

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Ancient conserved untranslated sequences

RRID:SCR_008130

Type: Tool

Proper Citation

Ancient conserved untranslated sequences (RRID:SCR_008130)

Resource Information

URL: <http://pbil.univ-lyon1.fr/acuts/ACUTS.html>

Proper Citation: Ancient conserved untranslated sequences (RRID:SCR_008130)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, Documented on August 12, 2014. Database that identifies new regulatory elements in untranslated regions of protein-coding genes (5 prime flanks, 5 prime UTRs, introns, 3 prime UTRs and 3 prime flanks). The analyses is focused on genes from metazoan species (essentially vertebrates, insects and nematodes). Information on highly conserved regions (sequences, alignments, annotations, bibliographic references) are compiled. Currently 176 out of 326 detected highly conserved regions (HCRs) have been analyzed and incorporated in the database. You can also access the list of annotated conserved elements and the list of conserved elements that remain to be processed. Their approach is based on comparative sequence analysis, for the identification of phylogenetic footprints.

Abbreviations: ACUTS

Resource Type: data or information resource, database

Defining Citation: [PMID:9204283](#)

Keywords: echinoderm, footprint, fragment, functional, gene, alignment, analysis, annotation, chordate, cis-element, coding, degradation, divergence, dna, dnase, highly conserved region, homologous, intron, metazoan, mrna, non-coding, nucleotide, phylogenetic, post-transcriptional, promoter, protein, region, regulatory, segment, sequence, structural, transcriptional repressor, translation, untranslated region

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Ancient conserved untranslated sequences

Resource ID: SCR_008130

Alternate IDs: nif-0000-20934

Record Creation Time: 20220129T080245+0000

Record Last Update: 20260905T133150+0000

Ratings and Alerts

No rating or validation information has been found for Ancient conserved untranslated sequences.

No alerts have been found for Ancient conserved untranslated sequences.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.