Cell Line Authentication Report

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科伯生物Cobioer

1. Sample information:

Sample No.: “HuH-6”

2. Methods:

a. Genomic DNA was extracted from the cell pellets provided by the customer.
b. Samples, together with positive and negative control were amplified using GenePrint System (Promega).
c. Amplified products were processed using the ABI3730xl Genetic Analyzer.
d. Data was analyzed by GeneMapper4.0 software and then compared with the ATCC, DSMZ or JCRB databases for reference matching.

3. Results:

STR profile

Marker	Sample				Database		
	Allele1	Allele2	Allele3	Allele4	Allele1	Allele2	Allele3
Amelogenin	X	Y			X	Y	
D5S818	8	11			8	11	
D13S317	8	12			8	12	
D7S820	11	12			11	12	
D16S539	10	11			10	11	
vWA	14	17			14	17	
TH01	7	8			7	8	
TPOX	8	8			8	8	
CSF1PO	10	12			10	12	
D19S433	12	12.2					
D21S11	29	30					
D18S51	13	21					
D6S1043	13	18					
D3S1358	14	17					
Penta D	9	13					
D2S441	10	12					
D8S1179	10	11					
Penta E	11	11					
D12S391	18	20					
D2S1338	18	18					
FGA	19	24					

4. Conclusion

A．The STR results show that there are no four alleles on the main nine locus, there is certainly no cross contamination of human cells in this cell line.

B．The match percent between the sample and the STR database profile is **100%**, the cell name is **HuH-6.**

Similarity	Cell line	Source	Shared	D5S818	D7S820	D13S317	D16S539	vWA	TH01	TPOX	CSF1PO	Amelogenin
		Your query		8, 11	11, 12	8, 12	10, 11	14, 17	7, 8	8, 8	10, 12	X, Y
100 %	HuH-6	RCB1367	9	8, 11	11, 12	8, 12	10, 11	14, 17	7, 8	8, 8	10, 12	X, Y

Note:

1. The STR profile data was compared with the ATCC, DSMZ or JCRB databases, if the cell line is not included in the three institutions, the results are not correct. More information you provided would be useful for Cell Line Authentication.
2. Based on the ANSI Standard, cell lines with 100% match are considered to be “identical”; cell lines with ≥80% but less than 100% match are considered to be “related”.

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Date：2023-12-01