## **Resource Summary Report**

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# XL-mHG

RRID:SCR\_016846 Type: Tool

**Proper Citation** 

XL-mHG (RRID:SCR\_016846)

### **Resource Information**

URL: https://github.com/flo-compbio/xlmhg

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**Description:** Software Python package as a semiparametric test for enrichment in ranked lists. Used for determining gene set enrichment.

Abbreviations: xlmhg

**Synonyms:** xlminimumhypergeometric, XL-minimum HyperGeometric test, XL-minimum HyperGeometric, xlmhg, XL-mHG

**Resource Type:** software application, software resource, data analysis software, data processing software

Keywords: semiparametric, test, enrichment, ranked, list, gene, bio.tools

#### **Funding:**

Availability: Free, Available for download, Freely available

Resource Name: XL-mHG

Resource ID: SCR\_016846

Alternate IDs: BioTools:XL-mHG, biotools:XL-mHG

Alternate URLs: https://bio.tools/XL-mHG, https://bio.tools/XL-mHG, https://bio.tools/XL-mHG

License: GNU General Public License v3.0

Record Creation Time: 20220129T080332+0000

Record Last Update: 20250517T060306+0000

### **Ratings and Alerts**

No rating or validation information has been found for XL-mHG.

No alerts have been found for XL-mHG.

### 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 <u>dkNET</u>.

Singer M, et al. (2016) A Distinct Gene Module for Dysfunction Uncoupled from Activation in Tumor-Infiltrating T Cells. Cell, 166(6), 1500.