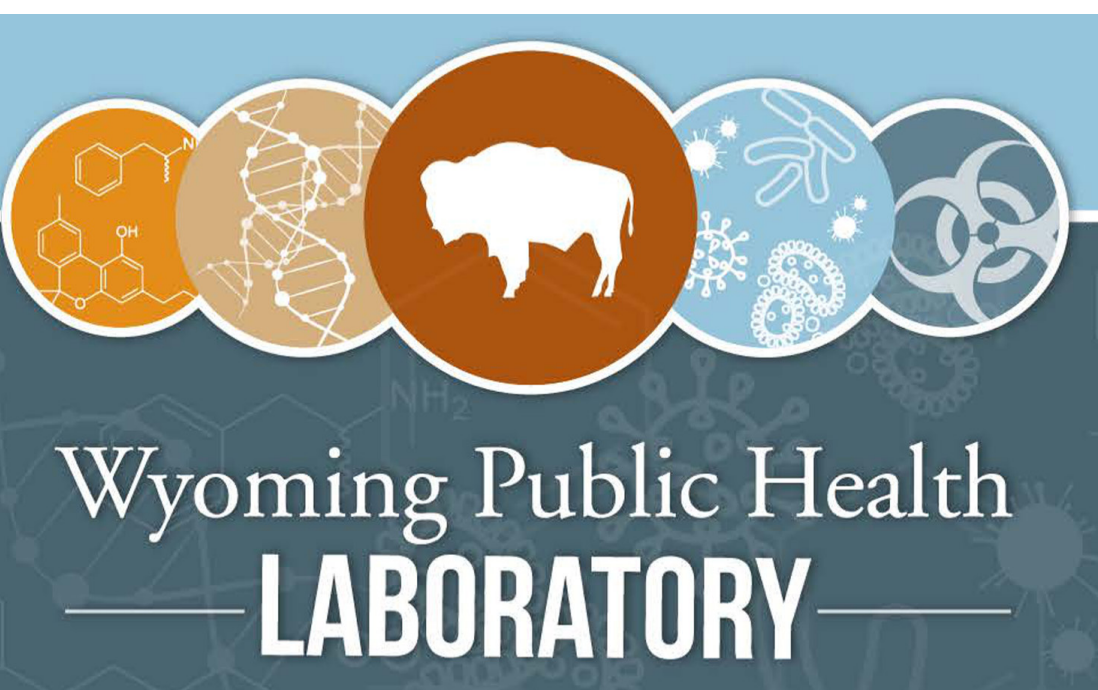


Supporting the bioinformatics community by supporting users of Bactopia



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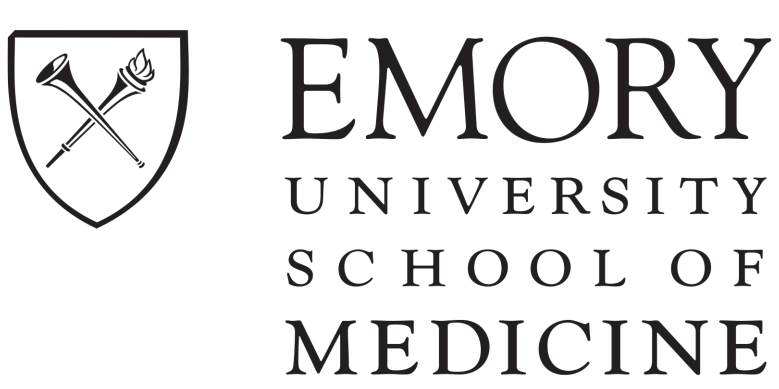
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[@rpetit3](https://twitter.com/rpetit3)

[@rpetit3](https://github.com/rpetit3)

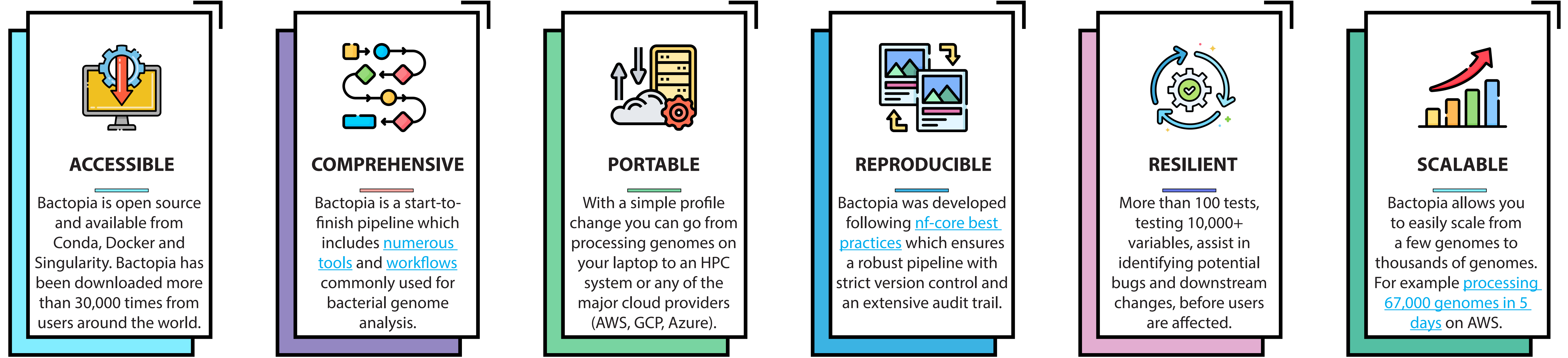
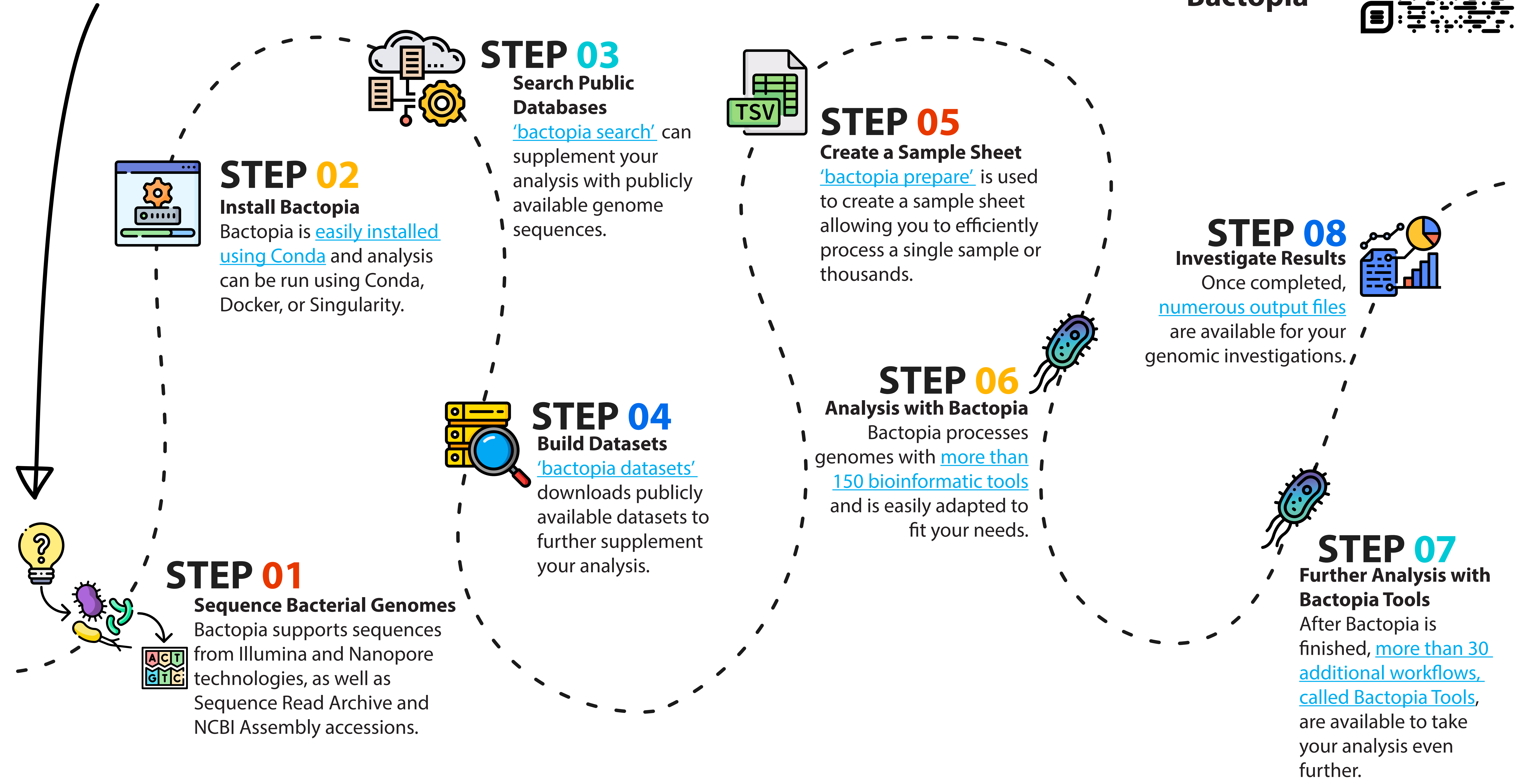
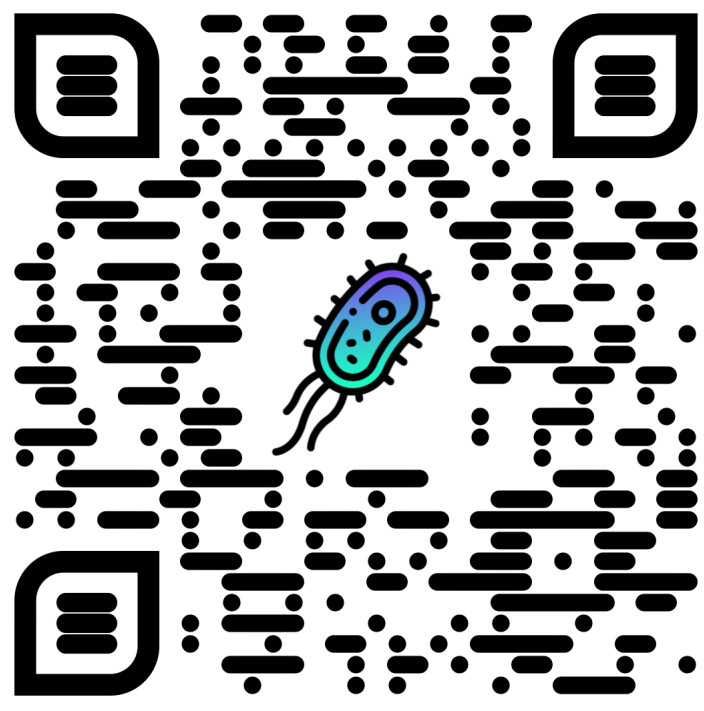
Bactopia Documentation: bactopia.github.io



What is Bactopia?

Bactopia is an extensive Nextflow pipeline for the complete analysis of bacterial genomes. To learn more, follow the step-by-step guide below.

Scan to get started with Bactopia

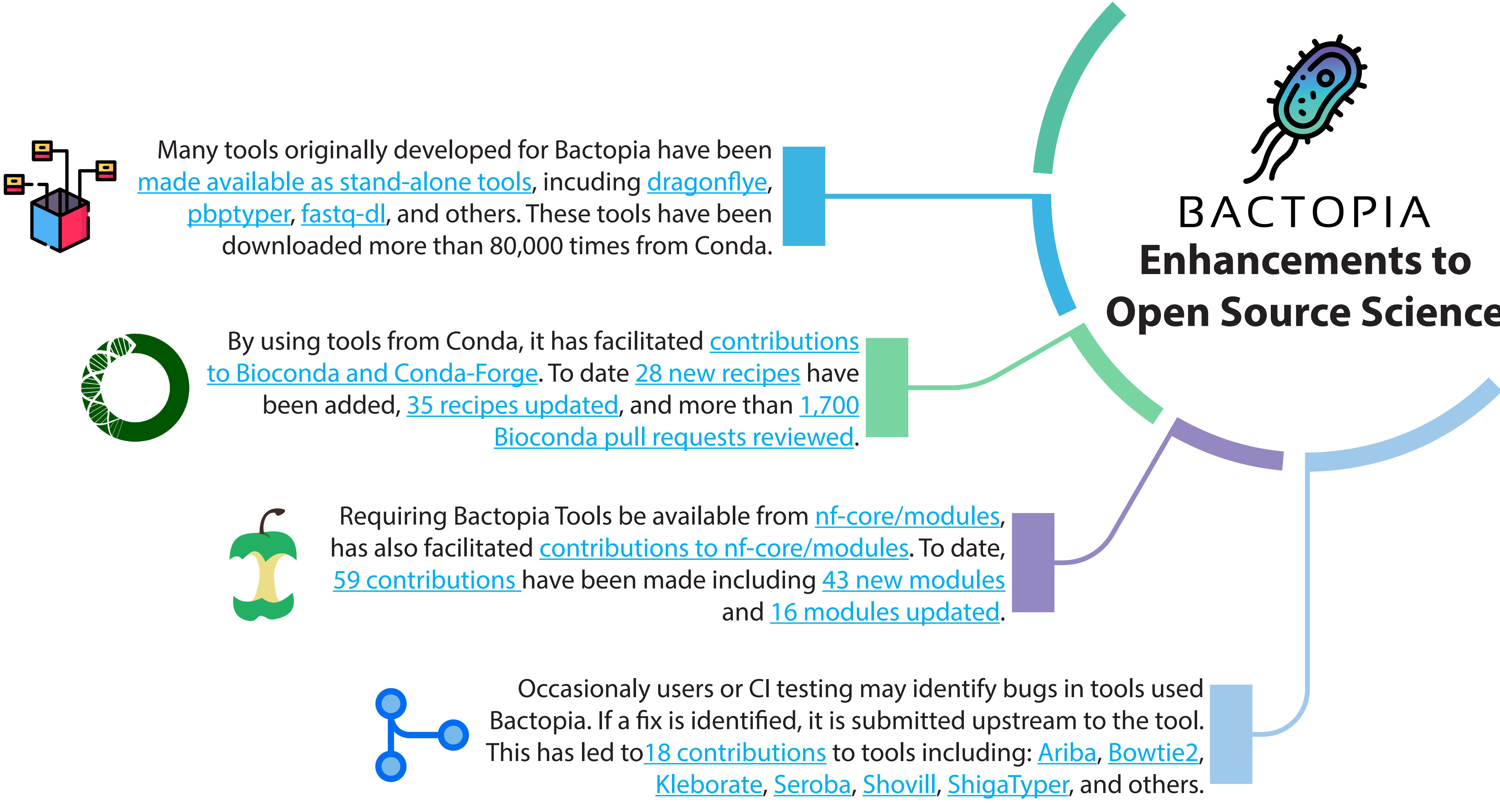


How supporting Bactopia users supports the Bioinformatics Community?

In order to reduce the burden of on-going maintenance and to better help users, a few design principles were adopted.

1. Tools must be free and open source
2. Tools must be available from Bioconda or Conda-Forge
3. Bactopia Tools must be available from nf-core/modules

By adopting these principles, we have been able to rapidly meet user's needs, while also opening pathways to contribute back to the wider bioinformatics community (described to the right).



Scan to see the full list of 150+ contributions

