

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

ProbeSelect

RRID:SCR_012965

Type: Tool

Proper Citation

ProbeSelect (RRID:SCR_012965)

Resource Information

URL: <http://sourceforge.net/projects/probeselect/>

Proper Citation: ProbeSelect (RRID:SCR_012965)

Description: Software for selecting probes in heterogenous transcriptional sets.

Abbreviations: ProbeSelect

Synonyms: ProbeSelect - Selecting probes in heterogenous transcriptional sets

Resource Type: software resource

Resource Name: ProbeSelect

Resource ID: SCR_012965

Alternate IDs: OMICS_00836

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260905T132730+0000

Ratings and Alerts

No rating or validation information has been found for ProbeSelect.

No alerts have been found for ProbeSelect.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Franklin A, et al. (2025) Cell-type-specific DNA methylation dynamics in the prenatal and postnatal human cortex. *Cell genomics*, 5(12), 101010.

Falaleeva M, et al. (2015) SNORD116 and SNORD115 change expression of multiple genes and modify each other's activity. *Gene*, 572(2), 266.

De Conti L, et al. (2015) TDP-43 affects splicing profiles and isoform production of genes involved in the apoptotic and mitotic cellular pathways. *Nucleic acids research*, 43(18), 8990.

Maslon MM, et al. (2014) The translational landscape of the splicing factor SRSF1 and its role in mitosis. *eLife*, 3, e02028.

Convertini P, et al. (2013) Genome wide array analysis indicates that an amyotrophic lateral sclerosis mutation of FUS causes an early increase of CAMK2N2 in vitro. *Biochimica et biophysica acta*, 1832(8), 1129.