

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

GenomeTraFaC

RRID:SCR_007686

Type: Tool

Proper Citation

GenomeTraFaC (RRID:SCR_007686)

Resource Information

URL: <http://genometrafac.cchmc.org>

Proper Citation: GenomeTraFaC (RRID:SCR_007686)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 29,2022. Whole genome resource for the detection of transcription factor binding site clusters associated with conventional and microRNA encoding genes conserved between mouse and human gene orthologs

Synonyms: GenomeTraFaC

Resource Type: data or information resource, database

Defining Citation: [PMID:17178752](#)

Keywords: gene orthologs, human mrna, mouse mrna

Availability: THIS RESOURCE IS NO LONGER IN SERVICE.

Resource Name: GenomeTraFaC

Resource ID: SCR_007686

Alternate IDs: nif-0000-02902

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260811T043318+0000

Ratings and Alerts

No rating or validation information has been found for GenomeTraFaC.

No alerts have been found for GenomeTraFaC.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Miller SJ, et al. (2009) Integrative genomic analyses of neurofibromatosis tumours identify SOX9 as a biomarker and survival gene. EMBO molecular medicine, 1(4), 236.

Kaiser S, et al. (2007) Transcriptional recapitulation and subversion of embryonic colon development by mouse colon tumor models and human colon cancer. Genome biology, 8(7), R131.