

# Resource Summary Report

Generated by RRID on Aug 5, 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:** 20260805T044003+0000

### Ratings and Alerts

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

No alerts have been found for ENTROPY BLOCKER.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.