|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 1 | **Sample ID:** | 3-95-NALM-6 | **DSMZ Best Match Cell Name:** | JJN-3 |
| **Database Best Match Score:** | | 76.92% | **DSMZ Best Match Cell Line No:** | ACC-541 |

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Methodology:** | Authenticity for all cell lines used in this study was performed using the Promega GenePrint 10 System and following the protocol described in ANSI/ATCC ASN-0002-2011. The STR alleles were searched on the **DSMZ** Database. | | | | | |
| **Marker Characterization** | | | | | | |
| **Marker** | | **Allele 1** | **Allele 2** | **Allele 3** | **Allele 4** | **DSMZ Best Match Profile** |
| D5S818 | | 11 | 12 |  |  | 11,12 |
| D13S317 | | 9 | 13 |  |  | 9 |
| D7S820 | | 8 | 11 |  |  | 8,9 |
| D16S539 | | 10 |  |  |  | 11,12 |
| vWA | | 15 |  |  |  | 15 |
| TH01 | | 8 | 9 |  |  | 9 |
| AMEL | | X | Y |  |  | X |
| TPOX | | 8 | 10 |  |  | 8,10 |
| CSF1PO | | 12 |  |  |  | 12 |
| D21S11 | | 27 | 29 |  |  |  |

| 1 | **Sample ID:** | 3-95-NALM-6 | **Sample's Electropherogram** |
| --- | --- | --- | --- |
| **Marker Raw Data** | | | |
|  | | | |

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 2 | **Sample ID:** | 3-96-RPMI | **ICLC 2 Best Match Cell Name:** | MDA-MB-436 |
| **Database Best Match Score:** | | 72.73% | **ICLC 2 Best Match Cell Line No:** | HTL11005 |

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Methodology:** | Authenticity for all cell lines used in this study was performed using the Promega GenePrint 10 System and following the protocol described in ANSI/ATCC ASN-0002-2011. The STR alleles were searched on the **ICLC 2** Database. | | | | | |
| **Marker Characterization** | | | | | | |
| **Marker** | | **Allele 1** | **Allele 2** | **Allele 3** | **Allele 4** | **ICLC 2 Best Match Profile** |
| D5S818 | | 12 | 13 |  |  | 13 |
| D13S317 | | 12 |  |  |  | 10 |
| D7S820 | | 8 | 13 |  |  | 10 |
| D16S539 | | 10 | 11 |  |  | 9,11 |
| vWA | | 14 | 20 |  |  | 14,20 |
| TH01 | | 9 | 9.3 |  |  | 9.3 |
| AMEL | | X |  |  |  | X |
| TPOX | | 8 |  |  |  | 8 |
| CSF1PO | | 11 | 12 |  |  | 12 |
| D21S11 | | 30 |  |  |  | 30,31.2 |

| 2 | **Sample ID:** | 3-96-RPMI | **Sample's Electropherogram** |
| --- | --- | --- | --- |
| **Marker Raw Data** | | | |
|  | | | |

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 3 | **Sample ID:** | dddd | **DSMZ Best Match Cell Name:** | Hs 274.T |
| **Database Best Match Score:** | | 71.43% | **DSMZ Best Match Cell Line No:** | CRL-7222 |

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Methodology:** | Authenticity for all cell lines used in this study was performed using the Promega GenePrint 10 System and following the protocol described in ANSI/ATCC ASN-0002-2011. The STR alleles were searched on the **DSMZ** Database. | | | | | |
| **Marker Characterization** | | | | | | |
| **Marker** | | **Allele 1** | **Allele 2** | **Allele 3** | **Allele 4** | **DSMZ Best Match Profile** |
| D5S818 | | 12 | 13 |  |  | 12,13 |
| D13S317 | | 12 |  |  |  | 11 |
| D7S820 | | 8 | 13 |  |  | 11 |
| D16S539 | | 10 | 11 |  |  | 11 |
| vWA | | 14 | 20 |  |  | 14,17 |
| TH01 | | 9 | 9.3 |  |  | 9,9.3 |
| AMEL | | X |  |  |  | X |
| TPOX | | 8 |  |  |  | 8,10 |
| CSF1PO | | 11 | 12 |  |  | 11,12 |
| D21S11 | | 30 |  |  |  |  |

| 3 | **Sample ID:** | dddd | **Sample's Electropherogram** |
| --- | --- | --- | --- |
| **Marker Raw Data** | | | |
|  | | | |