



Agilent Technologies

TapeStation Tips & Tricks





CHECKPOINTS BEFORE STARTING A SCREENTAPE RUN



Before the ScreenTape run begins...

The following quick and easy steps show how to set up a run and assure proper functionality

- Read the quick guide protocol and carefully follow the [Good Measurement Practices](#) 😊
- Let the reagents warm up to room temperature
- Remove bubbles from bottom of strip tubes/plate by centrifugation before placing in the instrument. Carefully remove tube strip-tube lids.
- Don't overload the ScreenTape. Have a reasonable idea of quantity before set up a run.
- Do NOT freeze any ScreenTape
- Do NOT mix match reagents and ScreenTape of different assays
- Flick screentapes briefly prior loading them
- Tip buckets must be emptied after every run.
- Load all 16 tips into the tip holder (2200). Load the TipRack into the holder (4200)
- Always remove lids from the 8 strip tubes!



Instrument Setup

- Only use included bundle PC for the operation of the instrument
- SW does not require a license. Data analysis software may be installed on additional PCs.
- Power up in the right path:
Laptop → Instrument → Controller
- Do not use wireless connections on TapeStation laptop to access company network. If necessary, store data files locally during data acquisition and move them on a server afterwards

4200 TapeStation System - Consumables

Advice:

- Only use Agilent accessories such as tips, 8-way strips, 96-well-plates and foil seal
- 96 well plates should be sealed with foil to avoid evaporation
- Be aware that tips and 96-well plates differ between 2200 and 4200 TapeStation system! No mix and match!

Possible issues with non-Agilent consumables:

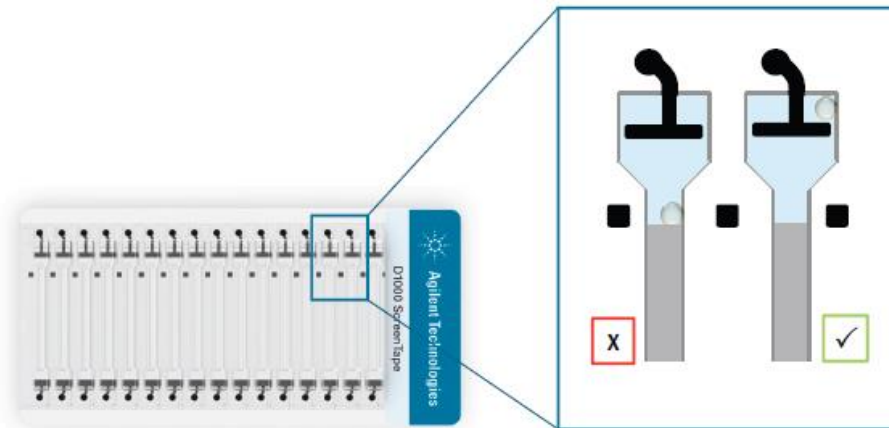
- Blank lines
- Bent tips
- Jammed ScreenTape
- **Instrument crash**



Before the ScreenTape run starts...

Flick the ScreenTape

- Bubbles can form in the buffer chamber of the ScreenTape & a loss of performance is observed if these are positioned at the gel/buffer interface
- Flicking the ScreenTape will move the bubbles to the top of the chamber



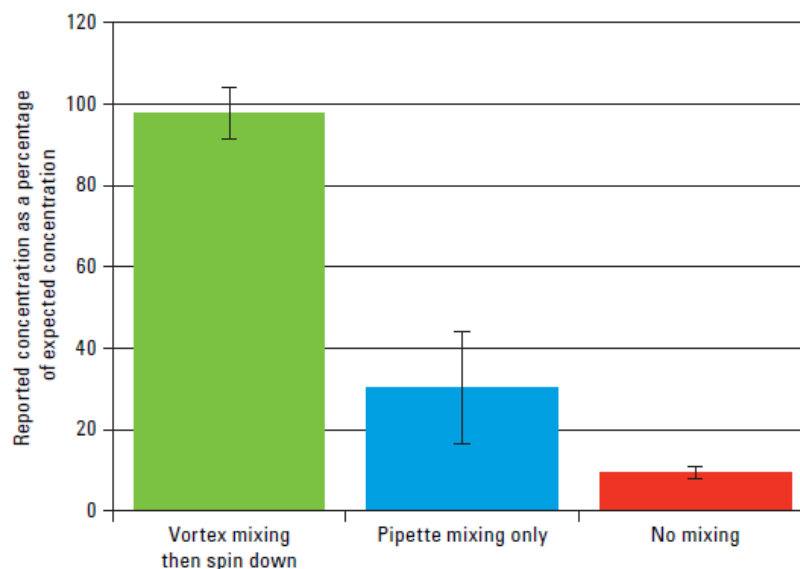
Vortexing: Critical for Concentration Calculation



- Discrepancies in sample quantification often caused by insufficient mixing
- Pipette mixing is often not sufficient!
- **Recommended to use IKA MS3 96-well plate vortexer using the default vortex settings (1 minute).**
- The graph below shows how poor mixing can lead to an underestimation of sample concentration.

MORE CONSISTENT MIXING
BETTER ALIGNMENT TO
BIOANALYZER

Mixing test - D1000

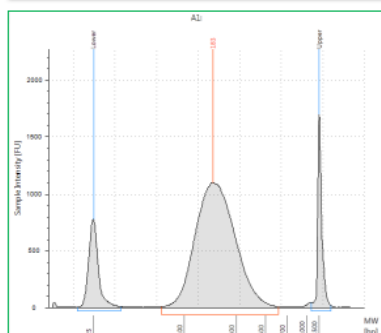


Cockpit error that will affect Sample Concentration

Peak Integration

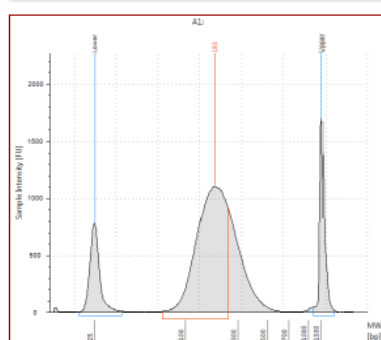
- Region functionality is the best practice for smear analysis
- Ensure the upper marker (UM) peak is proper integrated, this is used to determine quantification
- Sample peaks should be adjusted if required (see example below)

Sample A1 – correct
peak integration



Concentration – 42.2 ng/μl

Sample A1 – incorrect
peak integration



Concentration – 29.9 ng/μl



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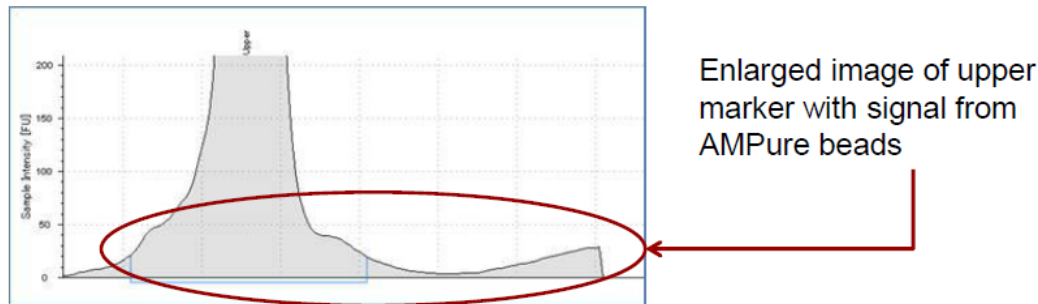
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Rev. 1 [March 9, 2016]

Other Issues

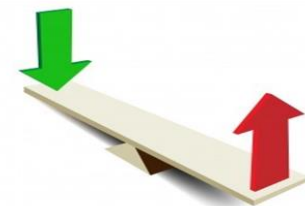
- Residual AMPure beads from SureSelect protocol can give signal which runs with the upper Marker (see picture below)
 - Any signal under the upper marker affects sample quantification
 - Removal of the beads removes the signal under the upper marker peak

Tip: Place the sample plate on a magnetic base when pipetting samples



- Over amplification can cause product to co-migrate with the upper marker peak

Ladder and Markers also play important role for sizing calculation



Lower & Upper Markers

- Always ensure that the upper and lower markers have been identified correctly
- The markers are used as internal reference to determine the molecular weight size of the sample
- Incorrect identification of the markers will lead to miscalculation in reported sizing values

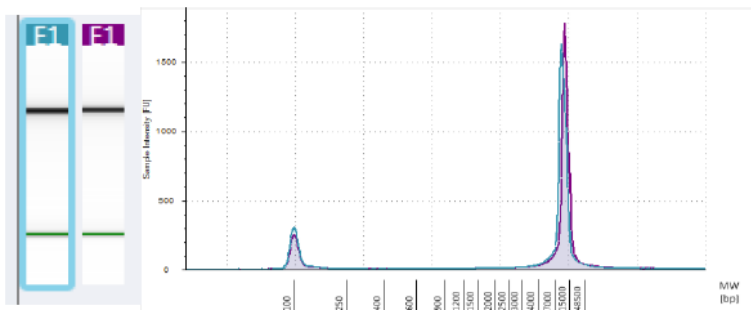
Ladder

- Its recommended to use the run ladder instead of electronic ladder for more accurate sizing; some assays (e.g. D5000, gDNA) do not have electronic ladders available.

Genomic DNA ScreenTape

Genomic DNA sizing

- Reagents must be equilibrated at room temperature for 30 minutes before use
- Failure to do so can affect sizing results
 - Cold reagents will be very difficult to pipette, due to high viscosity

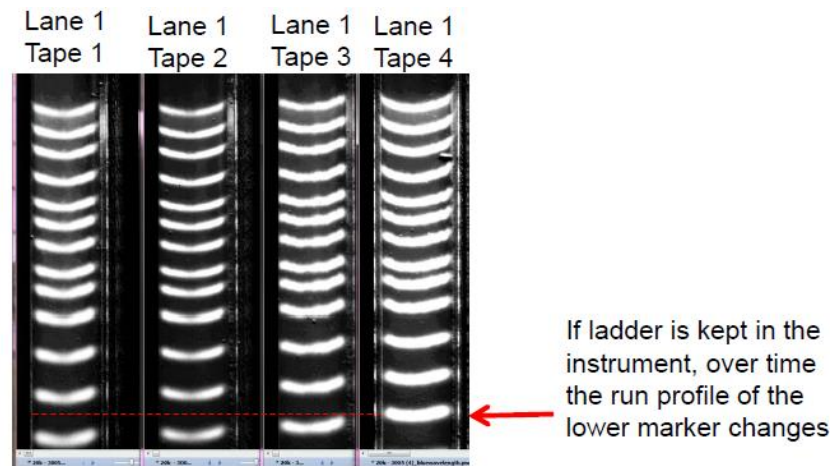


Size (bp)			
Sample	Reagents at 4°C (purple)	Reagents equilibrated to room temperature (green)	Difference
17k - 50ng/ul	24369 bp	19867 bp	4502 bp

Genomic DNA Sizing

Use Fresh Ladder:

- Ladder must be prepared fresh for each run & added to the first available position
- Run profile changes as ladder warms up and evaporates in the instrument: this will affect sizing results
- No software ladder available for genomic DNA



Molarity Calculation

- Molarity is determined from both size and concentration
- Errors in sizing and quantification will result in erroneous molarity calculations!
- Always ensure that the good measurement practices for sizing and quantification have been followed to confirm accuracy in molarity values

Good Measurement Practices for DNA analysis with the Agilent 2200 TapeStation system [5991-3187EN](#)





HARDWARE



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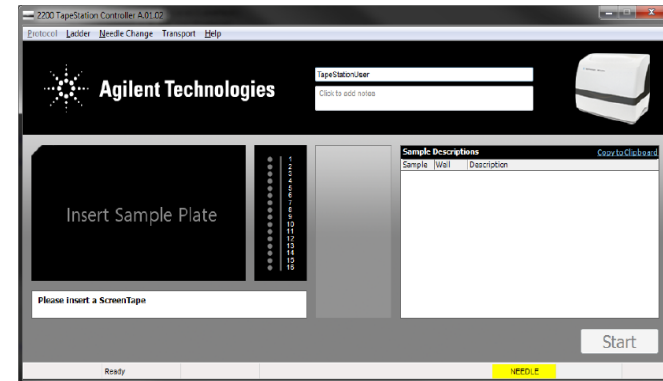
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Rev. 1 [March 9, 2016]



Instrument needle change

The controller software

- Warning message requesting needle change
 - Warning at 7680 pierces (2200 TapeStation)
 - Warning at 3840 pierces (4200 TapeStation)
 - *Needles need to be changed*
- 2200 TapeStation will stop after 8960 pierces
- 4200 TapeStation will stop after 4480 pierces



Needle cartridge can be replaced by users

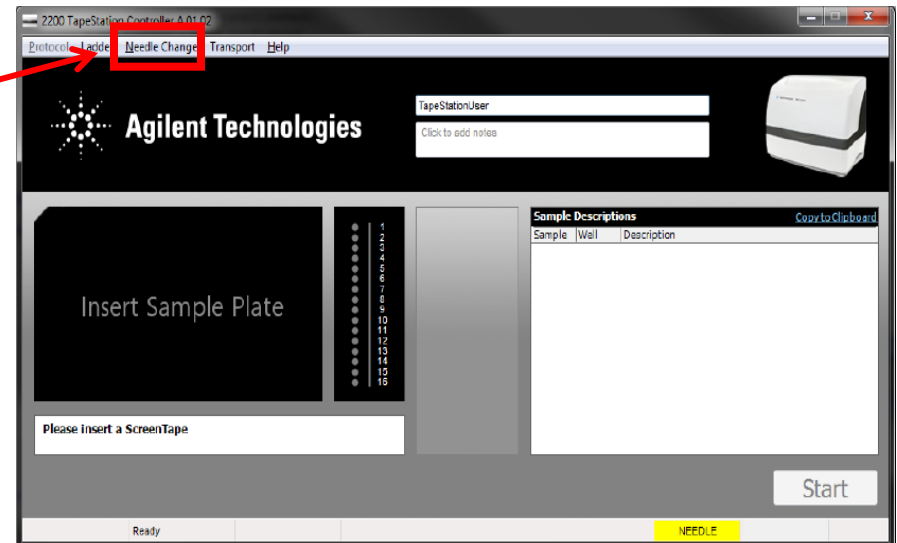
Needle cartridge part number 2200 TapeStation: **G2960-60063**

Needle cartridge part number 4200 TapeStation: **5067-5783**

Instrument needle change 2200 TapeStation System

How to change needles:

- Remove the sample plate and tip holder
- Remove the foil tab from the top of the needle cartridge
- Insert cartridge into the tip holder space (label facing right, printed arrow points to front of the instrument)
- Close the lid
- Select **Needle Change**
- Select **Run**

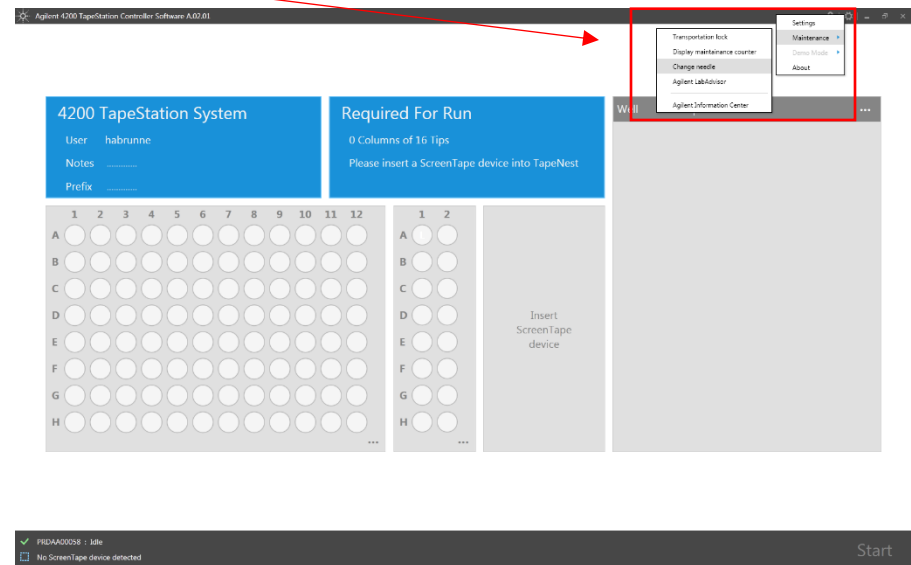
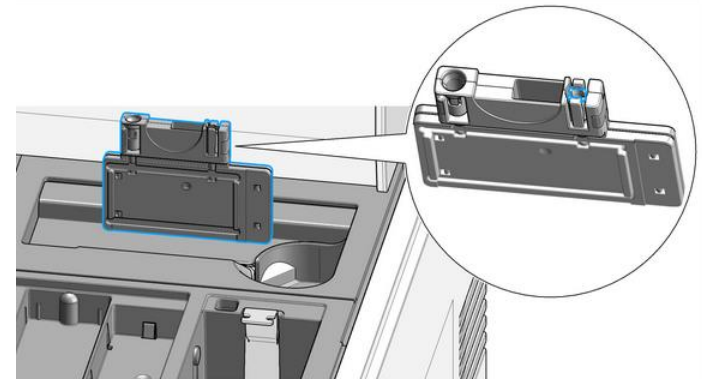


Instrument needle change 4200 TapeStation System

How to change needles:

- Prepare instrument by selecting **Change Needle** from the 4200 TapeStation Controller Software

- Insert Needle Cartridge into the ScreenTape nest
- Follow on-screen instructions to exchange the needle



- After needle change: remove needle change cartridge from the ScreenTape nest and discard
- Wait until Needle Calibration is complete

Preventive Maintenance (PM) Service

- TapeStation Preventive Maintenance service must be performed annually
- PM Service is provided by Agilent authorized engineer **only**
- **Includes:**
 - Fan filter replacement
 - Needle replacement
 - Electrophoresis probes replacement
 - Internal instrument inspection

NOTE: PM service is offered to guarantee good instrument performance.

There is no on-site repair offered for the TapeStation system.

Hardware troubleshooting



How to collect Log files:

The screenshot shows the TapeStation Analysis Software interface. The title bar reads "TapeStation Analysis Software A.02.01 SR1 - 2016-03-02 - 16 44 33.RNA [read-only]". The interface is divided into a left sidebar, a main content area, and a right sidebar.

Step 1: A red arrow points to the "File" menu icon in the top-left corner of the application window.

Step 2: A red arrow points to the "Help" option in the "File" menu, which is highlighted with a red box.

Step 3: A red arrow points to the "Export Log Files" option in the "Tools for working with TapeStation Software" section of the main content area, which is highlighted with a red box.

The main content area includes sections for "Support" (TapeStation Software Help, Contact Us, Agilent Information Center), "Tools for working with TapeStation Software" (Options, Export Log Files), and "About TapeStation Software" (Version: A.02.01 SR1, Copyright © 2011-2016 Agilent Technologies).



How to perform smear analysis?

1- File

2- Options

3- Define regions

4- Apply to File

5- ok

Run Properties

Analysis Software Version: 2.1.4.7682

Assay Options - D1000 High Sensitivity

	Valid	From [bp]	To [bp]	Region Comment
	✗			
	✓	100	300	
	✓	400	500	

Regions have been applied to active file.

Note: If you do not have a ladder assigned to the current file, the regions will not be visible.

☐ Don't show this message again

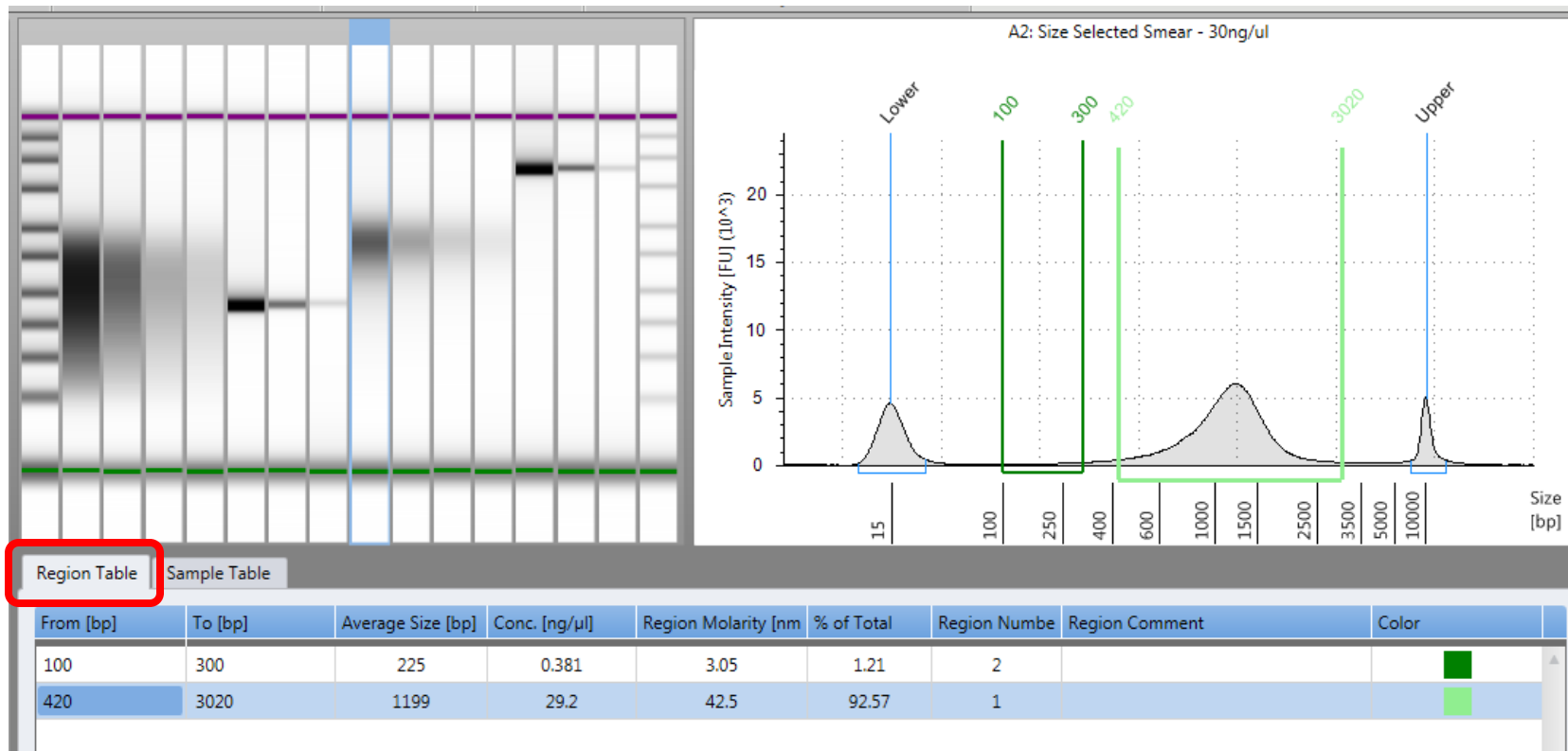
OK

Apply to File

OK Cancel Apply

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How to perform smear analysis?





2200 TapeStation System:

Important points to be aware...

- Flick the ScreenTape prior to use
- **REMOVE LIDS FROM 8 WAY STRIP TUBES**
- Tip buckets must be emptied after every run
- **Do not open the instrument lid during the run**
 - If so: Run will be aborted and data lost
 - Data is only saved to the computer at the end of the run.
- For 96 well plates => Plan ahead of time and ensure you will have the correct number of ScreenTape needed.
 - The controller software will NOT tell how many ScreenTape is required.
- For sizing accuracy: Run a fresh ladder ALWAYS – all assay type
- Do NOT freeze any ScreenTape
- Do NOT mix match reagents and ScreenTape
- Incorrect consumables, such as tips, tubes and seal foil will cause issues with Blank lane

