

Cell Line Authentication Service STR Profile Report

Sales Order: 240719A
Cell Line Designation: BEAS-2B
Date Sample Received: Jul 19th, 2024
Report Date: Jul 19th, 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: BEAS-2B		Database Profile: BEAS-2B	
Amelogenin	X	Y		
D3S1358	15	17	15	17
D5S818	12	13	12	13
D2S1338	22	23		
TPOX	6	11	6	11
CSF1PO	9	12	9	12
Penta D	2.2	13		
TH01	7	9.3	7	9.3
vWA	17	18	17	18
D7S820	10	13	10	13
D21S11	28	30	28	30
Penta E	5	8		
D10S1248	14	15		
D8S1179	13	15	13	15
D1S1656	14			
D18S51	18	19	18	19
D12S391	17	18		
D6S1043	12	18		
D19S433	13.2	15.2		
D16S539	12		12	
D13S317	13		13	
FGA	20	24	20	24

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 In order to protect the identity of the donor, **please do not publish** the allele calls from all the STR loci tested.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): BEAS-2B
- ☐ The submitted profile is similar to the following ExPASy human cell line(s):

e-Signature Technician:

Digitally signed by Xuekun Chen
DN: cn=Xuekun Chen, o=Genetic Testing
Biotechnology (Suzhou), ou=DNA Typing Section,
email=order@jsdna.org, c=CN
Date: 2024.07.19 15:44:05 +08'00'

e-Signature Reviewer:

Digitally signed by Xiankun Zhao
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Biotechnology (Suzhou), ou=Supervision Section,
email=service@jsdna.org, c=CN
Date: 2024.07.19 15:44:50 +08'00'



More information

Addendum: Electropherogram for the customer's sample set 1 of 1

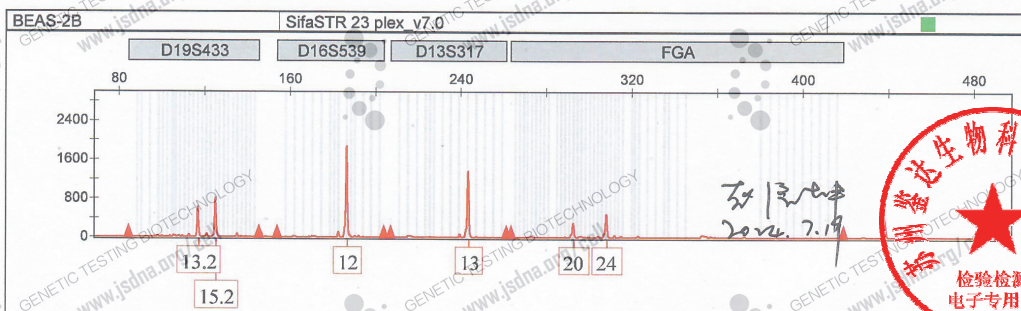
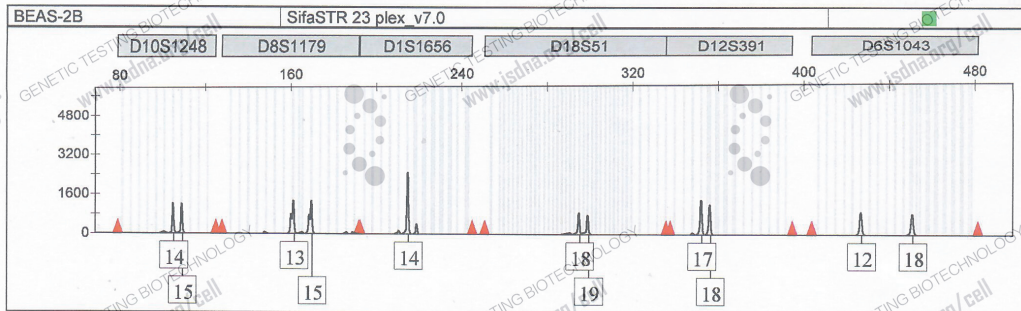
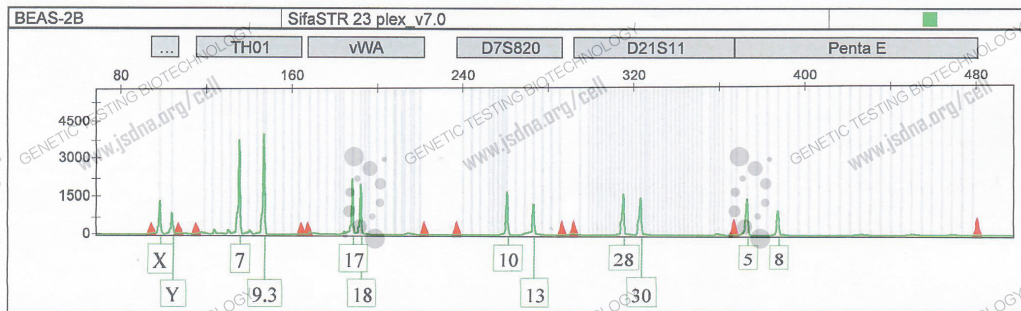
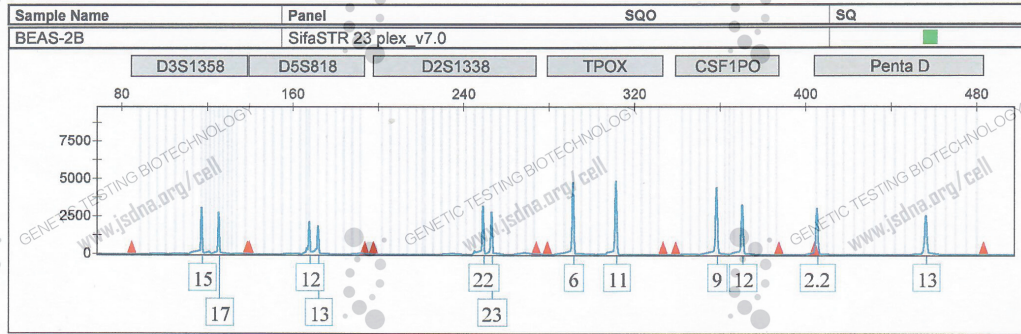


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Applied
Biosystems
GeneMapper ID v3.2

240719



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