## **Epigenomics**

## **Software Distribution Tools**

2017-EE-251	2017-EE-258	2017-EE-289	2017-EE-267
4C-ker	TRX-LOGOS	karyoploteR	genomation
Basic4Cseq	LeNup	tracktables	MUSIC
FourCSeq	SUSHI	gtrellis	MSR
fourSig	ATACseqQC	GenomeHubs	RECAP
HTSstation	HMMRATAC	AncestryView	Babelomics
HiFive	MMARGE	chromoMap	GOAtools
ShRec3D	DeepATAC	Squiggle	Categorizer
Sushi.R	chromVAR	MitoZ	myTAI
chromoHBM	cisTopic	ASCIIGenome	GOexpress
InfMod3DGen	Plate-scATAC-seq	kmerPyramid	mvGST
Mango	atacR	Paplot	CHRONOS
diffloop	ATAC-pipe	bulkVis	PathNet
ChiaSig	esATAC	genomeview	BrainCell
ChIP-nexus	ChIPseeker	PrediXcan	NetGSA
PeakXus	metaseq	PhenoNet	EnrichmentBrowser
CexoR	iTagPlot	Phenopredict	ToPASeq
Q-nexus	gplots	PhenotypeSeeker	topologyGSA
chromstaR	CanvasXpress	BaMM!motif	timeClip
hiHMM	Epigenomics in	RTFBSDB	ICoVeR
	JBrowse		
EpiCSeg	MeViz	hiddenDomains	MultiGPS
GenoSTAN	Circleator	Regmex	DChIPRep
Deopen	HiCBrowser	chequeR	SignalSpider
DeepDiff	caOmicsV	LOLA	DESeq2
AltAnalyze	Repliscan	RnBeads	MOABS
scLVM	MultiDataSet	DISMISS	MethylAction