

## STR Authentication Report For Cell Line

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1. Cell Name: IMR32

2. Test Method: DNA was extracted using the genome extraction kit (Axygen), amplified

using a 21- STR amplification protocol, the STR loci and gender gene Amelogenin were

|          | Genotype               | analysis res   | Ilts of STR and | Amelogenin l                   | oci in cells |         |  |
|----------|------------------------|----------------|-----------------|--------------------------------|--------------|---------|--|
|          |                        | rofile for Sam |                 | STR profile for Cell Bank cell |              |         |  |
| STR Loci | Cell name: STR24091208 |                |                 | Cell name: IMR-32              |              |         |  |
|          | Allele1                | Allele2        | Allele3         | Allele1                        | Allele2      | Allele3 |  |
| D5S818   | 11                     | 12             | - IBIC          | 11                             | 12           |         |  |
| D13S317  | 9                      | 9              |                 | 9                              | 9            |         |  |
| D7S820   | 9                      | 10             |                 | 9                              | 10           | BIGE    |  |
| D16S539  | 8                      | 8              |                 | 8                              | 8            |         |  |
| VWA      | 15                     | 15             |                 | 15                             | 15           | F       |  |
| TH01     | GF7                    | 9.3            |                 | 7                              | 9.3          |         |  |
| AMEL     | Х                      | Y              |                 | х                              | Y            |         |  |
| ТРОХ     | 11                     | 11             | 0               | -141                           | 11           |         |  |
| CSF1PO   | 11                     | 12             | UBIC            | 11                             | 12           |         |  |
| D12S391  | 23                     | 23             | 8               |                                |              | E       |  |
| FGA      | 21                     | 24             |                 |                                |              | BIGE    |  |
| D2S1338  | 23                     | 24             |                 |                                | 8            |         |  |
| D21S11   | 30                     | 31             |                 |                                |              |         |  |
| D18S51   | 12                     | 15             |                 |                                |              |         |  |
| D8S1179  | 13                     | 13             |                 |                                |              |         |  |
| D3S1358  | 16                     | 16             |                 | ENE                            |              |         |  |
| D6S1043  | 14                     | 18             | , BIC           |                                |              |         |  |

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

| D19S433 14 15    PENTAD 11 12     D1S1656 17 17.3 | PENTAE  | GE 7 | 15   |     |     |  |
|---|---------|------|------|-----|-----|--|
|   | D19S433 | 14   | 15   |     |     |  |
| D1S1656 17 17.3                                   | PENTAD  | 11   | 12   |     | ENE |  |
|   | D1S1656 | 17   | 17.3 | BIC |     |  |

4. Conclusion: The STR results of <u>IMR32</u> cells are consistent with the genotypes of <u>IMR32</u> cell lines in <u>DSMZ</u> database, the cell ID corresponded to <u>ACC-165</u>, and the STR results <u>completely matched</u>. In the test, <u>No multi allele was found</u> in this cell line.

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## 5. Attached Image



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