





# UCSC Trackhub Converter

The table below lists the public trackhubs made available by UCSC. Browse, search, and copy the converted Quickload (Q) and source UCSC Trackhub (U) URLs. Quickload URLs can be added as data sources in the Integrated Genome Browser to leverage its advanced visualization capabilities.

#	Hub Name	Description	Copy URLs	Genome Versions
1	ALFA Hub	NCBI's Allele Frequency Aggregator (ALFA) allele frequency for variants in dbGaP studies.	<b>Q</b> <b>U</b>	hg19 
2	ChIP-seq data track HUBs from MSC cells from GSE79815	ChIP-seq data from the publication "Epigenetic plasticity drives adipogenic and osteogenic differentiation of marrow-derived mesenchymal stem cells" (10.1074/jbc.M116.736538) from GEO	<b>Q</b> <b>U</b>	GCF_005870125.1 (Lonchura striata domestica) GCF_013377495.1 (Cygnus atratus) GCF_012460135.1 (Molothrus ater)  
3	Coloc segments	Colocalized segments of human genome for Roadmap cell types	<b>Q</b> <b>U</b>	hg19 
...				
...				

Set row height so as to fit both entire hub name and description.

Copy icon letters are larger than the rest of the row content and have colors matching UCSC and Bioviz themes

Show tooltip confirming copy action

Table is wider overall, to better allow for displaying of information by preventing many line breaks in each column.

Set the last copied URL to bold. Font choice WIP.

Users can expand the list of genome versions.