

Cell STR Certification Report



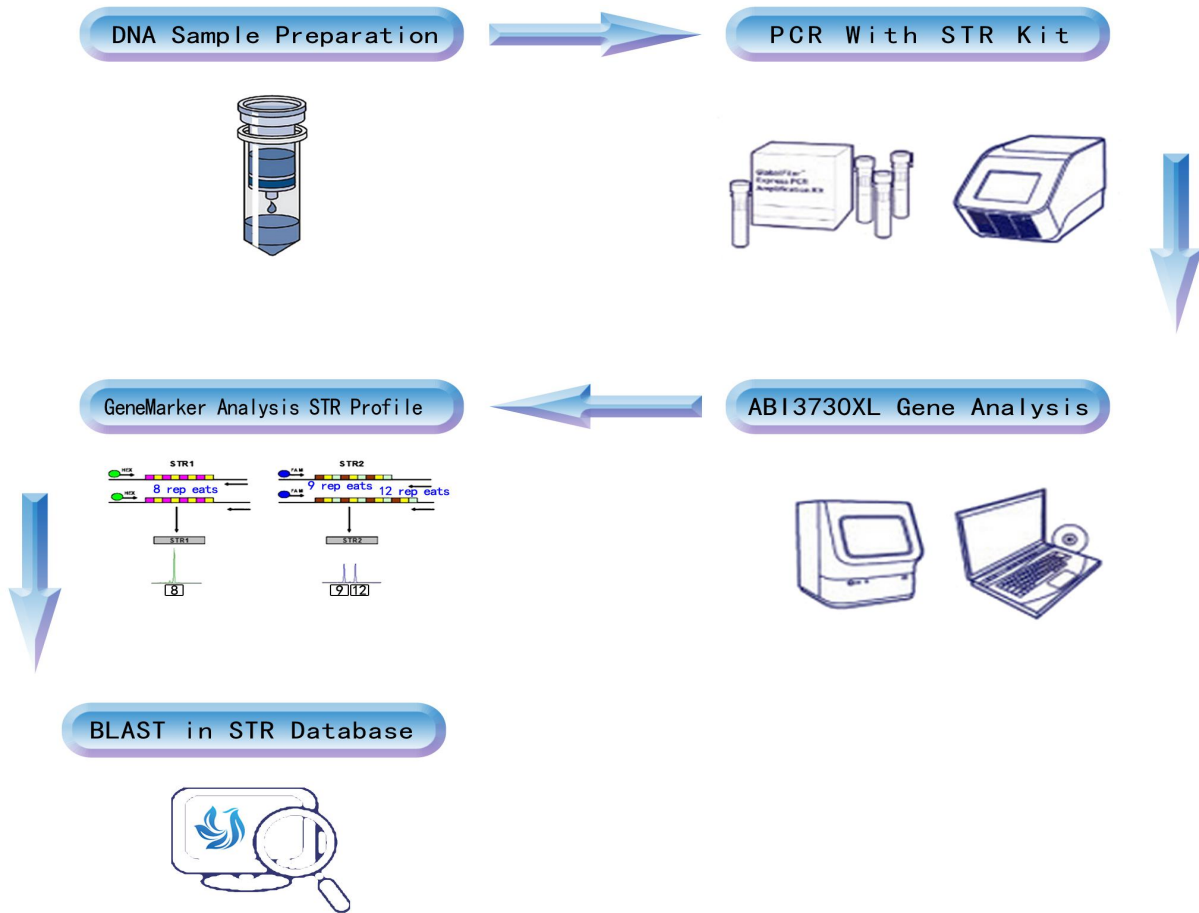
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- 1、 Sample ID: HepG2
- 2、 Original Material: Cell pellets
- 3、 Check time:2020-10-21
- 4、 Methods:



5、 Results:

Negative and positive test results are correct.

Amplification map of Genomic DNA clear, Genotyping results well.

STR Profile :

Genetic Site	Customer sample		ATCC	
	HepG2		HepG2	
Amelogenin	X	Y	X	Y
CSF1PO	10	11	10	11
D13S317	9	13	9	13
D16S539	12	12	12	12
D5S818	11	12	11	12
D7S820	10	10	10	10
THO1	9	9	9	9
TPOX	8	9	8	9
vWA	17	17	17	17



Percent match between the sample and the database profile: 100%

6、 Summary:

The result of STR profile showed no more than 2 distinct alleles were found ,the sample derived from a common ancestry(Figure 1); which matched 100% the reference cell line in the ATCC STR database, named HepG2 .

Notes:

- $P=100\% \times (2 \times M)/N$; $M=18$, $N=36$, $P=100\% \times (2 \times 18)/36=100\%$
M: number of the matching peaks; N: number of all peaks
- Based on ASN-0002-2011 Standard , cell lines with $\geq 80\%$ match are considered to be related ; i.e.,derived from a common ancestry. Cell lines with between a 55% to 80% match require futher profiling for authentication of relatedness.
- This data and analysis are for research use only.

Operator: Yuqiu Qin

Auditor:



Report time: 2020-10-23

Figure:

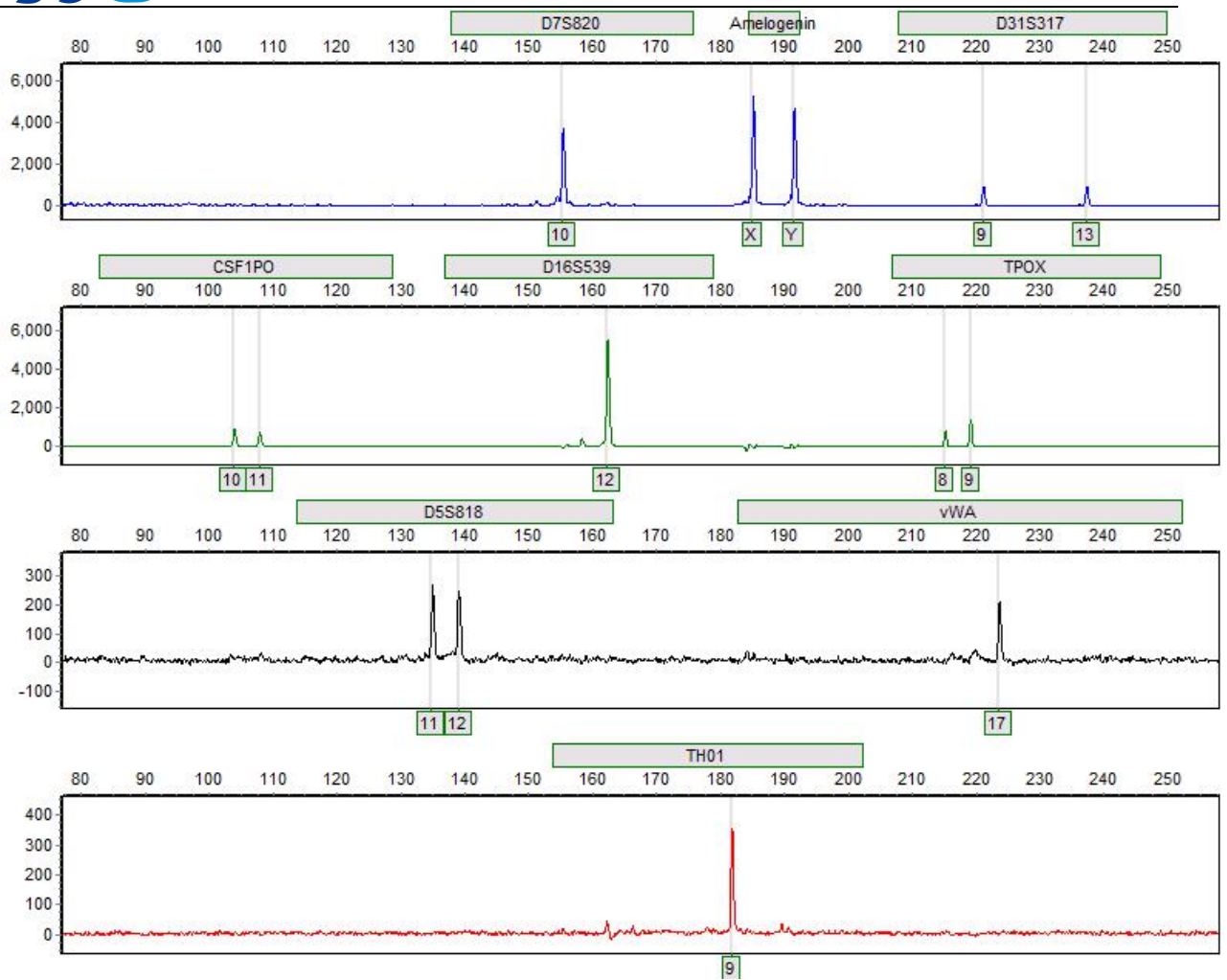


Figure 1.STR profiles of HepG2 cell line