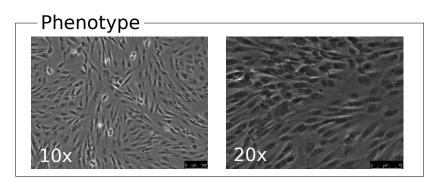
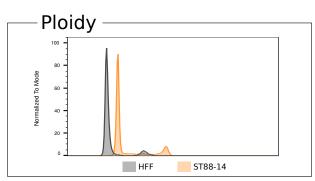
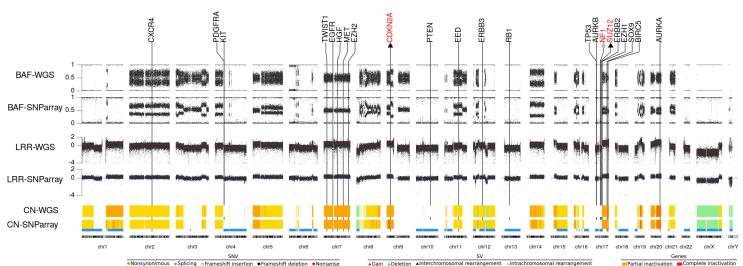
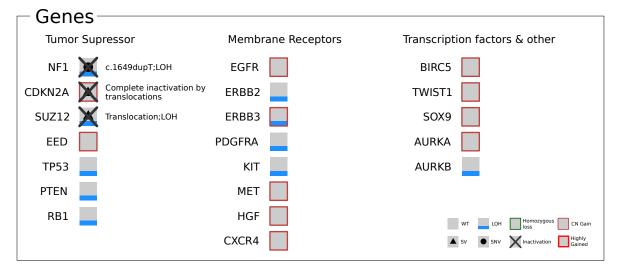
ST88-14

RRID:CVCL_8916









| ⊢STR − | |
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| • | |
| D8S1179 | 14 |
| D21S11 | 29, 32.3 |
| D7S820 | 8 |
| CSF1PO | 9, 12 |
| D3S1358 | 15, 18 |
| TH01 | 9 |
| D13S317 | 12 |
| D16S539 | 13 |
| D2S11338 | 17, 23 |
| D19S433 | 13, 14 |
| vWA | 16 |
| TPOX | 11, 12 |
| D18S51 | 12 |
| AMEL | X, Y |
| D5S818 | 12, 13 |
| FGA | 21 |

- Methylome class -MPNST - Marker expression-

SOX9: Positive SOX10: Negative S100B: Negative

Additional alterations:

Additional information: In some repositories and journal articles the NF1 germline mutation has been erroneously identified as c.910C>T

Aliases: ST88.14; ST 88-14; ST-8814; ST8814; 88-14; NF188-14

Original publication: Fletcher JA, Kozakewich HP, Hoffer FA, Lage JM, Weidner N, Tepper R, Pinkus GS, Morton CC, Corson JM. Diagnostic relevance of clonal cytogenetic aberrations in malignant soft-tissue tumors. N Engl J Med. 1991 Feb 14;324(7):436-42. https://doi.org/10.1056/nejm199102143240702