

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

University of Oxford; Oxford; United Kingdom

RRID:SCR_006361

Type: Tool

Proper Citation

University of Oxford; Oxford; United Kingdom (RRID:SCR_006361)

Resource Information

URL: <http://www.ox.ac.uk/>

Proper Citation: University of Oxford; Oxford; United Kingdom (RRID:SCR_006361)

Description: Collegiate research university in Oxford, England. Teaching as early as 1096, making it the oldest university in English speaking world and world second oldest university in continuous operation.

Abbreviations: Oxford

Synonyms: Oxford University

Resource Type: university

Resource Name: University of Oxford; Oxford; United Kingdom

Resource ID: SCR_006361

Alternate IDs: Crossref funder ID:501100000769, Wikidata:Q34433, ISNI:0000 0004 1936 8948, grid.4991.5, nlx_59631

Alternate URLs: <https://ror.org/052gg0110>

Record Creation Time: 20220129T080235+0000

Record Last Update: 20260725T190622+0000

Ratings and Alerts

No rating or validation information has been found for University of Oxford; Oxford; United Kingdom.

No alerts have been found for University of Oxford; Oxford; United Kingdom.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Unitt A, et al. (2025) Neisseria gonorrhoeae LIN codes provide a robust, multi-resolution lineage nomenclature. eLife, 14.

Gao Y, et al. (2022) Lipid-mediated phase separation of AGO proteins on the ER controls nascent-peptide ubiquitination. Molecular cell, 82(7), 1313.

Baker H, et al. (2022) Information sharing practices during the COVID-19 pandemic: A case study about face masks. PloS one, 17(5), e0268043.

McDonald A, et al. (2019) Paediatric neurodisability and sleep disorders: clinical pathways and management strategies. BMJ paediatrics open, 3(1), e000290.

De Maio N, et al. (2018) Bayesian reconstruction of transmission within outbreaks using genomic variants. PLoS computational biology, 14(4), e1006117.

Klebl BM, et al. (2004) Stacking up the armory against viruses. Drug discovery today, 9(4), 162.