

Resource Summary Report

Generated by [dkNET](#) on Sep 9, 2026

Compugen

RRID:SCR_004631

Type: Tool

Proper Citation

Compugen (RRID:SCR_004631)

Resource Information

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

Proper Citation: Compugen (RRID:SCR_004631)

Description: Drug discovery company providing pharma with protein and antibody therapeutics in the fields of Oncology and Immunology.

Abbreviations: CGEN

Synonyms: Compugen Ltd

Resource Type: commercial organization

Keywords: drug, diagnostic, protein therapeutics, target discovery, therapeutics, oncology, immunology, monoclonal antibody

Resource Name: Compugen

Resource ID: SCR_004631

Alternate IDs: Wikidata: Q5157220, grid.418916.3, nlx_63124, ISNI: 0000 0004 0506 4804

Alternate URLs: <https://ror.org/017vd4789>

Record Creation Time: 20220129T080225+0000

Record Last Update: 20260905T132522+0000

Ratings and Alerts

No rating or validation information has been found for Compugen.

No alerts have been found for Compugen.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 18 mentions in open access literature.

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

Chaudry M, et al. (2023) Simulating flexibility, variability and decentralisation with an integrated energy system model for Great Britain. Scientific reports, 13(1), 4772.

Blechter B, et al. (2023) Polygenic Risk Score, Environmental Tobacco Smoke, and Risk of Lung Adenocarcinoma in Never-Smoking Women in Taiwan. JAMA network open, 6(11), e2339254.

Zhou X, et al. (2022) Alcohol consumption, DNA methylation and colorectal cancer risk: Results from pooled cohort studies and Mendelian randomization analysis. International journal of cancer, 151(1), 83.

Peters BA, et al. (2022) Menopause Is Associated with an Altered Gut Microbiome and Estrobolome, with Implications for Adverse Cardiometabolic Risk in the Hispanic Community Health Study/Study of Latinos. mSystems, 7(3), e0027322.

Silva MDS, et al. (2022) NLRP3 inflammasome activation in human peripheral blood mononuclear cells induced by venoms secreted PLA2s. International journal of biological macromolecules, 202, 597.

Li J, et al. (2021) PVRIG is a novel natural killer cell immune checkpoint receptor in acute myeloid leukemia. Haematologica, 106(12), 3115.

Bi W, et al. (2019) A Fast and Accurate Method for Genome-wide Scale Phenome-wide G × E Analysis and Its Application to UK Biobank. American journal of human genetics, 105(6), 1182.

Nold-Petry CA, et al. (2017) Gp96 Peptide Antagonist gp96-II Confers Therapeutic Effects in Murine Intestinal Inflammation. Frontiers in immunology, 8, 1531.

Liu WW, et al. (2016) Nanodroplet-Vaporization-Assisted Sonoporation for Highly Effective Delivery of Photothermal Treatment. Scientific reports, 6, 24753.

Greenald D, et al. (2015) Genome-wide mapping of Hif-1? binding sites in zebrafish. BMC genomics, 16, 923.

Chao A, et al. (2015) BAI1-Associated Protein 2-Like 1 (BAIAP2L1) Is a Potential Biomarker in Ovarian Cancer. PloS one, 10(7), e0133081.

Gallardo VE, et al. (2013) Fluorescent activated cell sorting (FACS) combined with gene expression microarrays for transcription enrichment profiling of zebrafish lateral line cells. Methods (San Diego, Calif.), 62(3), 226.

Kosmehl T, et al. (2012) A combined DNA-microarray and mechanism-specific toxicity approach with zebrafish embryos to investigate the pollution of river sediments. *Reproductive toxicology* (Elmsford, N.Y.), 33(2), 245.

Kos M, et al. (2010) Yeast pre-rRNA processing and modification occur cotranscriptionally. *Molecular cell*, 37(6), 809.

Gardiner MR, et al. (2007) A global role for zebrafish *klf4* in embryonic erythropoiesis. *Mechanisms of development*, 124(9-10), 762.

Haines AN, et al. (2005) Immunoglobulins in the eggs of the nurse shark, *Ginglymostoma cirratum*. *Developmental and comparative immunology*, 29(5), 417.

Eisenberg E, et al. (2005) Identification of RNA editing sites in the SNP database. *Nucleic acids research*, 33(14), 4612.

Zhou Y, et al. (2003) Database and analyses of known alternatively spliced genes in plants. *Genomics*, 82(6), 584.