

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

ROSO

RRID:SCR_010967

Type: Tool

Proper Citation

ROSO (RRID:SCR_010967)

Resource Information

URL: <http://pbil.univ-lyon1.fr/roso/help.php>

Proper Citation: ROSO (RRID:SCR_010967)

Description: A software to design optimized oligonucleotide probes (size over 25 nucleotides) for microarrays.

Abbreviations: ROSO

Synonyms: ROSO - A friend for your design, Recherche et Optimisation de Sondes Oligonucléotidiques, Recherche et Optimisation de Sondes Oligonucleotidiques

Resource Type: software resource

Defining Citation: [PMID:14734320](#)

Availability: Free to non-profit users, On request

Resource Name: ROSO

Resource ID: SCR_010967

Alternate IDs: OMICS_00838

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260905T132651+0000

Ratings and Alerts

No rating or validation information has been found for ROSO.

No alerts have been found for ROSO.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Abot A, et al. (2016) CAZyChip: dynamic assessment of exploration of glycoside hydrolases in microbial ecosystems. BMC genomics, 17(1), 671.