

Resource Summary Report

Generated by T1D on Aug 1, 2026

EHPLUS

RRID:SCR_009159

Type: Tool

Proper Citation

EHPLUS (RRID:SCR_009159)

Resource Information

URL: <http://www.mrc-epid.cam.ac.uk/Personal/jinghua.zhao/software/>

Proper Citation: EHPLUS (RRID:SCR_009159)

Description: THIS RESOURCE IS NO LONGER IN SERVCE, documented September 22, 2016.

Abbreviations: EHPLUS

Synonyms: EH, FASTEHPPLUS

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c, unix, solaris, dec alpha, ms-windows

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EHPLUS

Resource ID: SCR_009159

Alternate IDs: nlx_154295

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260731T042809+0000

Ratings and Alerts

No rating or validation information has been found for EHPLUS.

No alerts have been found for EHPLUS.

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 [T1D](#) .

Razi F, et al. (2019) Haplotypes in vitamin D receptor gene encode risk in diabetic nephropathy. *Gene*, 683, 149.

Lu X, et al. (2009) Variants in the insulin-degrading enzyme gene are associated with metabolic syndrome in Chinese elders. *Metabolism: clinical and experimental*, 58(10), 1465.