



GenePhile X-Plex

Human X Chromosome STR Mapping Set

14 Powerful X-STR loci in a single PCR reaction

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The GenePhile X-Plex PCR Amplification Kit is a short tandem repeat (STR) multiplex assay that amplifies 13 X-STR loci and an amelogenin locus in a single PCR reaction. The X-Plex kit contains all the necessary reagents for the amplification of human DNA.

Each X-Plex kit contains materials sufficient to perform 50 reactions at a 25 μ L reaction volume.

GenePhile X-Plex PCR Amplification Kit

Kit Components	<ul style="list-style-type: none">• 5X PCR buffer 400μl• X-Plex Primer mix 125μl• GenePhile Taq Polymerase(5U/μl) 13μl• Control DNA K01(10ng/μl) 50μl• X-Plex Allelic Ladder 20μl
Number of Reactions	50
Reaction Volume	25 μ L
Loci in Kit	DXS8378, DXS9898, DXS8377, HPRTB, GATA172D05, DXS7423, DXS6809, DXS7132, DXS101, DXS6789, Amelogenin, DXS9902, DXS6807, DXS7424
Amplicon Allele Size Range	100-360 base pairs
Dyes Used	6-FAM™, VIC®, NED™, PET®, LIZ® dyes
Storage Conditions	<ul style="list-style-type: none">• 5X PCR buffer• X-Plex Primer mix• Control DNA K01 Store at 2 to 8°C.• GenePhile Taq Polymerase Store at -15 to -25°C• X-Plex Allelic Ladder Store at -15 to -25°C upon receipt, 2 to 8°C after initial use
Matrix and Size Standards	Matrix Standards: ThermoFisher DS-33 Matrix Standard kit(Dye Set G5) (PN 4345833) Size standards: ThermoFisher GeneScan™ 500 LIZ® dye Size Standard (PN 4322682) ThermoFisher GeneScan™ 600 LIZ® dye Size Standard (PN 4366589)

IMPORTANT! The fluorescent dyes attached to the primers are light sensitive. Protect the primer mix from light when not in use. Amplified DNA, Allelic Ladder, and GeneScan 500/600 LIZ Size Standards should also be protected from light. Keep freeze-thaw cycles to a minimum.

1. X-Plex kit loci and alleles

The following table shows the loci amplified by the X-Plex kit and the corresponding dyes used. The GenePhile X-Plex Kit Allelic Ladder is used to genotype the analyzed samples. The alleles contained in the allelic ladder and the genotype of the Control DNA 9947A and K01 are listed in the table.

Locus	Designation Alleles Included in X-Plex	Dye Label	9947A Genotype	Control DNA K01 Genotype
DXS8378	8-14	6-FAM™	10,11	11
DXS9898	8.3-16		12,15	12
DSX8377	33-60		45,47	51
HPRTB	6-17		14,14	15
GATA172D05	5-12	VIC®	10,10	11
DXS7423	12-18		14,15	14
DXS6809	27-40		31,34	34
DXS7132	8-19	NED™	12,12	16
DXS101	14-32		24,26	25
DXS6789	13-25		21,22	20
AMEL	X, Y	PET	X,X	X/Y
DXS9902	6-13		11,11	11
DXS6807	11-17		12,14	15
DXS7424	9-18		14,16	16

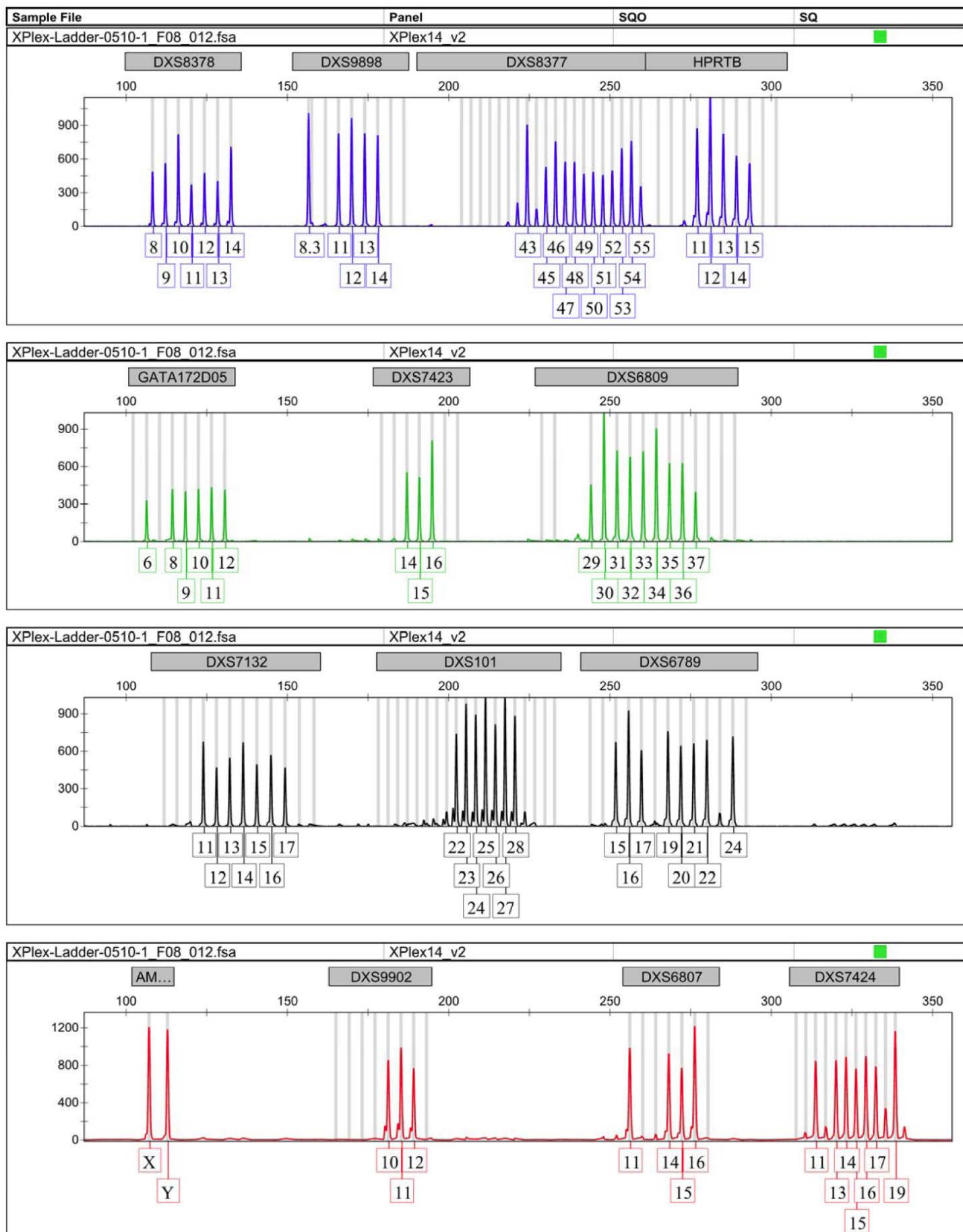
2. PCR reaction

5X PCR Buffer	5.0ul
X-Plex Primer Mix	2.5ul
GenePhile Taq Polymerase (5U/uL)	0.3ul
dd H ₂ O	14.7ul
<u>Genomic DNA(2-10ng)</u>	<u>2.5ul</u>
Total	25.0ul

3. PCR cycle

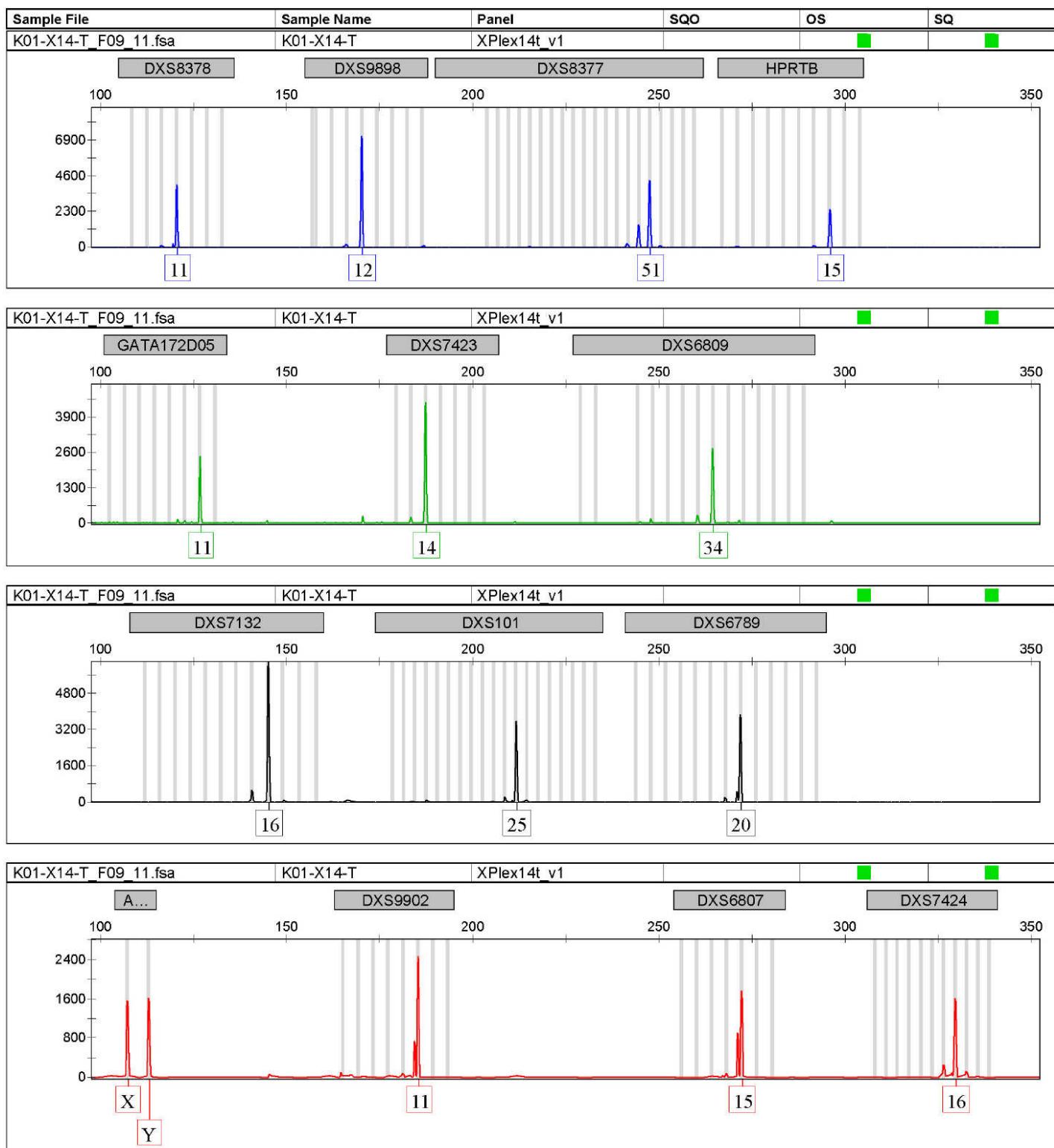
Initial Incubation Step	Cycle (30 cycles)			Final Extension	Final Hold
	Denature	Anneal	Extend		
95 °C	94 °C	60 °C	72 °C	60 °C	4 °C
10 min	1 min	1 min	1 min	45 min	Hold

4. XPlex Allelic Ladder Profile: (Load 1ul Ladder DNA per well, in 10ul Hi-Di-LIZ500/600 mixture)



5. Control DNA K01 Profile:

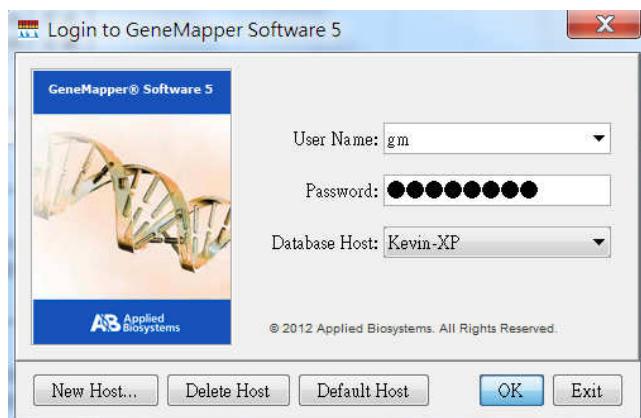
Control DNA K01 is a positive control for evaluating the efficiency of the amplification step and STR genotyping using the GenePhile X-Plex Kit Allelic Ladder.



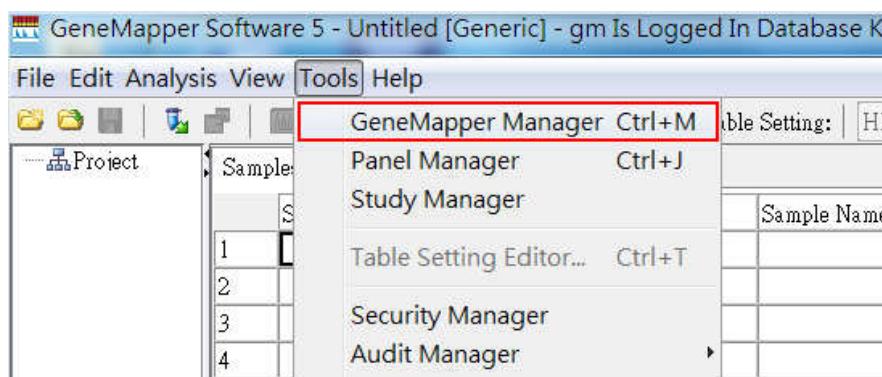
6. First-time installation of Xplex Panel & Binset: (For ABI GeneMapper users)

Download panel and bin set from website: <http://www.genephile.com.tw/products/XPlex.rar>

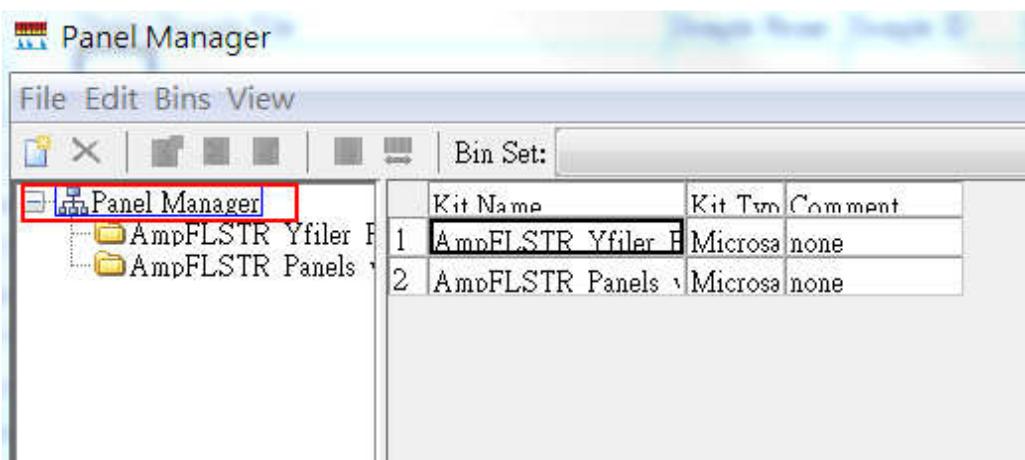
- a. Save and Unzip the **XPlex.rar** to the desktop on your PC.
- b. Run **GeneMapper** software
- c. Login with registered **User Name** and **Password**.



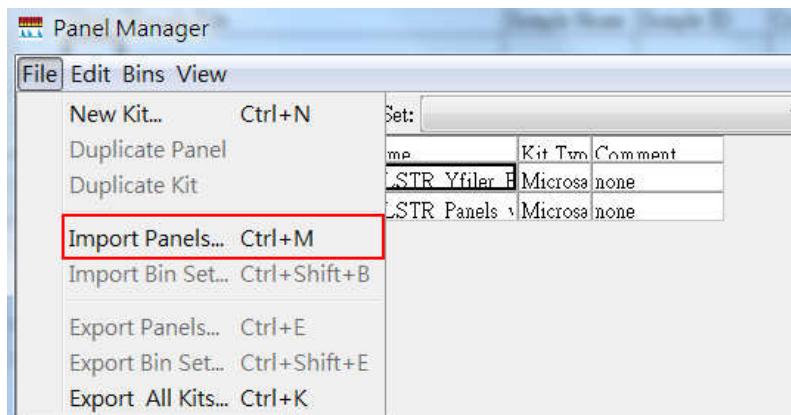
- d. Select '**Tools**' → '**Panel Manager**'



- e. Click on '**Panel Manager**'



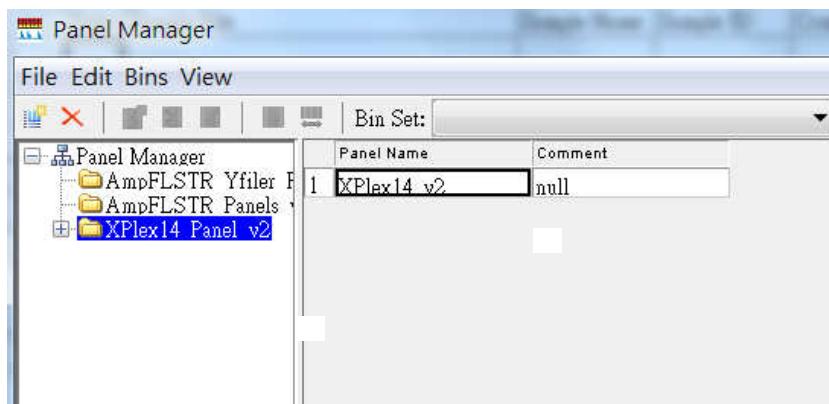
f. 'File' → 'Import Panels'



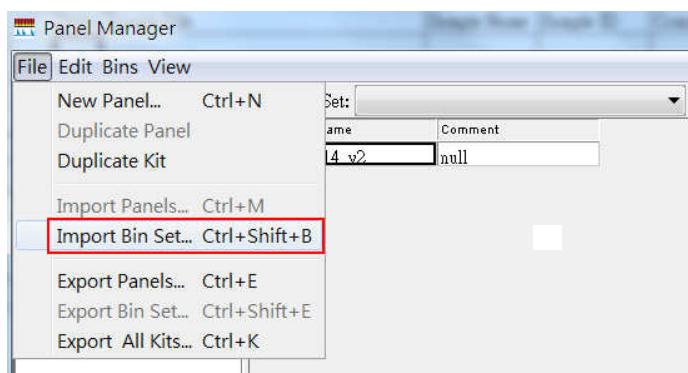
g. Direct to the file named '[XPlex14_Panel_v2.txt](#)'.

h. Click **Import**

i. Click on '[XPlex14_Panel_v2](#)'



j. 'File' → 'Import Bin Set', direct to the file named '[XPlex14_Bin_v2.txt](#)'.



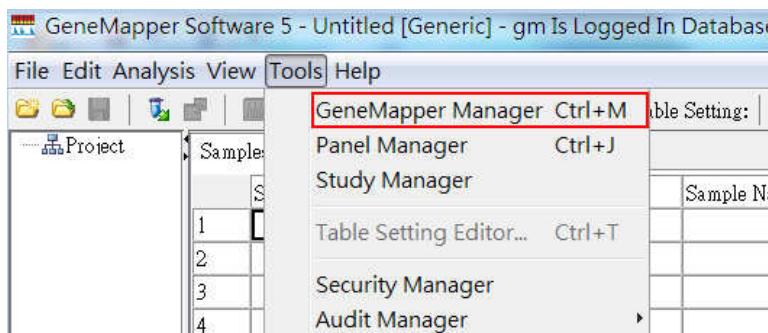
k. Click **Import**

l. Click '**Apply**' or '**OK**' and exit.

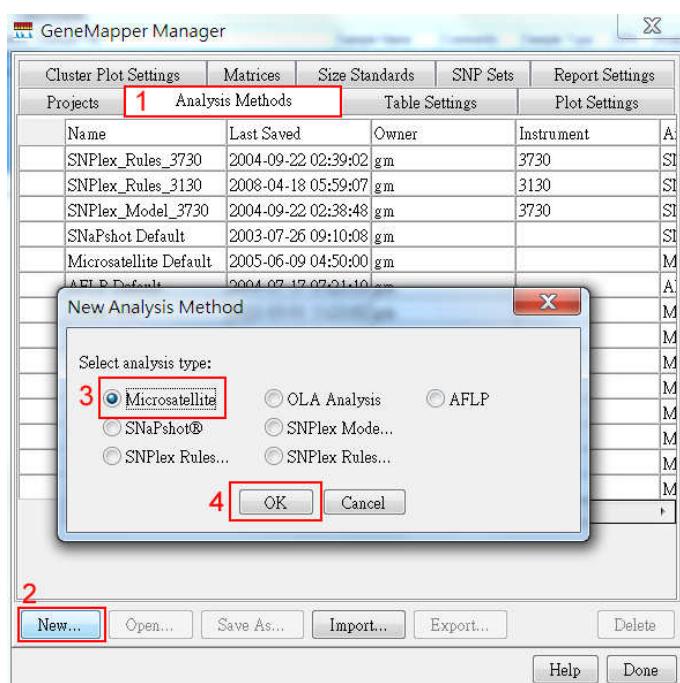


7. First-time initialization of XPlex Analysis method: (For ABI GeneMapper users)

- Start **GeneMapper** software, login with **User Name** and **Password**.
- Click 'Tools' → 'GeneMapper Manager'



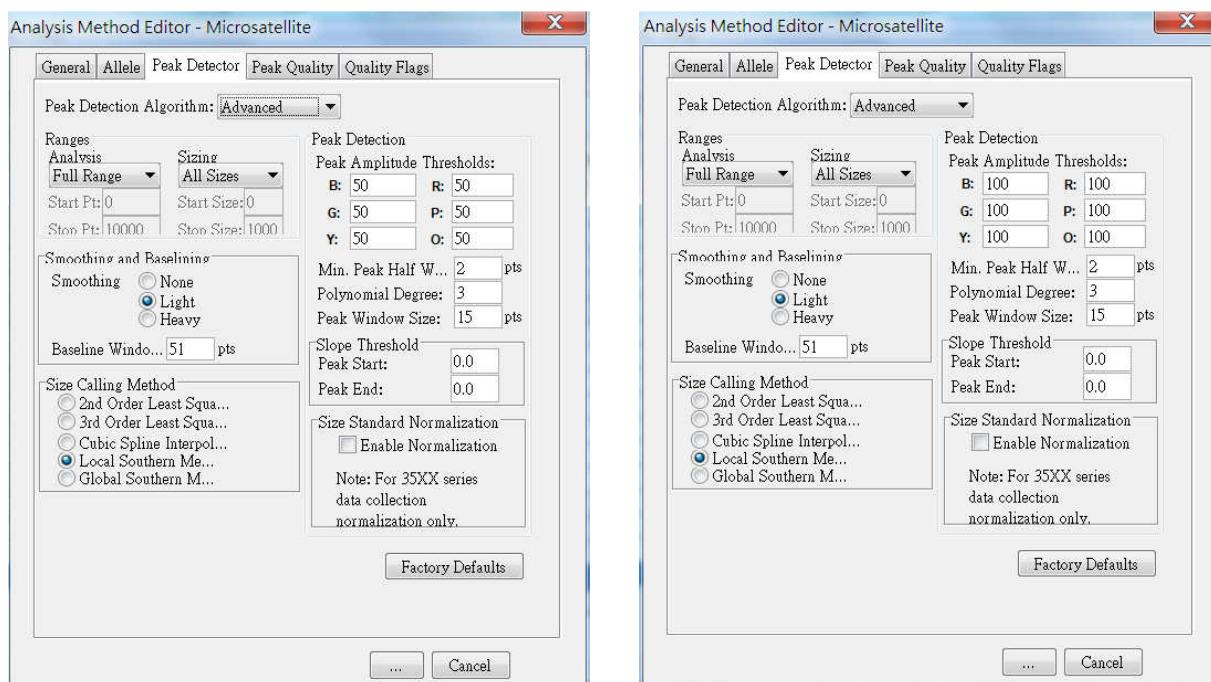
- Select **1** 'Analysis Methods' sheet → **2** 'New' → **3** 'Microsatellite' → **4** 'OK'



- Assign a name for this method: '**X-Plex**'. In the '**Allele**' sheet, select '**Xplex14_Binset_v2**'.

The screenshot shows two instances of the 'Analysis Method Editor - Microsatellite' dialog box. The left instance shows the 'General' tab with 'Name' set to 'X-Plex' and 'Analysis Type' set to 'Microsatellite'. The right instance shows the 'Allele' tab with 'Bin Set' set to 'None' and 'Marker Re...' dropdown set to 'Xplex14_Binset_v2'. Other tabs like Peak Detector, Peak Quality, and Quality Flags are visible at the top of both dialogs.

- e. In the 'Peak Detector' sheet, users can modify the peak amplitude thresholds to meet analytic requirements. Press 'OK' to finish a setting.



8. Perform a XPlex analysis:

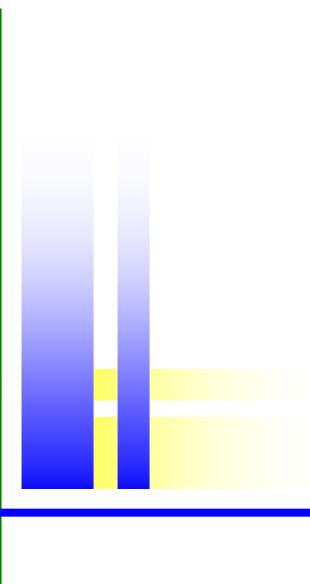
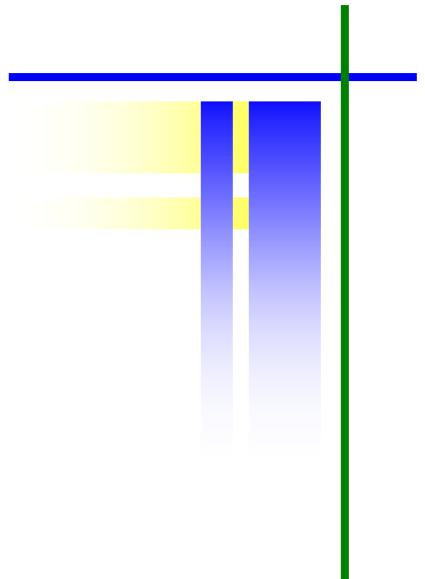
- In the sample list view, create a new project and import a "XPlex ladder" file and "sample" files.
(Note: XPlex ladder file and sample files must be placed in a same folder.)
- Assign sample types **1**, set Analysis Method to 'XPlex' and Panel to 'XPlex14_v2' **2**. Finally, press button to start an analysis **3**.

Status	Sample File	Sample Name	Comments	Sample Type	SFN	Analysis Method	Panel	Size Standard	Matrix
1	GPT190155-D-X_	GPT190155-D-X	None	Sample	NA	X-Plex	XPlex14_v2	GS500(-250)LIZ	
2	GPT190155-D2-X	GPT190155-D2-X	None	Sample	NA	X-Plex	XPlex14_v2	GS500(-250)LIZ	
3	GPT190155-M-X_	GPT190155-M-X	None	Sample	NA	X-Plex	XPlex14_v2	GS500(-250)LIZ	
	X-ladder-108040	X-ladder-1080401A	None	Allelic Ladder	NA	X-Plex	XPlex14_v2	GS500(-250)LIZ	

- Select samples and perform genotypes.

9. Publication:

- Hwa HL, Chang YY, Lee J CI, Yin HY, Tseng LH, Su YN, Ko TM. **Thirteen X-Chromosomal short tandem repeat loci multiplex data from Taiwanese.** *Int J Legal Med.* 2009 July;123:263-269.
- Hwa HL, Lee, J CI, Chang, YY, Yin HY, Chen YH., Tseng LH., Ko TM. **Genetic analysis of eight population groups living in Taiwan using a 13 X-chromosomal STR loci multiplex system.** *Int J Legal Med.* 2010; 125(1):33-37.



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