

## Clone-Based Functional Genomics Resources (ORFeomics)

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Our subcommittee “Clone-Based Functional Genomics Resources (ORFeomics)” has kept tracking progress made towards the production of full-length cDNAs and open reading frame (ORF) clones for all annotated Arabidopsis protein-coding genes, there are no new large-scale “ORFeomics” projects and no new activities in our subcommittee during 2020-2021. So, we think that we should dissolve our subcommittee.

However, we think that the production of full-length cDNAs and open reading frame (ORF) clones for all annotated Arabidopsis protein-coding genes is so important in the Arabidopsis community, and we hope that the goal will be completed in the future. As our subcommittee can be related with the subcommittee “Systems and Synthetic Biology”, that subcommittee will absorb our activity goal (towards the production of full-length cDNAs and open reading frame (ORF) clones for all annotated Arabidopsis protein-coding genes)