

Resource Summary Report

Generated by T1D on Aug 27, 2026

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: 20260815T182621+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 717 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Robinson E, et al. (2026) STK11 mutations and deletions define a distinct subtype of cervical adenocarcinoma. medRxiv : the preprint server for health sciences.

Oluwatola O, et al. (2026) Transdermal bicarbonate buffer therapy increases intratumoral pH and elicits antitumor responses in bladder cancer. Frontiers in immunology, 17, 1706250.

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.

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.

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

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.

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.

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

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).

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.

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).

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.

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

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.

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

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.

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.