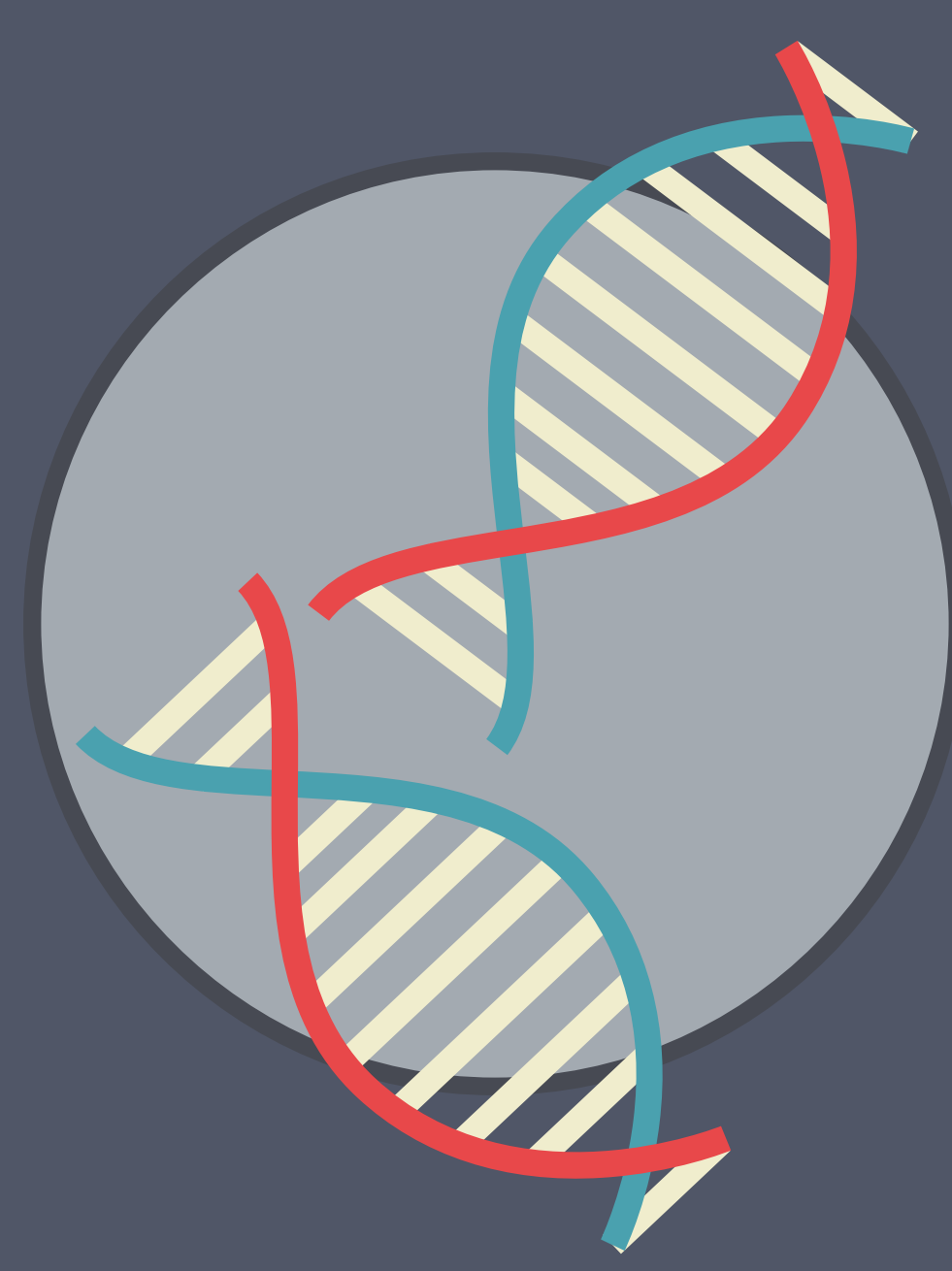


# PROFESSOR DES HIGGINS

## IMPACT JOURNEY



In 1988, Professor Des Higgins developed a computer programme called Clustal that could quickly compare sequences of genetic information.

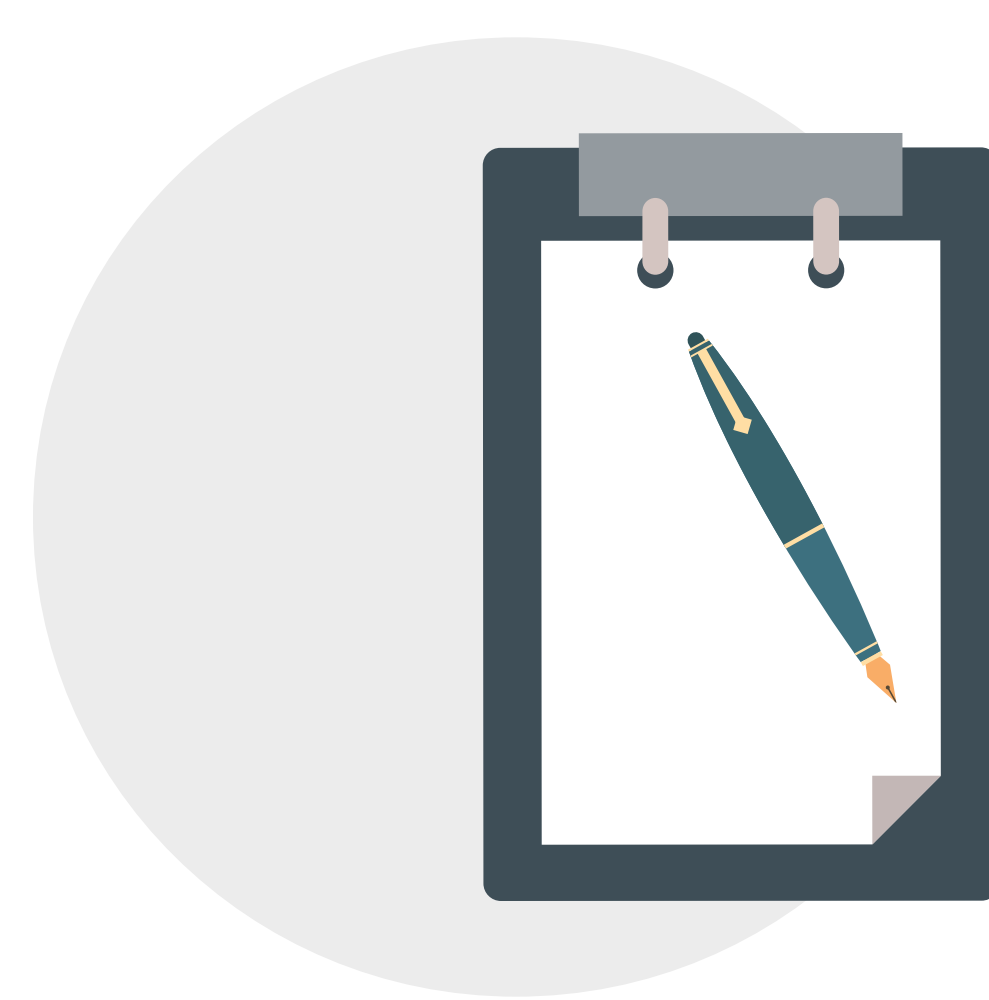
It is now a global standard, used hundreds of times a day by scientists addressing real-world challenges. As a result, Professor Higgins' Clustal publications are among the most highly cited in the world.

### INPUTS

01

Existing knowledge

Challenge: Tired of comparing sequences of genetic information by hand

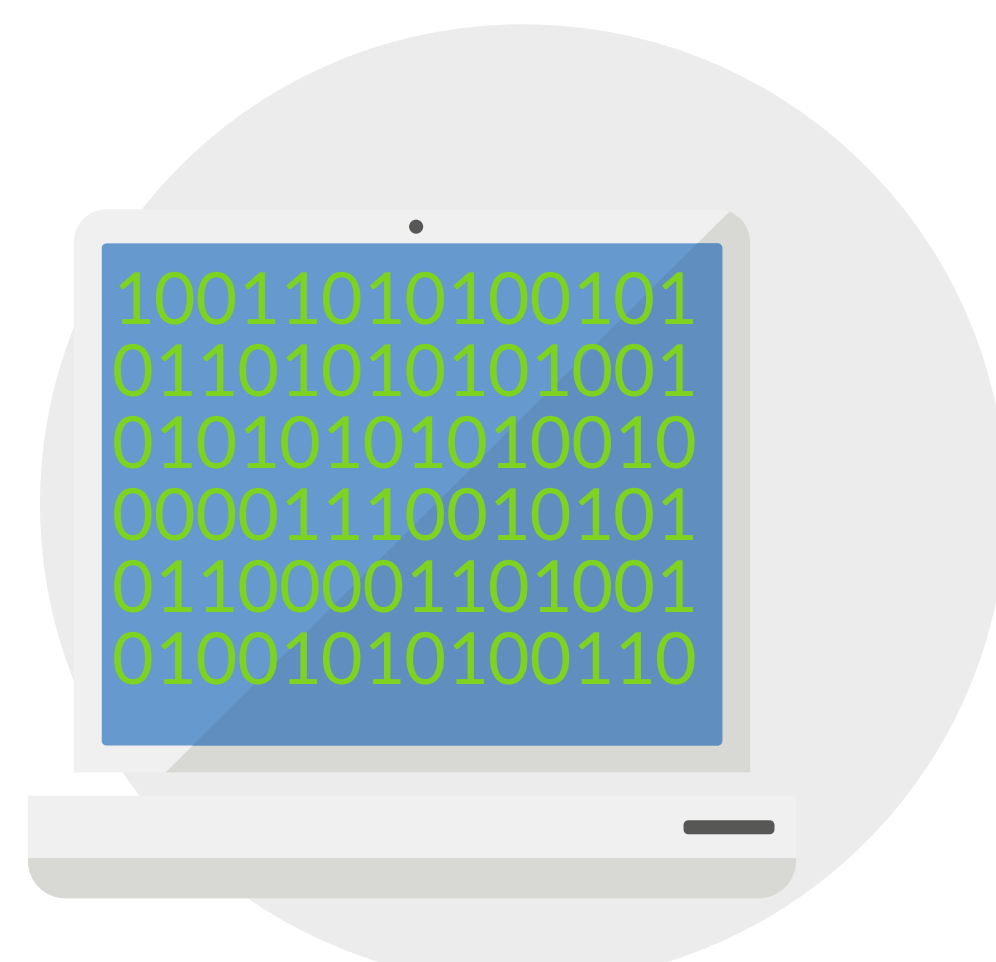


### ACTIVITIES

02

Developing 'Clustal', a computer programme able to quickly compare large amounts of genetic information

Collaborating with other researchers on more and more powerful versions of the programme

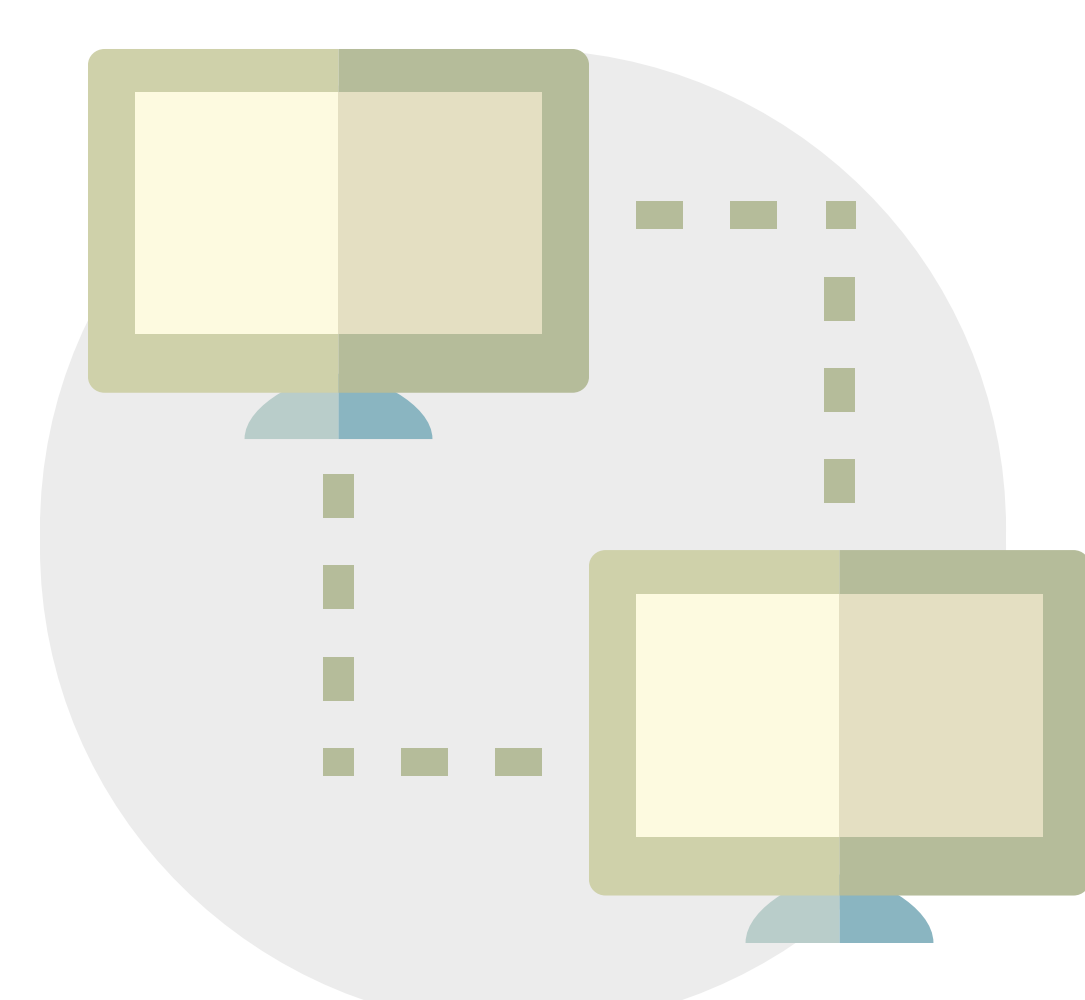


### OUTPUTS

03

Clustal software, made freely available to all

A series of academic publications describing different versions of the programme



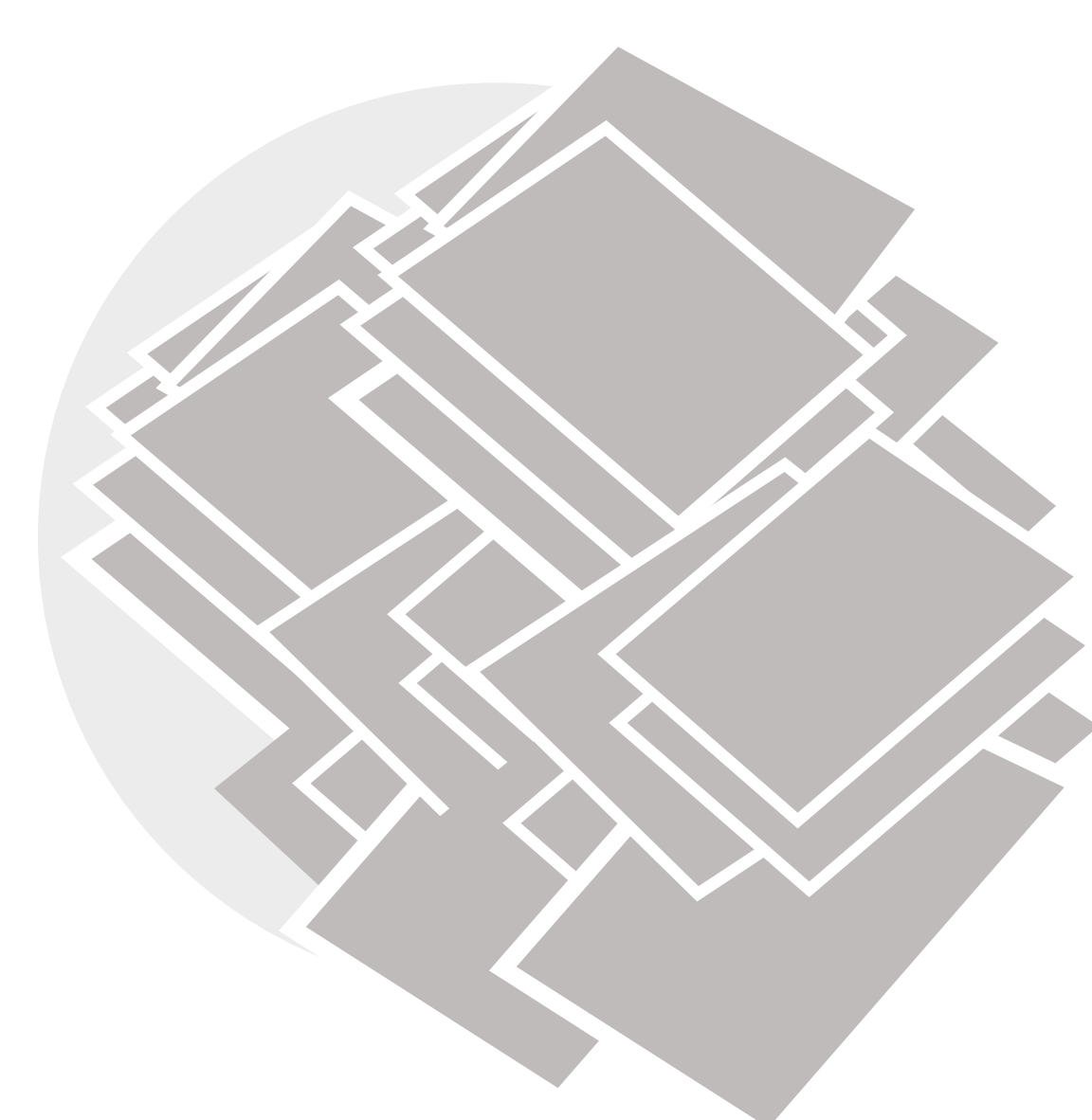
### OUTCOMES

04

Clustal shared widely among the scientific community, initially on floppy disk

Over 150,000 citations, with one paper in the top ten most cited of all time

Clustal hosted on large servers, like the one at the European Bioinformatics Institute



### IMPACTS

05



#### ACADEMIC

Clustal is a global standard, used hundreds of times a day



#### SOCIAL

Clustal used to address various real-world problems, including:

- Tracking infectious diseases
- Producing biofuels
- Creating disease-resistant plants



#### ECONOMIC

Clustal used by companies to make genetic comparison vastly more efficient

Cited in over 20,000 patent documents

[Read the full case study here](#)