

BluePippin™ User Guide

High-Pass™ DNA Size Selection

Sage Science Part Nos:

PAC-20KB

PAC-30KB

Library Size Selection Chart for PacBio® SMRTbell™ Templates

Cassette Kit PN	System	High Pass Threshold Range	Cassette Definition	PacBio Application	Validated by PacBio?	
PAC20KB	BluePippin	4-10kb	0.75% DF Marker S1 high-pass 4-10kb vs2	~20kb SMRTBell Templates	yes	1,2
PAC20KB	BluePippin	6-10kb	0.75% DF Marker S1 high-pass 6-10kb vs3	~20kb SMRTBell Templates	yes	
PAC20KB	BluePippin	15-20kb	0.75% DF Marker S1 high-pass 15-20kb	~20kb SMRTBell Templates	yes	
PAC30KB	BluePippin	30-40kb	0.75% DF Marker U1 high-pass 30-40kb v3	>30kb SMRTBell Templates	yes	3
BLF7510	BluePippin	NA	0.75% DF 2-6 kb Marker S1	Iso-Seq	yes	
BLF7510	BluePippin	NA	0.75% DF 3-10 kb Marker S1	Targeted Sequencing & Phasing	yes	
HPD7510	PippinHT	6-10kb	0.75% Agarose, 6-10kb High Pass, 75D	~20kb SMRTbell Templates	pending	4
HPE7510	PippinHT	15-20kb	0.75% Agarose, 15-20kb High Pass, 75E	~20kb SMRTbell Templates	pending	
HPF7510	PippinHT	30-40kb	0.75% Agarose, 30-40kb High Pass, 75F	>30kb SMRTbell Templates	pending	
ELD7510	SageELF	NA	0.75% 1 kb – 18 kb	Iso-Seq	yes	

Notes:

- 1 PAC20KB is the same kit as Sage PN BLF7510, except for the user quick guide.
- 2 The version2, 4-10kb, size selection accuracy can be affected by the DNA input amount. New users should use version 3, 6-10kb.
- 3 PAC30KB is the same kit as Sage PN BUF7510, except for the user quick guide.
- 4 The maximum input amount for the PippinHT is 1.5 ug. The PippinHT runs up to 22 samples per run.

Using this Guide

This guide is meant to provide a comprehensive workflow for using the BluePippin for High-Pass DNA size-selection. High-Pass size selection is a process by which user enter DNA size selection threshold value, in base pairs, into a software protocol. When the procol is run, The BluePippin will collect all fragments above that threshold. This allows users to filter out unwanted smaller DNA fragments from a sheared sample. The larger DNA fragments include those that are to are not resolved electrophoretically, and are compressed as band.

The instructions here provide special recommendations for software, hardware, and gel cassette usage High-Pass size selection. However, users should read the **BluePippin Operations Manual** to completely familiarize themselves with the system and features, and safety and warnings.

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A. Preparing DNA samples

1. Dilute up to 5 µg SMRTbell template into a final volume of 30 µl EB (PacBio Elution Buffer).

Note: For >20kb SMRTbell libraries, do not use less than 500 ng per lane
For >30kb SMRTbell libraries, do not use less than 2 µg per lane

2. Bring loading solution to room temperature and add 10 µl of loading solution to the 30 µl sample.

Note: Loading solution is viscous: pipet slowly to ensure complete transfer of the loading solution into the DNA sample.

3. Mix thoroughly by gentle pipetting. Do not vortex. Spin briefly to collect the contents at the bottom of the tube.

4. Bring the DNA marker to room temperature.

B. Creating a Protocol

The following instructions show how to program the size selection protocol for SMRTbell template preparation. Optional pre-programmed protocols may also be loaded onto a BluePippin, and are shown in the gray box on page 5.

1. In the BluePippin Software, click the Protocol Editor Tab.

2. Press the “NEW” button, to program a new protocol.

3. Click “Cassette Type” folder icon, and select the appropriate cassette definition from the list below.

Cassette Kit PN	High Pass Threshold Range	Cassette Definition
PAC20KB	6-10kb	0.75% DF Marker S1 high-pass 6-10kb vs3
PAC20KB	15-20kb	0.75% DF Marker S1 high-pass 15-20kb
PAC30KB	30-40kb	0.75% DF Marker U1 high-pass 30-40kb v3

For software steps, a small representation of the screen is provided, with a yellow circle that indicates the area of the screen that is being referred to in the text.



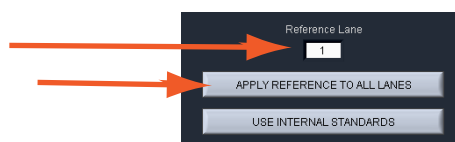
4. Select “Range” mode for the corresponding sample lanes to which the samples will be run. Do not select a collection mode for the sample lane to which the DNA marker will be added.



	Tight	Range	Time	Peak
5				
4				
3				
2				
1				

In this example, DNA reference marker will be loaded into lane 1, and the DNA samples will be loaded into lanes 2-5.

5. Enter the lane number to which the DNA marker will be loaded into the “Reference Lane” field, and press the “APPLY REFERENCE TO ALL LANES” button.



6. Enter a base pair threshold (BP Start) for DNA size selection:

For ~20kb SMRTbell libraries, enter a value between 4000 - 20000
 For >30kb SMRTbell libraries, enter a value of 30000 or 40000



7. If necessary, enter 50000 (for PAC-20KB) or 80000 (for PAC-30KB) into the (BP End) field.
 For software versions 6.2 and above, these values will auto-populate the protocol editor screen when the High-Pass cassette definition has been selected.

8. Enter Sample information into the “Sample ID” fields (optional).



9. Press “Save As” to name and save a new protocol.



B.1 Loading optional pre-set protocols

The protocols may be downloaded from www.sagescience.com/product-support/bluepippin. Use the typical software update instructions. See page 3 for more information.

1. Press the “LOAD” button.



2. Select the appropriate SMRTbell protocol from the protocol list.



3. The following protocol names will be found on the list:

SMRTbell 20kb, 4000bp High-Pass
 SMRTbell 20kb, 6000bp High-Pass v3
 SMRTbell 20kb, 7000bp High-Pass v3
 SMRTbell 20kb, 7000bp High-Pass
 SMRTbell 20kb, 8000bp High-Pass v3
 SMRTbell 20kb, 10000bp High-Pass v3
 SMRTbell 20kb, 15000bp High-Pass
 SMRTbell 20kb, 17000bp High-Pass
 SMRTbell 20kb, 20000bp High-Pass
 SMRTbell 30kb, 30000bp High-Pass v3
 SMRTbell 30kb, 40000bp High-Pass v3

4. The high pass size selection values will auto-populate the protocol editor, reserving lane 1 for the external marker :

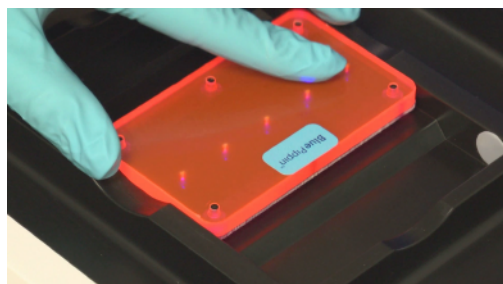
Sample Protocol: SMRTbell 20kb, 10000bp High-Pass v3

	Tight	Range	Time	Peak	Ref Lane	BP Target	BP Start	BP End	BP Pause
5					1	30000	10000	50000	0
4					1	30000	10000	50000	0
3					1	30000	10000	50000	0
2					1	30000	10000	50000	0
1					1	0	0	0	0



C. Calibrating the Optics with the Calibration Fixture

1. In the BluePippin Software, click the **Main Tab**.
2. Place the calibration fixture onto the optical nest. The dark side of the fixture must be down, and completely cover all LED detectors.



Correct placement of the calibration fixture.

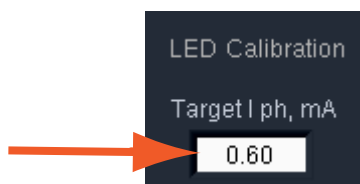
3. Close the Lid.
4. Press the **"CALIBRATE"** button.



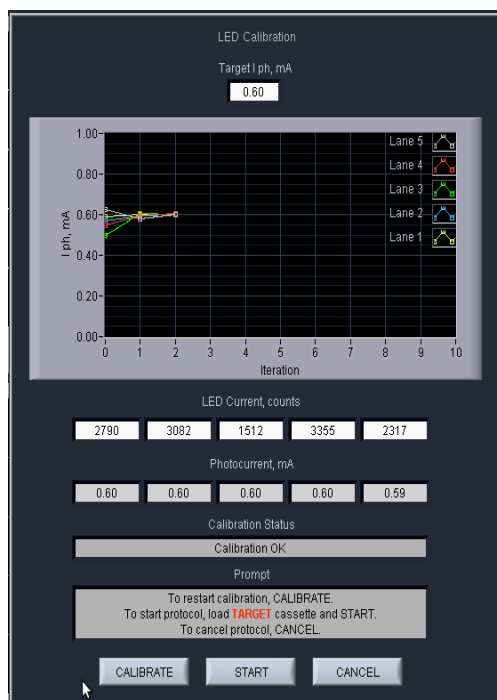
5. A calibration sub-window will launch.



6. Check that the calibration target box ("Target I ph, mA") at the top of the window reads **"0.60"**. If not, manually change the target value to **"0.60"**.



7. Press “CALIBRATE” button within the calibration window.



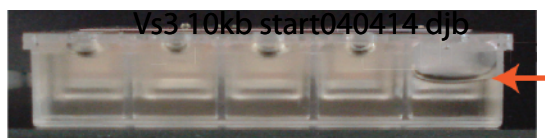
The Calibration sub-window after a successful calibration. If the status window reads “Calibration OK”, proceed to the next step.

Note: The optical calibration routine ensures that each LED light intensity is at the correct level (0.60 mA, in this case). The LEDs will adjust individual current (counts) values to achieve this intensity. If the count level is <200, the test will fail, likely due to light infiltration such as an open lid. The maximum count value is 4095, and if the LED cannot achieve the proper intensity at that level, the test will fail due to a faulty LED.

8. Press “EXIT” to return to the Main Screen.

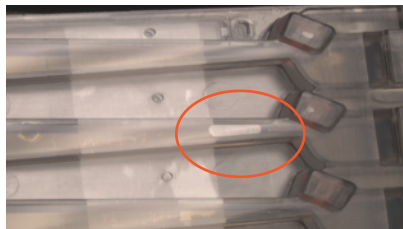
D. Inspecting the Gel Cassette

1. Remove the gel cassette from the foil packaging.
2. **Inspect the levels of buffer in all buffer reservoirs.** Reservoirs should be nearly full. If the buffer level in any reservoir appears lower, or less than 50% full, fill with spare electrophoresis buffer (provided with the BluePippin reagent kit).



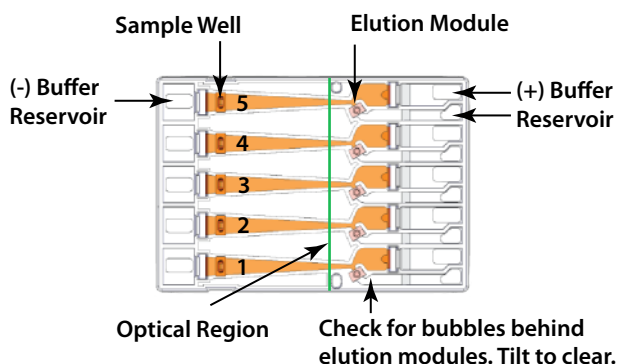
If a buffer reservoir is low, fill with spare buffer.

3. **Inspect for bubbles due to delamination of agarose from the bottom of the cassette in the region used for optical detection of DNA.** To inspect, turn the cassette upside down and gently tilt the cassette under a light source. If a bubble is detected, do not use the affected lane to run the DNA reference marker. Affected lanes can be used for size selection of sample DNA.



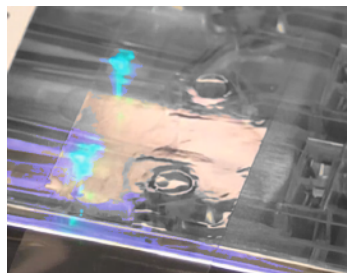
If the agarose has delaminated in the optical region, do not use that lane to run the reference marker.

E. Preparing the Cassette for Loading



1. **Dislodge air bubbles from behind the elution wells.** Tilt the cassette sample well side down, and tap gently to release the bubbles.
2. **Place the cassette into the optical nest.** Keep the cassette slightly tilted down so that the air bubbles do not return to the area behind the elution modules. Make sure the cassette is fully seated into the bottom of the nest to ensure proper optical alignment.
3. **Remove the white tabbed adhesive strips from the cassette.** Place one hand on the cassette, and hold it firmly in the nest. Grab the white tabs and pull the strips firmly and slowly toward the front of the instrument until they are removed.
4. **Completely remove all buffer from each elution module, and replace with 40 μ l of fresh electrophoresis buffer** (provided with the BluePippin reagent kit). When refilling, dispense the fresh buffer slowly, raising the pipette tip slowly from the bottom of the elution module, to avoid trapping air bubbles in the module.

5. **Re-seal the elution wells with an adhesive tape strip (supplied with cassette packaging).** Place the tape over the elution wells and rub firmly to seal the elution ports. The port should be tightly sealed without any wrinkles around the edges of the port.



Rub firmly to ensure a tightly sealed elution port.

6. **Close the lid.**

7. **Perform the Continuity Test.** Press the “Test” button. The test sub-window will open, and the test routine will automatically measure the current in each separation and elution channel. A successful continuity test will return a “PASS” indication. The cassette temperature must be above 17°C (62°F). If a lane fails due to low elution current (the affected lane will be highlighted with an orange color), refilling the elution module (see step 4 above) will usually resolve the problem.



Continuity Test Parameters

I separate max, mA
I elute max, mA

2.50

1.65

I separate min, mA
I elute min, mA

1.55

1.15

Actual Lane Currents, mA

	Separation	Elution
5	1.73	1.41
4	1.78	1.39
3	1.78	1.39
2	1.74	1.37
1	1.70	1.34

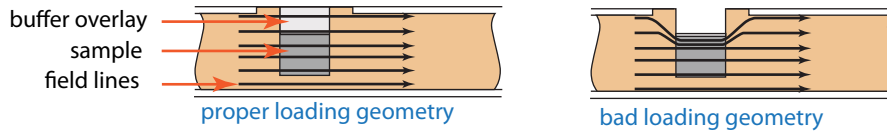
PASS

RETURN

A successful continuity test will return a “PASS” indication.

8. Press “Return” to close the Continuity Test sub-window.

F. Loading Samples

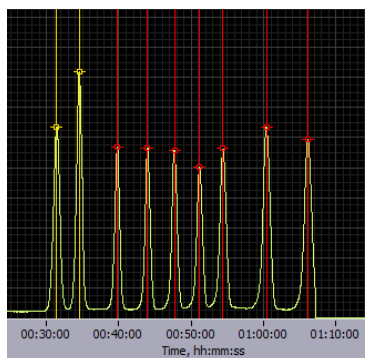


1. **Check the buffer levels in the sample wells.** Make sure that sample wells are completely full to the top with electrophoresis buffer. Fill with additional buffer if necessary.
2. **Remove 40 μ l of buffer from the first sample well, and load 40 μ l of sample or marker into that well.** Take care not to pierce the agarose with the pipette tip. There is gel on all sides and at the bottom of the sample well. In addition, there is an agarose “chimney” surrounding the top of the sample well that protrudes up through the cassette cover. When removing buffer, some users find it useful to immerse the pipette tip just below the surface of the buffer and follow the liquid level down with the tip as the buffer is removed. When buffer removal is completed, there will be ~30 μ l of buffer left in the well. When adding sample, place the tip of the pipette just below the surface of the buffer, and slowly eject the sample. Don’t be concerned if the sample well slightly overfills. Then density of the sample will allow it to sink before it can flow out of the well.
3. Repeat step 2 for the remaining four wells.

Note: Cassettes may be reused if 2-3 lanes are used (1-2 samples). Reseal the cassette with the adhesive tape provided and store at RT. However one remaining lane must again be loaded with the external marker for calibration. Unused lanes must have all fields deselected in the software protocol.

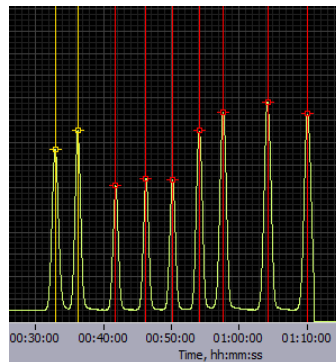
G. Running

1. Close the lid, go to the Main Tab, and make sure the proper protocol is loaded in the “Protocol Name” field.
2. Press “START”. The run will automatically stop when every collection is complete. Electrogram images of the DNA markers in the Mains screen should appear as shown below.



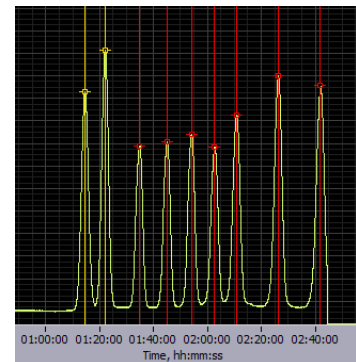
Marker S1

PAC-20KB High-Pass Size Selection
4-10kb thresholds
Approx run time : 2 hours



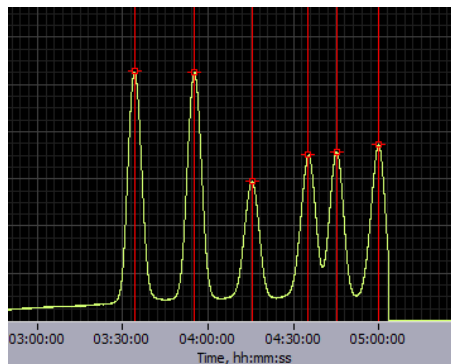
Marker S1

PAC-20KB High-Pass Size Selection
6-10kb thresholds
Approx run time : 2 hours



Marker S1

PAC-20KB High-Pass Size Selection
15-20kb thresholds
Approx run time : 5-6 hours



Marker U1

PAC-30KB High-Pass Size Selection
30-40kb thresholds
Approx run time : 9.5-10 hours

H. Collecting Fractions

1. For best recovery, wait at least 30-45 minutes after end of run before removing samples from the elution modules.
2. Remove samples using a standard 100-200 μ l pipette. Aspirate slowly to minimize breakage of large DNA fragments. Samples will be in a Tris-TAPS buffer at a volume of approximately 40 μ l.
3. **Optional: Additional 10-30% yield can often be achieved by rinsing the elution well with TE+0.1% Tween 20.** Tween solution is provided by Sage Science with cassette kits for HMW collections.
 - Add 40 μ l Tween solution (supplied with the cassettes) to the elution well
 - Wait 1 minute
 - Remove solution and pool with the original extracted sample, or process separately
4. **Remove the cassette and dispose of properly.** Do not keep used cassettes in the Pippin with the cover closed. Humidity from the cassette may cause damage to the electrodes.

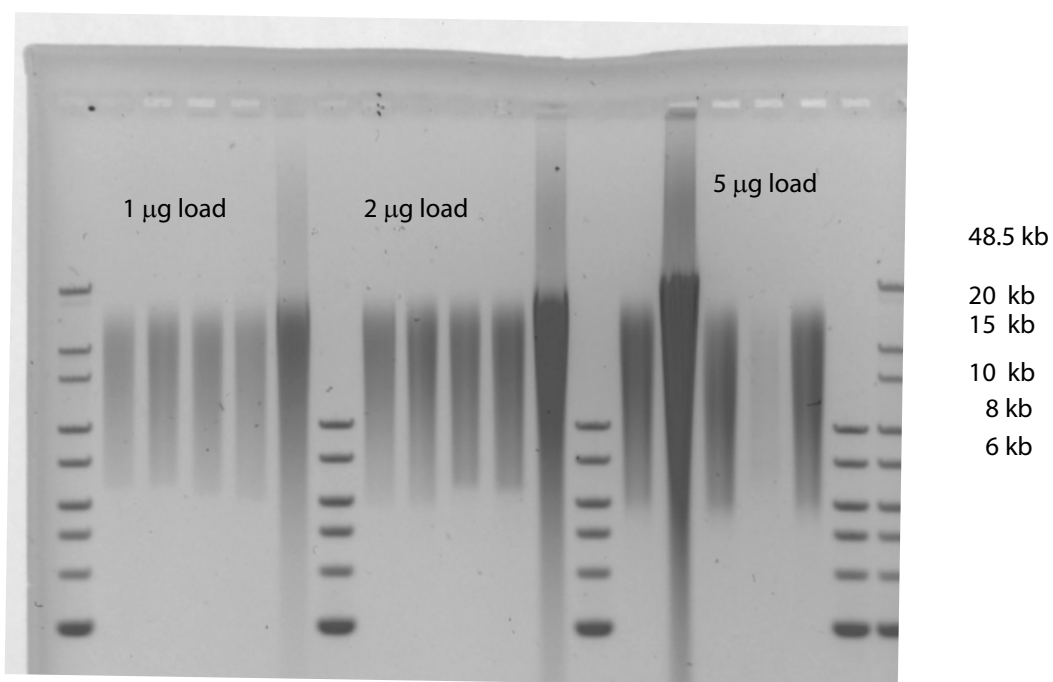


Important! Qubit fluorometry is recommended for determining DNA concentrations of eluted samples. UV absorbance-based methods (eg. Nanodrop) give inaccurate results on post-Pippin products: an artifact from the cassette manufacturing process interferes with accurate quantification.

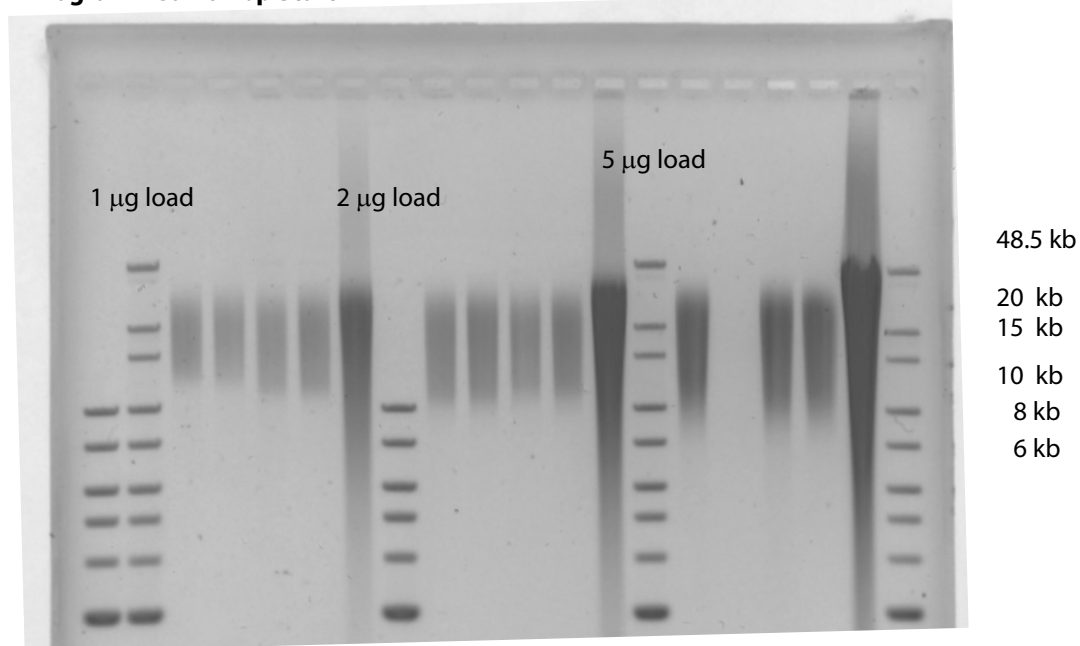
I. Gel Analysis Images of High-Pass Collections

The following pages show analytical gel images of sheared samples after size selection with High Pass protocols. Gels were run using the Pippin Pulse pulsed-field electrophoresis power supply (PPI0200).

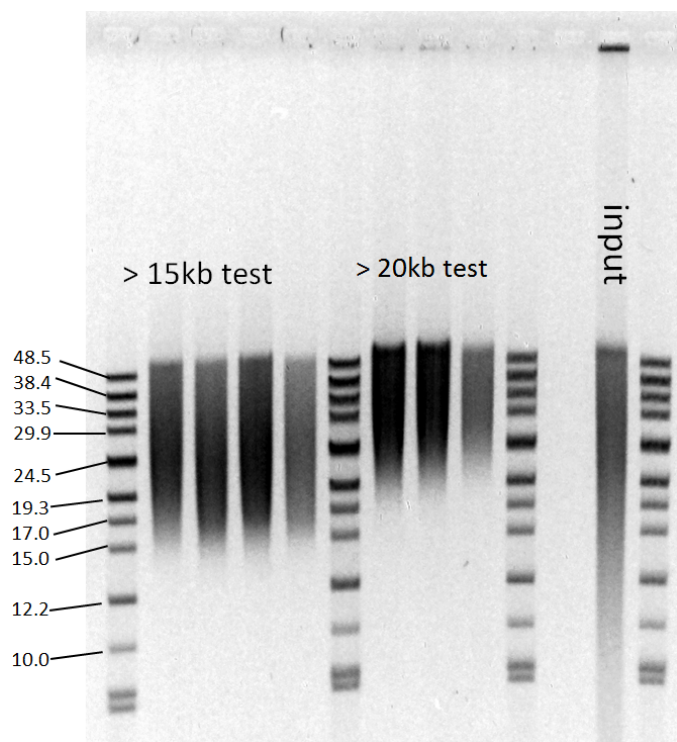
Programmed 6 kbp Start



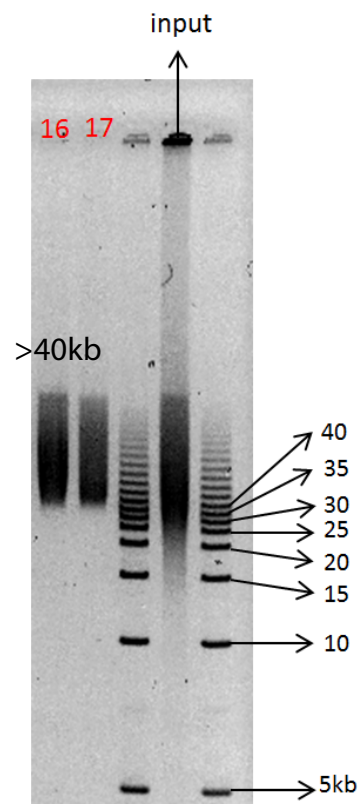
Programmed 10 kbp Start



Programmed 15 and 20 kbp Start (5 µg input)



Programmed 40 kbp Start (5 µg input)



J. DNA Control for High-Pass Collections

High Pass control DNA (**Product No. CHP-7504**) can be used to evaluate whether the High Pass protocol is working as expected. The control DNA is a ladder containing fragments ranging from 0.5kb – 48kb, formulated to load 1.25µg of DNA per sample lane.

