Table 5: Methylation databases

Name	URL	Brief description	Download of Data	Current status
ANCOGeneDB	https://bioinfo.uth.edu/ancogenedb/	Epigenomic, enhancers, and expression quantitative trait loci	Yes	Online
BECon*	https://redgar598.shinyapps.io/BECon/	Interpreting methylation findings from blood in the context of brain	Yes	Online
CMS	http://cbbiweb.uthscsa.edu/KMethylomes/	Analytic functions for cancer methylome datasets	No	Offline
DBCAT	http://dbcat.cgm.ntu.edu.tw/	Methylation profiles of DNA alteration in human cancer	Yes	Online
DiseaseMeth*	http://bio-bigdata.hrbmu.edu.cn/diseasemeth/	Aberrant methylomes of human diseases	No	Online
GED	http://gametsepi.nwsuaflmz.com/	Epigenetic modification of gametogenesis in mammals	Yes	Online
Geneimprint	http://www.geneimprint.com/site/home	Gene imprinting and which allele is expressed	Yes	Online
Lnc2Meth	http://bio-bigdata.hrbmu.edu.cn/Lnc2Meth/	Informs about RNAs and DNA methylation of transcripts	Yes	Online
MeInfoText	http://bws.iis.sinica.edu.tw:8081/MeInfoText2/	Gene methylation and cancers, protein-protein interactions, and biological pathways	Yes	Offline
MethHC	http://methhc.mbc.nctu.edu.tw/	Focuses on aberrant methylomes of human diseases	No	Offline
MethylomeDB	http://epigenomics.columbia.edu/methylomedb/index.html	DNA methylation profiles for human and mouse brains	Yes	Offline
mPod	www.genome.org	Genome-wide tissue-specific DNA methylation profiles	No	Online
PhenoScanner *	http://www.phenoscanner.medschl.cam.ac.uk/	Methylation and human genotype-phenotype associations	Yes	Online
ROADMAP*	https://egg2.wustl.edu/roadmap/web_portal/index.html	Epigenetic modifications and mRNA expression of human cell types and tissues	Yes	Online
TCGA*	https://www.cancer.gov/	Cancer methylation and expression	Yes	Online
TSGene	http://bioinfo.mc.vanderbilt.edu/TSGene/	Methylation status of tumor suppressor genes	No	Offline

^{*}Top tools ranked by the authors.