The Cancer Center Amsterdam (CCA) with 400 cancer researchers is part of the academic hospital VU University Medical Center. Given the rise of multiple omics technologies and their clinical potential we are currently establishing a Drylab that empowers both researchers and clinicians with state-of-the-art bioinformatics solutions. Galaxy-based analysis workflows are implemented usable by non-bioinformatics researchers. We promote Sustainable Software Development of internal tools and their integration in Galaxy. We train users in bioinformatics through seminars, tutorials and coaching; give them power to do more analyses themselves.

References
1. www.bioconductor.org package QDNAseq
2. www.biobix.be/proteoformer

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