

Resource Summary Report

Generated by [NIF](#) on Aug 14, 2026

R/FEST

RRID:SCR_013347

Type: Tool

Proper Citation

R/FEST (RRID:SCR_013347)

Resource Information

URL: <http://folk.uio.no/thoree/FEST/>

Proper Citation: R/FEST (RRID:SCR_013347)

Description: An R package for simulations and likelihood calculations of pair-wise family relationships using DNA marker data. (entry from Genetic Analysis Software)

Synonyms: FEST

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, r

Resource Name: R/FEST

Resource ID: SCR_013347

Alternate IDs: nlx_154111, SCR_000830, nlx_154582

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260813T055046+0000

Ratings and Alerts

No rating or validation information has been found for R/FEST.

No alerts have been found for R/FEST.

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 [NIF](#) .

Apavaloaei A, et al. (2025) Tumor antigens preferentially derive from unmutated genomic sequences in melanoma and non-small cell lung cancer. Nature cancer, 6(8), 1419.

Kina E, et al. (2024) Breast cancer immunopeptidomes contain numerous shared tumor antigens. The Journal of clinical investigation, 134(1).