## O UBIGENE **STR Authentication Report For Cell Line**

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1. Cell Name: NIH/3T3

2. Test Method: DNA was extracted using the genome extraction kit (Axygen), amplified using a 20- STR amplification protocol, the STR loci and gender gene Amelogenin were detected on the ABI 3730XL Genetic Analyzer.

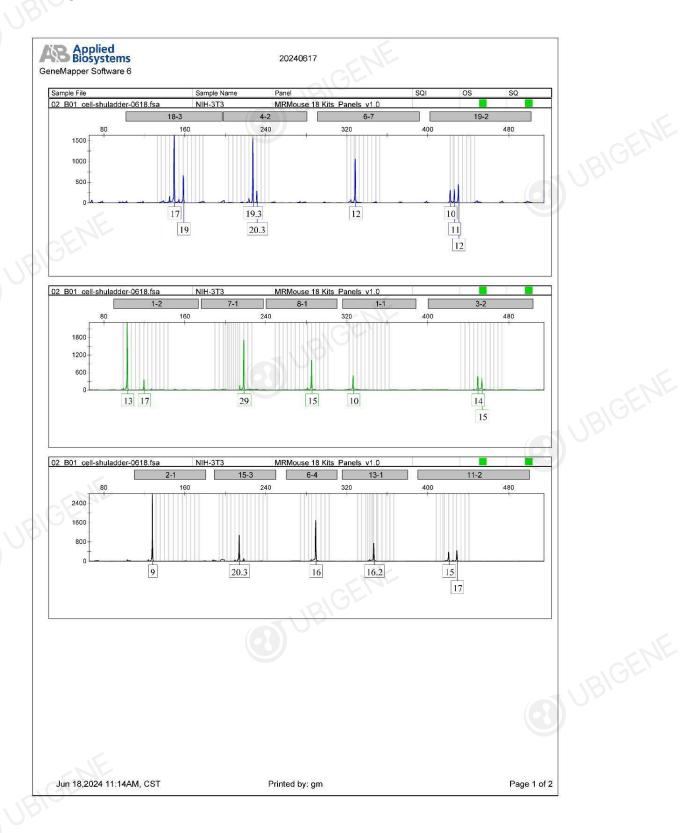
	Genoty	ype analysis	results of S	ΓR and Amel	ogenin loci	in cells	
STR Loci	GENE	STR profile fo	or Sample cel	STR profile for Cell Bank cell Cell name: NIH 3T3			
	0,	Cell name	e: NIH-3T3				
	Allele1	Allele2	Allele3	Allele4	Allele1	Allele2	Allele3
18-3	17.0	19.0		IBIO	17.0	19.0	
4-2	19.3	20.3	(3)		19.3	20.3	
6-7	12.0				12.0		181GK
19-2	10.0	11.0	12.0		11.0	12.0	0
1-2	13.0	17.0			13.0	17.0	
7-1	29.0				29.0		
8-1	15.0				15.0		
1-1	10.0			GEN	10.0		
3-2	14.0	15.0		JBIO	14.0	15.0	
2-1	9.0				9.0		
15-3	20.3				20.3		IBIO
6-4	16.0				15.3	(3)	
13-1	16.2				16.2		
11-2	15.0	17.0			15.0	17.0	
17-2	13.0	14.0			13.0	14.0	
12-1	20.0			EN	20.0		
5-5	14.0	15.0		BIOL	14.0	15.0	

X-1	25.0			25.0	
TH01	0.				
D5S818			CENT		

Note: The cell lines were compared with the STR data of cell lines from ATCC, DSMZ, JCRB and RIKEN databases, the cell lines not included in the above cell banks could not be matched. D4S2408 and TH01 in the above sites is a human site, which is used to detect whether the cell is contaminated by human sources.

4. Conclusion: This cell line is identified as a mouse cell line. The STR results of NIH/3T3 cells are consistent with the genotypes of NIH/3T3 cell lines in EXPASY database, the cell ID corresponded to CVCL 0594, and the STR results basically matched. In the test, No multiple alleles were found, no cross-contamination, no human contamination in this cell line.

## 5. Attached Image





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