

Cell Line Authentication Service STR Profile Report

Sales Order: 240625D-1
Cell Line Designation: NCI-H358
Date Sample Received: Jun 25th, 2024
Report Date: Jul 11th, 2024

Methodology: Twenty-one short tandem repeat (STR) loci plus the Amelogenin locus were amplified using the commercially available SiFaSTR™ 23 plex Kit. The cell line sample was processed using the ABI Prism® 3130 XL Genetic Analyzer. Data were analyzed using GeneMapper® ID v3.2 software (Applied Biosystems). Appropriate positive and negative controls were run and confirmed for each sample submitted.

Data Interpretation: Cell lines were authenticated using Short Tandem Repeat (STR) analysis as described in 2021 in ANSI Standard (ASN-0002) by the ATCC Standards Development Organization (SDO) and in Jamie L. Almeida et al., Authentication of Human and Mouse Cell Lines by Short Tandem Repeat (STR) DNA Genotype Analysis. Assay Guidance Manual. PMID: 23805434. Bookshelf ID: NBK144066.

GTB™ performs STR Profiling following ISO 9001:2008 and ISO/IEC 17025:2005 quality standards.

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NOTE: According to the recommendations of *IJC* on cell line authentication, the report is valid for 3 years since the issue date.

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Test Results for Submitted Sample			ExPASy Reference Database Profile	
Loci	Query Profile: NCI-H358		Database Profile: NCI-H358	
Amelogenin	X	Y		
D3S1358	14	18	14	18
D5S818	10	12	10	12
D2S1338	17	23		
TPOX	8	9	8	9
CSF1PO	11	12	11	12
Penta D	10	13		
TH01	6		6	
vWA	17		17	
D7S820	10	11	10	11
D21S11	28	30	28	30
Penta E	18			
D10S1248	13	14		
D8S1179	13	14	13	14
D1S1656	15	15.3		
D18S51	14		14	
D12S391	20			
D6S1043	14			
D19S433	13	14		
D16S539	12	13	12	13
D13S317	8	12	8	12
FGA	20	21	20	21

The allele match algorithm compares the 13 core STR loci only, even though alleles from all loci will be reported when available.

Note: Loci highlighted in grey (13 core STR loci) can be made public to verify cell identity. In order to protect the identity of the donor, **please do not publish** the allele calls from all the STR loci tested.

The sample match is based on the reference data available at the time of comparison.

Explanation of Test Results

Cell lines with $\geq 80\%$ match are derived from the same donor. Cell lines with between a 70% to 79% match require further profiling for authentication of relatedness. Cell lines with $< 70\%$ match are very unlikely to be from the same donor.

- ☐ The submitted sample profile is human, but not a match for any profile in the ExPASy STR database.
- ☒ The submitted profile is an exact match for the following human cell line(s) in the ExPASy STR database (13 core STR loci): NCI-H358
- ☐ The submitted profile is similar to the following ExPASy human cell line(s):

e-Signature Technician:

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DN: cn=Xuekun Chen, o=Genetic Testing
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More information

Addendum: Electropherogram for the customer's sample set 1 of 1
For Research Use ONLY



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