

Resource Summary Report

Generated by [RRID](#) on Aug 7, 2026

Akoya Biosciences: Vectra 3 Automated Quantitative Pathology Imaging System

RRID:SCR_025828

Type: Tool

Proper Citation

Akoya Biosciences: Vectra 3 Automated Quantitative Pathology Imaging System
(RRID:SCR_025828)

Resource Information

URL: <https://www.akoyabio.com/phenoimager/instruments/vectra-3-0/>

Proper Citation: Akoya Biosciences: Vectra 3 Automated Quantitative Pathology Imaging System (RRID:SCR_025828)

Description: Vectra 3 automated quantitative pathology imaging system, together with the inForm software, provides multispectral and automated multi-slide imaging of up to 6 slides to detect and measure weakly expressed and overlapping biomarkers in H and E, IHC or IF intact FFPE tissue sections or TMAs within familiar digital pathology workflow.

Synonyms: Vectra 3 Automated Quantitative Pathology Imaging System

Resource Type: instrument resource

Keywords: multispectral and automated multi-slide imaging, slide, imaging,

Specification URL: https://raw.githubusercontent.com/SciCrunch/RRID-Instruments/main/PDF/SCR_025828.pdf

Resource Name: Akoya Biosciences: Vectra 3 Automated Quantitative Pathology Imaging System

Resource ID: SCR_025828

Alternate URLs: <https://www.akoyabio.com/wp-content/uploads/2020/03/Vectra-3-Product-Note-1.pdf>

Record Creation Time: 20241001T053245+0000

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Ratings and Alerts

No rating or validation information has been found for Akoya Biosciences: Vectra 3 Automated Quantitative Pathology Imaging System.

No alerts have been found for Akoya Biosciences: Vectra 3 Automated Quantitative Pathology Imaging System.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Savage G, et al. (2026) Phenotypic Characterization and Prognostic Impact of CD103+ Tissue-Resident Memory T Cells in Diffuse Large B-cell Lymphoma. Cancer immunology research, 14(2), 291.

Nguyen TT, et al. (2026) NN-01-195, a Novel Conjugate of HSP90 and AURKA Inhibitors, Effectively Targets Solid Tumors. Molecular cancer therapeutics, 25(8), 1231.

Li C, et al. (2026) ARID1A deficiency unleashes centromeric RNA transcription to drive chromosomal instability and boosts PKMYT1 inhibitor efficacy via RNA sensing. bioRxiv : the preprint server for biology.

Garg SK, et al. (2026) Potentiating dendritic cell immunotherapy by interrupting the Semaphorin 4D-induced immune-suppressive barrier. Cell reports, 45(3), 117050.

Chen K, et al. (2026) Identification of a distinct cluster of LY6E+ macrophages in esophageal squamous cell carcinoma: functional phenotype, spatial interaction, and prognostic significance. British journal of cancer, 135(3), 382.

Garcia GL, et al. (2025) Aged and BRCA-Mutated Stromal Cells Drive Epithelial Cell Transformation. Cancer discovery, 15(6), 1203.