|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 3 | **Sample ID:** | 4-95-24143$pdx13 | **ICLC 2 Best Match Cell Name:** | RPMI 7932 |
| **Database Best Match Score:** | | 66.67% | **ICLC 2 Best Match Cell Line No:** | HTL96012 |

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Methodology:** | Authenticity for all cell lines used in this study was performed using the Promega GenePrint 24 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 |  |  |  | 11,12 |
| D13S317 | | 11 |  |  |  | 11 |
| D7S820 | | 11 |  |  |  | 11,13 |
| D16S539 | | 5 | 12 |  |  | 12 |
| vWA | | 18 | 19 |  |  | 17 |
| TH01 | | 9 |  |  |  | 9,9.3 |
| AMEL | | X |  |  |  | x |
| TPOX | | 8 | 10 |  |  | 8 |
| CSF1PO | | 10 |  |  |  | 10 |
| D21S11 | | 27 | 30 |  |  | 29,30 |
| D10S1248 | | 14 |  |  |  |  |
| D12S391 | | 18 | 22 |  |  |  |
| D18S51 | | 17 |  |  |  |  |
| D19S433 | | 13 | 14 |  |  |  |
| D1S1656 | | 15 | 17.3 |  |  |  |
| D22S1045 | | 11 |  |  |  |  |
| D2S1338 | | 20 |  |  |  |  |
| D2S441 | | 11 | 14 |  |  |  |
| D3S1358 | | 17 |  |  |  |  |
| D8S1179 | | 11 | 14 |  |  |  |
| DYS391 | |  |  |  |  |  |
| FGA | | 25 |  |  |  |  |
| Penta D | | 9 | 13 |  |  |  |
| Penta E | | 13 | 15 |  |  |  |

| 3 | **Sample ID:** | 4-95-24143$pdx13 | **Sample's Electropherogram** |
| --- | --- | --- | --- |
| **Marker Raw Data** | | | |
|  | | | |