

## STR Authentication Report For Cell Line

O) UBIGENE

1. Cell Name: H22

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

	Genot	ype analysis	results of ST	R and Ame	ogenin loci	in cells	
	GENE	STR profile fo	or Sample cell	STR profile for Cell Bank cell Cell name: H22			
STR Loci	0,	Cell name: S	TR25012026				
	Allele1	Allele2	Allele3	Allele4	Allele1	Allele2	Allele3
18-3	11.0	17.0		BIOL	11.0	17.0	
4-2	19.3	20.3			19.3	20.3	
6-7	13.0				13.0		BIGE
19-2	13.0						
1-2	18.0	19.0					
7-1	24.2						
8-1	17.0	18.0			e.		
1-1	15.0	16.0		GEN	Ĺ		
3-2	13.0	14.0		BIU			
2-1	9.0						a C
15-3	19.3	20.3			19.3	20.3	BIOL
6-4	15.3	16.3			15.3	16.3	9
13-1	16.2						
11-2	16.0						
17-2	13.0	15.0					
12-1	17.0			EN	17.0		
5-5	14.0			BIOL	14.0		

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Gene-editing cell lines | CRISPR Library Microorganisms | EZ-editor™ series products

X-1	25.0			25.0	
TH01	0.				
D5S818			CEN	Ĺ	

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 <u>H22</u> cells are consistent with the genotypes of <u>H22</u> cell lines in <u>EXPASY</u> database, the cell ID corresponded to <u>CVCL\_H613</u>, and the STR results <u>basically matched</u>. In the test, <u>Multiple alleles were found, no cross-contamination, no human contamination</u> in this cell line.

## 5. Attached Image

