

June 2016

Important Note

Dear GeneRead® Link customer,

We are writing to inform you of some restrictions that apply to the GeneRead Link software version 1.1.0.

In **Clonal Amplification**, GeneRead Link proposes a different way to dilute the sample than the handbook. Both ways lead to the same result.

The GeneRead Clonal Amp Q Kit handbook proposes to measure the **bead density** before preparing the flow cell. GeneRead Link does not guide you through this step, but offers to log the measured values in the “Comment” field for each sample. You will be prompted by GeneRead Link to note down the **bead density** in the **Prepare Flow Cell** step.

For **Clonal Amplification** and **Sequencing**, the **Process Report** only includes results for the final kit used in these processes (i.e. kit box 3 of 3). For information relating to results for all kits used during these steps you can refer to the run reports of the individual instruments used.

If you are using **Internet Explorer** and have many samples waiting for approval in **Target Enrichment**, the performance of your browser might be slow. If this happens, use Mozilla Firefox® or Google Chrome™ as an alternative browser.

For any changes in the kit handbooks, please check whether the procedures proposed by GeneRead Link still apply. In case of doubts, follow the procedures from the latest handbook.

If you have further questions, please contact QIAGEN Technical Services (see the back cover of our handbooks or visit www.qiagen.com for contact information).

Best regards,

Your GeneRead Link team

Trademarks: QIAGEN®, Sample to Insight™, GeneRead® QuantiMIZE™ (QIAGEN Group), Chrome™ (Google), Firefox® (Mozilla). Registered names, trademarks, etc. used in this document, even when not specifically marked as such, are not to be considered unprotected by law.
HB-2153-001 1103378 06/2016 © 2016 QIAGEN, all rights reserved.