A Galaxy-based framework to analyze raw sequencing data from microbiota
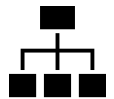
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A framework built on the shoulders of giants

To develop a modular, accessible, redistributable, sharable and user-friendly framework for scientists working with microbiota data

With a comprehensive set of microbiota related tools

With predefined and tested workflows

With documentation and training

GitHub: http://github.com/asaim/framework

Doc: http://asaim.readthedocs.io

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