

STR Profile Report

Sample Submitted By: Dr. Ping Wang

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Sales Order: 170408B
Cell Line Designation: A549

Date Sample Received: Apr 8th, 2017 **Report Date:** Apr 10th, 2017

Methodology: Nineteen short tandem repeat (STR) loci plus the gender determining locus,

Amelogenin, were amplified using the commercially available EX20 Kit from AGCU. The cell line sample was processed using the ABI Prism® 3500 Genetic Analyzer. Data were analyzed using GeneMapper® ID-X v1.4 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 2012 in ANSI Standard (ASN-0002) by the ATCC Standards Development Organization (SDO) and in Capes-Davis et al., Match criteria for human cell line authentication: Where do we draw the line? Int J Cancer.

2013;132(11):2510-9.

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

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| Test Results for Submitted Sample | | | | | | DSMZ Reference Database Profile | | | |
|-----------------------------------|----|-----------|------------|--|------------------------|---------------------------------|--|--|--|
| Loci | | Query Pro | file: A549 | | Database Profile: A549 | | | | |
| Amelogenin | Χ | Υ | | | Χ | Υ | | | |
| D3S1358 | 16 | | | | | | | | |
| D13S317 | 11 | | | | 11 | | | | |
| D7S820 | 8 | 11 | | | 8 | 11 | | | |
| D16S539 | 11 | 12 | <u>13</u> | | 11 | 12 | | | |
| Penta E | 7 | 11 | | | | | | | |
| TPOX | 8 | 11 | | | 8 | 11 | | | |
| TH01 | 8 | 9.3 | | | 8 | 9.3 | | | |
| D2S1338 | 24 | | | | | | | | |
| CSF1PO | 10 | 12 | | | 10 | 12 | | | |
| Penta D | 9 | | | | | | | | |
| D19S433 | 13 | | _ | | | | | | |
| vWA | 14 | | | | 14 | | | | |
| D21S11 | 29 | | | | | | | | |
| D18S51 | 14 | 17 | | | | | | | |
| D6S1043 | 11 | 13 | | | | | | | |
| D8S1179 | 13 | 14 | | | | | | | |
| D5S818 | 11 | | | | 11 | | | | |
| D12S391 | 18 | 19 | | | | | | | |
| FGA | 23 | | | | | | | | |

The allele match algorithm compares the 8 core loci plus amelogenin only, even though alleles from all loci will be reported when available.

Note: Loci highlighted in grey (8 core STR loci plus Amelogenin) 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 In order to protect the identity of the donor, **please do not publish** the allele calls from all the STR loci tested.

Explanation of Test Results

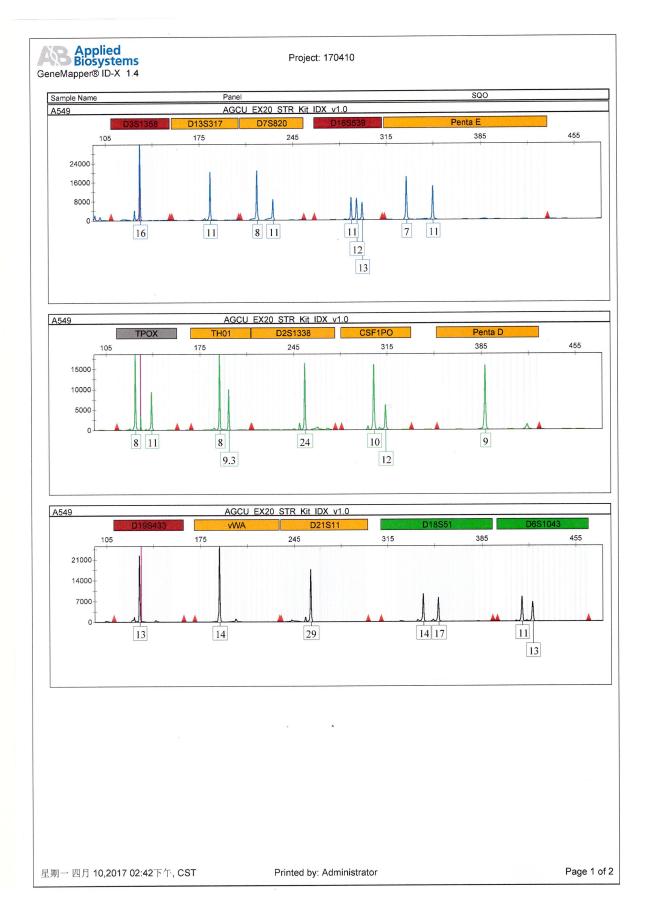
Cell lines with ≥80% match are considered to be related; i.e., derived from a common ancestry. Cell lines with between a 55% to 80% match require further profiling for authentication of relatedness.

| | | • | • | J | | | |
|---|---|-------------------|----------|-------------------|--------------------|------------------|----------|
| | The submitted sample | e profile is huma | n, but r | not a match for a | ny profile in the | e DSMZ STR data | abase. |
| | The submitted profile (8 core loci plus Ame | | h for th | e following huma | ın cell line(s) ir | n the DSMZ STR | database |
| • | The submitted profile | is similar to the | followin | ng DSMZ human | cell line(s): A5 | 49 (97% match) | |
| | nature, Technician: nature, Reviewer: | | | | _ | | |
| | | | | | | More information | |

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



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