

Resource Summary Report

Generated by [ASWG](#) on Sep 1, 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: 20260829T182913+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 [ASWG](#) .

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.