



Dear CCGP Investigators,

We would like to provide two updates regarding the whole genome re-sequencing component of your CCGP projects.

1. Amplify-on index options at QB3 at UC Berkeley. QB3 is phasing out their internal usage of dual-unique Amplify-on indexes in library preparations, as they are moving to a Normalase protocol.

- They have offered to continue to act as a distribution hub for Amplify-on indexes for CCGP projects and will be able to pool your libraries with other CCGP projects to optimize sequencing costs. However, this means that these indexes will be ordered specially for us. Therefore, labs that intend to acquire indexes from QB3 should budget for an approximate additional cost of \$1.39 per sample from their original budget.
- If you wish to have QB3 prepare your WGS libraries, they will use this new Normalase library prep protocol. Please contact them for more information on the specifics of the protocol. Learn more about QB3 [here](#).

2. The CCGP Mini-Core is fully open for business.

- We have been able to source enough reagents and disposables during the COVID-related supply slowdown to now be fully operational. We appreciate your patience! Please visit [the Mini-Core section](#) of the CCGP website for more information on the services we provide, costs, and submission guidelines. For Mini-Core questions, please contact Dan Oliveira (danrodriguesolive@gmail.com).
- We have received feedback that sample collection for some species has been slowed by both COVID and the bad rain year across the state. In order to get data flowing and projects moving, you may now submit samples to the Mini-Core in up to two batches per species-project, rather than waiting until your full set of samples is collected

Please let us know if you have any questions regarding any of the above.

The CCGP Team

California Conservation Genomics Project

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