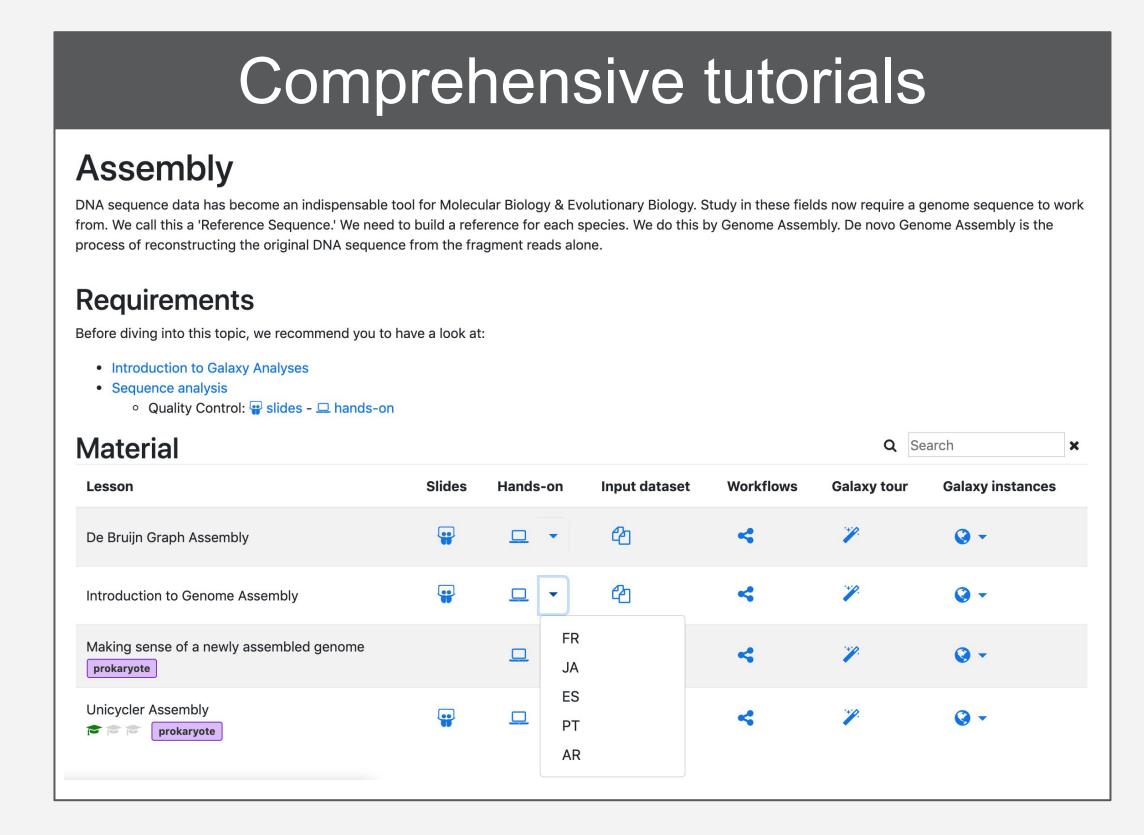
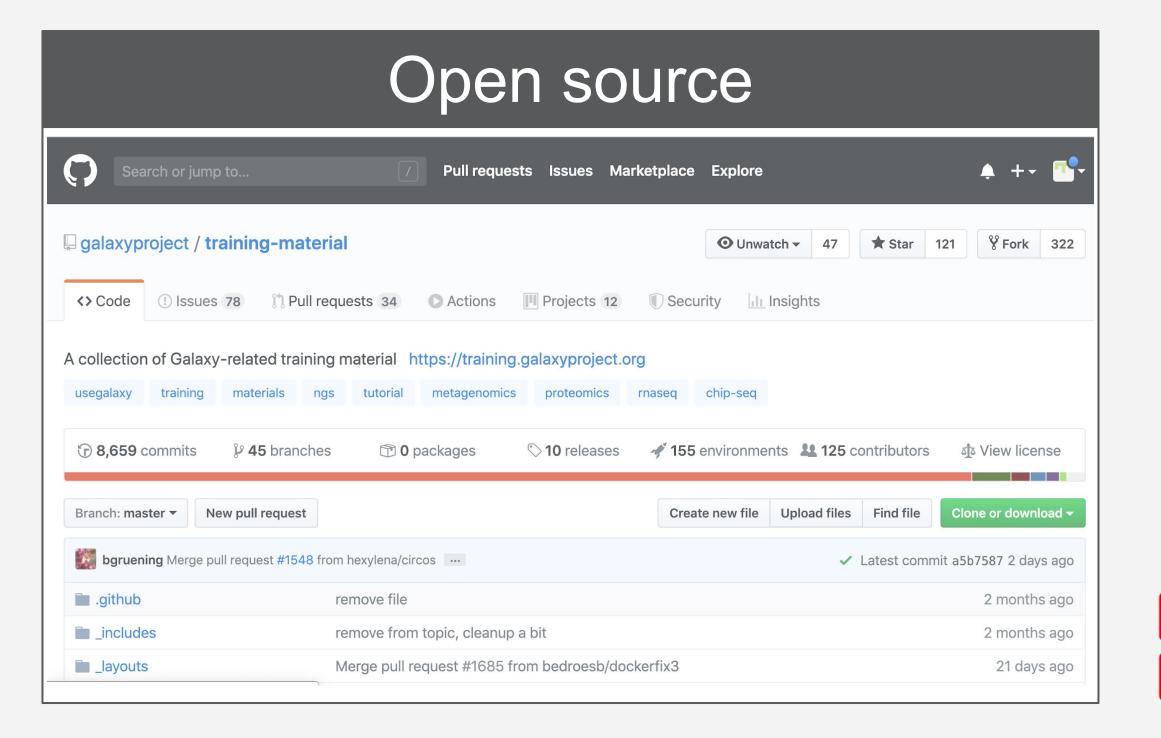
21 topics, +160 tutorials **Galaxy for Scientists Galaxy Tips & Tricks Tutorials Tutorials** Introduction to Galaxy Analyses **Data Manipulation** User Interface and Features Computational chemistry **Galaxy for Developers and Admins Epigenetics** 31 **Galaxy Server administration** Genome Annotation 13 **Development in Galaxy** Imaging How to contribute? Metabolomics First off, thanks for taking the time to contribute! Metagenomics You can report mistakes or errors, create more contents, etc. Whatever is Proteomics your background, there is probably a way to do it: via the GitHub website, via command-line. If you feel it is too much, you can even write it with any Sequence analysis text editor and contact us: we will work together to integrate it. Statistics and machine learning To get you started, check our dedicated tutorials or our Frequently Asked Transcriptomics 21 Galaxy for Contributors and Instructors Variant Analysis Tutorials Visualisation Contributing to the Galaxy Training Material Teaching and Hosting Galaxy training





Community GTN GFESTS Online, every 3 months Online, every 3 months Online, every 3 months Fig. 1. Sept. 1



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