

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

[PEpiD](#)

RRID:SCR_000235

Type: Tool

Proper Citation

PEpiD (RRID:SCR_000235)

Resource Information

URL: <http://wukong.tongji.edu.cn/pepid>

Proper Citation: PEpiD (RRID:SCR_000235)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 31,2025. A database to store the curated epigenetic data from studies of prostate cancer retrieved by literature mining. The Prostate Epigenetic Database (PEpiD) is meant as a resource for finding previous studies of prostate cancer in humans, mice and rats. Searches can be targeted through the categories of DNA methylation, histone modification, and microRNA.

Synonyms: Prostate Epigenetic Database

Resource Type: data or information resource, database

Defining Citation: [PMID:23696878](#)

Keywords: epigenetic, prostate cancer, dna methylation, histone modification, micro rna, mrna

Related Condition: Prostate Cancer

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: PEpiD

Resource ID: SCR_000235

Alternate IDs: OMICS_01845

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260801T190904+0000

Ratings and Alerts

No rating or validation information has been found for PEpiD.

No alerts have been found for PEpiD.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Singh AN, et al. (2017) Identification of key pathways and genes with aberrant methylation in prostate cancer using bioinformatics analysis. *OncoTargets and therapy*, 10, 4925.

Shi J, et al. (2013) PEpiD: a prostate epigenetic database in mammals. *PloS one*, 8(5), e64289.