

Resource Summary Report

Generated by [NIF](#) on Aug 18, 2026

[XPN](#)

RRID:SCR_008845

Type: Tool

Proper Citation

XPN (RRID:SCR_008845)

Resource Information

URL: <https://genome.unc.edu/xpn/>

Proper Citation: XPN (RRID:SCR_008845)

Description: Merging Two Gene Expression Studies via Cross Platform Normalization.

Abbreviations: XPN

Resource Type: software resource

Keywords: bio.tools

Resource Name: XPN

Resource ID: SCR_008845

Alternate IDs: OMICS_00863, biotools:xpn

Alternate URLs: <https://bio.tools/xpn>

Record Creation Time: 20220129T080249+0000

Record Last Update: 20260815T182353+0000

Ratings and Alerts

No rating or validation information has been found for XPN.

No alerts have been found for XPN.

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 [NIF](#) .

Feldman S, et al. (2024) Comparison and development of cross-study normalization methods for inter-species transcriptional analysis. PloS one, 19(9), e0307997.

Taminau J, et al. (2012) Unlocking the potential of publicly available microarray data using inSilicoDb and inSilicoMerging R/Bioconductor packages. BMC bioinformatics, 13, 335.