

# Resource Summary Report

Generated by [RRID](#) on Aug 6, 2026

## [XPN](#)

RRID:SCR\_008845

Type: Tool

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### Proper Citation

XPN (RRID:SCR\_008845)

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### 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:** 20260801T190354+0000

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### Ratings and Alerts

No rating or validation information has been found for XPN.

No alerts have been found for XPN.

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### 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 [RRID](#) .

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.