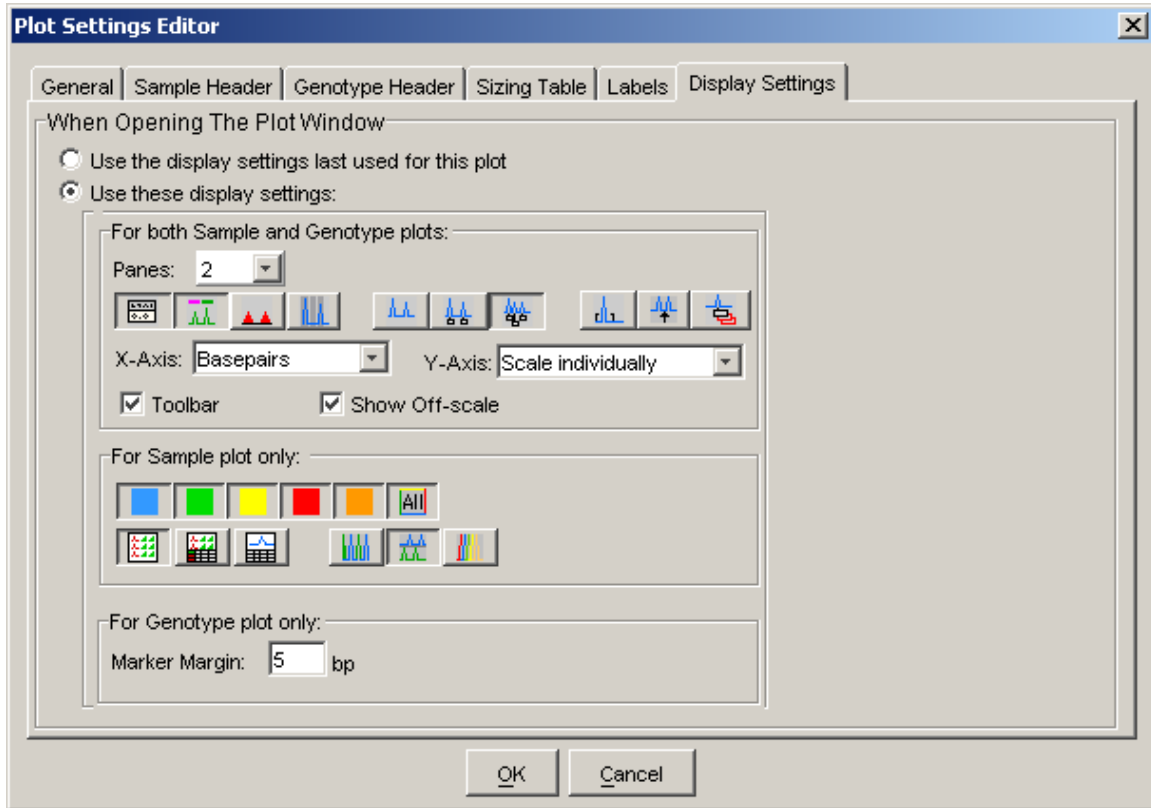
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### Analysis View: Display Settings



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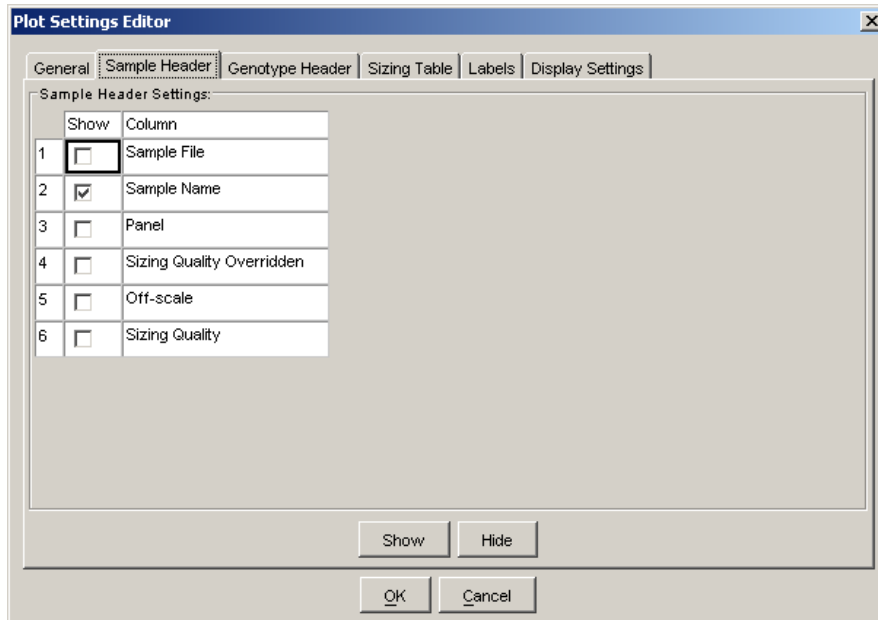
# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

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### PLOT SETTINGS: PRINT – IDENTIFILER ALLELIC LADDER

#### Print – Identifiler Allelic Ladder: Sample Header



#### Print – Identifiler Allelic Ladder: Genotype Header

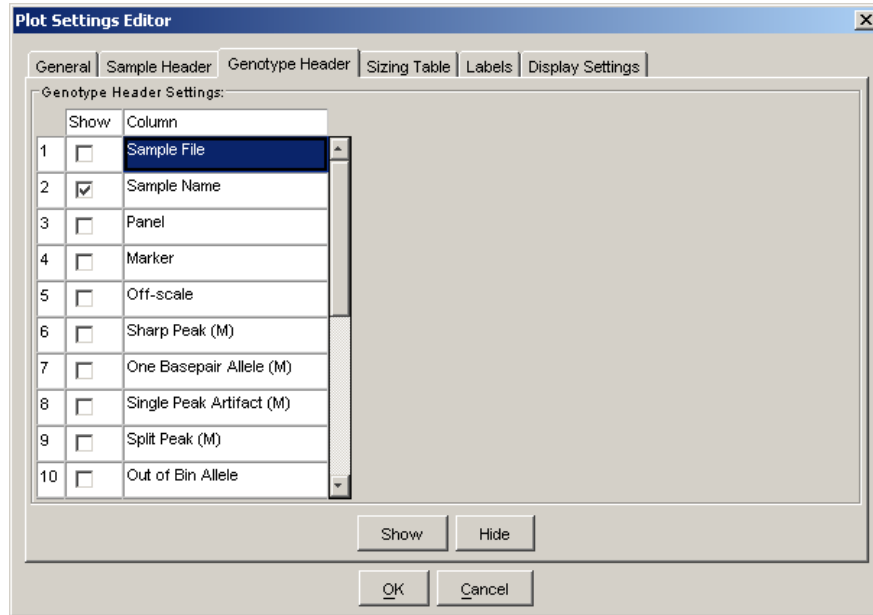
# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

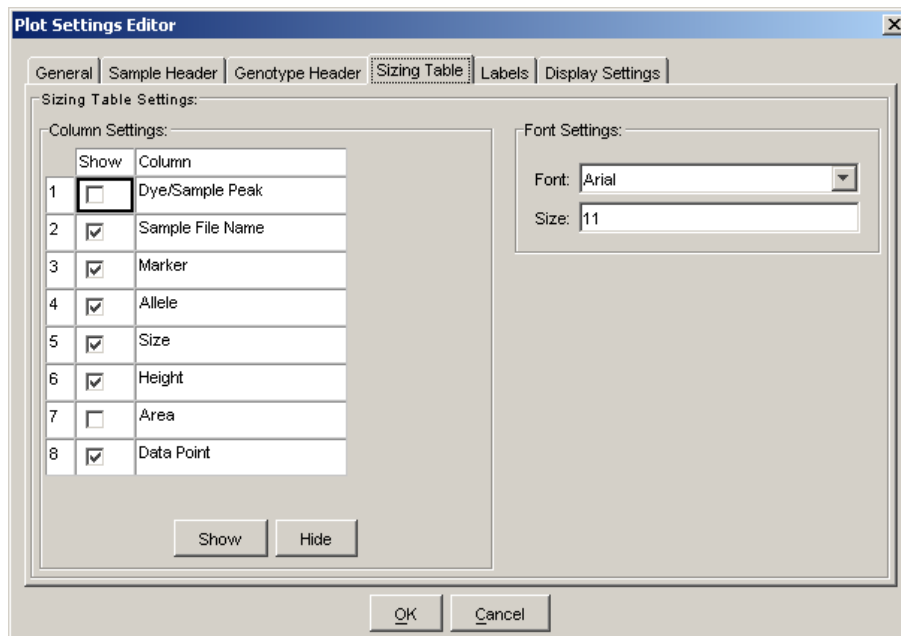
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**Boxes 3 – 20 are unchecked**  
**Print – Identifiler Allelic Ladder: Sizing Table**



**Print – Identifiler Allelic Ladder: Labels**

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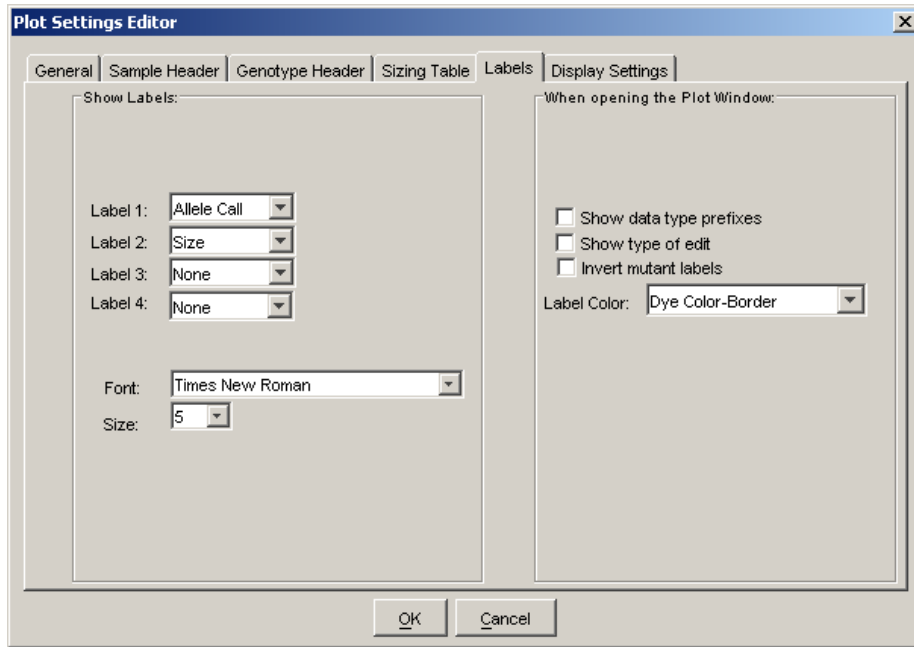
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### Print – Identifier Allelic Ladder: Display Settings

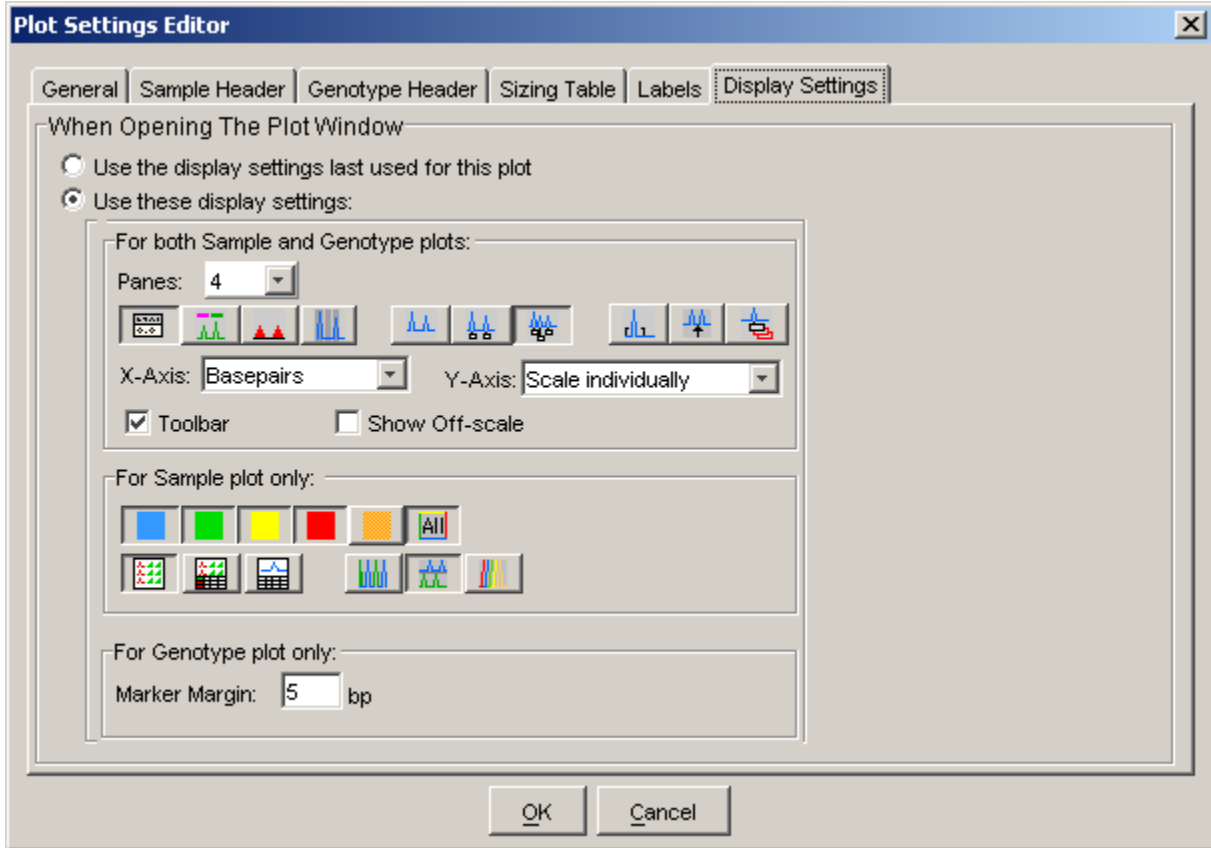
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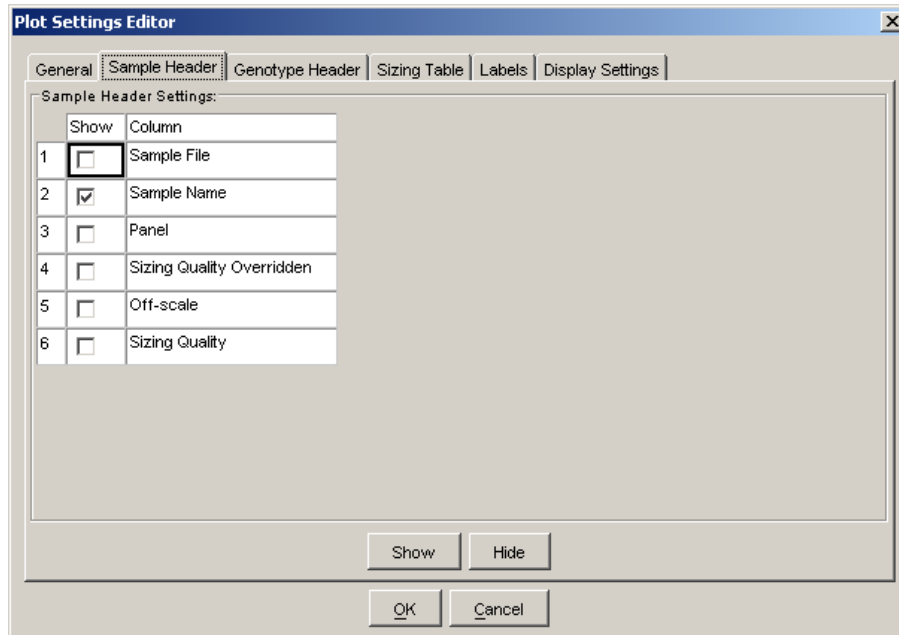
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### PLOT SETTINGS: PRINT – IDENTIFILER 28 CONTROLS

#### Print – Identifiler 28 Controls: Sample Header

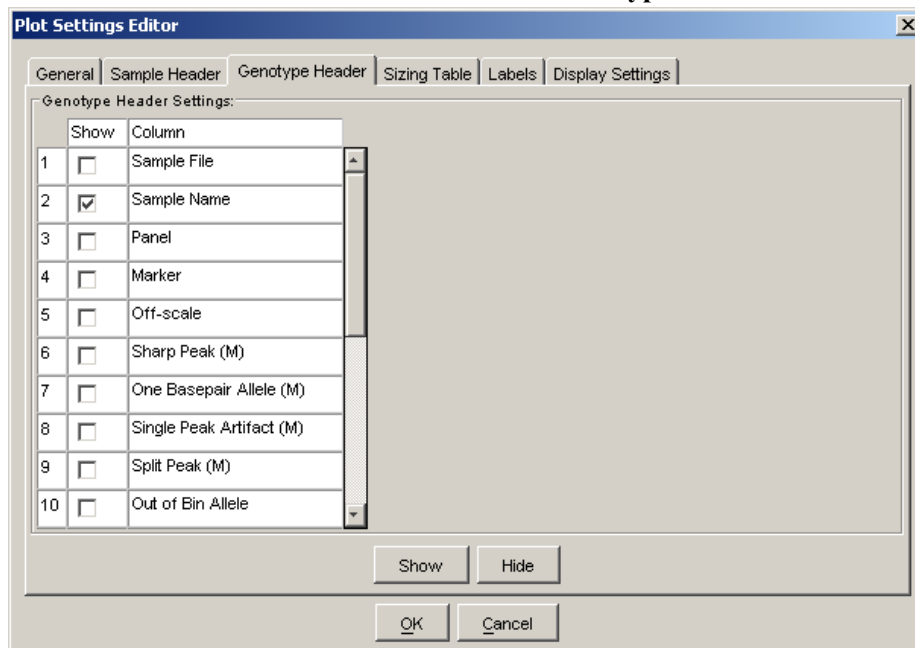


The screenshot shows the 'Plot Settings Editor' window with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Sizing Quality Overridden
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sizing Quality

Buttons at the bottom: Show, Hide, OK, Cancel.

#### Print – Identifiler 28 Controls: Genotype Header



The screenshot shows the 'Plot Settings Editor' window with the 'Genotype Header' tab selected. The 'Genotype Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Marker
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sharp Peak (M)
7	<input type="checkbox"/>	One Basepair Allele (M)
8	<input type="checkbox"/>	Single Peak Artifact (M)
9	<input type="checkbox"/>	Split Peak (M)
10	<input type="checkbox"/>	Out of Bin Allele

Buttons at the bottom: Show, Hide, OK, Cancel.

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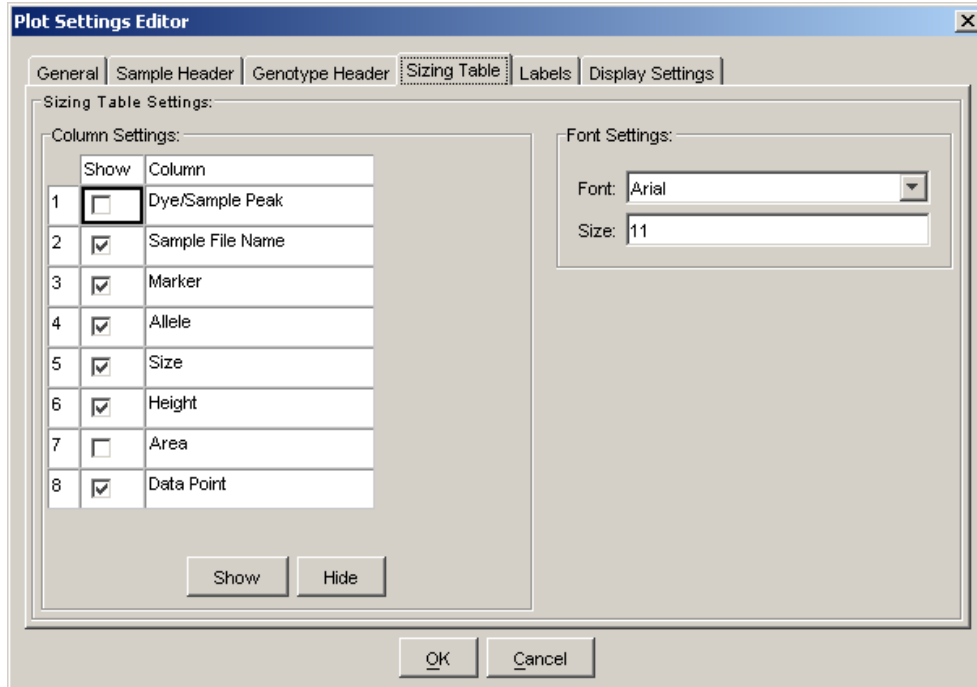
**Boxes 3 – 20 are unchecked**

# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

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### Print – Identifiler 28 Controls: Sizing Table



### Print – Identifiler 28 Controls: Labels

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**Plot Settings Editor**

General | Sample Header | Genotype Header | Sizing Table | **Labels** | Display Settings

Show Labels:

Label 1: Allele Call

Label 2: Size

Label 3: Height

Label 4: None

Font: Times New Roman

Size: 5

When opening the Plot Window:

Show data type prefixes

Show type of edit

Invert mutant labels

Label Color: Dye Color-Border

OK Cancel

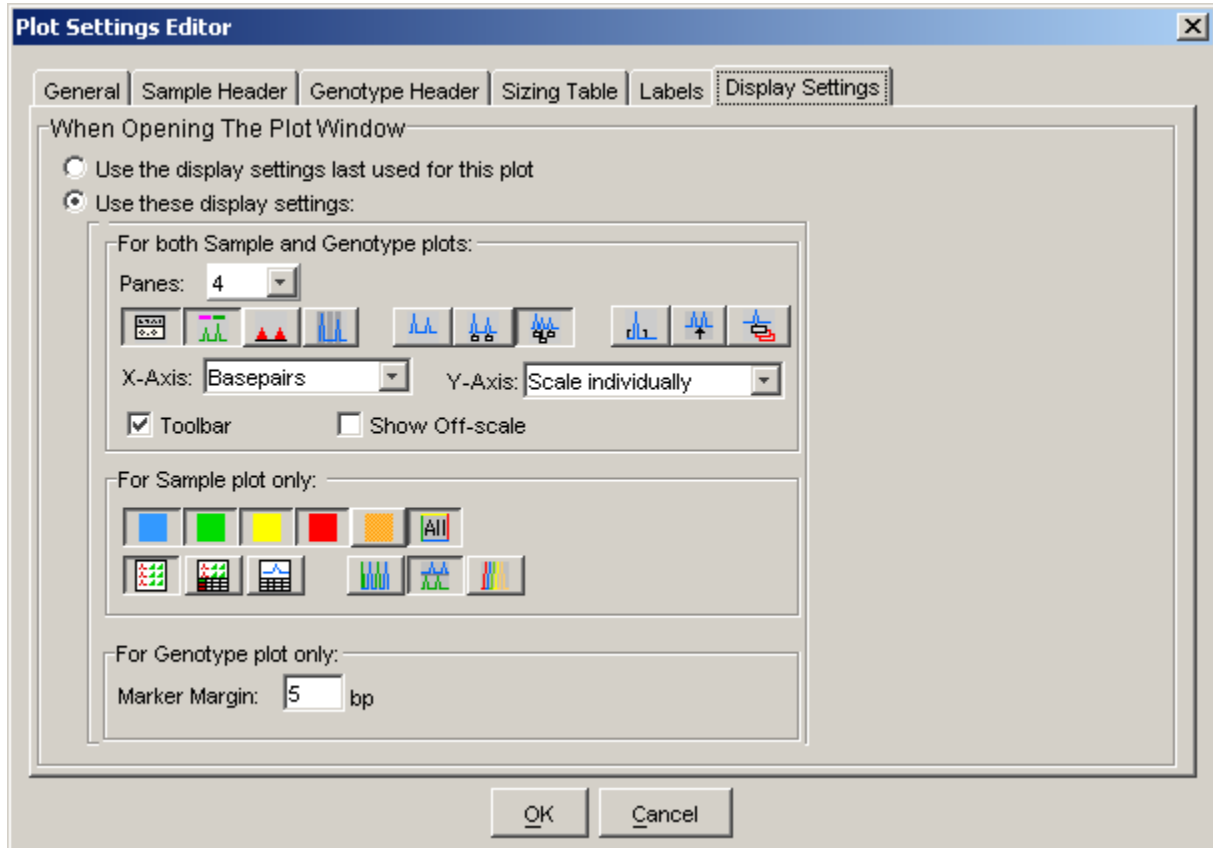
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### Print – Identifiler 28 Controls: Display Settings



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## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### PLOT SETTINGS: PRINT – IDENTIFILER 28 SAMPLES

#### Print – Identifiler28 Samples: Sample Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sample Header' tab selected. The dialog has a title bar with a close button (X). Below the title bar are tabs for 'General', 'Sample Header', 'Genotype Header', 'Sizing Table', 'Labels', and 'Display Settings'. The 'Sample Header' tab contains a table with the following settings:

Show	Column
1 <input type="checkbox"/>	Sample File
2 <input checked="" type="checkbox"/>	Sample Name
3 <input type="checkbox"/>	Panel
4 <input type="checkbox"/>	Sizing Quality Overridden
5 <input type="checkbox"/>	Off-scale
6 <input type="checkbox"/>	Sizing Quality

At the bottom of the dialog are buttons for 'Show', 'Hide', 'OK', and 'Cancel'.

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### Print – Identifier28 Samples: Genotype Header

Plot Settings Editor

General | Sample Header | Genotype Header | Sizing Table | Labels | Display Settings

Genotype Header Settings:

Show	Column
<input type="checkbox"/>	Sample File
<input checked="" type="checkbox"/>	Sample Name
<input type="checkbox"/>	Panel
<input type="checkbox"/>	Marker
<input type="checkbox"/>	Off-scale
<input type="checkbox"/>	Sharp Peak (M)
<input type="checkbox"/>	One Basepair Allele (M)
<input type="checkbox"/>	Single Peak Artifact (M)
<input type="checkbox"/>	Split Peak (M)
<input type="checkbox"/>	Out of Bin Allele

Show Hide

OK Cancel

**Boxes 3 – 20 are unchecked**

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## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### Print – Identifiler28 Samples: Sizing Table

Plot Settings Editor

General | Sample Header | Genotype Header | **Sizing Table** | Labels | Display Settings

Sizing Table Settings:

	Show	Column
1	<input type="checkbox"/>	Dye/Sample Peak
2	<input checked="" type="checkbox"/>	Sample File Name
3	<input checked="" type="checkbox"/>	Marker
4	<input checked="" type="checkbox"/>	Allele
5	<input checked="" type="checkbox"/>	Size
6	<input checked="" type="checkbox"/>	Height
7	<input type="checkbox"/>	Area
8	<input checked="" type="checkbox"/>	Data Point

Font Settings:

Font: Arial

Size: 11

Show Hide

OK Cancel

### Print – Identifiler28 Samples: Labels

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## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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**Plot Settings Editor**

General | Sample Header | Genotype Header | Sizing Table | **Labels** | Display Settings

Show Labels:

Label 1: Allele Call

Label 2: Size

Label 3: Height

Label 4: None

Font: Times New Roman

Size: 5

When opening the Plot Window:

Show data type prefixes

Show type of edit

Invert mutant labels

Label Color: Dye Color-Border

OK Cancel

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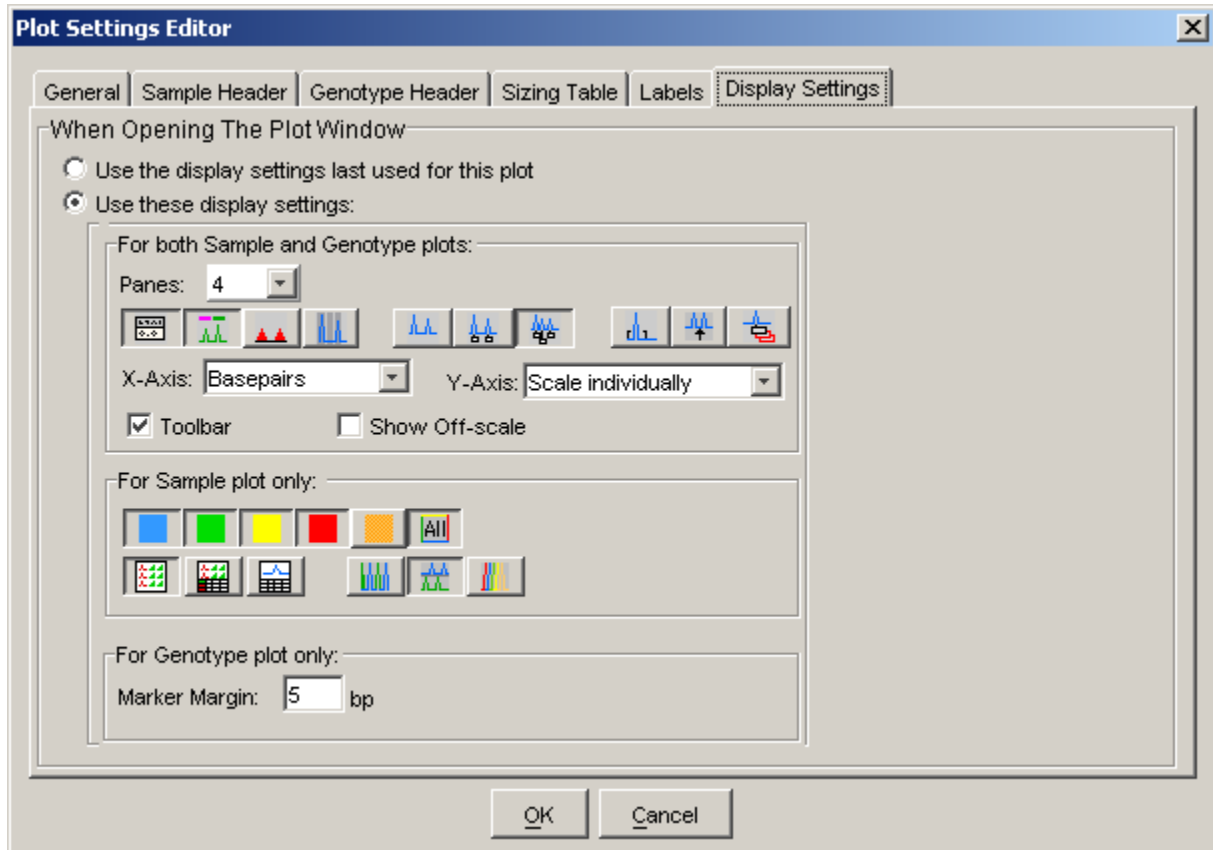
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## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### Print – Identifiler28 Samples: Display Settings



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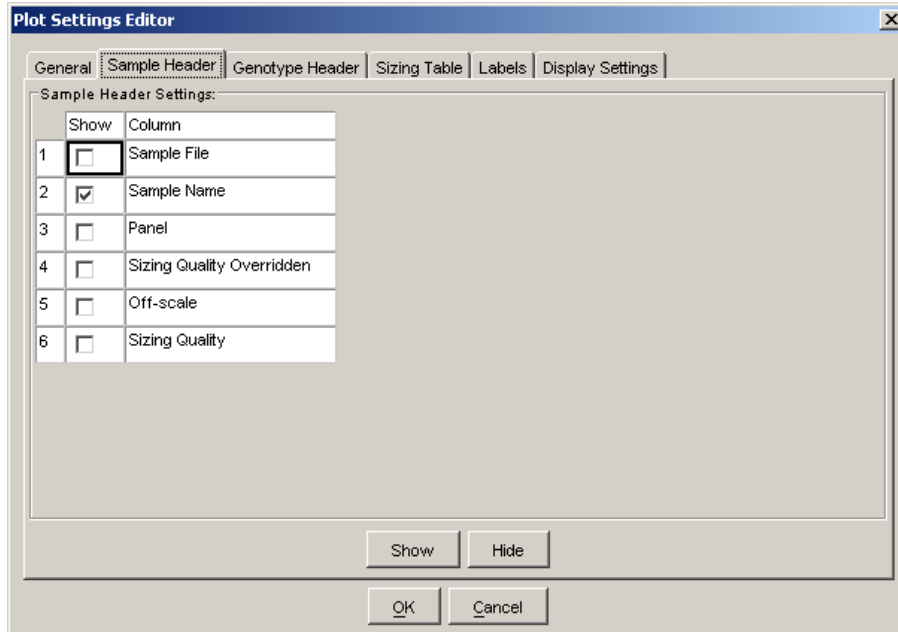
# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### PLOT SETTINGS: PRINT – IDENTIFILER 31 POSITIVE CONTROL (PE) AND SAMPLES

#### Print – ID 31 PE and Samples: Sample Header



#### Print – ID 31 PE and Samples: Genotype Header

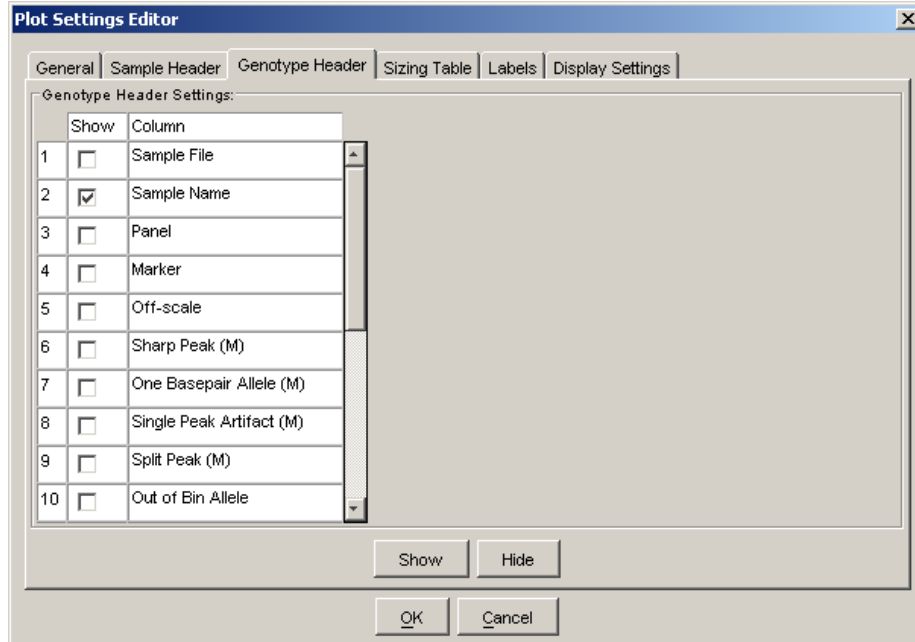
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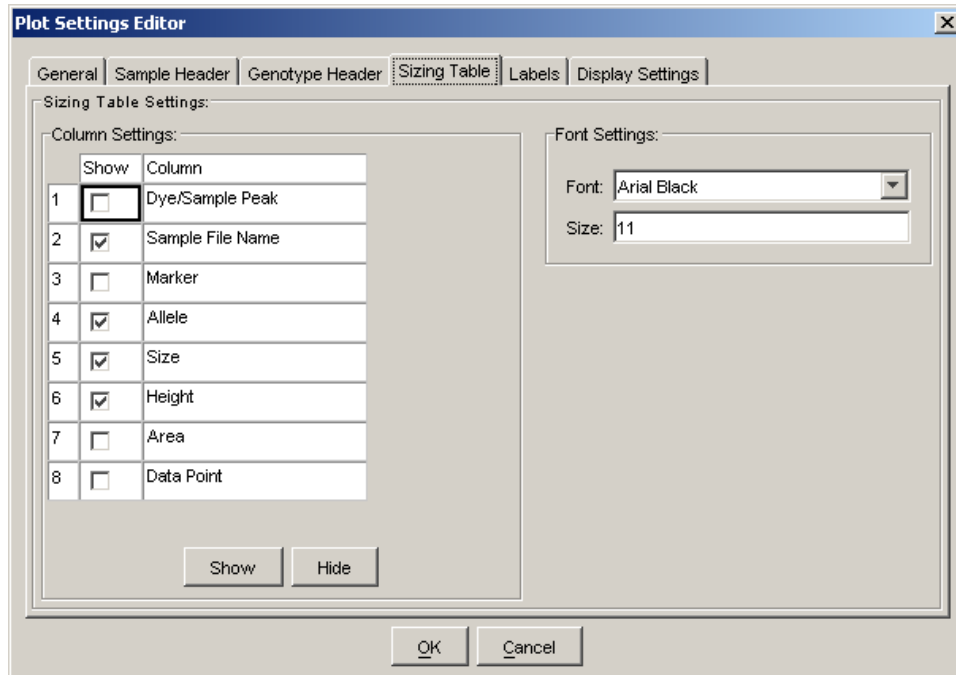
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### Print – ID 31 PE and Samples: Sizing Table



### Print – ID 31 PE and Samples: Labels

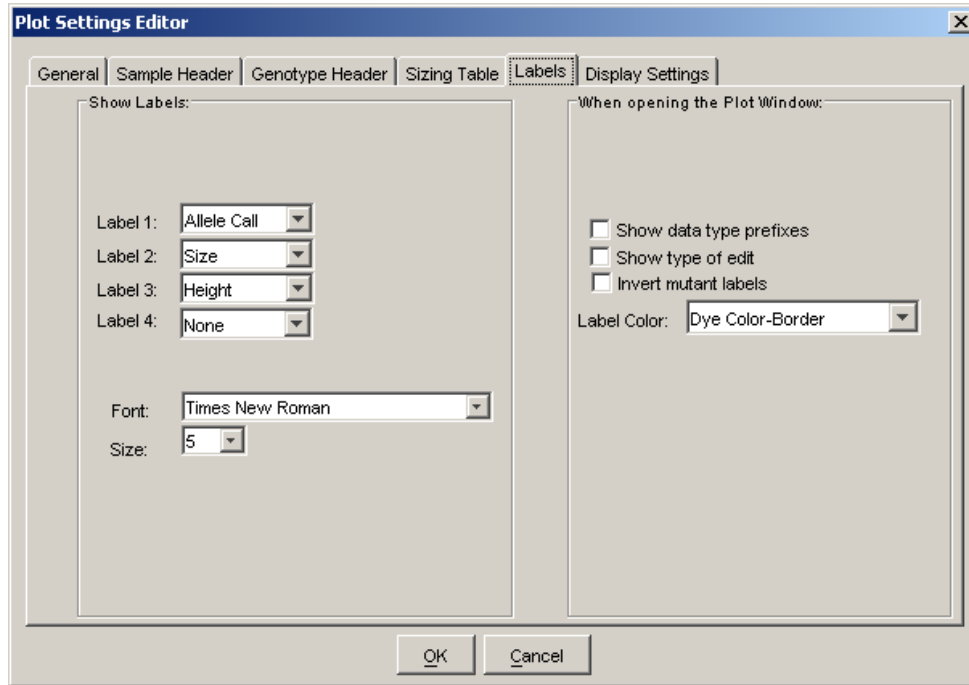
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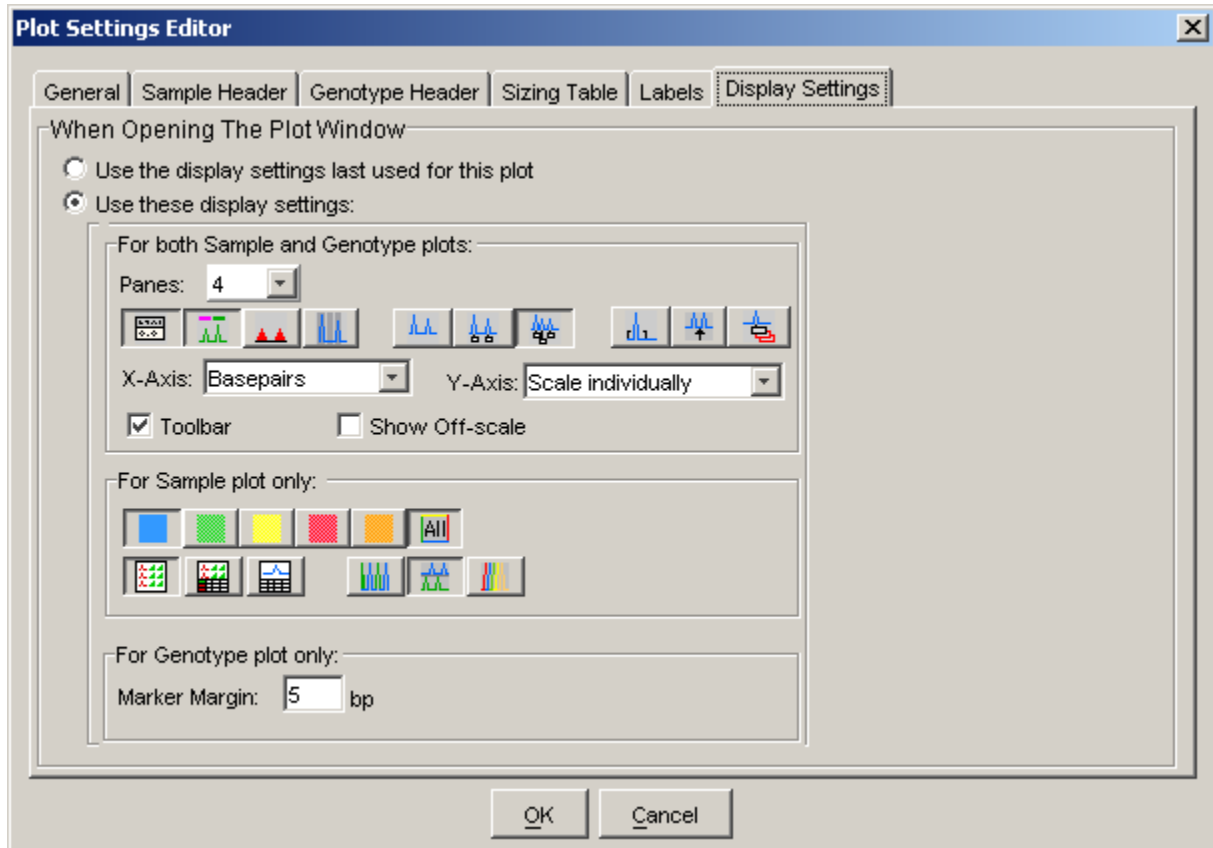


# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### Print – ID 31 PE and Samples: Display Settings



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## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### PLOT SETTINGS: PRINT – IDENTIFILER 31 NEGATIVE CONTROLS

#### Print – ID 31 Negative Controls: Sample Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Sizing Quality Overridden
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sizing Quality

Buttons at the bottom: Show, Hide, OK, Cancel.

#### Print – ID 31 Negative Controls: Genotype Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Genotype Header' tab selected. The 'Genotype Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Marker
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sharp Peak (M)
7	<input type="checkbox"/>	One Basepair Allele (M)
8	<input type="checkbox"/>	Single Peak Artifact (M)
9	<input type="checkbox"/>	Split Peak (M)
10	<input type="checkbox"/>	Out of Bin Allele

Buttons at the bottom: Show, Hide, OK, Cancel.

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**Boxes 3 – 20 are unchecked**

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### Print – ID 31 Negative Controls: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a table for column settings and font settings.

Show	Column
<input type="checkbox"/>	Dye/Sample Peak
<input checked="" type="checkbox"/>	Sample File Name
<input checked="" type="checkbox"/>	Marker
<input checked="" type="checkbox"/>	Allele
<input checked="" type="checkbox"/>	Size
<input checked="" type="checkbox"/>	Height
<input type="checkbox"/>	Area
<input checked="" type="checkbox"/>	Data Point

Font Settings:  
Font: Arial  
Size: 11

Buttons: Show, Hide, OK, Cancel

### Print – ID 31 Negative Controls: Labels

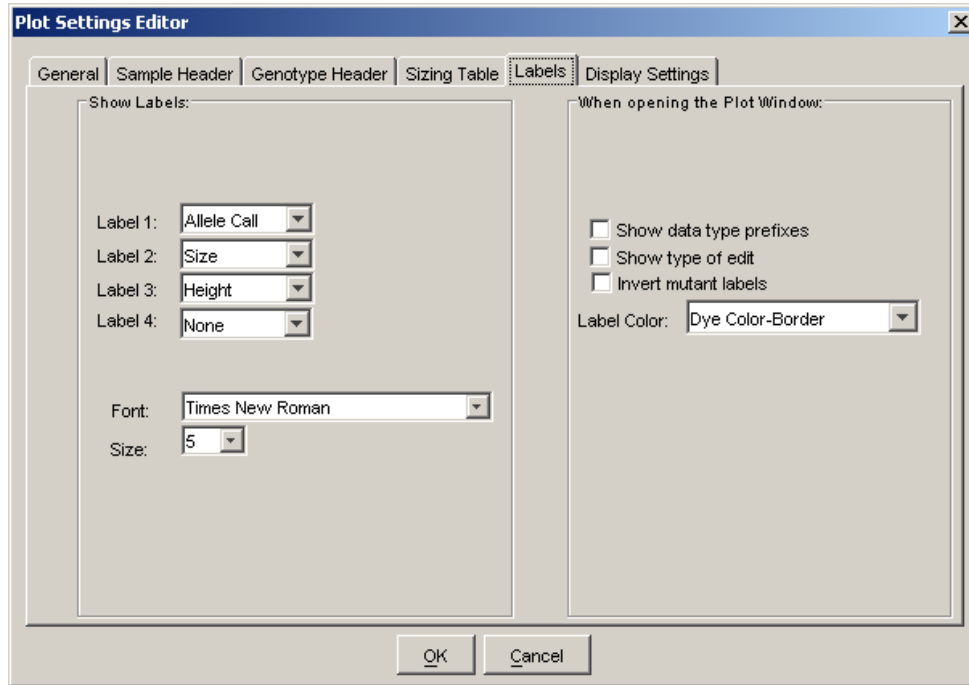
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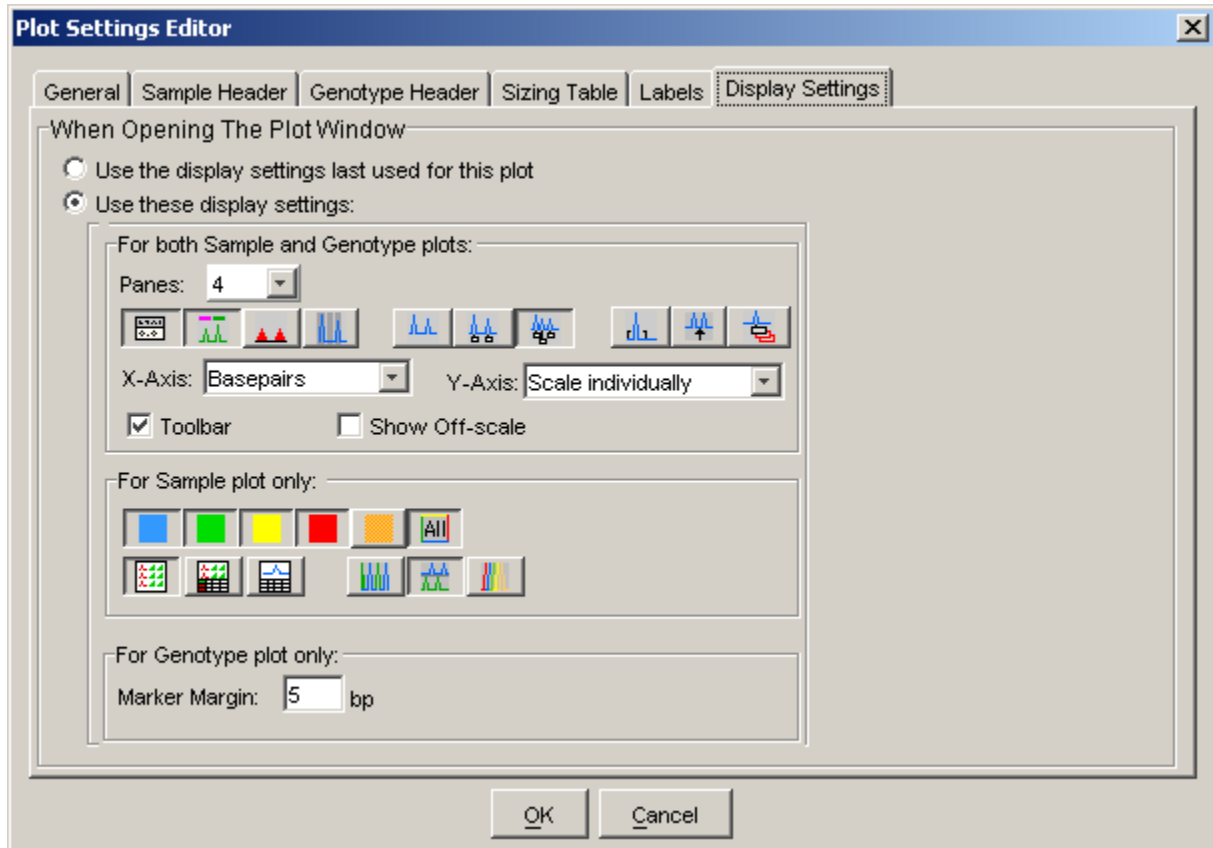
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### Print – ID 31 Negative Controls: Display Settings



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### PLOT SETTINGS: PRINT – YFILER ALLELIC LADDER

#### Print – YFiler Allelic Ladder: Sample Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sample Header' tab selected. The 'Sample Header Settings' section contains a table with the following items:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Sizing Quality Overridden
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sizing Quality

Buttons at the bottom include 'Show', 'Hide', 'OK', and 'Cancel'.

#### Print – YFiler Allelic Ladder: Genotypes Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Genotype Header' tab selected. The 'Genotype Header Settings' section contains a table with the following items:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Marker
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sharp Peak (M)
7	<input type="checkbox"/>	One Basepair Allele (M)
8	<input type="checkbox"/>	Single Peak Artifact (M)
9	<input type="checkbox"/>	Split Peak (M)
10	<input type="checkbox"/>	Out of Bin Allele

Buttons at the bottom include 'Show', 'Hide', 'OK', and 'Cancel'.

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**Boxes 3 – 20 are unchecked**



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### Print – YFiler Y Allelic Ladder: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a 'Column Settings' table and 'Font Settings'.

	Show	Column
1	<input type="checkbox"/>	Dye/Sample Peak
2	<input checked="" type="checkbox"/>	Sample File Name
3	<input checked="" type="checkbox"/>	Marker
4	<input checked="" type="checkbox"/>	Allele
5	<input checked="" type="checkbox"/>	Size
6	<input checked="" type="checkbox"/>	Height
7	<input type="checkbox"/>	Area
8	<input checked="" type="checkbox"/>	Data Point

Font Settings:  
Font: Arial  
Size: 11

Buttons: Show, Hide, OK, Cancel

### Print – YFiler Allelic Ladder: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section contains dropdown menus for Label 1, Label 2, Label 3, and Label 4, along with font settings. The 'When opening the Plot Window' section contains checkboxes for 'Show data type prefixes', 'Show type of edit', and 'Invert mutant labels', and a 'Label Color' dropdown.

Show Labels:  
Label 1: Allele Call  
Label 2: Size  
Label 3: None  
Label 4: None  
Font: Times New Roman  
Size: 5

When opening the Plot Window:  
 Show data type prefixes  
 Show type of edit  
 Invert mutant labels  
Label Color: Dye Color-Border

Buttons: OK, Cancel

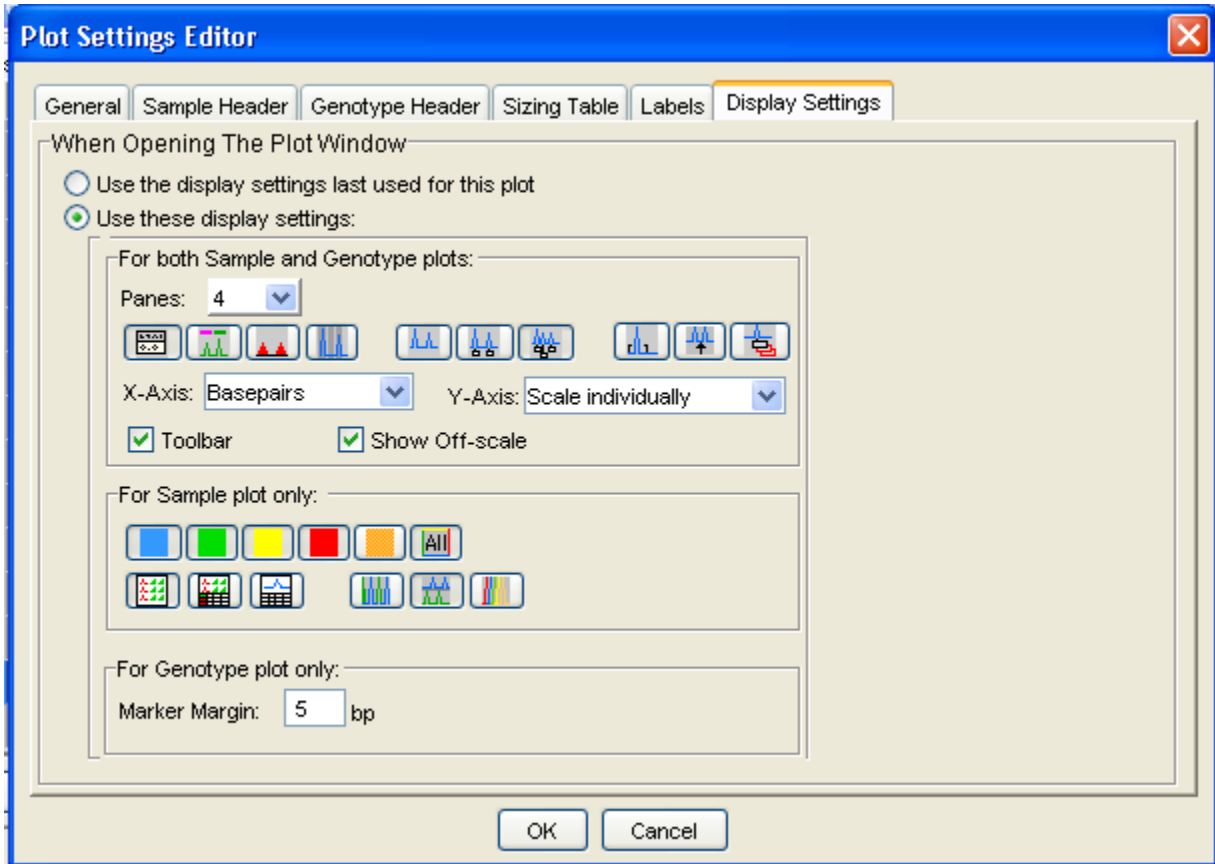
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### Print – Yfiler Allelic Ladder: Display Settings



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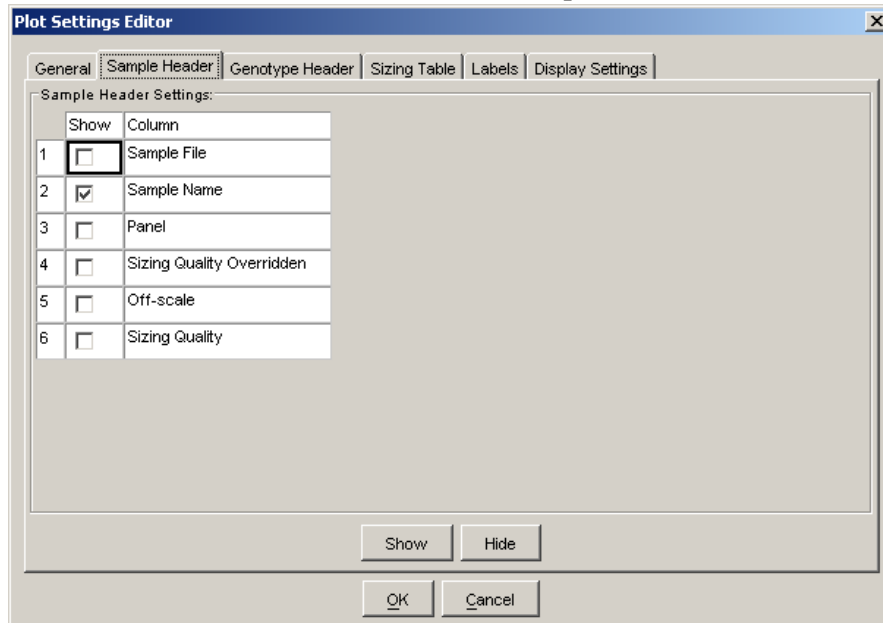
# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### PLOT SETTINGS: PRINT – YFILER CONTROLS

#### Print – YFiler Controls: Sample Header



#### Print – YFiler Controls: Genotypes Header

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The screenshot shows the 'Plot Settings Editor' dialog box with the 'Genotype Header' tab selected. The 'Genotype Header Settings' section contains a table with 10 rows. The first row, 'Sample File', has a checked checkbox. The remaining rows, from 'Sample Name' to 'Out of Bin Allele', have unchecked checkboxes. Below the table are 'Show' and 'Hide' buttons, and at the bottom are 'OK' and 'Cancel' buttons.

	Show	Column
1	<input checked="" type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Marker
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sharp Peak (M)
7	<input type="checkbox"/>	One Basepair Allele (M)
8	<input type="checkbox"/>	Single Peak Artifact (M)
9	<input type="checkbox"/>	Split Peak (M)
10	<input type="checkbox"/>	Out of Bin Allele

**Boxes 3 – 20 are unchecked**

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### Print – YFiler Controls: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a table for column visibility and font settings.

	Show	Column
1	<input type="checkbox"/>	Dye/Sample Peak
2	<input checked="" type="checkbox"/>	Sample File Name
3	<input checked="" type="checkbox"/>	Marker
4	<input checked="" type="checkbox"/>	Allele
5	<input checked="" type="checkbox"/>	Size
6	<input checked="" type="checkbox"/>	Height
7	<input type="checkbox"/>	Area
8	<input checked="" type="checkbox"/>	Data Point

Font Settings:  
Font: Arial  
Size: 11

Buttons: Show, Hide, OK, Cancel

### Print – YFiler Controls: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section has dropdown menus for label types and font settings. The 'When opening the Plot Window' section has checkboxes for display options and a label color dropdown.

Show Labels:  
Label 1: Allele Call  
Label 2: Size  
Label 3: Height  
Label 4: None  
Font: Times New Roman  
Size: 5

When opening the Plot Window:  
 Show data type prefixes  
 Show type of edit  
 Invert mutant labels  
Label Color: Dye Color-Border

Buttons: OK, Cancel

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### Print – Yfiler Controls: Display Settings



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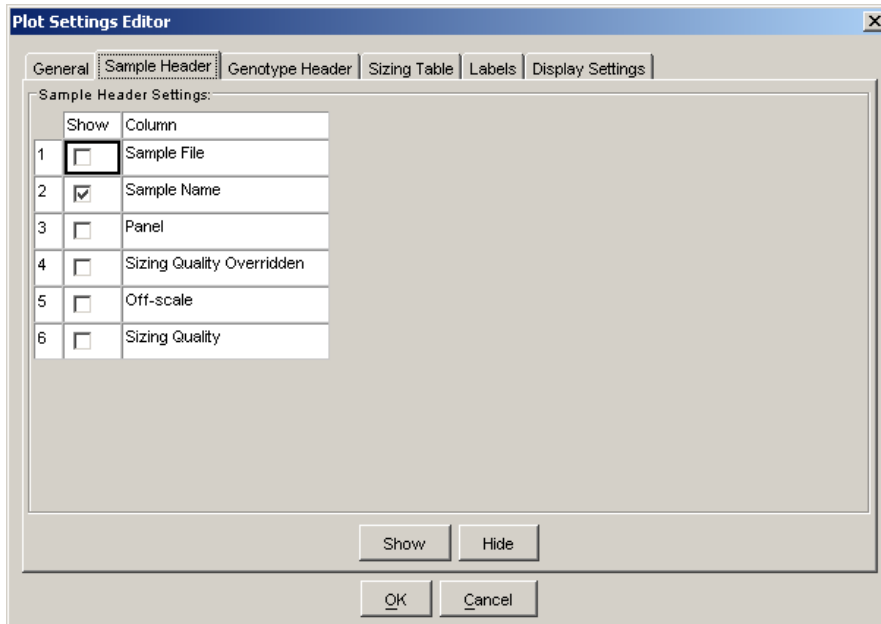
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## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### PLOT SETTINGS: PRINT – YFiler SAMPLES

#### Print – YFiler Samples: Sample Header

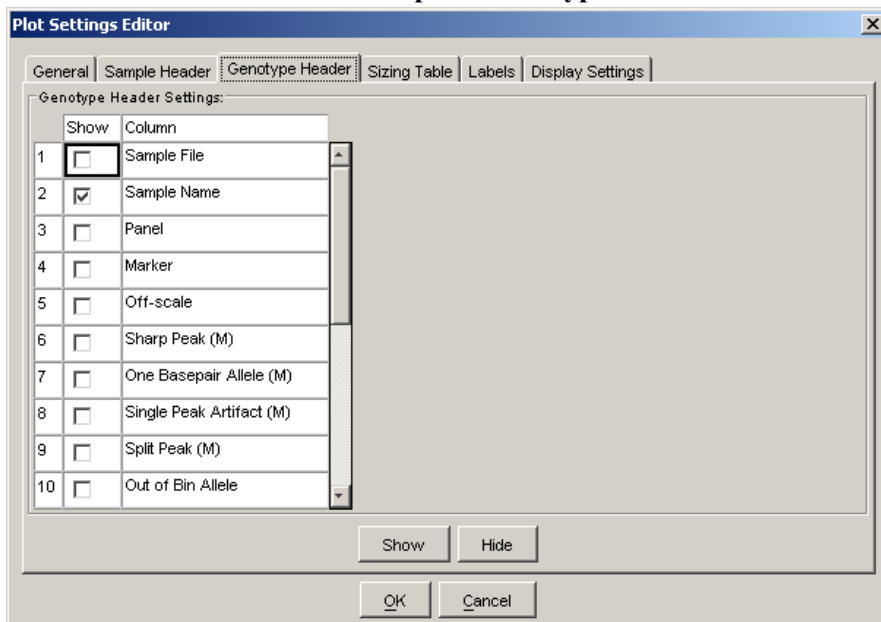


The screenshot shows the 'Plot Settings Editor' window with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Sizing Quality Overridden
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sizing Quality

Buttons at the bottom: Show, Hide, OK, Cancel.

#### Print – YFiler Samples: Genotypes Header



The screenshot shows the 'Plot Settings Editor' window with the 'Genotype Header' tab selected. The 'Genotype Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Marker
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sharp Peak (M)
7	<input type="checkbox"/>	One Basepair Allele (M)
8	<input type="checkbox"/>	Single Peak Artifact (M)
9	<input type="checkbox"/>	Split Peak (M)
10	<input type="checkbox"/>	Out of Bin Allele

Buttons at the bottom: Show, Hide, OK, Cancel.

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**Boxes 3 – 20 are unchecked**

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### Print – Yfiler Samples: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a 'Column Settings' table and 'Font Settings'.

	Show	Column
1	<input type="checkbox"/>	Dye/Sample Peak
2	<input checked="" type="checkbox"/>	Sample File Name
3	<input checked="" type="checkbox"/>	Marker
4	<input checked="" type="checkbox"/>	Allele
5	<input checked="" type="checkbox"/>	Size
6	<input checked="" type="checkbox"/>	Height
7	<input type="checkbox"/>	Area
8	<input checked="" type="checkbox"/>	Data Point

Font Settings:  
Font: Arial  
Size: 11

Buttons: Show, Hide, OK, Cancel

### Print – Yfiler Samples: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section has four dropdown menus for Label 1 through Label 4, and font settings. The 'When opening the Plot Window' section has three checkboxes and a label color dropdown.

Show Labels:  
Label 1: Allele Call  
Label 2: Size  
Label 3: Height  
Label 4: None  
Font: Times New Roman  
Size: 5

When opening the Plot Window:  
 Show data type prefixes  
 Show type of edit  
 Invert mutant labels  
Label Color: Dye Color-Border

Buttons: OK, Cancel

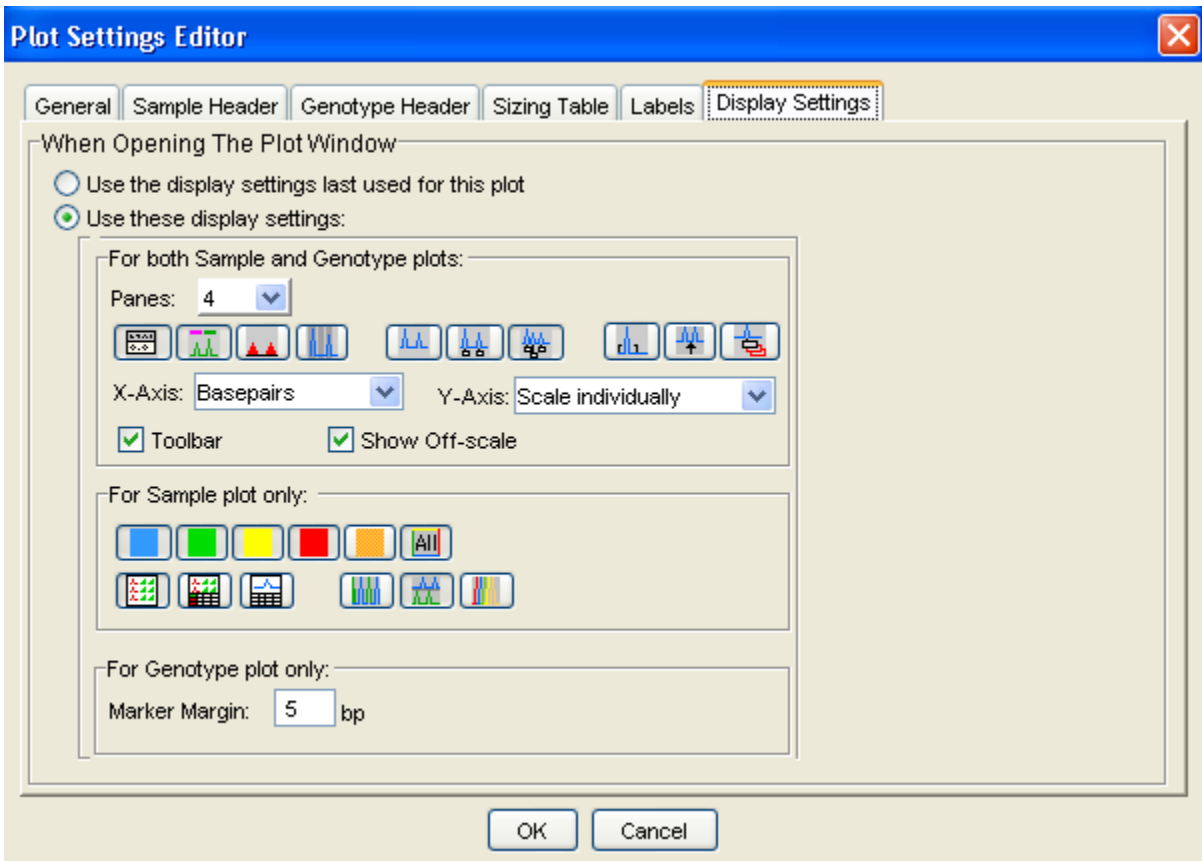
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### Print – Yfiler Samples: Display Settings



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### PLOT SETTINGS: MINIFILER ANALYSIS VIEW

#### MiniFiler Analysis View: Sample Header

The image shows a screenshot of the 'Plot Settings Editor' dialog box, specifically the 'Sample Header' tab. The dialog box has a title bar with 'Plot Settings Editor' and a close button. Below the title bar are several tabs: 'General', 'Sample Header' (which is selected), 'Genotype Header', 'Sizing Table', 'Labels', and 'Display Settings'. The 'Sample Header Settings' section contains a table with the following data:

	Show	Column
1	<input checked="" type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input checked="" type="checkbox"/>	Sizing Quality Overridden
5	<input checked="" type="checkbox"/>	Off-scale
6	<input checked="" type="checkbox"/>	Sizing Quality

Below the table are two buttons: 'Show' and 'Hide'. At the bottom of the dialog box are two buttons: 'OK' and 'Cancel'.

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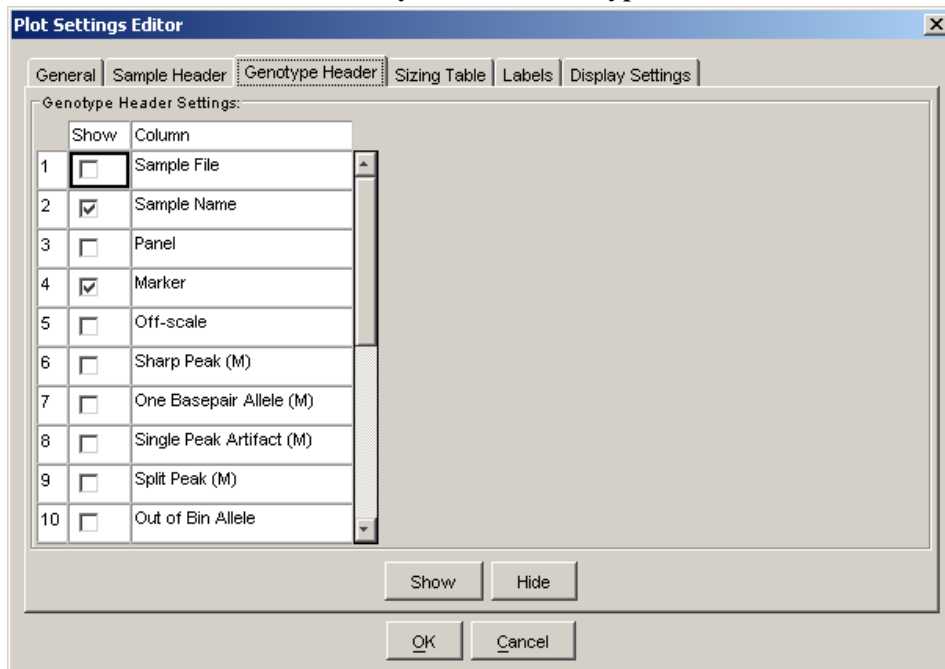
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# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### MiniFiler Analysis View: Genotype Header



5-20 are blank

### MiniFiler Analysis View: Sizing Table

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**Plot Settings Editor**

General | Sample Header | Genotype Header | **Sizing Table** | Labels | Display Settings

Sizing Table Settings:

Column Settings:

	Show	Column
1	<input checked="" type="checkbox"/>	Dye/Sample Peak
2	<input checked="" type="checkbox"/>	Sample File Name
3	<input checked="" type="checkbox"/>	Marker
4	<input checked="" type="checkbox"/>	Allele
5	<input checked="" type="checkbox"/>	Size
6	<input checked="" type="checkbox"/>	Height
7	<input checked="" type="checkbox"/>	Area
8	<input checked="" type="checkbox"/>	Data Point

Font Settings:

Font: Arial

Size: 11

Show Hide

OK Cancel

### MiniFiler Analysis View: Labels

**Plot Settings Editor**

General | Sample Header | Genotype Header | Sizing Table | **Labels** | Display Settings

Show Labels:

Label 1: Allele Call

Label 2: Size

Label 3: Height

Label 4: AE Comment

Font: Times New Roman

Size: 10

When opening the Plot Window:

Show data type prefixes

Show type of edit

Invert mutant labels

Label Color: Dye Color-Border

OK Cancel

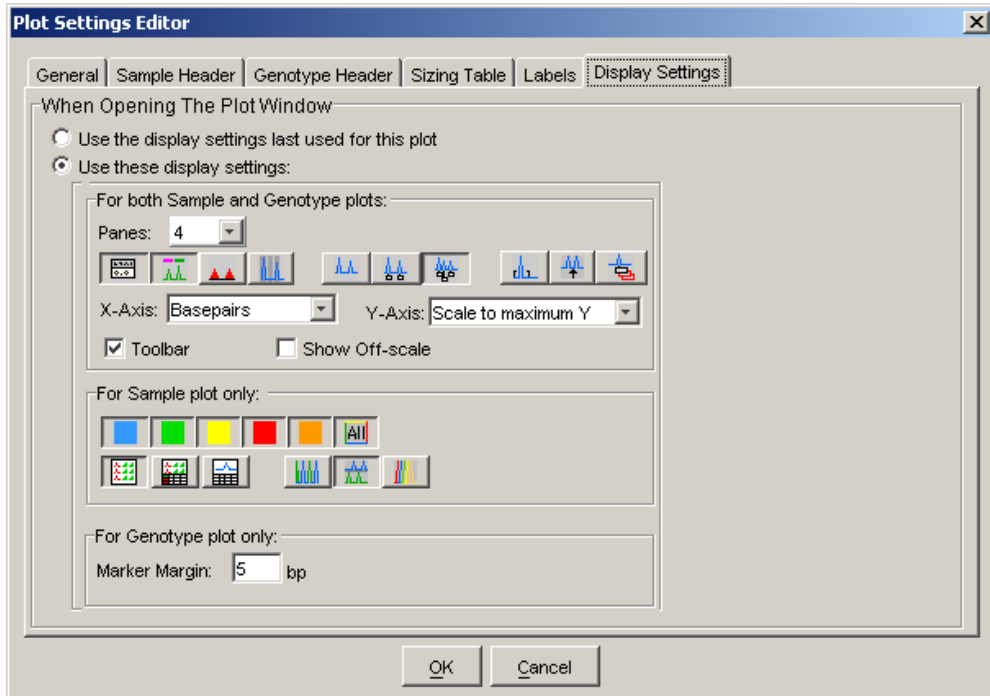
### MiniFiler Analysis View: Display Settings

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### PLOT SETTINGS: PRINT – MINIFILER ALLELIC LADDER

#### Print – MiniFiler Allelic Ladder: Sample Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

Show	Column
<input checked="" type="checkbox"/>	Sample File
<input checked="" type="checkbox"/>	Sample Name
<input type="checkbox"/>	Panel
<input type="checkbox"/>	Sizing Quality Overridden
<input type="checkbox"/>	Off-scale
<input type="checkbox"/>	Sizing Quality

Buttons at the bottom: Show, Hide, OK, Cancel.

#### Print – MiniFiler Allelic Ladder: Genotype Header

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Genotype Header' tab selected. The 'Genotype Header Settings' table is as follows:

Show	Column
<input type="checkbox"/>	Sample File
<input checked="" type="checkbox"/>	Sample Name
<input type="checkbox"/>	Panel
<input checked="" type="checkbox"/>	Marker
<input type="checkbox"/>	Off-scale
<input type="checkbox"/>	Sharp Peak (M)
<input type="checkbox"/>	One Basepair Allele (M)
<input type="checkbox"/>	Single Peak Artifact (M)
<input type="checkbox"/>	Split Peak (M)
<input type="checkbox"/>	Out of Bin Allele

Buttons at the bottom: Show, Hide, OK, Cancel.

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# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

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### Print – MiniFiler Allelic Ladder: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a table with 8 columns and 8 rows. The 'Show' column has checkboxes for each row, all of which are checked. The 'Column' column lists the following items: Dye/Sample Peak, Sample File Name, Marker, Allele, Size, Height, Area, and Data Point. To the right of the table is a 'Font Settings' section with a 'Font' dropdown set to 'Arial' and a 'Size' input field set to '8'. At the bottom of the dialog are 'OK' and 'Cancel' buttons.

Show	Column
<input checked="" type="checkbox"/>	Dye/Sample Peak
<input checked="" type="checkbox"/>	Sample File Name
<input checked="" type="checkbox"/>	Marker
<input checked="" type="checkbox"/>	Allele
<input checked="" type="checkbox"/>	Size
<input checked="" type="checkbox"/>	Height
<input checked="" type="checkbox"/>	Area
<input checked="" type="checkbox"/>	Data Point

### Print – MiniFiler Allelic Ladder: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section contains four 'Label' dropdown menus: Label 1 is set to 'Allele Call', Label 2 is set to 'Size', Label 3 is set to 'None', and Label 4 is set to 'None'. Below these are 'Font' and 'Size' dropdown menus, with 'Font' set to 'Times New Roman' and 'Size' set to '5'. The 'When opening the Plot Window' section contains three checkboxes: 'Show data type prefixes' (unchecked), 'Show type of edit' (unchecked), and 'Invert mutant labels' (unchecked). A 'Label Color' dropdown is set to 'Dye Color-Border'. At the bottom of the dialog are 'OK' and 'Cancel' buttons.

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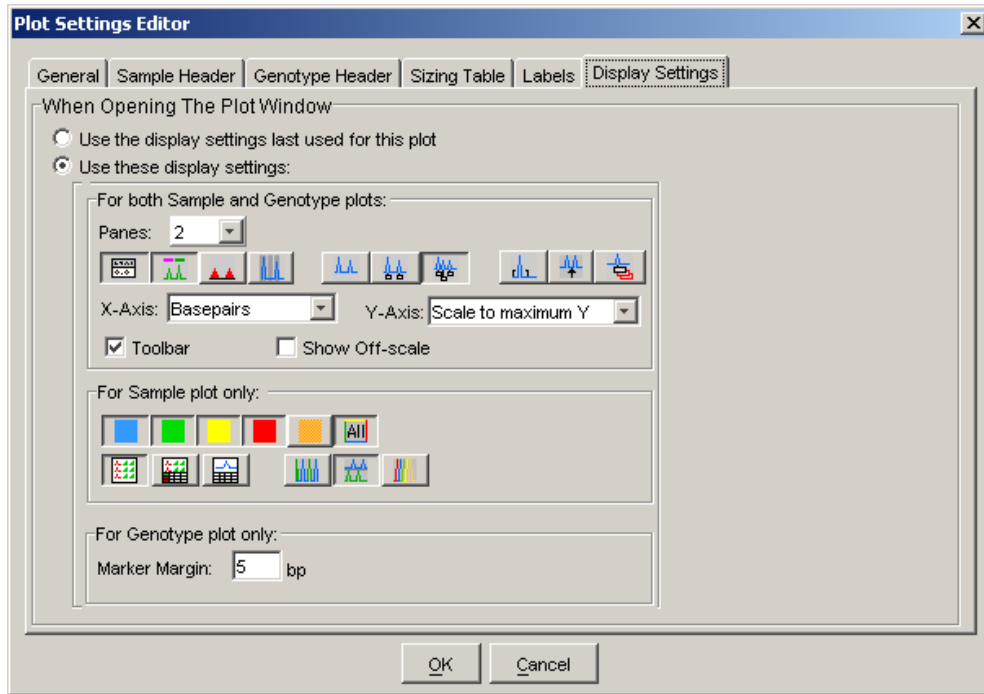
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### Print – MiniFiler Allelic Ladder: Display Settings



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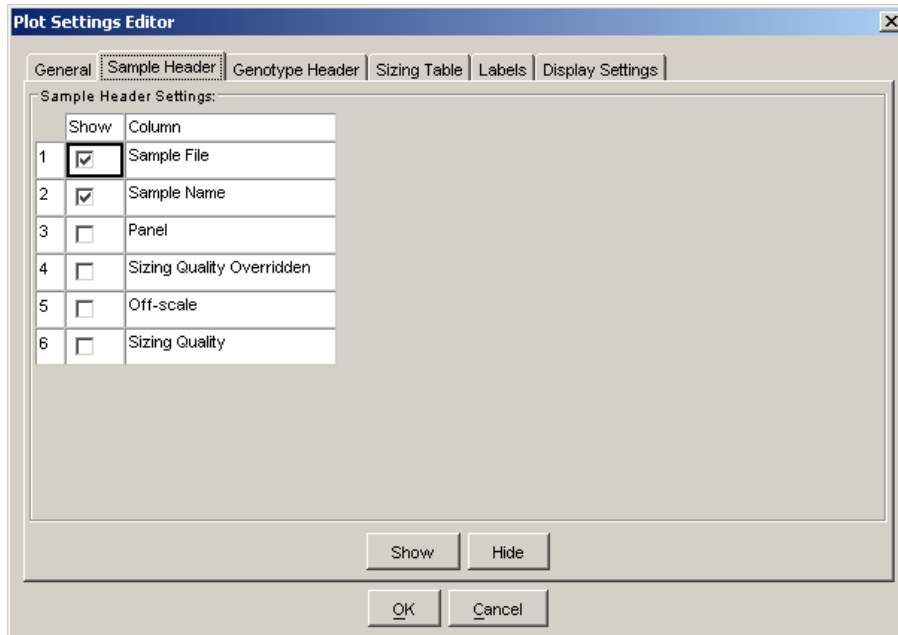
# FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

## GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

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### PLOT SETTINGS: PRINT – MINIFILER CONTROLS

#### Print – MiniFiler Controls: Sample Header

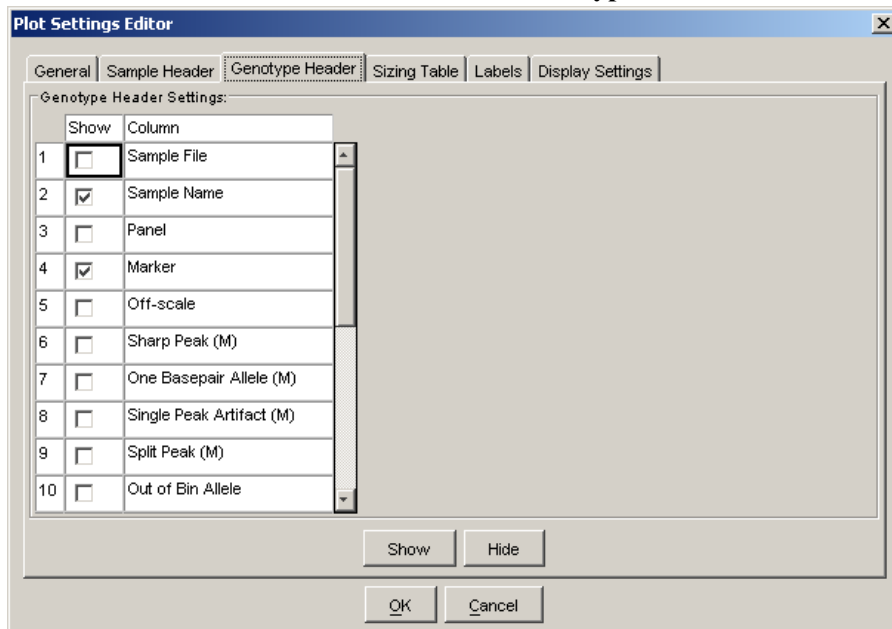


The screenshot shows the 'Plot Settings Editor' window with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

	Show	Column
1	<input checked="" type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input type="checkbox"/>	Sizing Quality Overridden
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sizing Quality

Buttons at the bottom: Show, Hide, OK, Cancel.

#### Print – MiniFiler Controls: Genotype Header



The screenshot shows the 'Plot Settings Editor' window with the 'Genotype Header' tab selected. The 'Genotype Header Settings' table is as follows:

	Show	Column
1	<input type="checkbox"/>	Sample File
2	<input checked="" type="checkbox"/>	Sample Name
3	<input type="checkbox"/>	Panel
4	<input checked="" type="checkbox"/>	Marker
5	<input type="checkbox"/>	Off-scale
6	<input type="checkbox"/>	Sharp Peak (M)
7	<input type="checkbox"/>	One Basepair Allele (M)
8	<input type="checkbox"/>	Single Peak Artifact (M)
9	<input type="checkbox"/>	Split Peak (M)
10	<input type="checkbox"/>	Out of Bin Allele

Buttons at the bottom: Show, Hide, OK, Cancel.

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### Print – MiniFiler Controls: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a table with 8 rows and 2 columns: 'Show' and 'Column'. All 'Show' checkboxes are checked. The 'Font Settings' section shows 'Font' set to 'Arial' and 'Size' set to '8'. 'Show' and 'Hide' buttons are at the bottom of the table, and 'OK' and 'Cancel' buttons are at the bottom of the dialog.

Show	Column
<input checked="" type="checkbox"/>	Dye/Sample Peak
<input checked="" type="checkbox"/>	Sample File Name
<input checked="" type="checkbox"/>	Marker
<input checked="" type="checkbox"/>	Allele
<input checked="" type="checkbox"/>	Size
<input checked="" type="checkbox"/>	Height
<input checked="" type="checkbox"/>	Area
<input checked="" type="checkbox"/>	Data Point

### Print – MiniFiler Controls: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section has four dropdown menus: 'Label 1' (Allele Call), 'Label 2' (Size), 'Label 3' (Height), and 'Label 4' (None). The 'Font' is set to 'Times New Roman' and 'Size' is '5'. The 'When opening the Plot Window' section has three unchecked checkboxes: 'Show data type prefixes', 'Show type of edit', and 'Invert mutant labels'. The 'Label Color' is set to 'Dye Color-Border'. 'OK' and 'Cancel' buttons are at the bottom.

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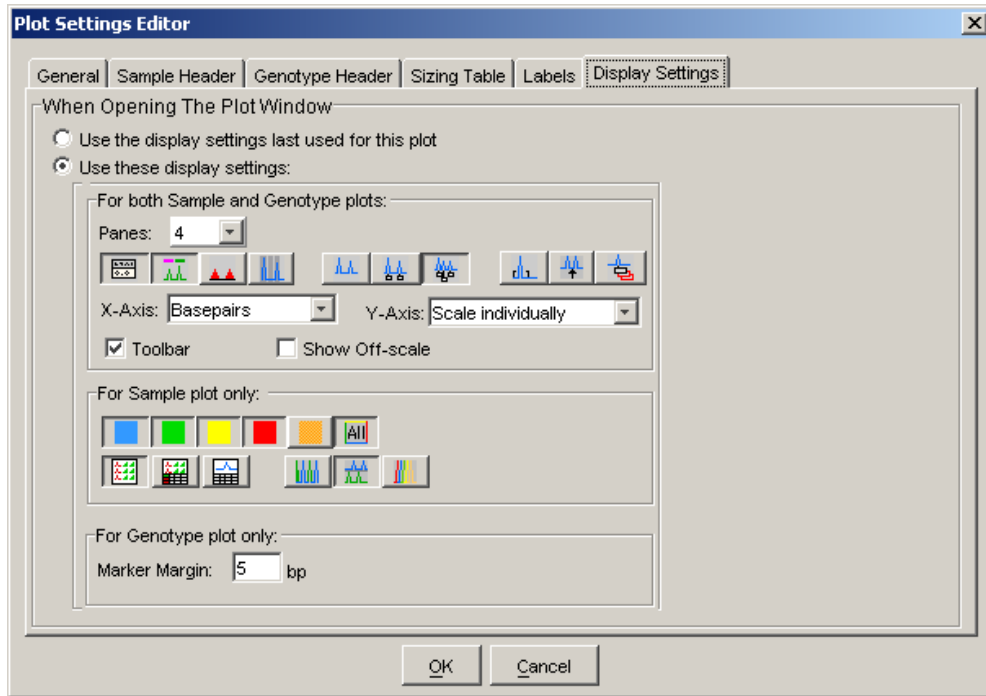
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### Print – MiniFiler Controls: Display Settings



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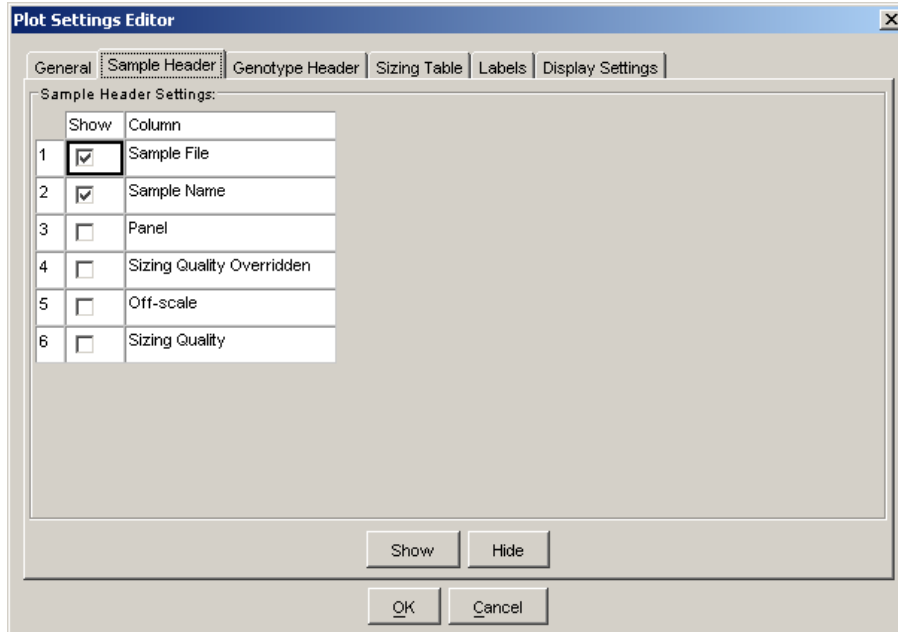
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### PLOT SETTINGS: PRINT – MINIFILER SAMPLES

#### Print – MiniFiler Samples: Sample Header



#### Print – MiniFiler Samples: Genotype Header

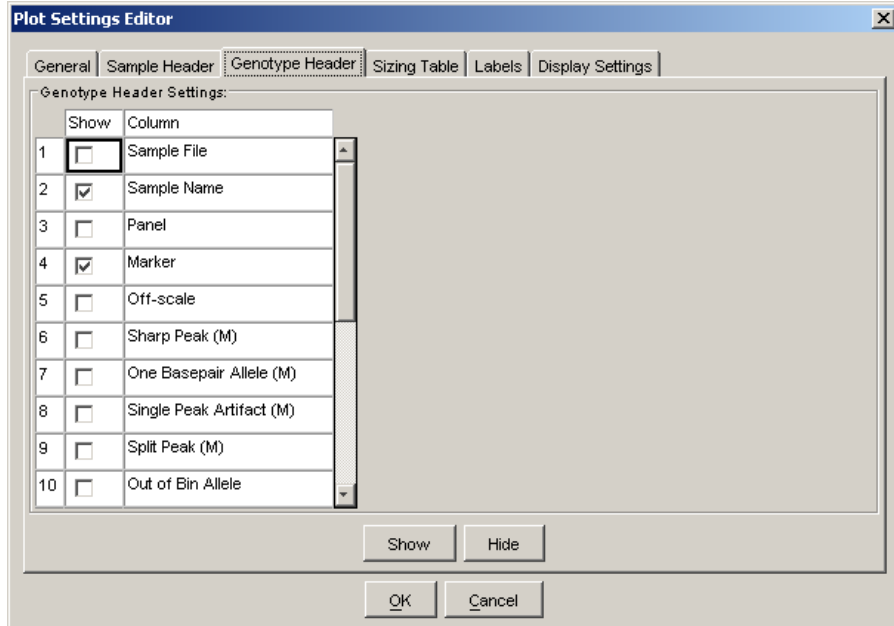
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### Print – MiniFiler Samples: Sizing Table

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Sizing Table' tab selected. The 'Sizing Table Settings' section contains a 'Column Settings' table and 'Font Settings'.

Show	Column
<input checked="" type="checkbox"/>	Dye/Sample Peak
<input checked="" type="checkbox"/>	Sample File Name
<input checked="" type="checkbox"/>	Marker
<input checked="" type="checkbox"/>	Allele
<input checked="" type="checkbox"/>	Size
<input checked="" type="checkbox"/>	Height
<input checked="" type="checkbox"/>	Area
<input checked="" type="checkbox"/>	Data Point

Font Settings:  
Font: Arial  
Size: 8

Buttons: Show, Hide, OK, Cancel

### Print – MiniFiler Samples: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section has four dropdown menus and font settings. The 'When opening the Plot Window' section has three checkboxes and a label color dropdown.

Show Labels:  
Label 1: Allele Call  
Label 2: Size  
Label 3: Height  
Label 4: None  
Font: Times New Roman  
Size: 5

When opening the Plot Window:  
 Show data type prefixes  
 Show type of edit  
 Invert mutant labels  
Label Color: Dye Color-Border

Buttons: OK, Cancel

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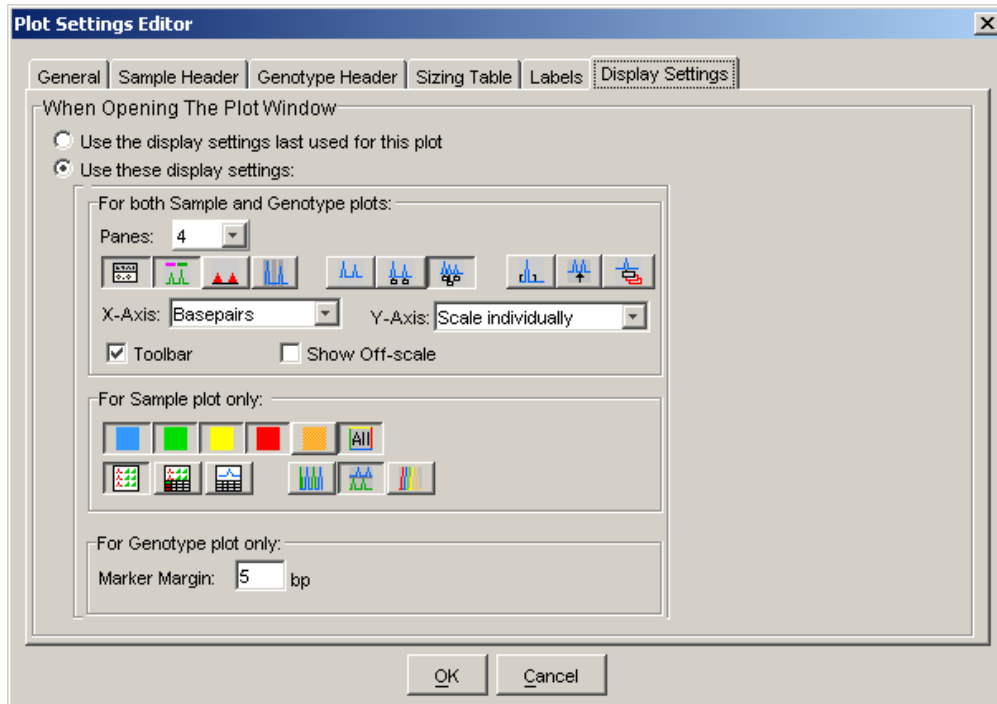
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