Wellcome Trust Sanger Institute; Hinxton; United Kingdom

RRID:SCR_011784
Type: Tool

Proper Citation

Wellcome Trust Sanger Institute; Hinxton; United Kingdom (RRID:SCR_011784)

Resource Information

**URL:** [http://www.sanger.ac.uk/](http://www.sanger.ac.uk/)

**Proper Citation:** Wellcome Trust Sanger Institute; Hinxton; United Kingdom (RRID:SCR_011784)

**Description:** Non profit research organization for genome sequences to advance understanding of biology of humans and pathogens in order to improve human health globally. Provides data which can be translated for diagnostics, treatments or therapies including over 100 finished genomes, which can be downloaded. Data are publicly available on limited basis, and provided more extensively upon request.

**Abbreviations:** WTSI, Sanger

**Synonyms:** The Wellcome Sanger Institute, Sanger Institute, Wellcome Trust Sanger Institute, Genome Research Limited, Wellcome Trust Sanger Institute Genome Research Limited

**Resource Type:** institution

**Keywords:** research, genome, sequence, human, health, project, global, data, treatment, therapy

**Funding Agency:** Wellcome Trust

**Resource Name:** Wellcome Trust Sanger Institute; Hinxton; United Kingdom

**Resource ID:** SCR_011784
Alternate IDs: nlx_91258, grid.10306.34, ISNI: 0000 0004 0606 5382, Wikidata: Q1142544

Alternate URLs: https://ror.org/05cy4wa09

Ratings and Alerts

No rating or validation information has been found for Wellcome Trust Sanger Institute; Hinxton; United Kingdom.

No alerts have been found for Wellcome Trust Sanger Institute; Hinxton; United Kingdom.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 504 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

Kotler O, et al. (2023) SUMOylation of NaV1.2 channels regulates the velocity of backpropagating action potentials in cortical pyramidal neurons. eLife, 12.


Zou X, et al. (2022) Mammalian splicing divergence is shaped by drift, buffering in, and a scaling law. Life science alliance, 5(4).


Matsushita K, et al. (2021) The role of Sp140 revealed in IgE and mast cell responses in Collaborative Cross mice. JCI insight, 6(12).

Candler T, et al. (2021) DNA methylation at a nutritionally sensitive region of the gene is associated with thyroid volume and function in Gambian children. Science advances, 7(45), eabj1561.


