

Resource Summary Report

Generated by SPARC SAWG on Aug 11, 2026

GenScript

RRID:SCR_002891

Type: Tool

Proper Citation

GenScript (RRID:SCR_002891)

Resource Information

URL: <http://www.genscript.com>

Proper Citation: GenScript (RRID:SCR_002891)

Description: Commercial organization which provides life science services and products to researchers. They specialize in gene synthesis, peptide, protein, antibody and preclinical drug development service.

Synonyms: GenScript USA, GenScript USA Inc

Resource Type: commercial organization

Keywords: gene synthesis service, drug development, reagent provider

Resource Name: GenScript

Resource ID: SCR_002891

Alternate IDs: nlx_152374

Alternate URLs: <http://www.dddmag.com/company-profiles/genscript-usa-inc>

Record Creation Time: 20220129T080216+0000

Record Last Update: 20260808T185748+0000

Ratings and Alerts

No rating or validation information has been found for GenScript.

No alerts have been found for GenScript.

Data and Source Information

Usage and Citation Metrics

We found 7134 mentions in open access literature.

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

Martedì R, et al. (2026) De novo cysteine biosynthesis in *Pseudomonas aeruginosa*: Characterization of the two main cysteine synthase isoforms. *iScience*, 29(1), 114304.

Boudjadi S, et al. (2026) Involvement of the FGF8/FGF Receptor Signaling Pathway in the Maintenance and Progression of Fusion-Positive Rhabdomyosarcoma. *Molecular cancer therapeutics*, 25(3), 493.

Rock RR, et al. (2026) Female-enriched *Eggerthella lenta* drives neuroinflammation and IFN- γ via host receptor TLR2. *bioRxiv* : the preprint server for biology.

Bontekoe E, et al. (2026) FOXM1-Specific TCR-Engineered T Cells Target Non-Small Cell Lung Cancer. *Cancer immunology research*, 14(6), 958.

Peirano E, et al. (2026) Trio analysis in dystonia identifies de novo KLC1 variants in a kinesinopathy with distinct motor and neurodevelopmental features. *EBioMedicine*, 129, 106358.

Søndreli KL, et al. (2026) Transcriptomic and functional analyses uncover a conserved effector driving genotype-dependent virulence in the *Sphaerulina musiva*-*Populus trichocarpa* interaction. *mBio*, 17(5), e0312025.

Johnson-Lynn S, et al. (2026) Use of TaqMan array card to investigate bacterial isolates in diabetic foot ulcers. *Journal of medical microbiology*, 75(4).

Preira CMF, et al. (2026) Sporozoite egress from *Plasmodium* oocysts requires a trimeric NF-Y-like complex. *Communications biology*, 9(1).

Altay K, et al. (2026) Application of LAMP and TaqMan qPCR for the rapid diagnosis of *Anaplasma Capra* (an emerging tick-borne zoonotic pathogen) and comparison with Nested-PCR. *Veterinary research communications*, 50(2), 153.

Banna HA, et al. (2026) Structures of the eIF4G-binding RNA domains among picornaviral IRES types are topologically conserved. *Nature communications*, 17(1).

Freiherr von Scholley GL, et al. (2026) Structural analysis of the flexibility of the Ubl2 domain within the papain-like protease of SARS-CoV-2. *Acta crystallographica. Section F, Structural biology communications*, 82(Pt 6), 222.

Ran T, et al. (2026) Genome-Wide Identification of the PLATZ Transcription Factor Family in *Populus euphratica* Oliv. and Functional Characterisation of PePLATZ8 in Drought Tolerance. *Plants (Basel, Switzerland)*, 15(7).

Alharbi S, et al. (2026) Biophysical insights into recombinant Zeocin binding protein: conformational stability and folding dynamics across pH and temperature. *Frontiers in molecular biosciences*, 13, 1748036.

Sharifpour MF, et al. (2026) A New Conditional Transcription Factor for Use in *Toxoplasma Gondii*. *Biotechnology journal*, 21(5), e70240.

Chakraborty S, et al. (2026) Cortisol-resistant CAR-NK cells overcome steroid-induced immunosuppression in lung cancer. *Signal transduction and targeted therapy*, 11(1).

Singh RR, et al. (2026) Transcription activation mechanism of a noncanonical DNA damage response pathway by the WYL-activator, DriD. *Science advances*, 12(12), eaec6337.

Jobe TO, et al. (2026) Multiomics analyses of mutants for *Marchantia polymorpha* FERONIA and MARIS reveal a link between cell wall integrity and abscisic acid responses. *Plant & cell physiology*, 67(6), 959.

Lv J, et al. (2026) Enhancing heritable genome editing in soybean by optimizing promoter combinations for the LbCas12a system. *BMC plant biology*, 26(1).

Wei B, et al. (2026) Dual-mechanism therapy for chronic rhinosinusitis via siRNA targeting of bacterial 16S rRNA and disruption of CFTR-pathogen interactions. *Brazilian journal of otorhinolaryngology*, 92(5), 101851.

Jaswal AP, et al. (2025) Targeting Tumor-Infiltrating Immune Cells for Targeted Alpha Therapy in Gliomas: Optimization of [225Ac]Ac-DOTA- α CD11b Dosing through PET Imaging. *Molecular cancer therapeutics*, 24(12), 1948.