SFD-10HT
Performance Data Analysis

SFD-10HT allows for the progressive elimination of short DNA fragments less than 10 kb. Eliminating these fragments decreases the number of reads below 10 kb, resulting in an improved mean read length. This enhancement leads to better sequence assembly and alignment outcomes, as longer reads provide more context and overlap, facilitating more accurate genomic sequence reconstruction. Additionally, SFD-10HT exhibits a high tolerance to a wide range of concentrations (10–150 ng/µL) and demonstrates an improved DNA Quality Number (DQN).

A. Progressive Elimination of <10 kb DNA fragments:

SFD-10HT's Tolerance to a wide range of DNA concentrations

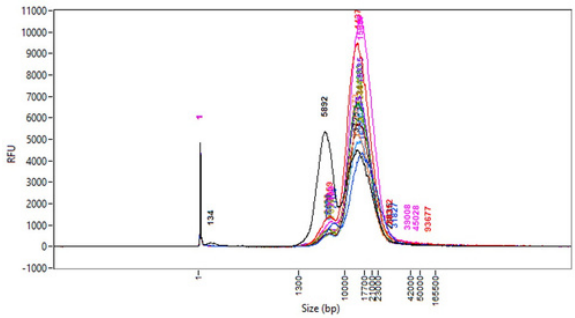

Figure 1: Femtopulse traces - Sheared DNA was diluted to different concentrations and 0.4x bead ratio of SFD-10HT was used to eliminate DNA fragments less than 10 kb.

SFD-10HT offers consistent efficiency in eliminating fragments less than 10 kb across a wide range of DNA concentrations

B. Improvement of DNA Quality of Sheared/Fragmented HMW DNA (15-20 kb)

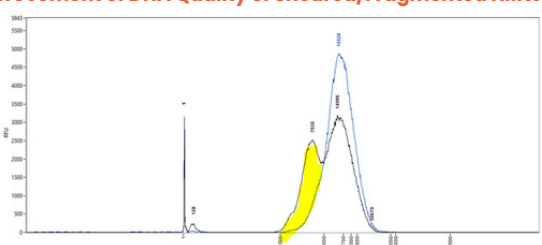

Figure 2: Sheared DNA with fragments >10 kb (55.8%)-see smear analysis Table 1. After elimination of fragments <10 kb with SFD-10HT, the sample quality was improved **from 55.8% to 84.4%**. The DQN was also improved from **5.6 to 8.4**.

Table 1: Smear Analysis-Sheared HMW DNA (10-15 kb)

Sample ID	Range	% Total	Avg. Size	%CV	Size Threshold (b.p.)	DQN
Sheared HMW DNA -post short fragments removal	10000 bp to 300000 bp	84.4	17180	25.19	10000	**8.4**
Control input-Sheared HMW DNA	10000 bp to 300000 bp	55.8	16837	25.64	10000	**5.6**

SFD-10HT provides a significant enhancement in input DNA quality, optimizing samples for long-read sequencing