

Resource Summary Report

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Highly Expressed Genes Database (HEG-DB)

RRID:SCR_007704

Type: Tool

Proper Citation

Highly Expressed Genes Database (HEG-DB) (RRID:SCR_007704)

Resource Information

URL: <https://omictools.com/heg-db-tool>

Proper Citation: Highly Expressed Genes Database (HEG-DB) (RRID:SCR_007704)

Description: Genomic database that includes prediction of which genes are highly expressed in prokaryotic complete genomes under strong translational selection.

Synonyms: HEG-DB

Resource Type: data or information resource, database

Keywords: gene expression prediction, prokaryote, prokaryotic genome

Resource Name: Highly Expressed Genes Database (HEG-DB)

Resource ID: SCR_007704

Alternate IDs: nif-0000-02948

Old URLs: <http://genomes.urv.cat/HEG-DB>

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260811T043315+0000

Ratings and Alerts

No rating or validation information has been found for Highly Expressed Genes Database (HEG-DB).

No alerts have been found for Highly Expressed Genes Database (HEG-DB).

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Sharma V, et al. (2014) Analysis of tetra- and hepta-nucleotides motifs promoting -1 ribosomal frameshifting in Escherichia coli. Nucleic acids research, 42(11), 7210.