

STR Authentication Report For Cell Line

O) UBIGENE

1. Cell Name: B16-F10

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

| | | | | | | (A) | UBIC |
|----------|---------|---------------|----------------|------------------------------------------------------|-------------|----------|---------|
| | Genoty | vpe analysis | results of ST | R and Ame | ogenin loci | in cells | |
| . < | GENS | TR profile fo | or Sample cell | STR profile for Cell Bank cell Cell name: B16-F10 | | | |
| STR Loci | | Cell name | e: B16-F10 | | | | |
| | Allele1 | Allele2 | Allele3 | Allele4 | Allele1 | Allele2 | Allele3 |
| 18-3 | 15.0 | 16.0 | | BIGE | 15.0 | 16.0 | |
| 4-2 | 20.3 | 21.3 | | 0 | 20.3 | 21.3 | |
| 6-7 | 15.0 | | | | 15.0 | | BIGE |
| 19-2 | 13.0 | 14.0 | | | 13.0 | 14.0 | 04 |
| 1-2 | 19.0 | 20.0 | | | 19.0 | 20.0 | |
| 7-1 | 26.2 | | | | 26.2 | | |
| 8-1 | 16.0 | 17.0 | | | 16.0 | 17.0 | |
| 1-1 | 17.0 | 18.0 | 19.0 | GEN | 17.0 | 18.0 | 19.0 |
| 3-2 | 14.0 | 15.0 | | JBIOS | 14.0 | 15.0 | |
| 2-1 | 16.0 | | 8 | | 16.0 | | |
| 15-3 | 22.3 | 23.3 | | | 22.3 | 23.3 | BIGE |
| 6-4 | 18.0 | 19.0 | | | 18.0 | 19.0 | 0. |
| 13-1 | 17.1 | 18.1 | | | 17.1 | 18.1 | |
| 11-2 | 16.0 | 17.0 | | | 16.0 | 17.0 | |
| 17-2 | 15.0 | 16.0 | 17.0 | | 15.0 | 16.0 | 17.0 |
| 12-1 | 17.0 | 18.0 | | - N | 17.0 | 18.0 | |
| 5-5 | 16.0 | 20.0 | | BIGE | 16.0 | 20.0 | |

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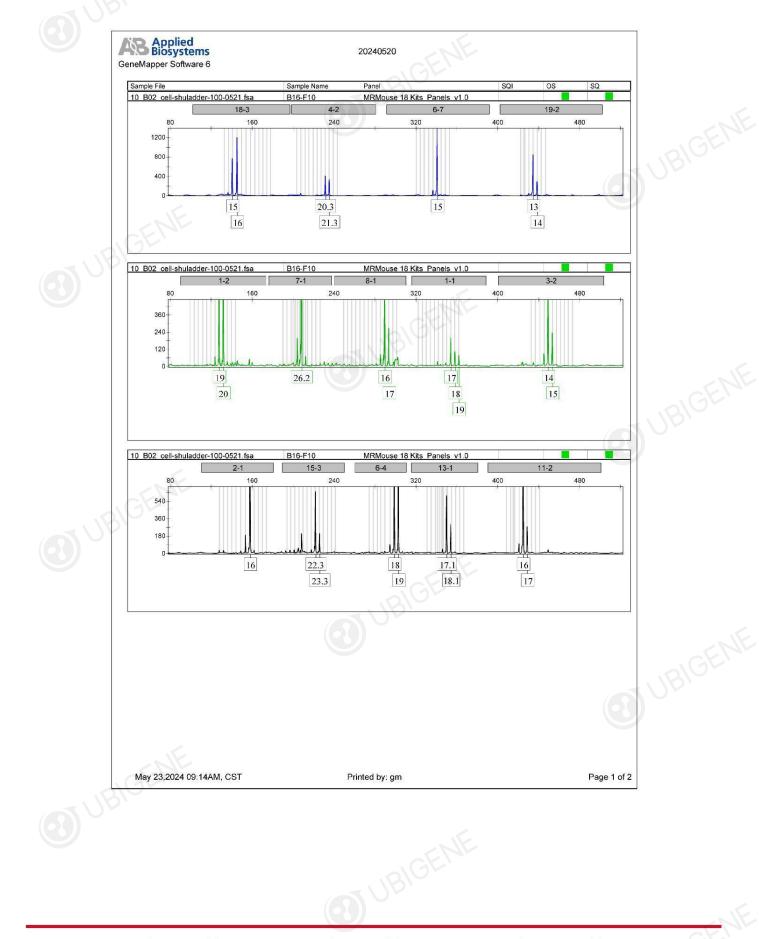
Gene-editing cell lines | CRISPR Library Microorganisms | EZ-editor™ series products

| X-1 | 28.0 | | | 28.0 | |
|--------|------|--|-----|------|--|
| TH01 | 0. | | | | |
| D5S818 | | | CEN | 1 | |

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 **B16-F10** cells are consistent with the genotypes of **B16-F10** cell lines in **EXPASY** database, the cell ID corresponded to **CVCL_0159**, and the STR results **completely 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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