

Resource Summary Report

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Agilent: TapeStation Laptop

RRID:SCR_019547

Type: Tool

Proper Citation

Agilent: TapeStation Laptop (RRID:SCR_019547)

Resource Information

URL: <https://www.agilent.com/en/product/automated-electrophoresis/tapestation-systems/tapestation-instruments/tapestation-laptop-345933>

Proper Citation: Agilent: TapeStation Laptop (RRID:SCR_019547)

Description: Standardize your data acquisition and analysis by using the Agilent validated TapeStation Laptop together with the TapeStation instrument. The laptop comes with the TapeStation Software and required drivers pre-installed and offers maximum convenience and security.

Resource Type: instrument resource

Keywords: Agilent, TapeStation System, Instrument Equipment, USEDit,

Resource Name: Agilent: TapeStation Laptop

Resource ID: SCR_019547

Record Creation Time: 20220129T080345+0000

Record Last Update: 20260801T190633+0000

Ratings and Alerts

No rating or validation information has been found for Agilent: TapeStation Laptop.

No alerts have been found for Agilent: TapeStation Laptop.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 715 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Lozada JR, et al. (2025) Expression Patterns of DLL3 across Neuroendocrine and Non-neuroendocrine Neoplasms Reveal Broad Opportunities for Therapeutic Targeting. *Cancer research communications*, 5(2), 318.

Lucero K, et al. (2025) CTCF-RNA interactions orchestrate cell-specific chromatin loop organization. *bioRxiv : the preprint server for biology*.

Starble RM, et al. (2025) Epigenetic priming promotes acquisition of tyrosine kinase inhibitor resistance and oncogene amplification in human lung cancer. *bioRxiv : the preprint server for biology*.

Ollinger TL, et al. (2025) Genetic interaction analysis of *Candida glabrata* transcription factors CST6 and UPC2A in the regulation of respiration and fluconazole susceptibility. *Antimicrobial agents and chemotherapy*, 69(2), e0129424.

Kuroha K, et al. (2025) Abnormal H3K27me3 underlies degenerative spermatogonial stem cells in cryptorchid testis. *Development (Cambridge, England)*, 152(2).

Toupenet Marchesi L, et al. (2025) Transcriptomic analysis reinforces the implication of spatacsin in neuroinflammation and neurodevelopment. *Scientific reports*, 15(1), 2370.

Choucair K, et al. (2025) Molecular and immune landscape of tumours in geriatric patients with non-small cell lung cancer, melanoma and renal cell carcinoma. *BMJ oncology*, 4(1), e000551.

Solomon A, et al. (2025) Early and delayed STAT1-dependent responses drive local trained immunity of macrophages in the spleen. *eLife*, 13.

Li J, et al. (2025) Biomolecular condensation of human IDRs initiates endogenous transcription via intrachromosomal looping or high-density promoter localization. *Nucleic acids research*, 53(4).

Parafati M, et al. (2025) Microgravity Accelerates Skeletal Muscle Degeneration: Functional and Transcriptomic Insights from a Muscle Lab-on-Chip Model Onboard the ISS. *bioRxiv : the preprint server for biology*.

Rekhtman N, et al. (2025) Chromothripsis-Mediated Small Cell Lung Carcinoma. *Cancer discovery*, 15(1), 83.

Metzner E, et al. (2025) Multiome Perturb-seq unlocks scalable discovery of integrated perturbation effects on the transcriptome and epigenome. *Cell systems*, 16(1), 101161.

Zawidzka EM, et al. (2025) Tumor-specific CD8⁺ T cells from the bone marrow resist exhaustion and exhibit increased persistence in tumor-bearing hosts as compared with tumor-infiltrating lymphocytes. *Journal for immunotherapy of cancer*, 13(2).

Liu JF, et al. (2025) Decoding m 6 Am by simultaneous transcription-start mapping and methylation quantification. *bioRxiv : the preprint server for biology*.

Spealman P, et al. (2025) Multilevel Gene Expression Changes in Lineages Containing Adaptive Copy Number Variants. *Molecular biology and evolution*, 42(2).

Zhang Y, et al. (2025) EHMT2-mediated R-loop formation promotes the malignant progression of prostate cancer via activating Aurora B. *Clinical and translational medicine*, 15(1), e70164.

Fox CR, et al. (2025) Resistance to complement-mediated lysis of parainfluenza virus 5-infected cells is acquired after transition from acute to persistent infection. *Journal of virology*, 99(2), e0189524.

Estevam GO, et al. (2025) Mapping kinase domain resistance mechanisms for the MET receptor tyrosine kinase via deep mutational scanning. *eLife*, 13.

Funasaki S, et al. (2025) Protocol for transcriptomic and epigenomic analyses of tip-like endothelial cells using scRNA-seq and ChIP-seq. *STAR protocols*, 6(1), 103326.

Zheng W, et al. (2025) Single-cell analysis reveals Mycobacterium tuberculosis ESX-1-mediated accumulation of permissive macrophages in infected mouse lungs. *Science advances*, 11(3), eadq8158.