


Public UCSC Trackhub Converter

This is a tool for browsing and converting public UCSC Trackhubs into Quickload data sources that can be loaded in the Integrated Genome Browser.




























Converted Trackhub URL

Use the convert icon () in the table below to convert a UCSC Trackhub to an IGB Quickload.

Browse public UCSC track hubs





#	Hub Name	Actions 	Genome Versions
1	ALFA Hub	 	hg38, hg19
2	Bird assemblies	 	GCF_005870125.1 (Lonchura striata domestica); GCF_013377495.1 (Cygnus atratus); GCF_012460135.1 (Molothrus ater); GCF_000691785.1 (Leptosomus discolor); GCF_003342905.1 (Dromaius novaehollandiae); GCF_000708925.1 (Phalacrocorax carbo); GCF_015832195.1 (Motacilla alba alba); GCF_000691405.1 (Haliaeetus albicilla); GCF_000698965.1 (Struthio camelus australis); GCF_001577835.2 (Coturnix japonica); GCF_004027225.1 (Strigops habroptila); GCF_000385455.1 
3	Blueprint Hub	 	hg38
4	BrainEpigenomeHub	 	hg19
5	Breast Cancer lncRNA	 	hg19
6	Broad Improved Canine Annotation v1	 	canFam3
7	C_elegans_isolates	 	CB4856Princeton_JR-contig, CB4856Princeton_Contigs, CB4856Princeton_cycle20 (Caenorhabditis elegans)
8	CADD	 	hg38, hg19
9	Cancer Genomics Tracks	 	hg19
10	CEMT (CEEHRC)	 	hg19, hg38
11	CESAR Gene Mappings	 	gadMor1, macFas5, susScr3, geoFor1, ailMel1, triMan1, dasNov3, rheMac3, sarHar1, bosTau7, oviAri3, eriEur2, ochPri3, chrPic1, petMar2, tetNig2, melUnd1, oryCun2, hetGla2, vicPac2, musFur1, saiBol1, fr3, speTri2, macEug2, papHam1, gasAcu1, allMis1, ornAna1, danRer7, pteVam1, calJac3, monDom5, anoCar2, otoGar3, equCab2, loxAfr3, nomLeu3, cerSim1, galGal4, taeGut2, melGal1, cavPor3, gorGor3, turTru2, echTel2, rn5, sorAra2, panTro4, felCat5, oreNil2, criGri1, ponAbe2, mm10, 
12	ChIP-seq data track HUBs from MSC cells	 	mm9