EWAS Catalog Eligibility Criteria

# PubMed search terms

"epigenome-wide" OR "epigenome wide" OR "EWAS" OR "genome-wide AND methylation" OR "genome wide AND methylation"

# Study eligibility

## Study inclusion

Studies are eligible for inclusion in the EWAS Catalog if:

* Include at least 100,000 CpG sites in the analysis.
* Have a sample size of at least 100 individuals.
* Have studied DNA methylation data.

## Study exclusion

Studies are not eligible for inclusion in the EWAS Catalog if:

* The DNA methylation data studied was not genome-wide.
* The study does not include any new EWAS data (e.g. a review of EWAS on smoking).
* Not used human samples.

# PubMed results

The PubMed search results are shown in an excel spreadsheet with the columns: PMID, Author (first author), Year (year of publication), Date (date of publication), Title, Abstract, Include.

Include column key:

* Y (Yes) – extract data and include in the catalog.
* M (Maybe) – paper is relevant but does not satisfy all of the inclusion/exclusion criteria (e.g. N<100).
* R (Review) – the paper is a relevant review but is unlikely to contain novel results.
* N (No) – paper is not relevant to the catalog.