

resDNASEQ® MDCK Residual DNA Quantitation System

Integrated sample preparation and real-time PCR assay for the quantitation of Madin-Darby canine kidney (MDCK) host cell DNA

- Highly sensitive quantitation using proven TaqMan® real-time qPCR technology (Figure 1)
- Manual and automated sample preparation, optimized for quantitative recovery from complex sample matrices (Table 1)
- Consistent performance across the expected range of DNA fragment sizes (Figure 2)
- Integrated system from sample to results, with sample preparation, master mix, TaqMan® primer/probe mix, and genomic DNA standard



The resDNASEQ® MDCK Residual DNA Quantitation System is a quantitative PCR (qPCR)-based system for the detection of host cell DNA from Madin-Darby canine kidney (MDCK) cells, an expression system commonly used for the production of vaccines. Reliable and rapid, the system enables sensitive (LOQ = 1.5 pg DNA per mL of test sample, Figure 1) and specific

(Figure 3) quantitation of MDCK cell DNA, typically in less than 4 hours. This performance helps ensure a high degree of confidence in quantitation data obtained from a broad range of sample types—from in-process samples to final product—whether the sample contains high molecular weight or sheared DNA (Figure 2).

Table 1. DNA recovery using the manual PrepSEQ® sample preparation protocol.

Assay performance data from 1 pg MDCK genomic DNA spiked into 6 test samples.

| | Mean recovery | Mean CV |
|------|---------------|---------|
| MDCK | 83% | 3% |

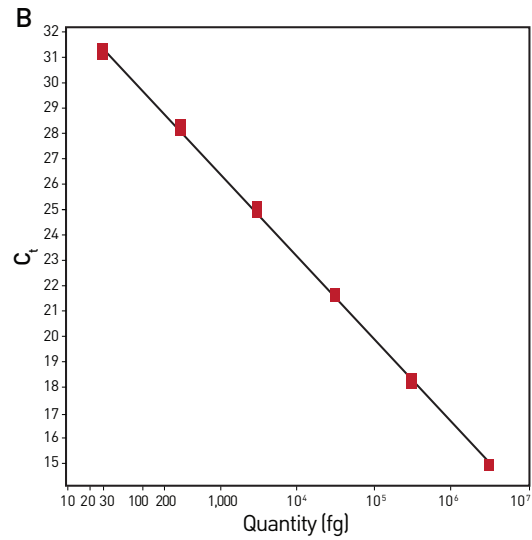
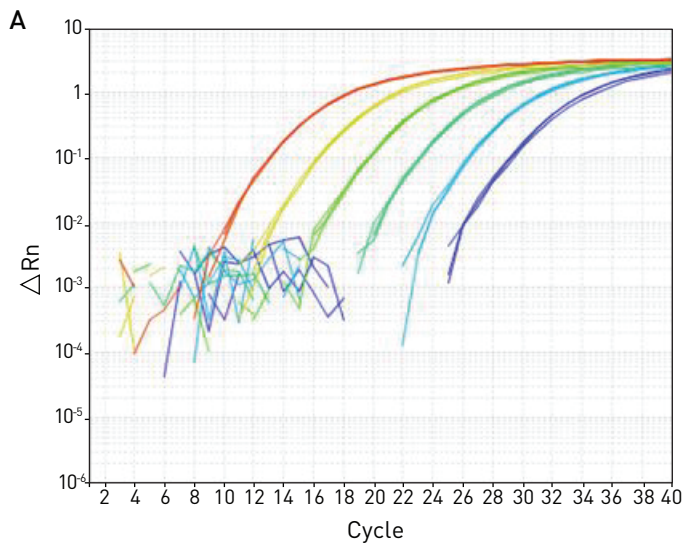


Figure 1. High sensitivity and broad dynamic range using the resDNASEQ[®] MDCK Residual DNA Quantitation System. (A) The amplification plots were generated using 10-fold serial dilutions [containing 3 ng to 30 fg] of MDCK genomic DNA, provided in the kit. (B) Standard curve of the 10-fold dilution series. Data were analyzed using AccuSEQ[®] Real-Time Detection Software.

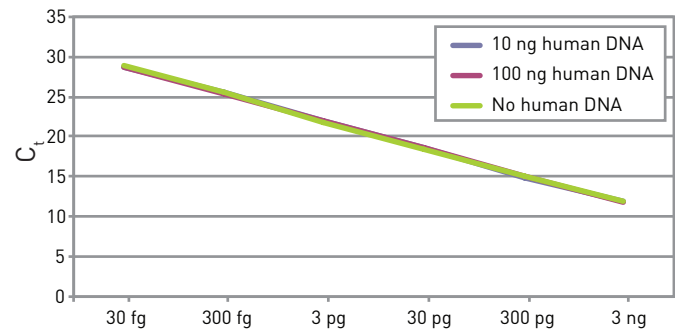
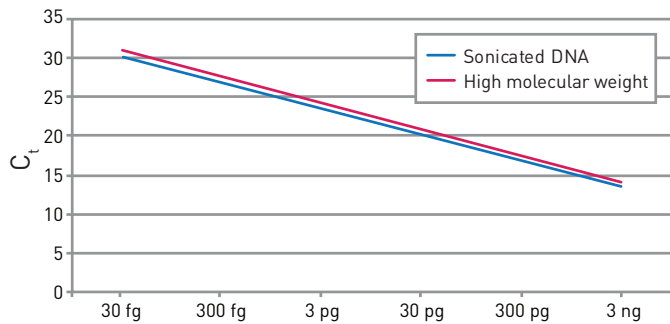


Figure 2. Consistent quantitation across a broad range of fragment sizes. Standard curves were generated using 10-fold serial dilutions of high molecular weight or fragmented DNA, from 3 ng to 30 fg. Fragmented DNA was generated by sonicating total MDCK genomic DNA. Fragmentation of the DNA was confirmed by agarose gel analysis.

Figure 3. Assay specificity. Standard curves were generated using 10-fold serial dilutions of MDCK genomic DNA in the presence of 100 ng, 10 ng, or no human DNA.

Ordering information

| Product | Cat. No. |
|---|----------|
| resDNASEQ[®] MDCK Residual DNA Quantitation System | |
| resDNASEQ [®] MDCK Residual DNA Kit, 100 rxns, without protocol and Quick Reference Card | 4464335 |
| PrepSEQ[®] Residual DNA Sample Preparation Kit | |
| PrepSEQ [®] Residual DNA Sample Preparation Kit, 100 rxns, with protocol and Quick Reference Card | 4415414 |
| PrepSEQ [®] Residual DNA Sample Preparation Kit, 100 rxns, without protocol and Quick Reference Card | 4413686 |
| AccuSEQ[®] Real-Time PCR Software | |
| AccuSEQ [®] Real-Time PCR Software v1.0 | 4443420 |

For Research Use Only. Not intended for any animal or human therapeutic or diagnostic use.

© 2011 Life Technologies Corporation. All rights reserved. The trademarks mentioned herein are the property of Life Technologies Corporation or their respective owners. TaqMan[®] is a registered trademark of Roche Molecular Systems, Inc., used under permission and license. Printed in the USA. C024250 0111