

FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 1 OF 70
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Default Table and Plot Settings

TABLE SETTINGS – ANALYSIS VIEW: SAMPLES SETTINGS

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 2 OF 70
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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 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

Show Hide

OK Cancel

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 3 OF 70
------------------------------	---	-----------------

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

Font Settings:

Font: Arial

Size: 11

Show Hide

Allele Settings

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

OK Cancel

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 4 OF 70
------------------------------	---	-----------------

TABLE SETTINGS – ANALYSIS VIEW: GENOTYPES SETTINGS (continued)

Table Setting Editor

General | Samples | **Genotypes**

Genotypes Table Settings:

Column Settings:

	Show	Column	Filtering	Content
20	<input type="checkbox"/>	Sharp Peak (M)	Show All Records	N/A
21	<input type="checkbox"/>	One Basepair Allele (M)	Show All Records	N/A
22	<input type="checkbox"/>	Single Peak Artifact (M)	Show All Records	N/A
23	<input type="checkbox"/>	Split Peak (M)	Show All Records	N/A
24	<input checked="" type="checkbox"/>	Out of Bin Allele	Show All Records	N/A
25	<input checked="" type="checkbox"/>	Peak Height Ratio	Show All Records	N/A
26	<input type="checkbox"/>	Low Peak Height	Show All Records	N/A
27	<input checked="" type="checkbox"/>	Spectral Pull-up	Show All Records	N/A
28	<input checked="" type="checkbox"/>	Allele Number	Show All Records	N/A
29	<input checked="" type="checkbox"/>	Broad Peak	Show All Records	N/A
30	<input type="checkbox"/>	Double Peak (SNP)	Show All Records	N/A
31	<input type="checkbox"/>	Narrow Bin (SNP)	Show All Records	N/A
32	<input checked="" type="checkbox"/>	Control Concordance	Show All Records	N/A
33	<input checked="" type="checkbox"/>	Overlap (HID)	Show All Records	N/A
34	<input type="checkbox"/>	Cross Talk	Show All Records	N/A
35	<input type="checkbox"/>	Genotype Quality	Show All Records	N/A
36	<input type="checkbox"/>	User Defined Column 1	Show All Records	
37	<input type="checkbox"/>	User Defined Column 2	Show All Records	
38	<input type="checkbox"/>	User Defined Column 3	Show All Records	

Font Settings:

Font: Arial

Size: 11

Show Hide

Allele Settings

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

OK Cancel

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 5 OF 70
------------------------------	---	-----------------

TABLE SETTINGS – CASEWORK VIEW: SAMPLES SETTINGS

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 6 OF 70
------------------------------	---	-----------------

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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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 7 OF 70
------------------------------	---	-----------------

TABLE SETTINGS – CASEWORK 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 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 checked="" 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 checked="" type="checkbox"/>	AE Comment	Show All Records	
15	<input type="checkbox"/>	Integration Comments	Show All Records	
16	<input 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 type="checkbox"/>	Off-scale	Show All Records	N/A

Font Settings:

Font: Arial

Size: 11

Allele Settings

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

Show Hide

OK Cancel

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 8 OF 70
------------------------------	---	-----------------

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 for column settings and font settings.

Column Settings:	Font Settings:																																																																																
<table border="1" style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th>Show</th> <th>Column</th> <th>Filtering</th> <th>Content</th> </tr> </thead> <tr><td><input type="checkbox"/></td><td>Sharp Peak (M)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>One Basepair Allele (M)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Single Peak Artifact (M)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Split Peak (M)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Out of Bin Allele</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Peak Height Ratio</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Low Peak Height</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Spectral Pull-up</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Allele Number</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Broad Peak</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Double Peak (SNP)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Narrow Bin (SNP)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Control Concordance</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Overlap (HID)</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Cross Talk</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>Genotype Quality</td><td>Show All Records</td><td>N/A</td></tr> <tr><td><input type="checkbox"/></td><td>User Defined Column 1</td><td>Show All Records</td><td></td></tr> <tr><td><input type="checkbox"/></td><td>User Defined Column 2</td><td>Show All Records</td><td></td></tr> <tr><td><input type="checkbox"/></td><td>User Defined Column 3</td><td>Show All Records</td><td></td></tr> </table>	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: Arial Size: 11
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																																																																															

Below the table settings, there are 'Show' and 'Hide' buttons. At the bottom of the window, the 'Allele Settings' section includes a text box for 'Number of Alleles' set to 15, a checkbox for 'Keep Allele, Size, Height, Area, Data Point, Mutation and Comment together' (which is unchecked), and 'OK' and 'Cancel' buttons.

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 9 OF 70
------------------------------	---	-----------------

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

Show Hide

OK Cancel

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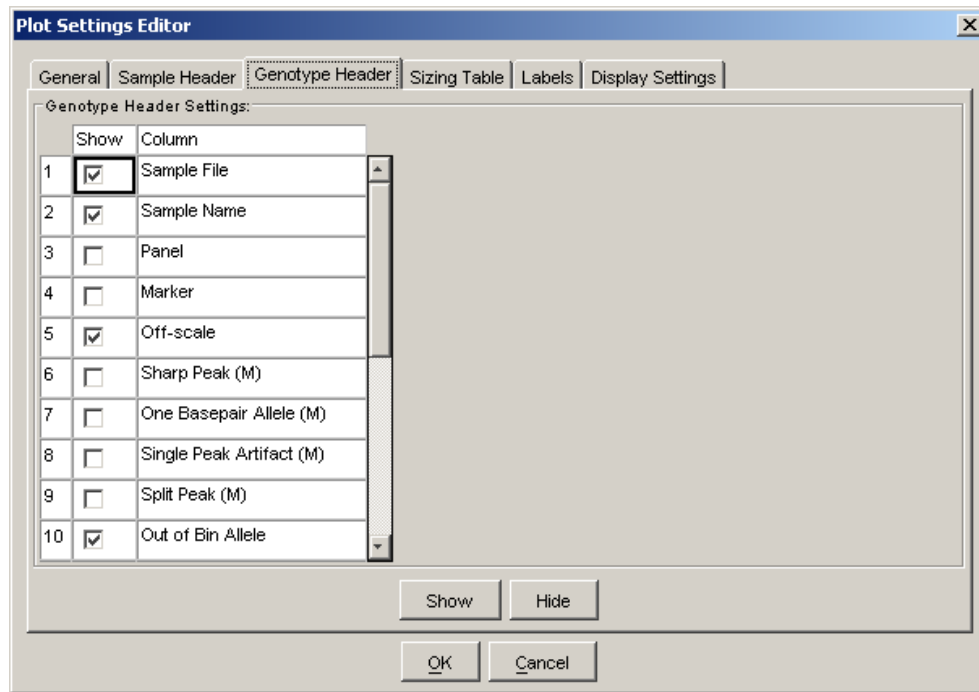
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 10 OF 70
------------------------------	---	------------------

Analysis View: Genotype Header



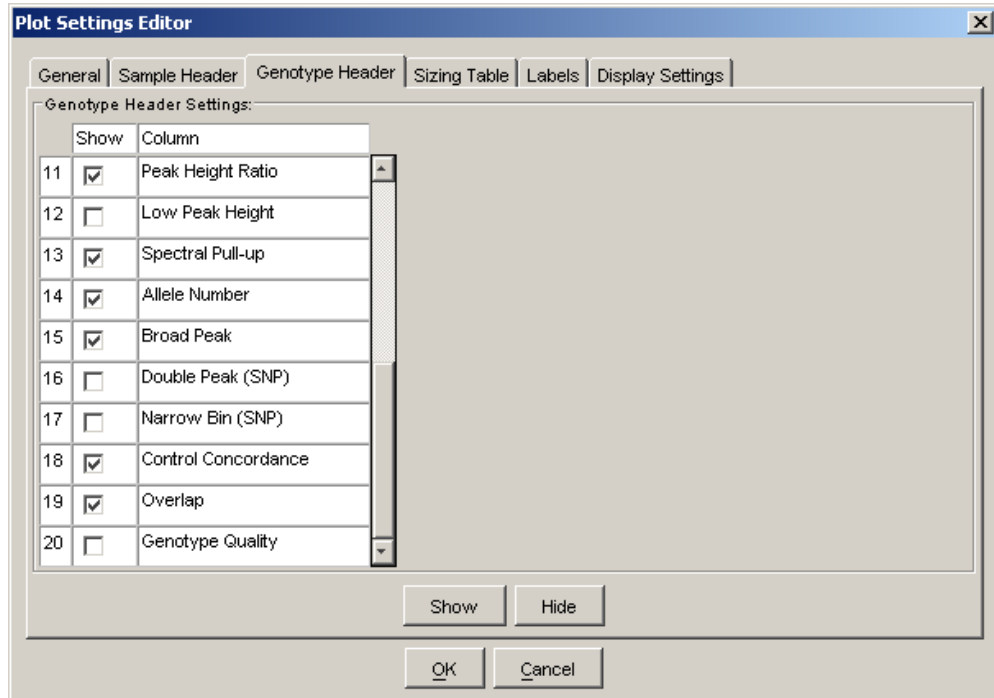
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 11 OF 70
------------------------------	---	------------------



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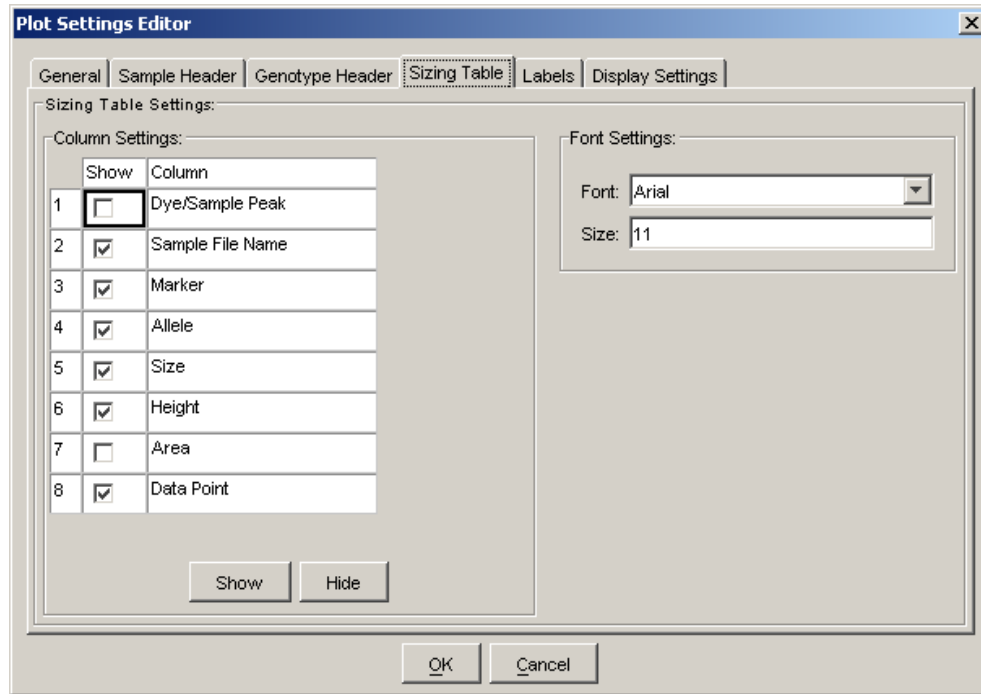
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 12 OF 70
------------------------------	---	------------------

Analysis View: Sizing Table



Analysis View: Labels

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FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 13 OF 70
------------------------------	---	------------------

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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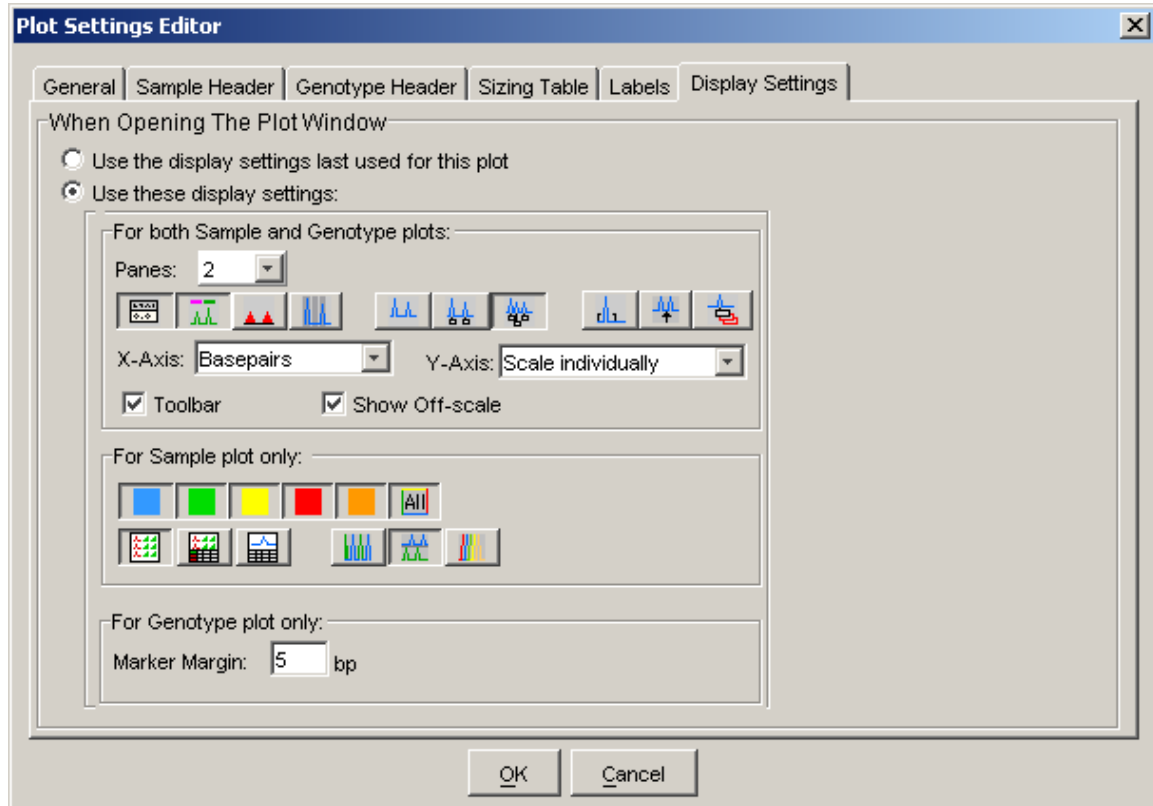
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 14 OF 70
------------------------------	---	------------------

Analysis View: Display Settings



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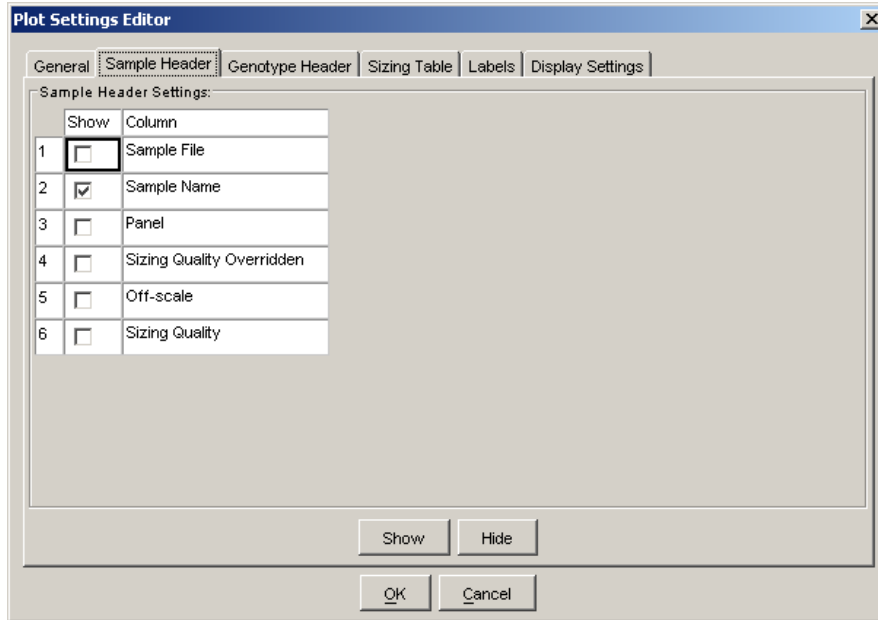
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 15 OF 70
------------------------------	---	------------------

PLOT SETTINGS: PRINT – IDENTIFILER ALLELIC LADDER

Print – Identifiler Allelic Ladder: Sample Header



Print – Identifiler Allelic Ladder: Genotype Header

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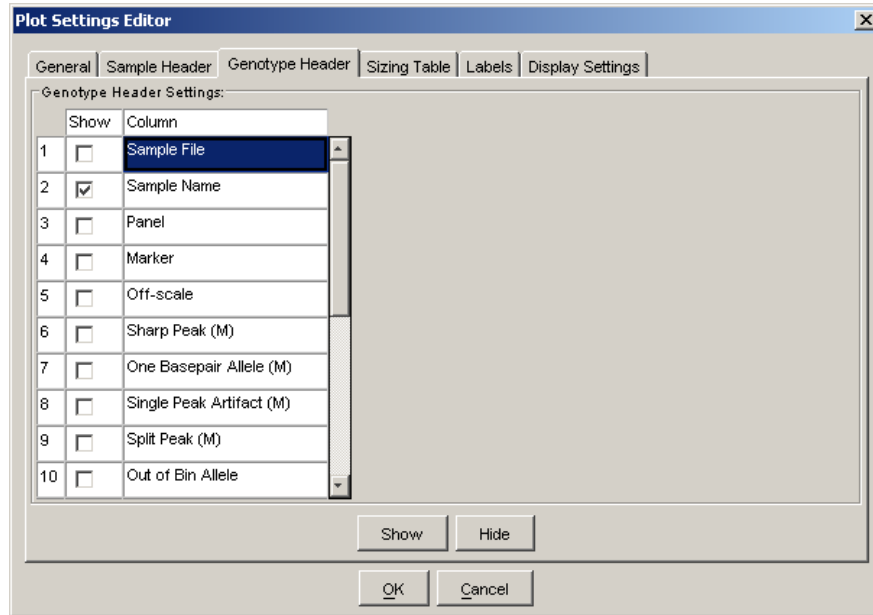
FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

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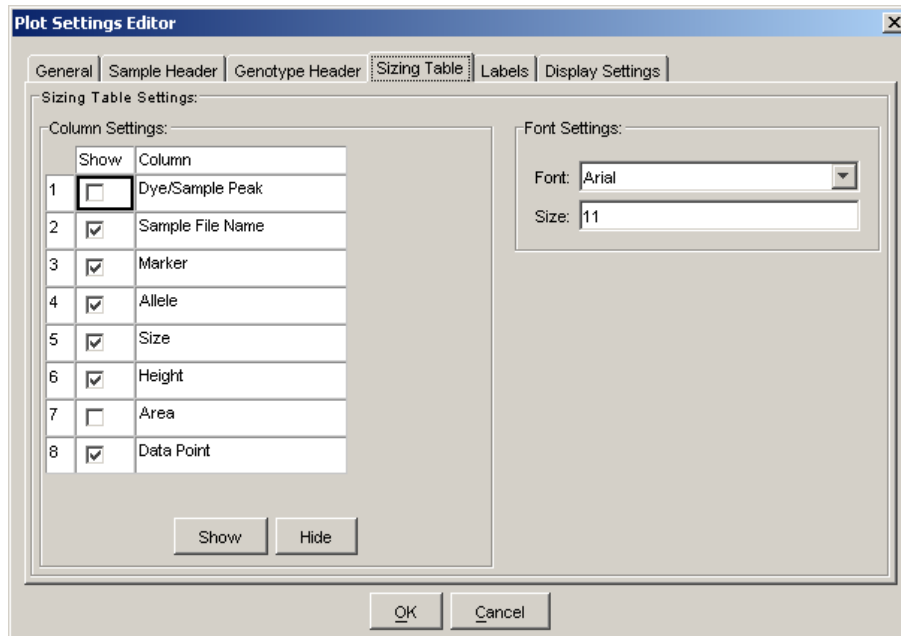
DATE EFFECTIVE
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PAGE
16 OF 70



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



Print – Identifiler Allelic Ladder: Labels

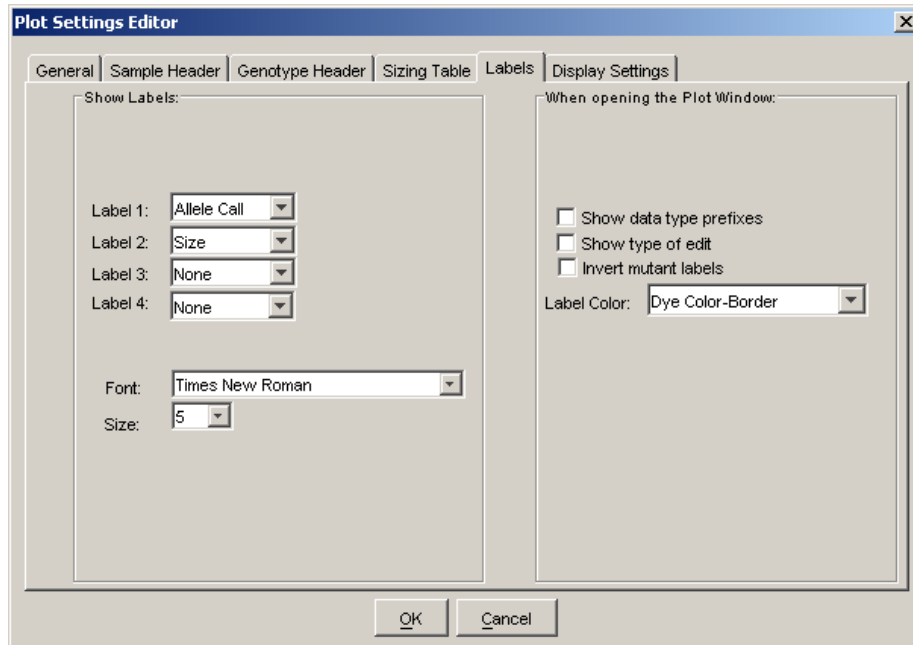
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 17 OF 70
------------------------------	---	------------------



Print – Identifier Allelic Ladder: Display Settings

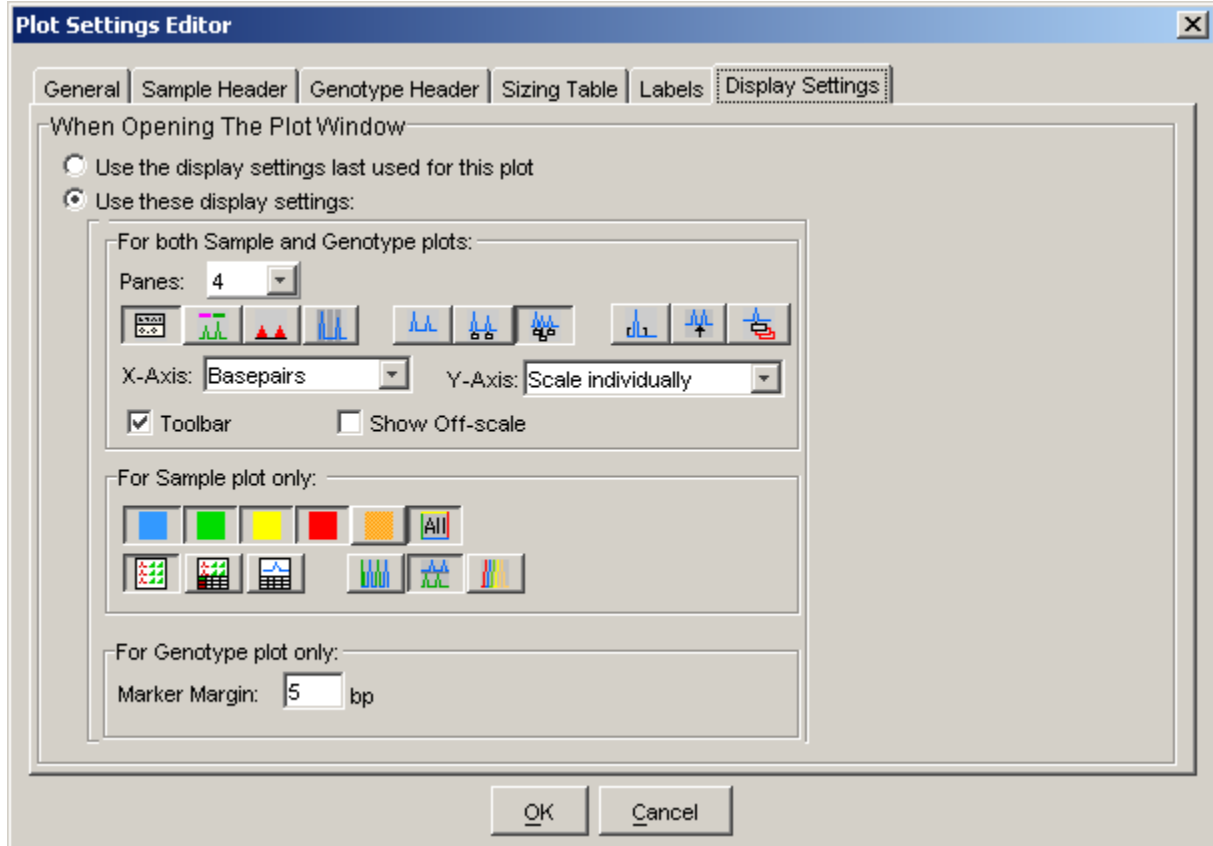
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 18 OF 70
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 19 OF 70
------------------------------	---	------------------

PLOT SETTINGS: PRINT – IDENTIFILER 28 CONTROLS

Print – Identifiler 28 Controls: 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 – Identifiler 28 Controls: Genotype 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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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 20 OF 70
------------------------------	---	------------------

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 21 OF 70
------------------------------	---	------------------

Print – Identifiler 28 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 '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 – Identifiler 28 Controls: Labels

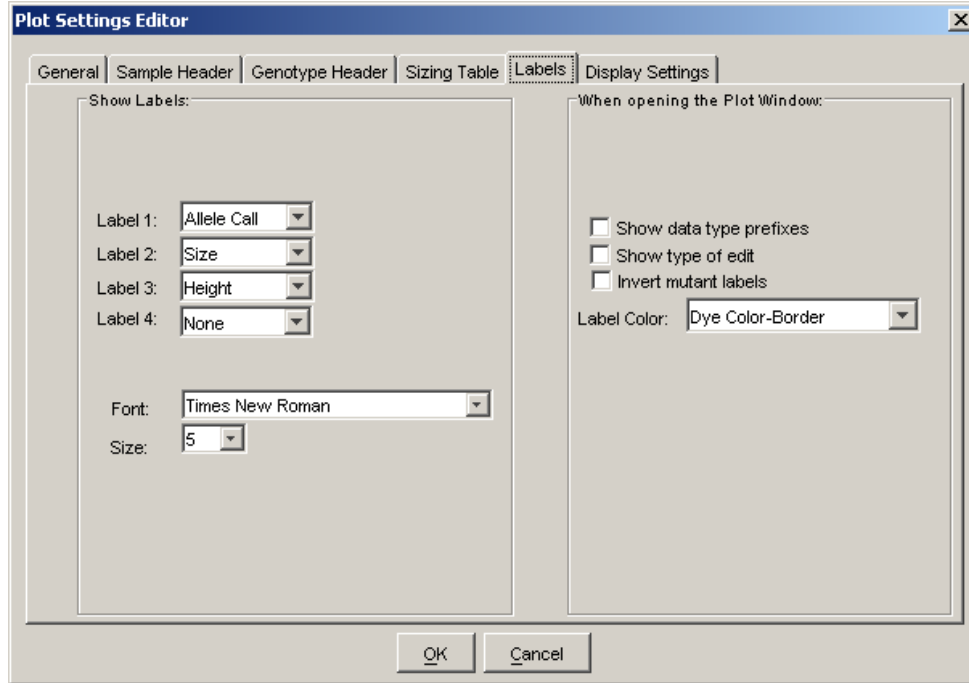
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 22 OF 70
------------------------------	---	------------------



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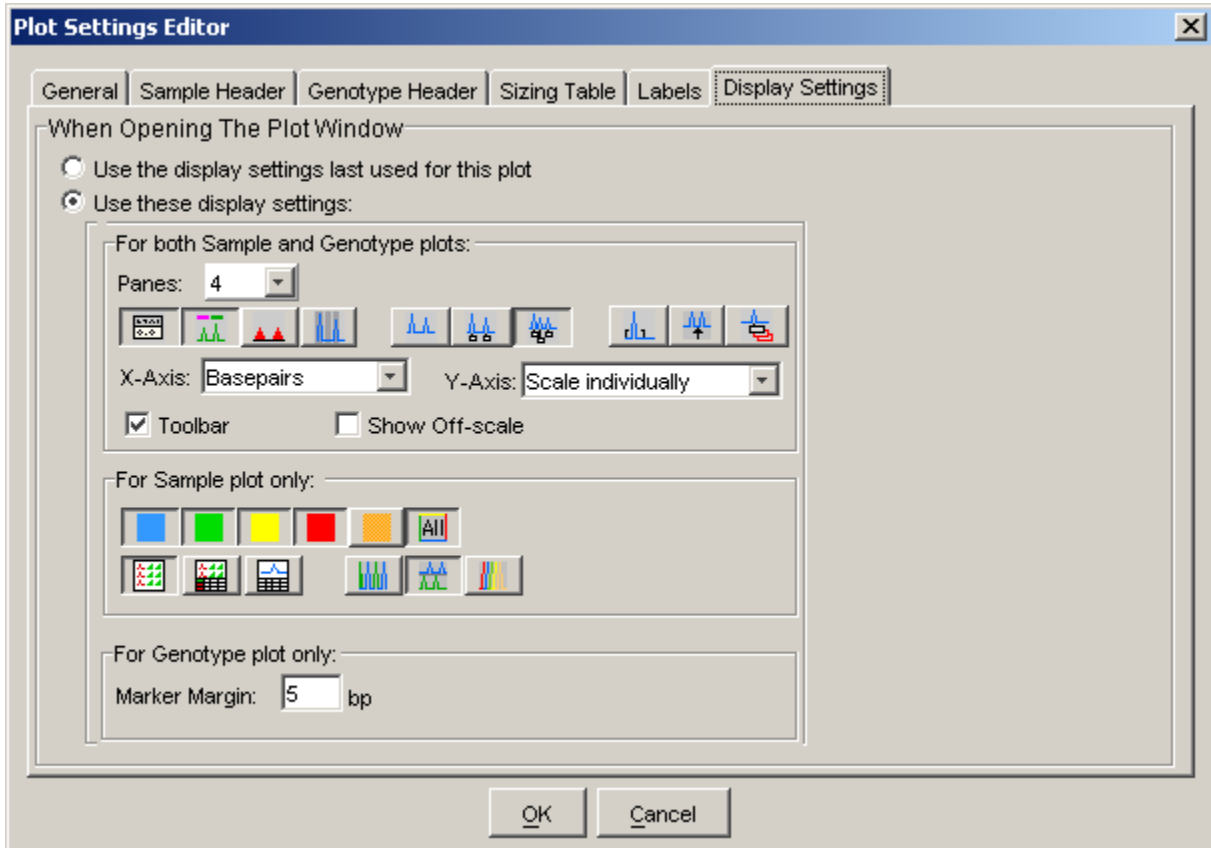
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 23 OF 70
------------------------------	---	------------------

Print – Identifiler 28 Controls: Display Settings



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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 24 OF 70
------------------------------	---	------------------

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 data:

Show	Column
<input 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

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

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 25 OF 70
------------------------------	---	------------------

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

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 26 OF 70
------------------------------	---	------------------

Print – Identifiler28 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 – Identifiler28 Samples: Labels

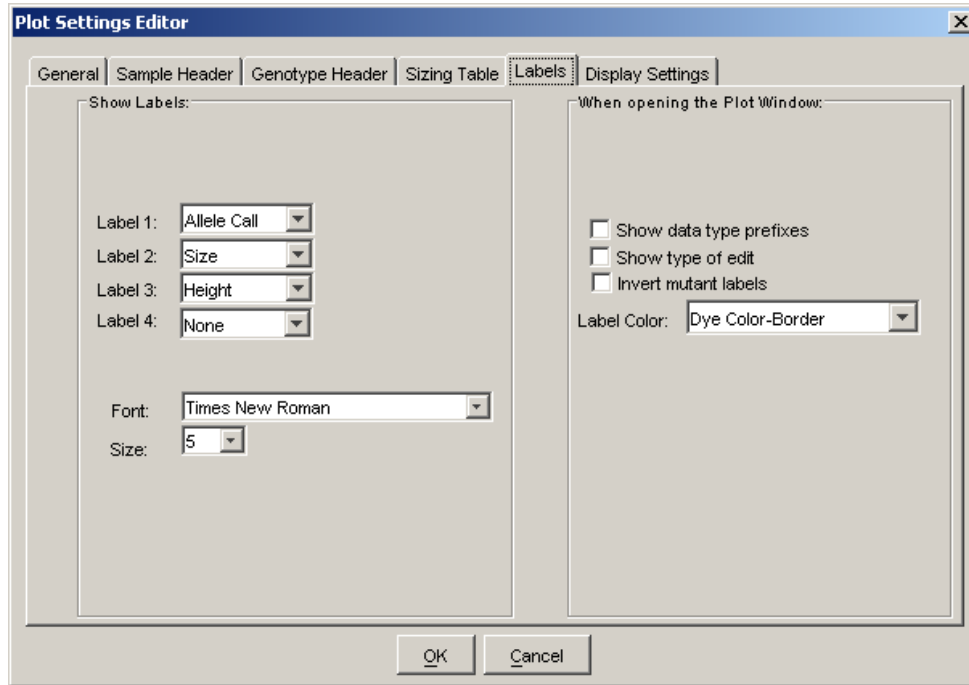
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 27 OF 70
------------------------------	---	------------------



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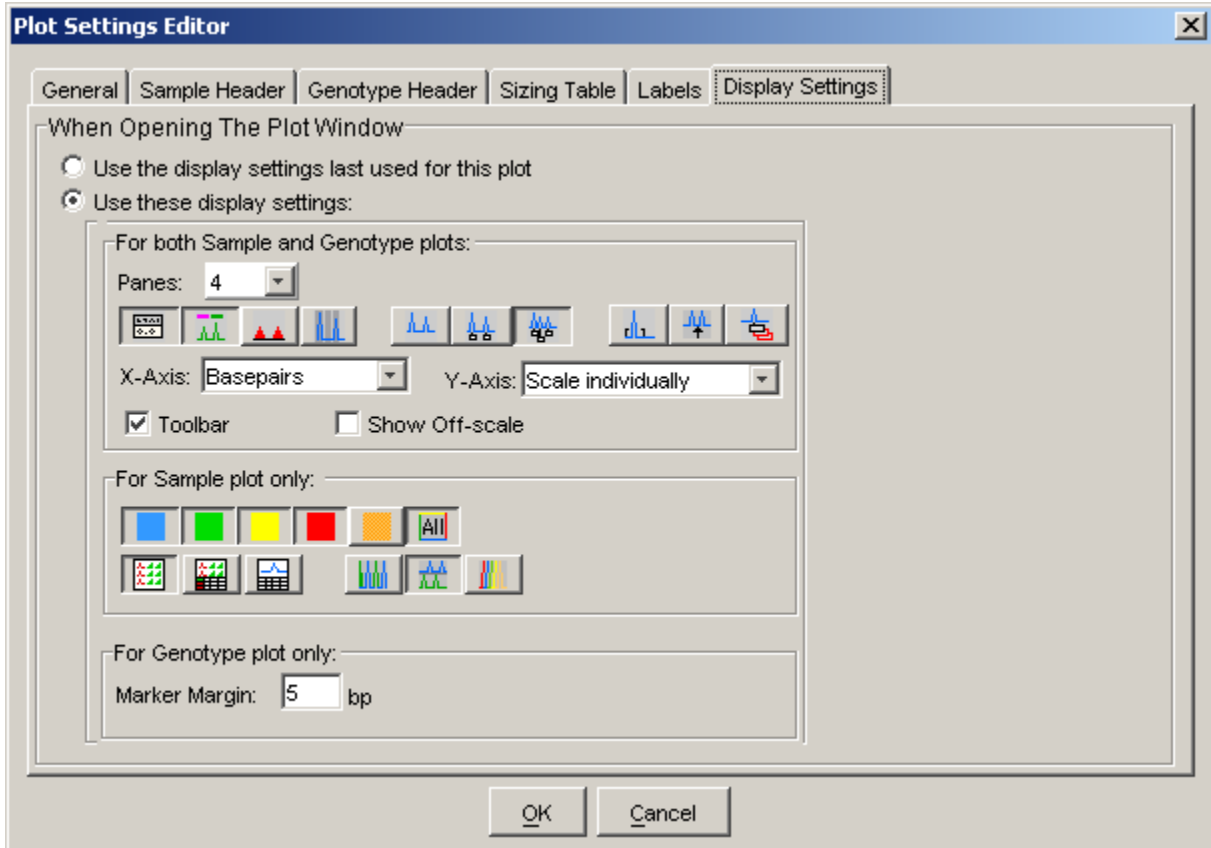
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 28 OF 70
------------------------------	---	------------------

Print – Identifiler28 Samples: Display Settings



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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 29 OF 70
------------------------------	---	------------------

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

Print – ID 31 PE and Samples: 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 data:

	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 box, there are buttons for 'Show', 'Hide', 'OK', and 'Cancel'.

Print – ID 31 PE and Samples: Genotype Header

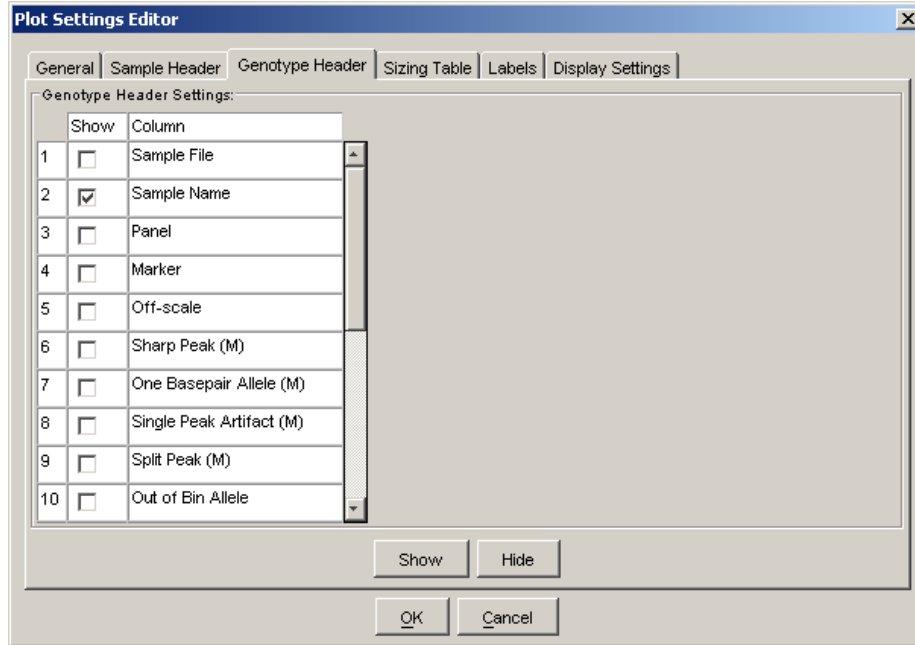
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 30 OF 70
------------------------------	---	------------------



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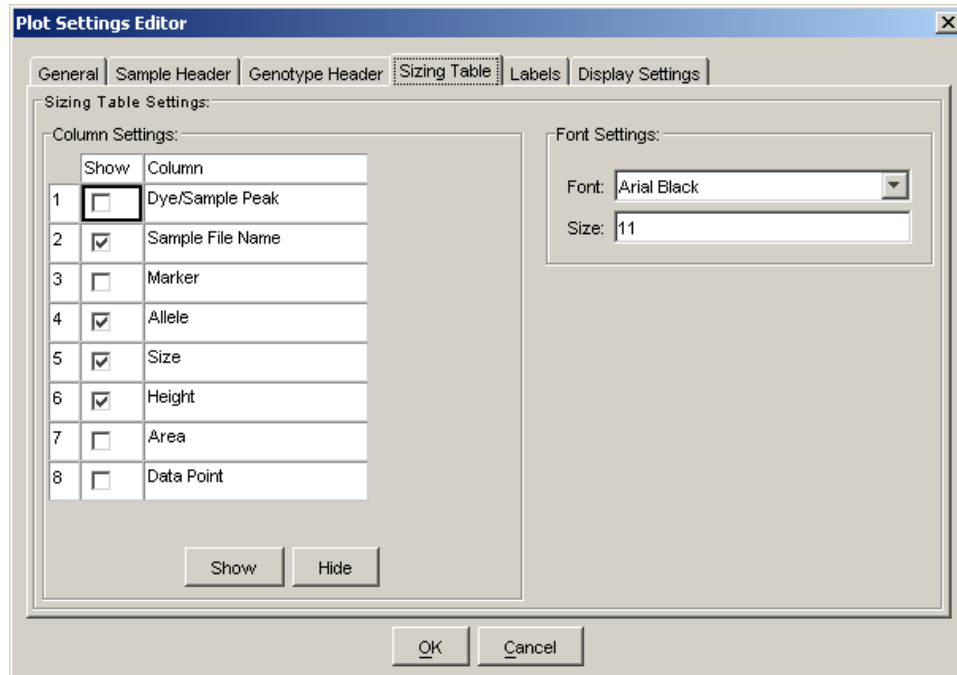
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 31 OF 70
------------------------------	---	------------------

Print – ID 31 PE and Samples: Sizing Table



Print – ID 31 PE and Samples: Labels

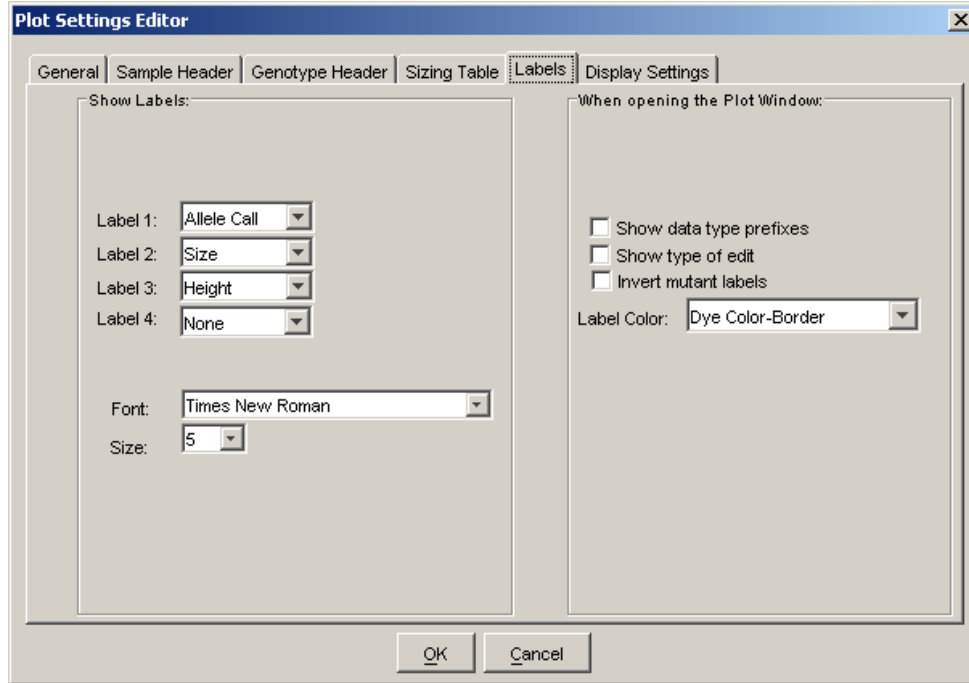
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 32 OF 70
------------------------------	---	------------------



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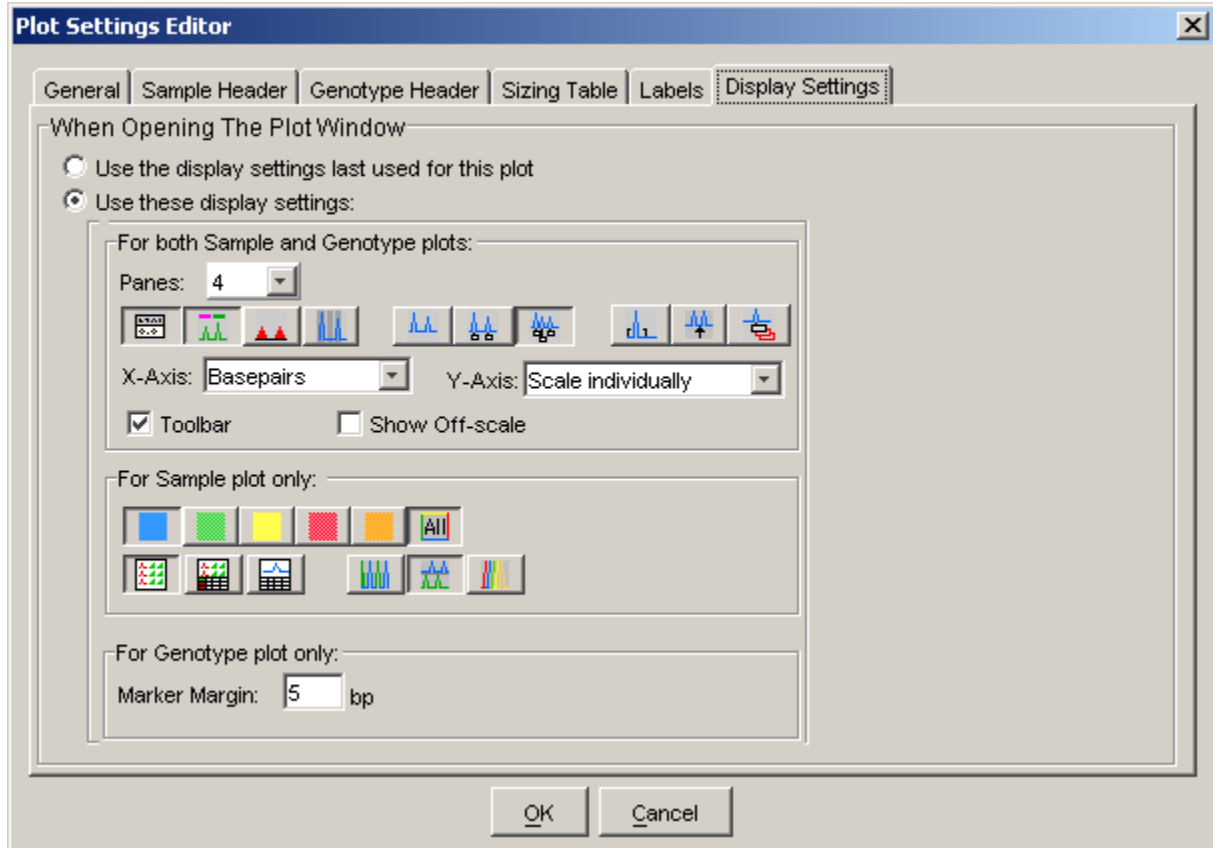
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 33 OF 70
------------------------------	---	------------------

Print – ID 31 PE and Samples: Display Settings



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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 34 OF 70
------------------------------	---	------------------

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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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 35 OF 70
------------------------------	---	------------------

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 36 OF 70
------------------------------	---	------------------

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

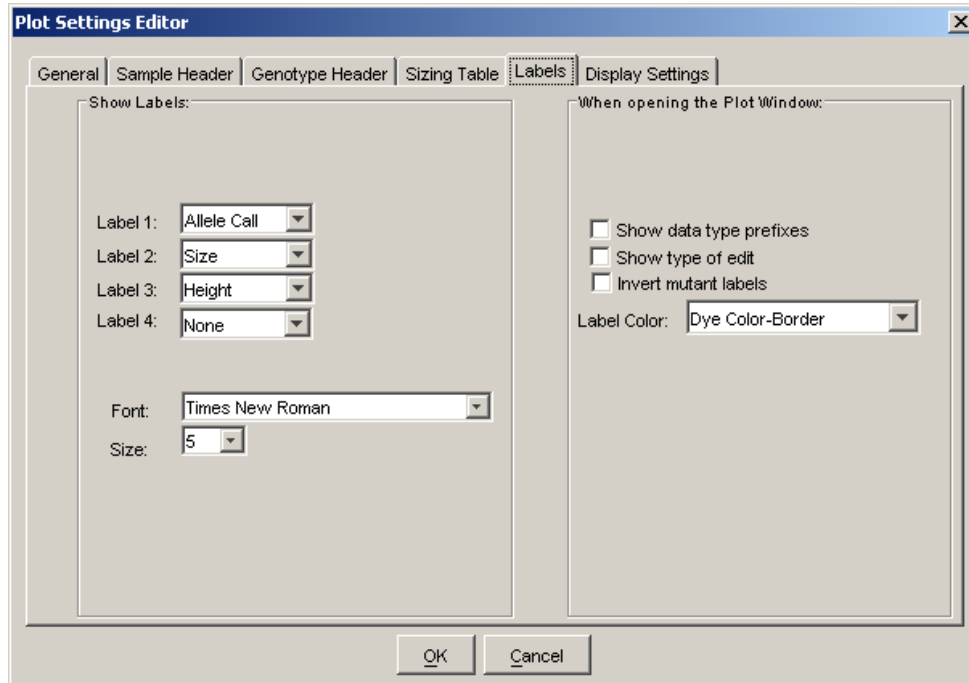
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 37 OF 70
------------------------------	---	------------------



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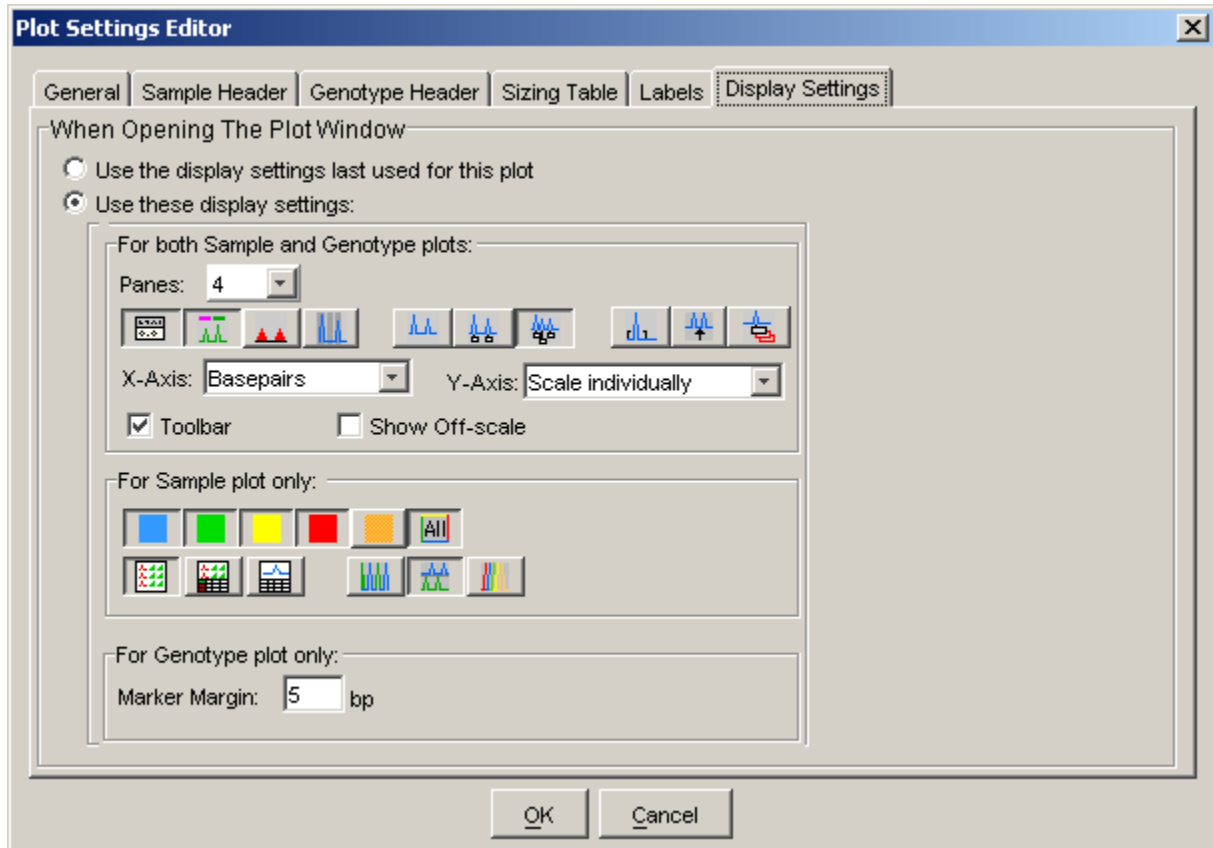
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 38 OF 70
------------------------------	---	------------------

Print – ID 31 Negative Controls: Display Settings



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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 39 OF 70
------------------------------	---	------------------

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' 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 Allelic Ladder: Genotypes 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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------------------------------	---	------------------

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 41 OF 70
------------------------------	---	------------------

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 table for column settings 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 settings and font settings.

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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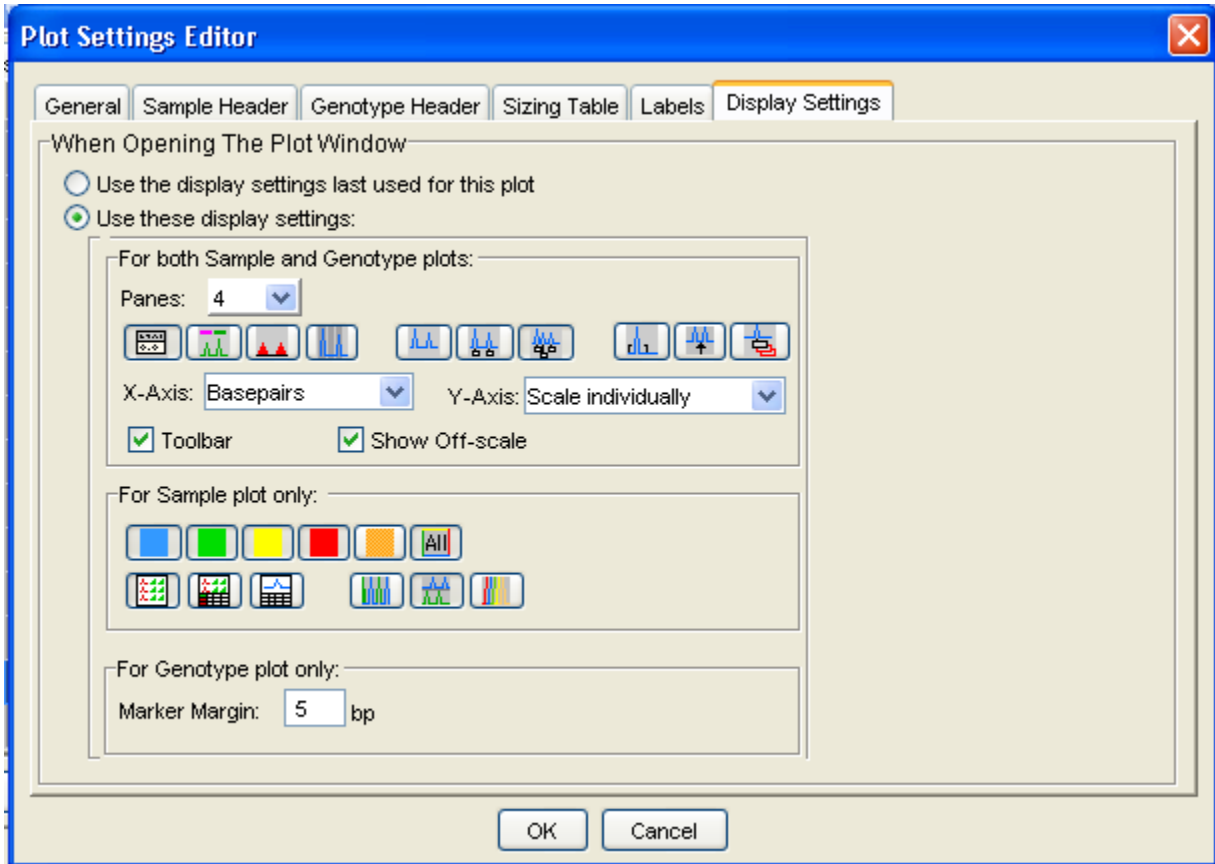
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 42 OF 70
------------------------------	---	------------------

Print – Yfiler Allelic Ladder: Display Settings



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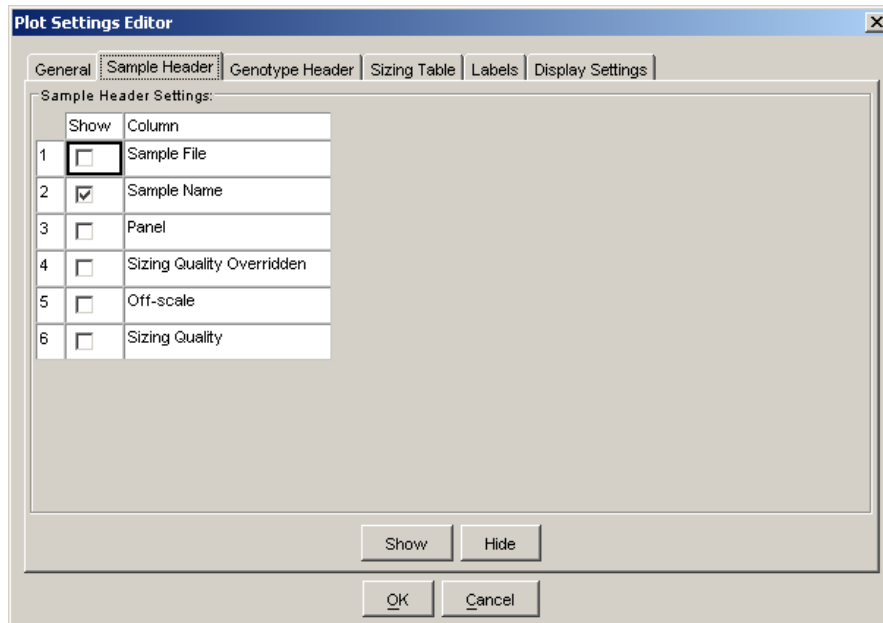
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 43 OF 70
------------------------------	---	------------------

PLOT SETTINGS: PRINT – YFILER CONTROLS

Print – YFiler Controls: Sample Header



Print – YFiler Controls: Genotypes Header

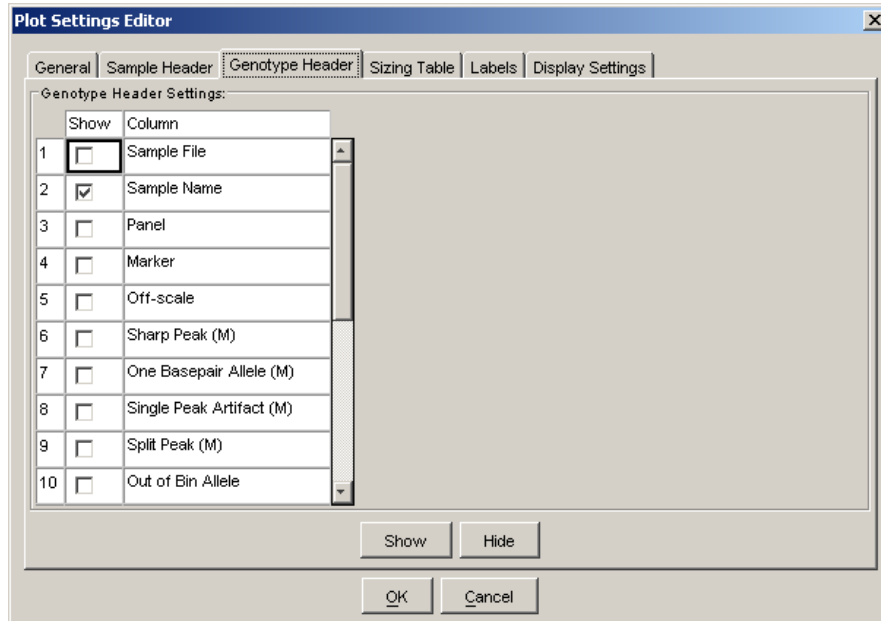
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 44 OF 70
------------------------------	---	------------------



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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 settings 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 contains dropdown menus for label types and font settings.

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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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 46 OF 70
------------------------------	---	------------------

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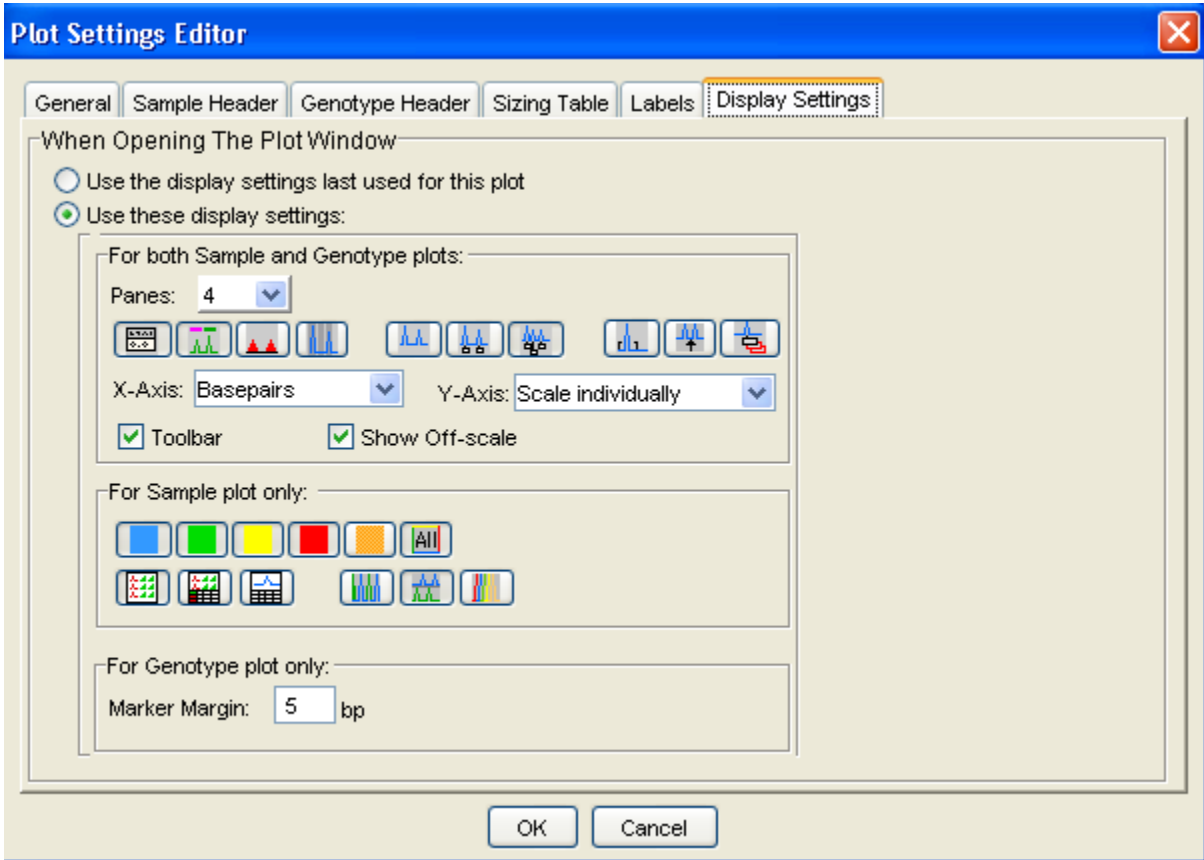
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 47 OF 70
------------------------------	---	------------------

Print – Yfiler Controls: Display Settings



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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 48 OF 70
------------------------------	---	------------------

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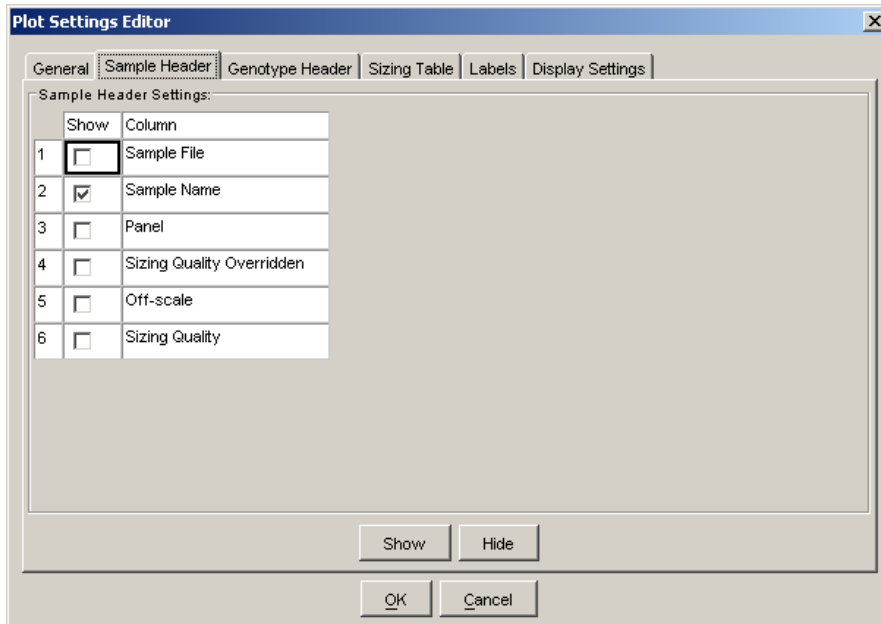
FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR TYPING

GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 49 OF 70
------------------------------	---	------------------

PLOT SETTINGS: PRINT – YFiler SAMPLES

Print – YFiler Samples: Sample Header

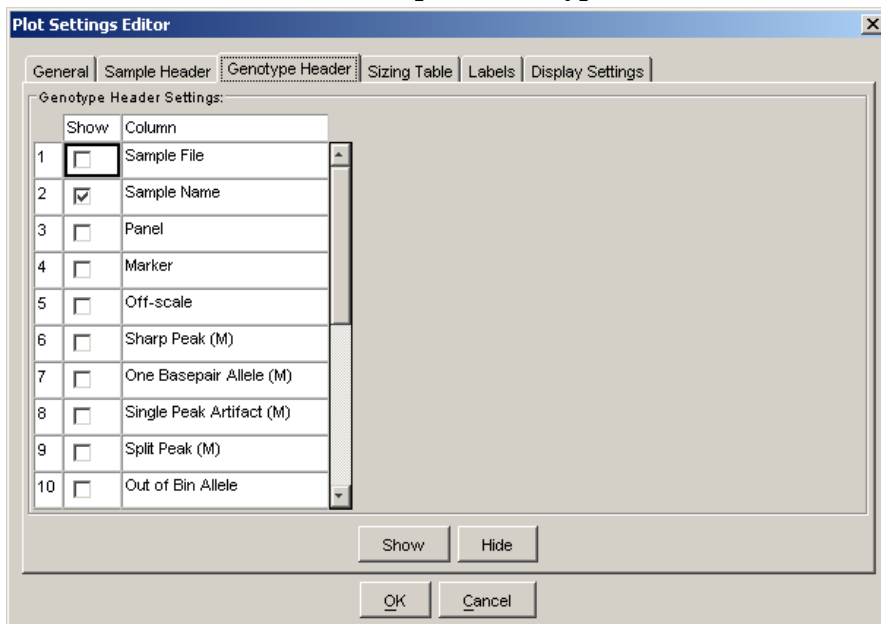


The screenshot shows the 'Plot Settings Editor' window with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

	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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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 50 OF 70
------------------------------	---	------------------

Boxes 3 – 20 are unchecked

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 51 OF 70
------------------------------	---	------------------

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 table for column settings 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 labels 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

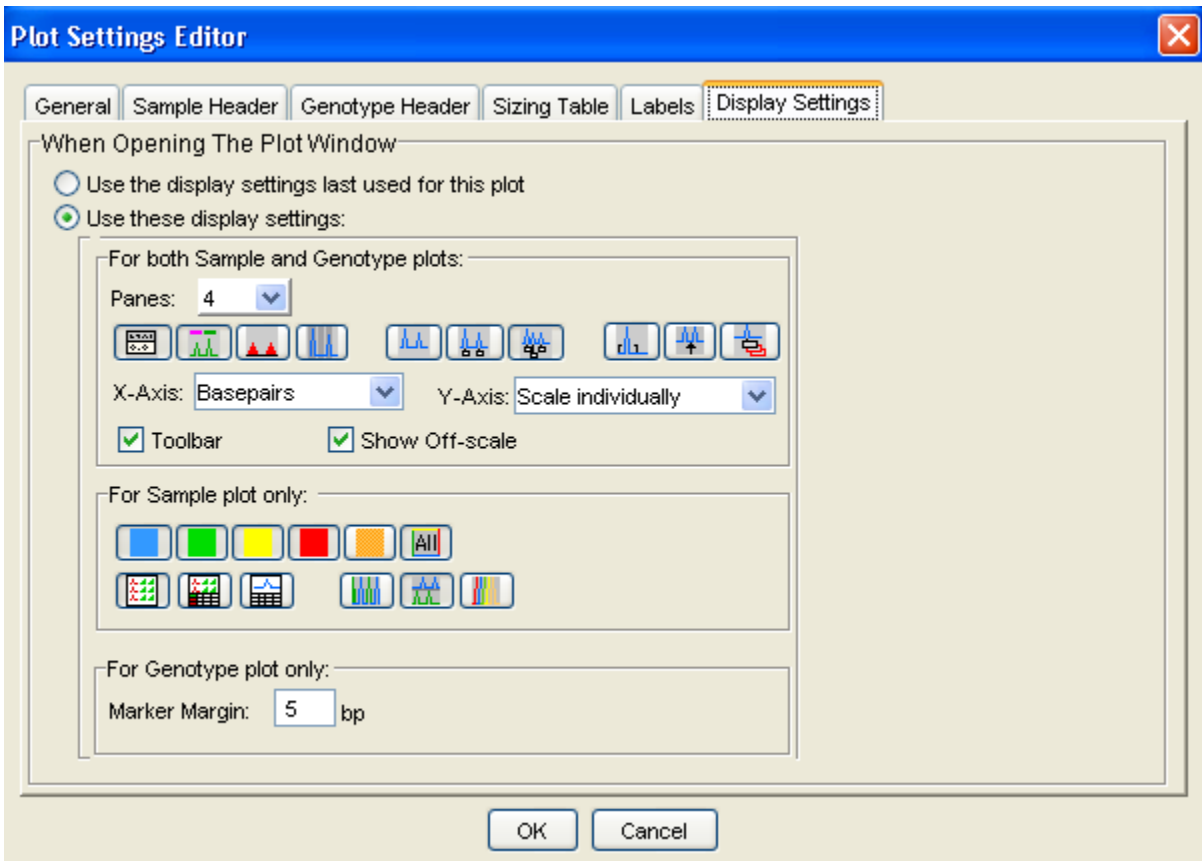
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 52 OF 70
------------------------------	---	------------------

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 53 OF 70
------------------------------	---	------------------

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 has a title bar with a close button (X) and a tabbed interface with the following tabs: 'General', 'Sample Header' (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 are two buttons: 'OK' and 'Cancel'.

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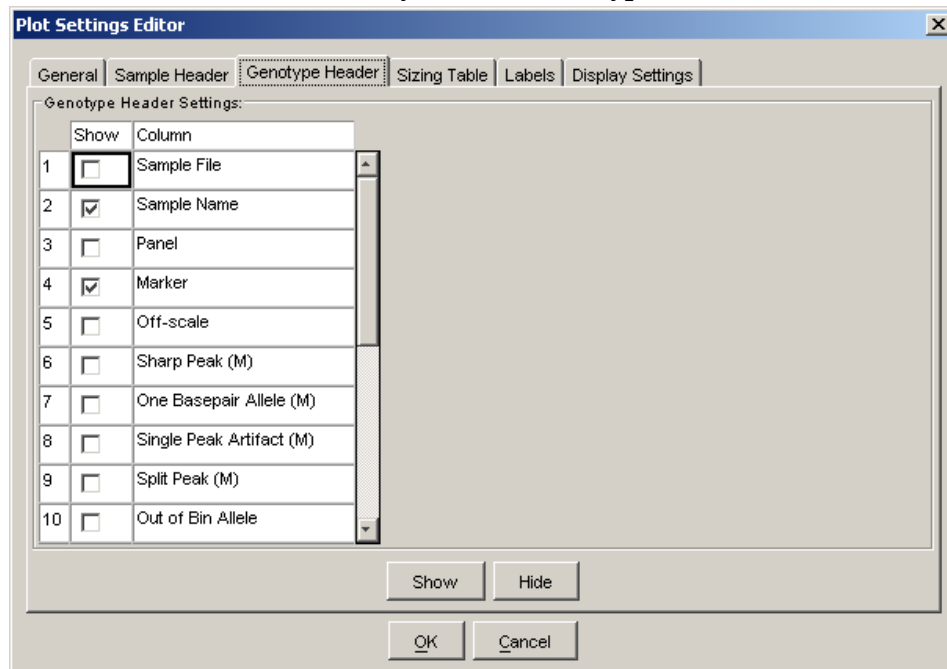
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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 54 OF 70
------------------------------	---	------------------

MiniFiler Analysis View: Genotype Header



5-20 are blank

MiniFiler Analysis View: Sizing Table

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 55 OF 70
------------------------------	---	------------------

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

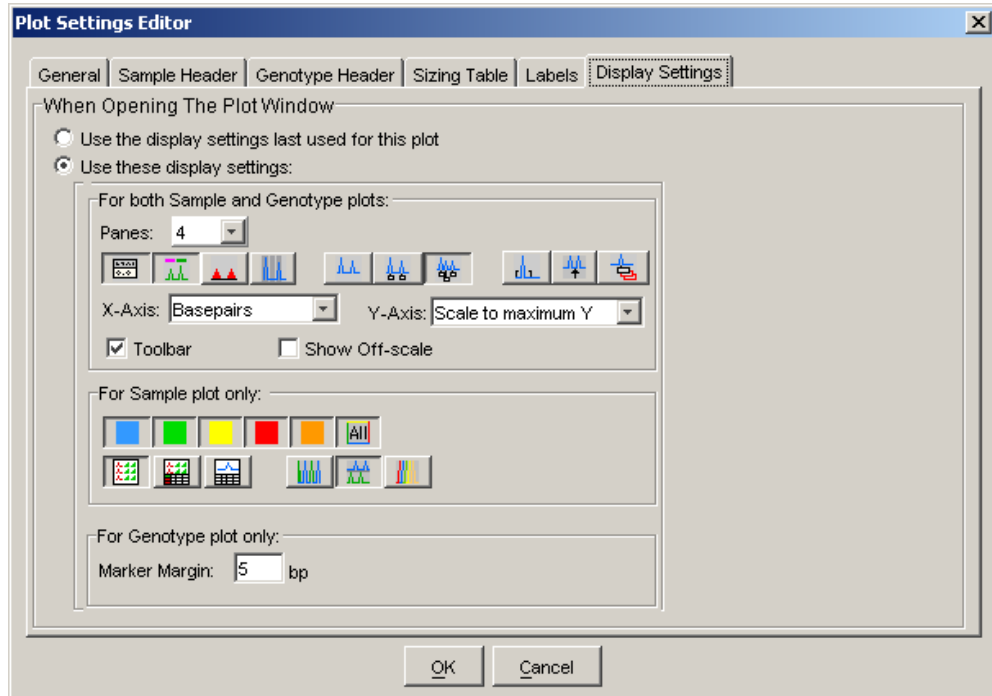
MiniFiler Analysis View: Display Settings

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 56 OF 70
------------------------------	---	------------------



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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 57 OF 70
------------------------------	---	------------------

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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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 58 OF 70
------------------------------	---	------------------

5-20 are blank

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 59 OF 70
------------------------------	---	------------------

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. All columns are checked in the 'Show' column. The 'Font Settings' section shows 'Font: Arial' and 'Size: 8'. There are 'Show' and 'Hide' buttons at the bottom of the table settings, and 'OK' and 'Cancel' buttons 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 Allelic Ladder: Labels

The screenshot shows the 'Plot Settings Editor' dialog box with the 'Labels' tab selected. The 'Show Labels' section has four labels: Label 1 (Allele Call), Label 2 (Size), Label 3 (None), and Label 4 (None). The font is set to 'Times New Roman' and size to '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'. There are 'OK' and 'Cancel' buttons at the bottom of the dialog.

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 60 OF 70
------------------------------	---	------------------

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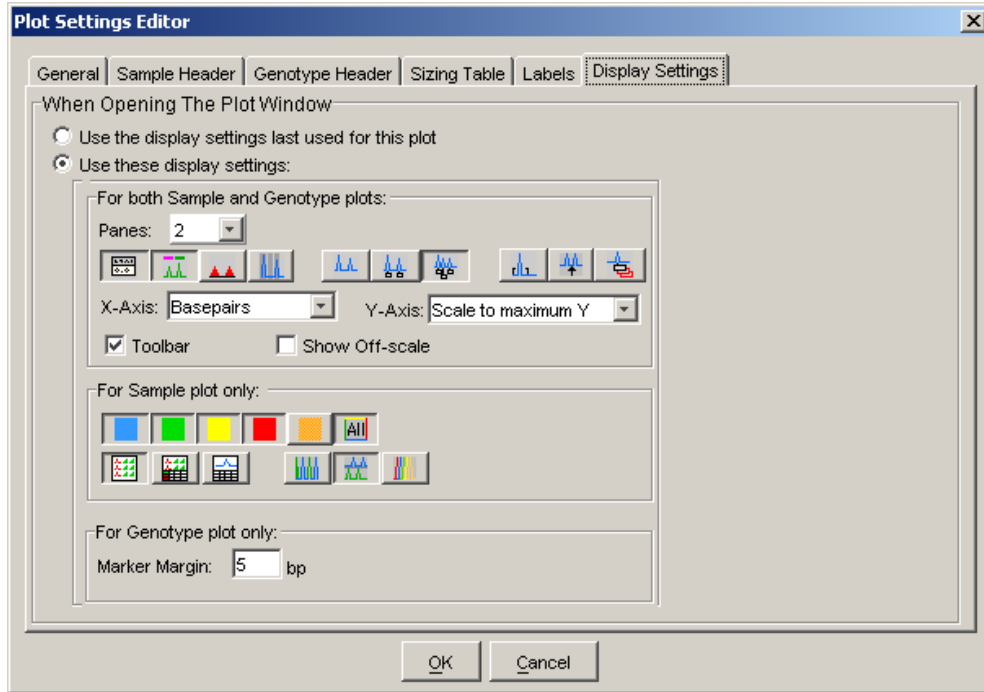
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 61 OF 70
------------------------------	---	------------------

Print – MiniFiler Allelic Ladder: Display Settings



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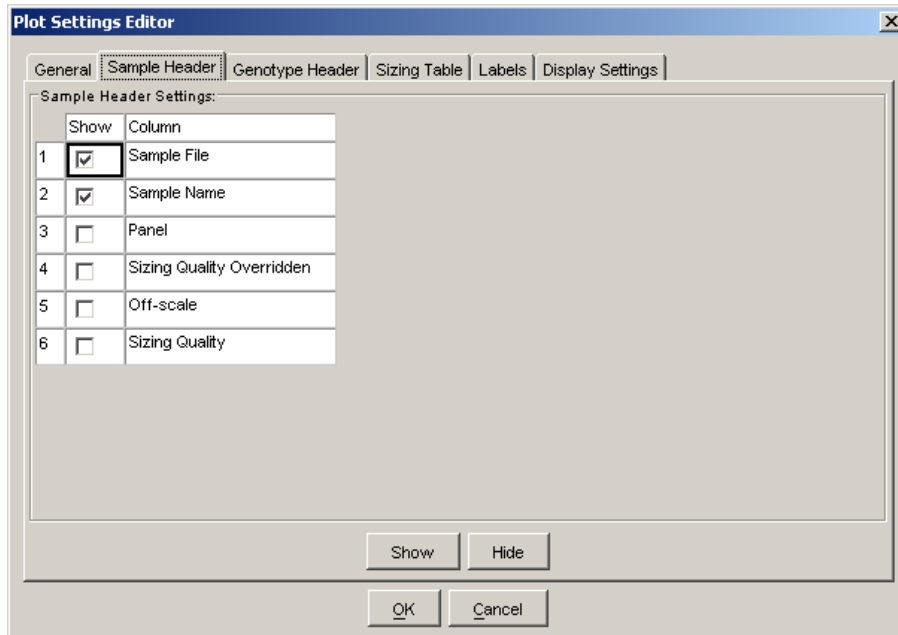
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 62 OF 70
------------------------------	---	------------------

PLOT SETTINGS: PRINT – MINIFILER CONTROLS

Print – MiniFiler Controls: Sample Header

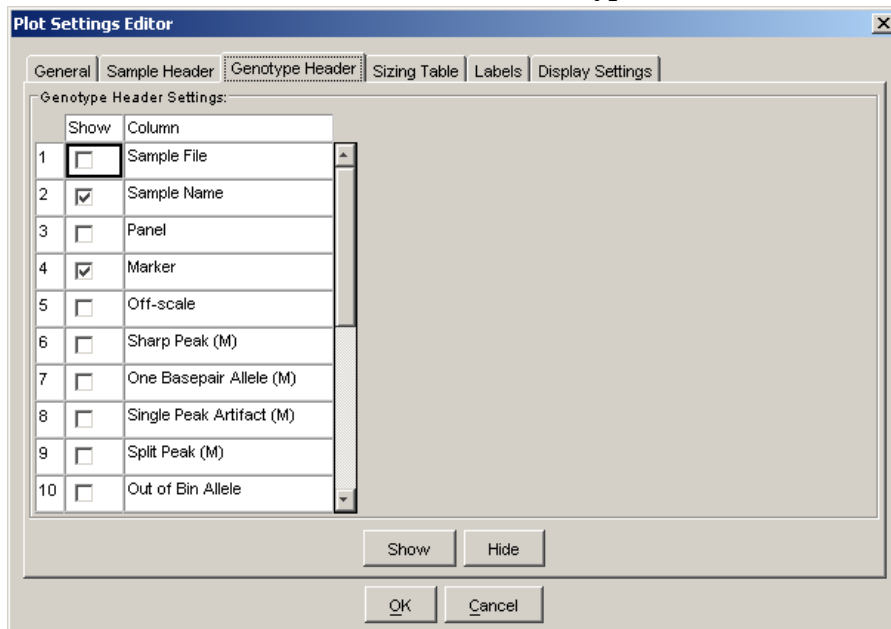


The screenshot shows the 'Plot Settings Editor' window with the 'Sample Header' tab selected. The 'Sample Header Settings' table is as follows:

	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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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 63 OF 70
------------------------------	---	------------------

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DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 64 OF 70
------------------------------	---	------------------

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 columns and 8 rows. The first column is labeled 'Show' and the second is 'Column'. All checkboxes in the 'Show' column are checked. The 'Font Settings' section shows 'Font' set to 'Arial' and 'Size' set to '8'. There are 'Show' and 'Hide' buttons at the bottom of the table, and 'OK' and 'Cancel' buttons 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'. There are 'OK' and 'Cancel' buttons at the bottom of the dialog.

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 65 OF 70
------------------------------	---	------------------

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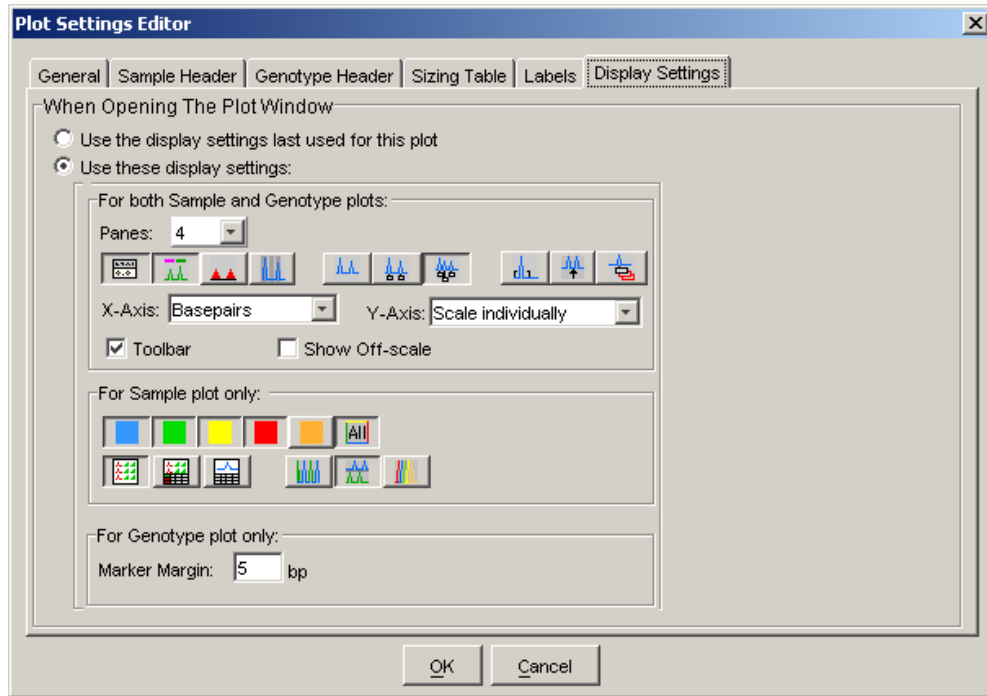
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 66 OF 70
------------------------------	---	------------------

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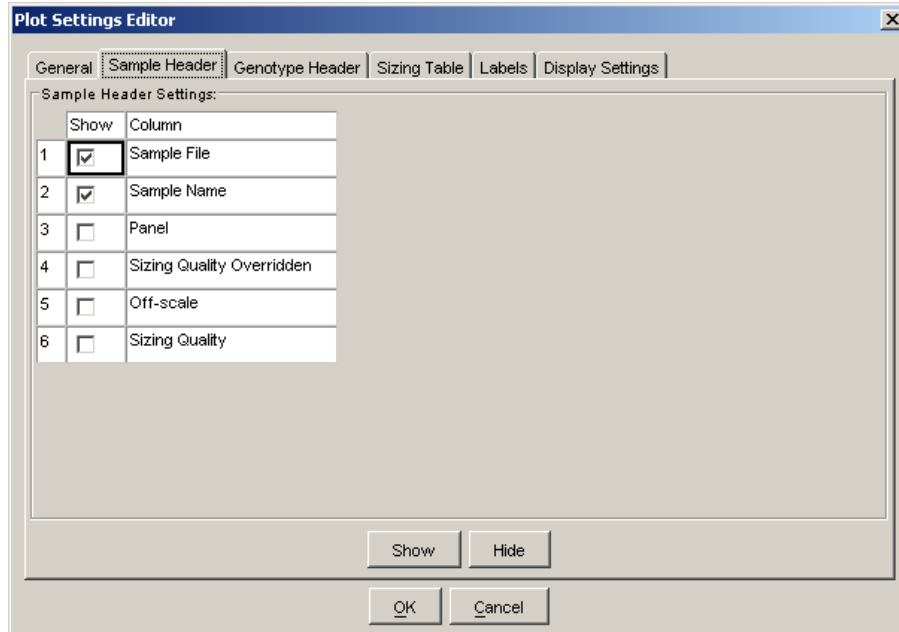
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 67 OF 70
------------------------------	---	------------------

PLOT SETTINGS: PRINT – MINIFILER SAMPLES

Print – MiniFiler Samples: Sample Header



Print – MiniFiler Samples: Genotype Header

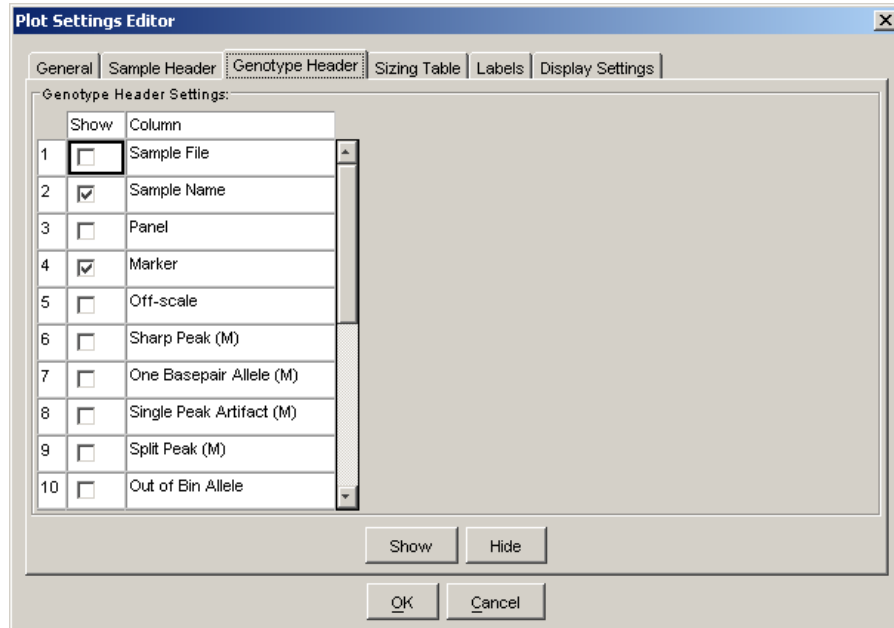
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 68 OF 70
------------------------------	---	------------------



5-20 are blank

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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 69 OF 70
------------------------------	---	------------------

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 dropdown menu.

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

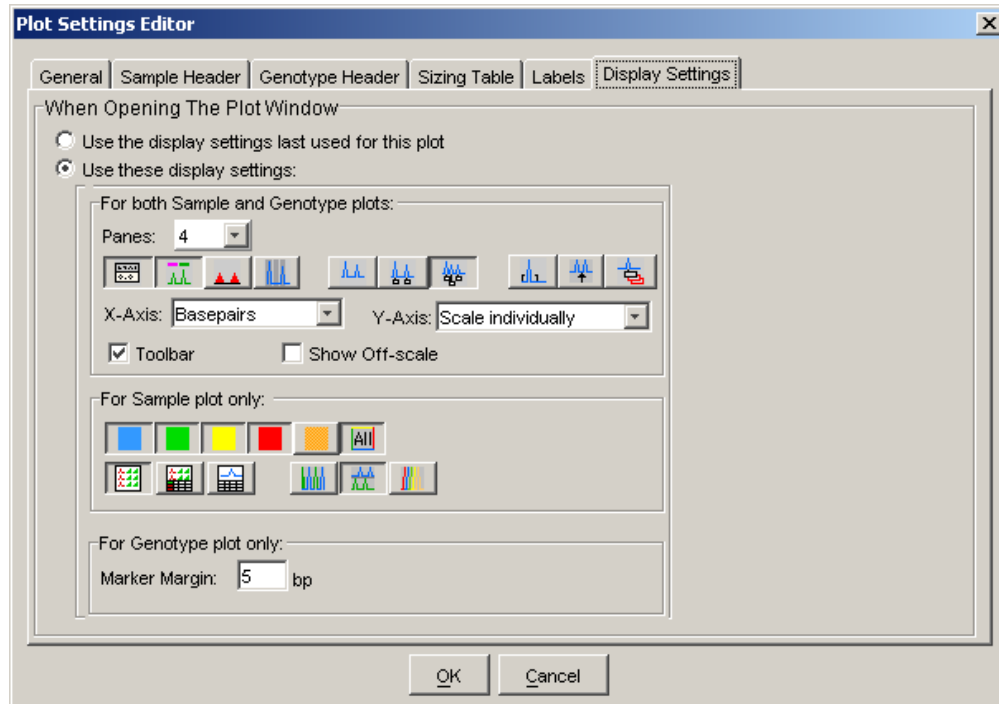
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GENEMAPPER ID – DEFAULT TABLE AND PLOT SETTINGS

DATE EFFECTIVE 06-20-2016	APPROVED BY NUCLEAR DNA TECHNICAL LEADER	PAGE 70 OF 70
------------------------------	---	------------------

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