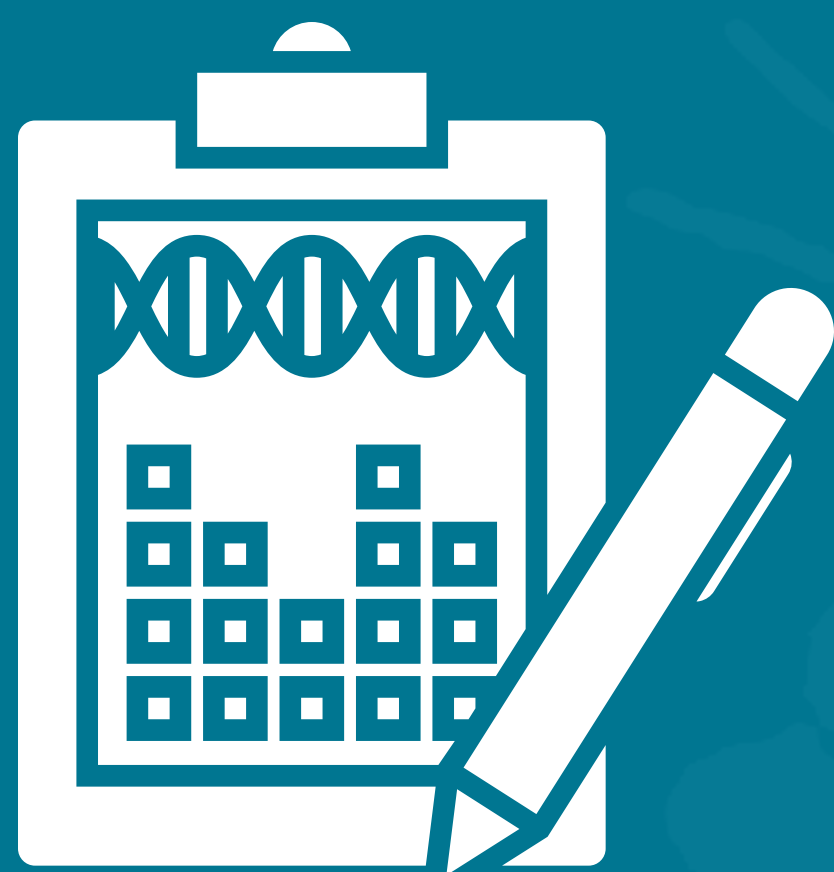


Automated review



The **Wellcome Sanger Institute** sequences each genome and collates information to form the basis of a Genome Note.



A report is created about each genome sequence including **key metrics on genome continuity, base accuracy and functional completeness**.



Both the Genome Note and metrics report are sent to **Wellcome Open Research** as an XML file.



The file is checked by the editorial team and is published as a **Genome Note**. The metrics report is published alongside the article as an automated review.

Community review



Peer reviewers are invited to review the **Genome Notes**. They can use the **automated review** to assess the article's quality.



Peer reviewers submit their reviewer report.



A **Genome Note** has passed peer review when it has one **automated review** and one **“Approved”** peer reviewer report.