UniSTS

RRID:SCR_006843
Type: Tool

Proper Citation

UniSTS (RRID:SCR_006843)

Resource Information


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Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented August 22, 2016. Database of sequence tagged sites (STSs) derived from STS-based maps and other experiments. STSs are defined by PCR primer pairs and are associated with additional information such as genomic position, genes, and sequences. Chromosome maps are labeled by name of the originating organism, the map title, total markers, total UniSTSSs and links to view maps as well as research documents available through PubMed, another NCBI database. The search functions within UniSTS allow the user to search by gene marker, chromosome, gene symbol and gene description terms to locate markers on specified genes. A representation of the UniSTS datasets is available by ftp. NOTE: All data from this resource have been moved to the Probe database, http://www.ncbi.nlm.nih.gov/probe. You can retrieve all UniSTS records by searching the probe database using the search term unists(properties). (use brackets instead of parenthesis). Additionally, legacy data remain on the NCBI FTP Site in the UniSTS Repository (ftp://ftp.ncbi.nih.gov/pub/ProbeDB/legacy_unists).

Abbreviations: UniSTS

Synonyms: Entrez UniSTS, NCBI UniSTS, UniSTS: Integrating Markers and Maps

Resource Type: data or information resource, database

Keywords: marker, primer sequence, mapping, sequence tagged site, genomic position, gene, sequence, nucleotide, nucleotide sequence, chromosome, gold standard
Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: UniSTS

Resource ID: SCR_006843

Alternate IDs: nif-0000-03614

Ratings and Alerts

No rating or validation information has been found for UniSTS.

No alerts have been found for UniSTS.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 39 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.


