



Welcome to Cromwell

Cromwell is an open-source Workflow Management System for bioinformatics. Licensing is BSD 3-Clause.

The Cromwell documentation has a dedicated site.

First time to Cromwell? Get started with Tutorials.

Community

Thinking about contributing to Cromwell? Get started by reading our Contributor Guide.

Cromwell has a growing ecosystem of community-backed projects to make your experience even better! Check out our Ecosystem page to learn more.

Talk to us: - Join the Cromwell Slack workspace to discuss the Cromwell workflow engine. - Join the OpenWDL Slack workspace to discuss the evolution of the WDL language itself. - More information about WDL is available in that project's repository.

Capabilities and roadmap

Many users today run their WDL workflows in Terra, a managed cloud bioinformatics platform with built-in WDL support provided by Cromwell. See [here](#) for a quick-start guide.

Users with specialized needs who wish to install and maintain their own Cromwell instances can download a JAR or Docker image. The development team accepts reproducible bug reports from self-managed instances, but cannot feasibly provide direct support.

Cromwell's backends receive development resources proportional to user demand. The team is actively developing for Google Cloud and Microsoft Azure (see [Cromwell on Azure](#)). Maintenance of other backends is primarily community-based.

Cromwell supports the WDL workflow language. Cromwell version 80 and above no longer support CWL.

CWL will be re-introduced at a later date in the Terra platform, using a solution other than Cromwell. See the blog post "[Terra's roadmap to supporting more workflow languages](#)" for details.

Security reports

If you believe you have found a security issue please contact infosec@broadinstitute.org.

Issue tracking

Need to file an issue? Head over to Github Issues.

If you previously filed an issue in JIRA, the link is [here](#). New signups are no longer accepted.

