

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

ENTROPY BLOCKER

RRID:SCR_000123

Type: Tool

Proper Citation

ENTROPY BLOCKER (RRID:SCR_000123)

Resource Information

URL: <http://wpicr.wpic.pitt.edu/WPICCompGen/blocks.htm>

Proper Citation: ENTROPY BLOCKER (RRID:SCR_000123)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 31,2025. Software application aiming at identifying haplotype blocks. The likelihood of the data is calculated minus the model complexity. The resulting blocks have very low diversity and the linkage disequilibrium with SNP's outside the blocks is low. (entry from Genetic Analysis Software)

Abbreviations: ENTROPY BLOCKER

Synonyms: R/ENTROPY BLOCKER, R/ENTROPY_BLOCKER

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, r, ms-windows, linux

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ENTROPY BLOCKER

Resource ID: SCR_000123

Alternate IDs: nlx_154300, nlx_154581, SCR_007247

Record Creation Time: 20220129T080159+0000

Record Last Update: 20260919T195625+0000

Ratings and Alerts

No rating or validation information has been found for ENTROPY BLOCKER.

No alerts have been found for ENTROPY BLOCKER.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.