

# Resource Summary Report

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## CREMOFAC: A web-database of Chromatin Remodeling Factors

RRID:SCR\_007613

Type: Tool

### Proper Citation

CREMOFAC: A web-database of Chromatin Remodeling Factors (RRID:SCR\_007613)

### Resource Information

**URL:** <http://www.jncasr.ac.in/cremofac/>

**Proper Citation:** CREMOFAC: A web-database of Chromatin Remodeling Factors (RRID:SCR\_007613)

**Description:** CREMOFAC is a database for chromatin remodeling factors has been developed. The database harbors 64 types of remodeling factors from 49 different organisms reported in literature and facilitates a comprehensive search for them. In addition, it also provides in-depth information for the factors reported in the three widely studied mammals namely, human, mouse and rat. Further, information on literature, pathways, and phylogenetic relationships has also been covered.

**Synonyms:** CREMOFAC

**Resource Type:** data or information resource, database

**Keywords:** chromatin, chromatin remodeling factor, human chromatin, mouse chromatin, rat chromatin

**Resource Name:** CREMOFAC: A web-database of Chromatin Remodeling Factors

**Resource ID:** SCR\_007613

**Alternate IDs:** nif-0000-02697

**Record Creation Time:** 20220129T080242+0000

**Record Last Update:** 20260829T183003+0000

### Ratings and Alerts

No rating or validation information has been found for CREMOFAC: A web-database of Chromatin Remodeling Factors.

No alerts have been found for CREMOFAC: A web-database of Chromatin Remodeling Factors.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Ignatieva EV, et al. (2016) A compendium of human genes regulating feeding behavior and body weight, its functional characterization and identification of GWAS genes involved in brain-specific PPI network. BMC genetics, 17(Suppl 3), 158.

Milanowska K, et al. (2011) Databases and bioinformatics tools for the study of DNA repair. Molecular biology international, 2011, 475718.