

Resource Summary Report

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Dragon Database for Exploration of Ovarian Cancer Genes

RRID:SCR_007621

Type: Tool

Proper Citation

Dragon Database for Exploration of Ovarian Cancer Genes (RRID:SCR_007621)

Resource Information

URL: <http://apps.sanbi.ac.za/ddoc/>

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Description: DDOC provides a comprehensive compilation of the published research related to the genes associated with ovarian cancer. DDOC provides details of the cell line, tissue or cell type, expression status, disease stage, tumor grade, OC type and laboratory method provided in the literature. The links to the relevant sources of data used to extract information related to genes are also included. Many aspects of the information provided in the DDOC were curated by biologists, which increases its accuracy. DDOC is freely accessible for academic and non-profit users.

Synonyms: DDOC

Resource Type: data or information resource, database

Keywords: ovarian cancer, ovarian cancer cell line, ovarian cancer gene, ovarian cancer stages, ovarian tissue

Resource Name: Dragon Database for Exploration of Ovarian Cancer Genes

Resource ID: SCR_007621

Alternate IDs: nif-0000-02742

Record Creation Time: 20220129T080242+0000

Record Last Update: 20260821T193825+0000

Ratings and Alerts

No rating or validation information has been found for Dragon Database for Exploration of Ovarian Cancer Genes.

No alerts have been found for Dragon Database for Exploration of Ovarian Cancer Genes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Xiong C, et al. (2020) ASDmiR: A Stepwise Method to Uncover miRNA Regulation Related to Autism Spectrum Disorder. *Frontiers in genetics*, 11, 562971.

Jin D, et al. (2015) A computational approach to identifying gene-microRNA modules in cancer. *PLoS computational biology*, 11(1), e1004042.

Delfino KR, et al. (2013) Transcription factor-microRNA-target gene networks associated with ovarian cancer survival and recurrence. *PloS one*, 8(3), e58608.

Essack M, et al. (2013) Information exploration system for sickle cell disease and repurposing of hydroxyfasudil. *PloS one*, 8(6), e65190.

Kaur M, et al. (2011) In Silico discovery of transcription factors as potential diagnostic biomarkers of ovarian cancer. *BMC systems biology*, 5, 144.