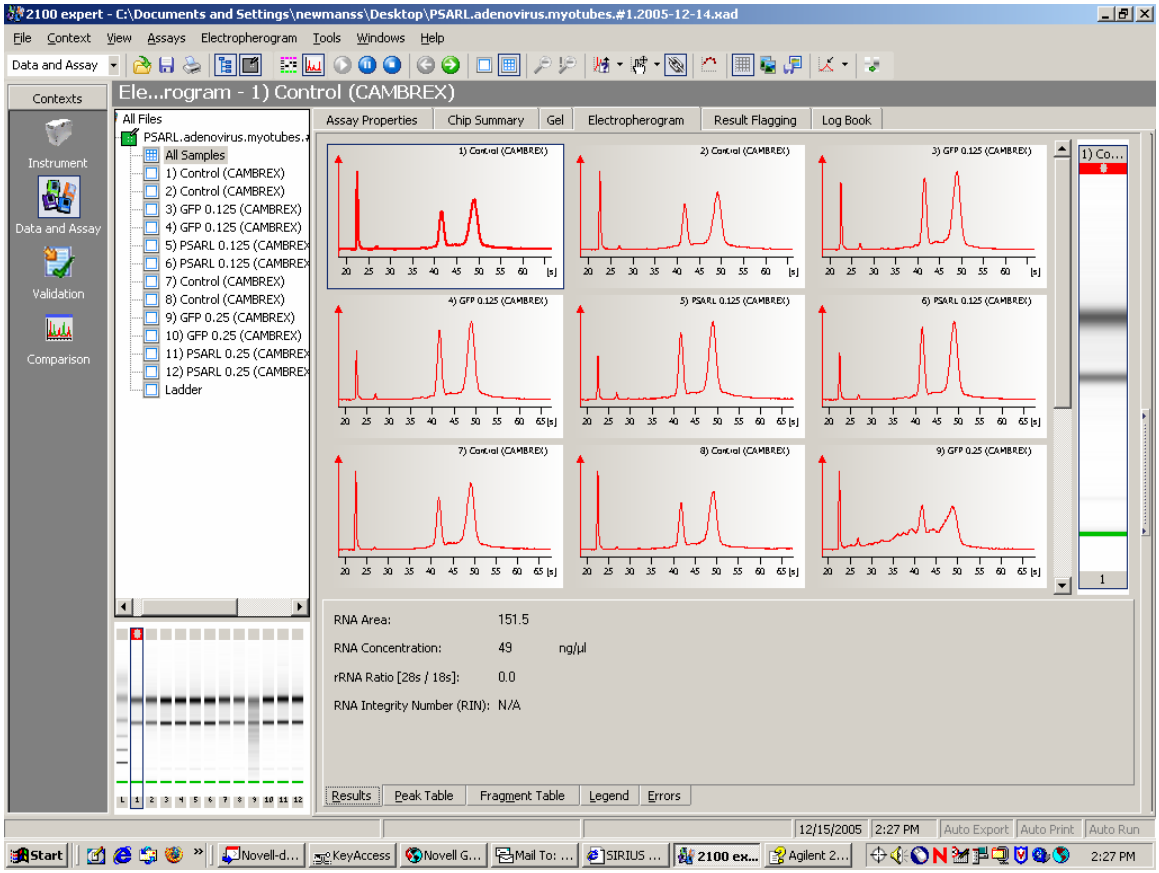
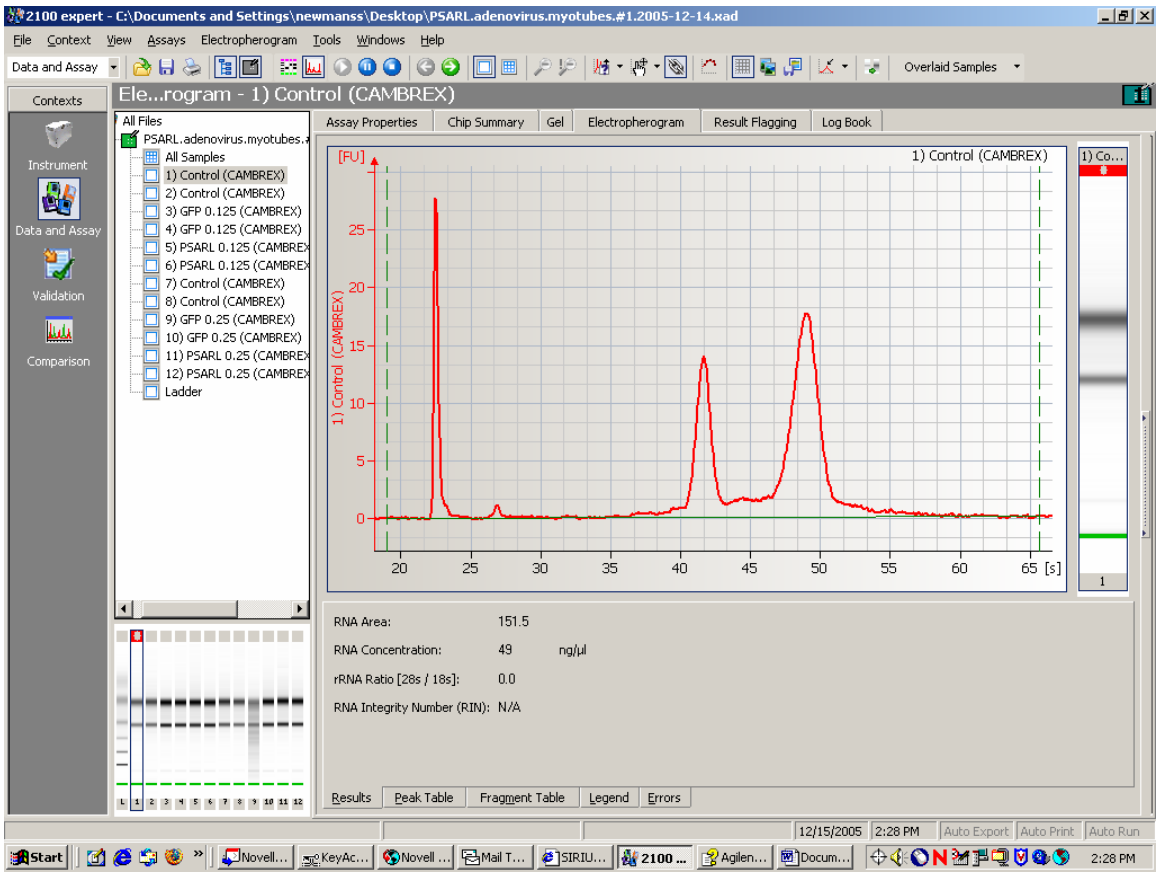
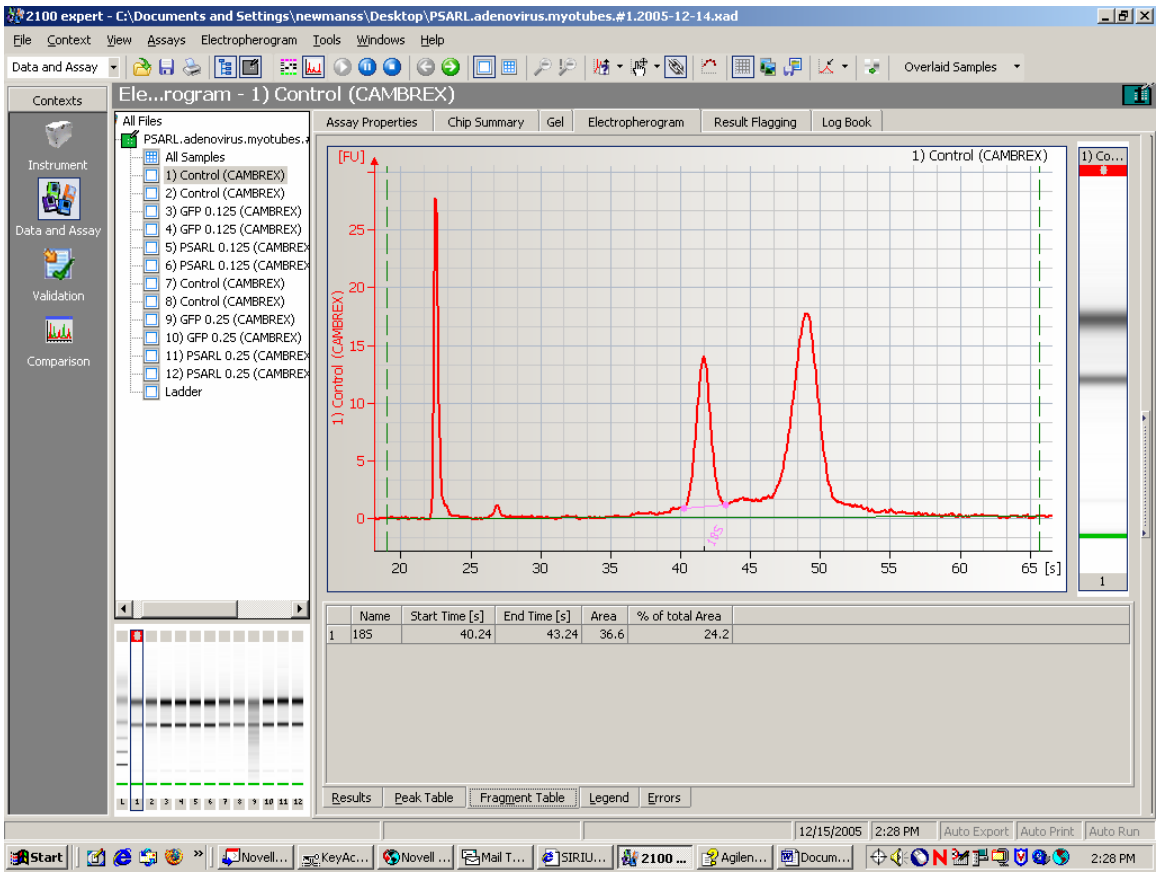
Correcting Missed or Incorrect Peak Calls



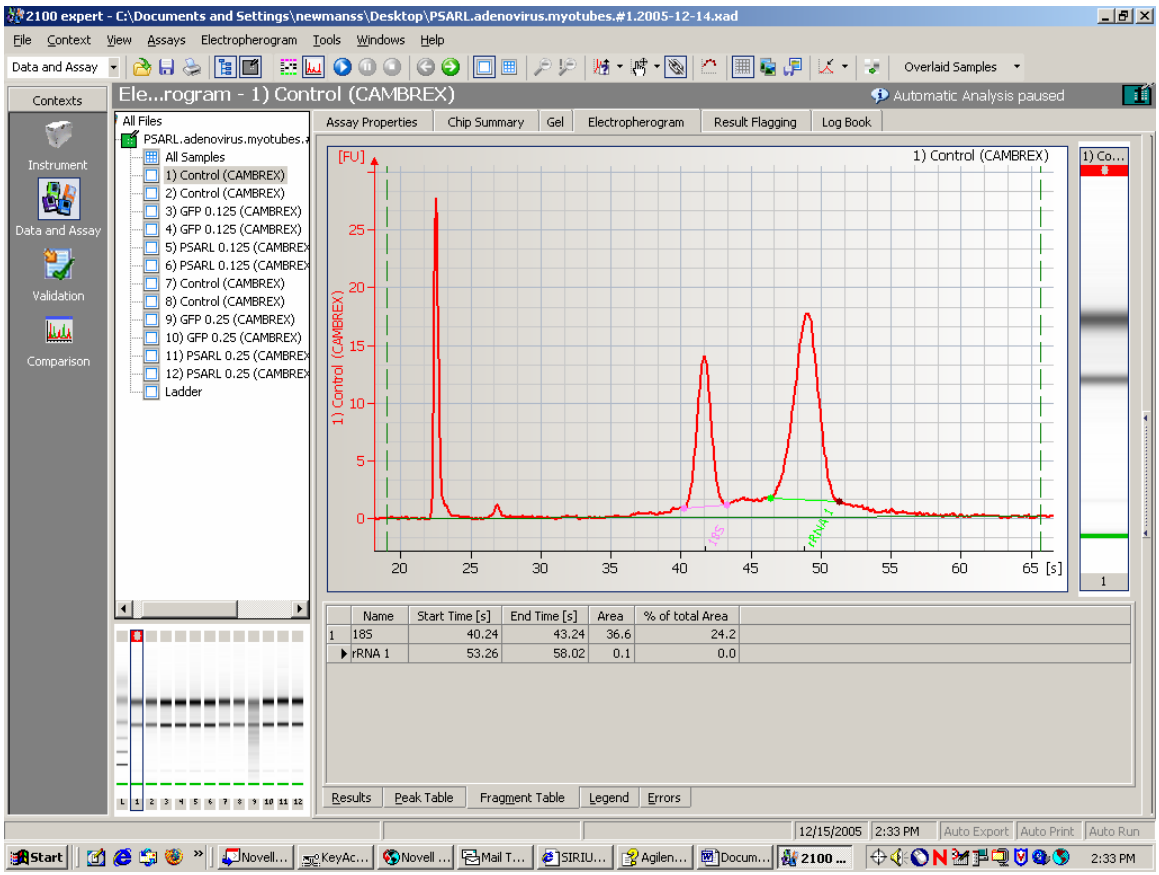
Double click on the electropherogram in question to select it.



Select the Fragment Table tab at the bottom of the screen.

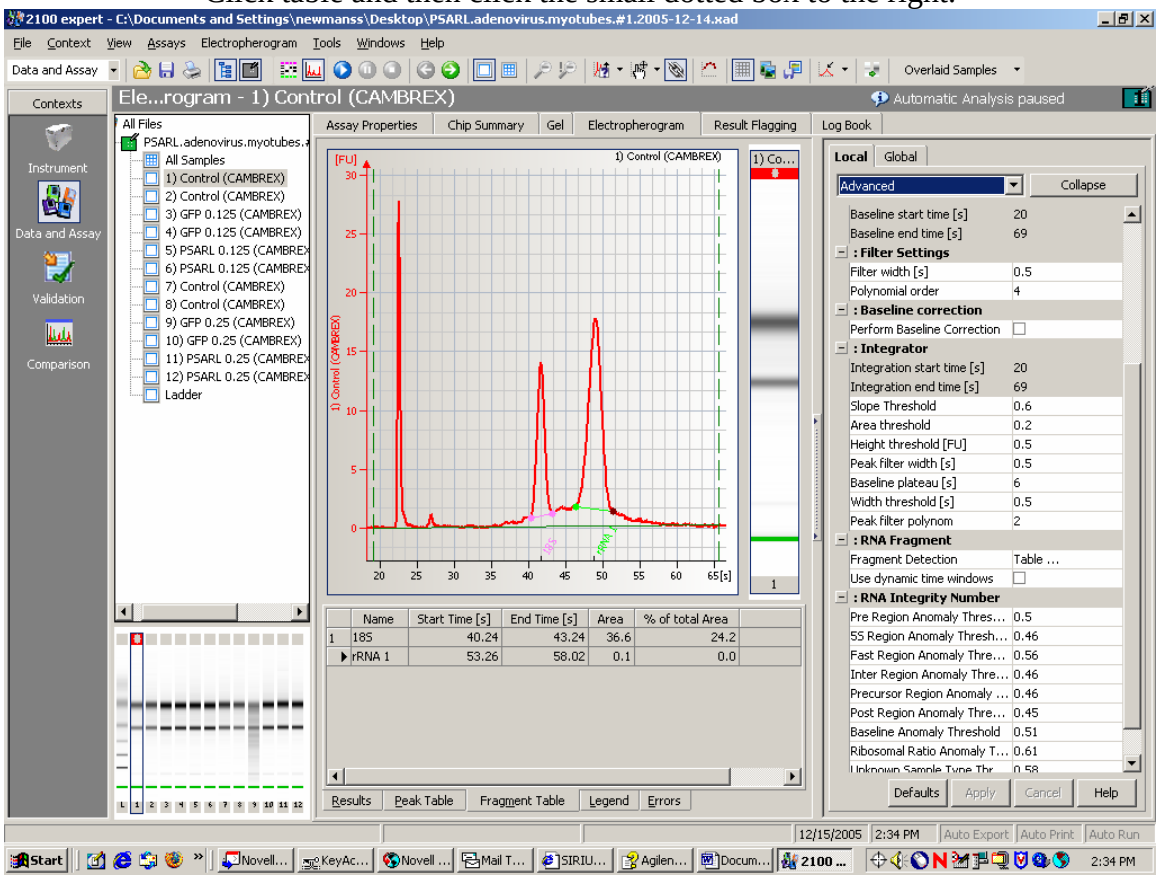


Right click on the graph and select “Add Fragment” in the popup window.
On the menu at the top of the screen select Electropherogram>Pause Analysis.
A new green bar will be added to the electropherogram. Click on each end of the bar and drag the bar-ends to the correct locations at the beginning and end of the fragment.

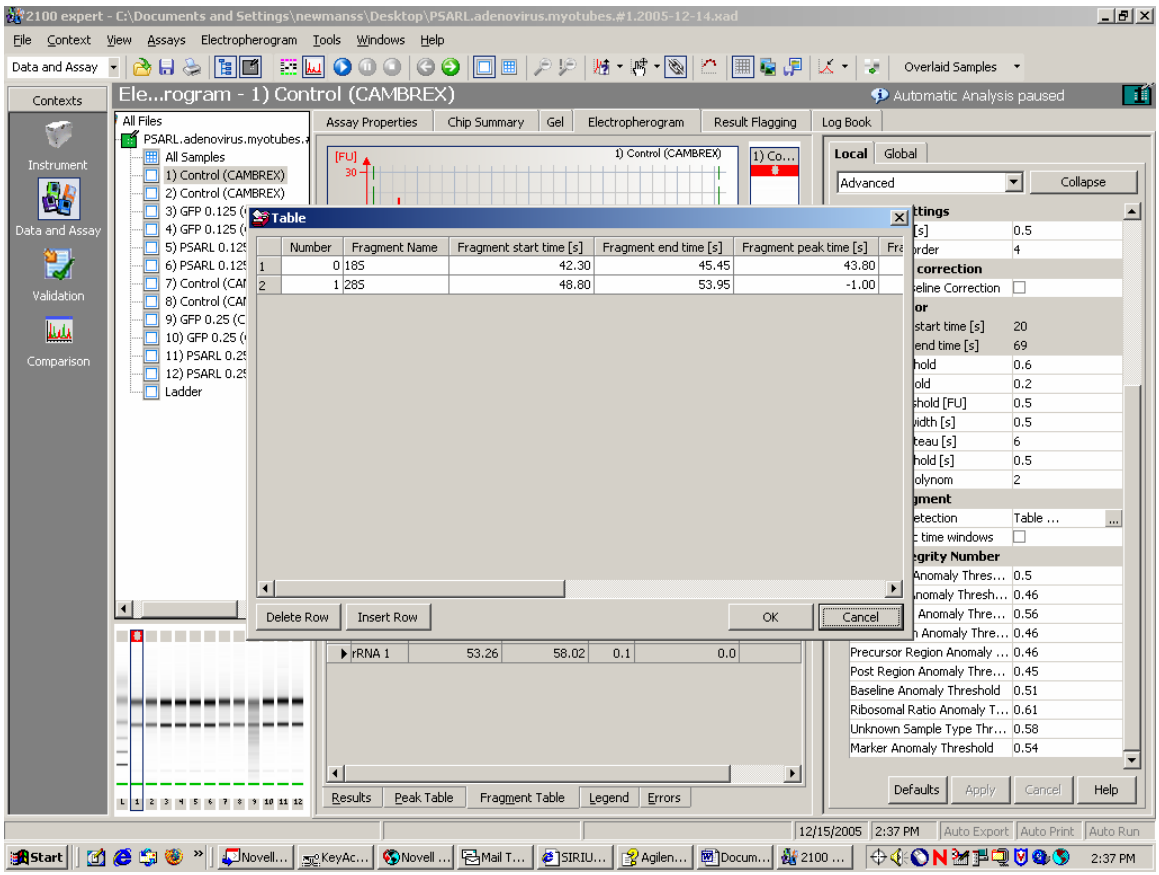


To open the “Setpoint Explorer”, double click on the dotted line found on the bar at the right edge of your screen. In the Setpoint Explorer, select the Local tab. Select “Advanced” from the drop down menu. Scroll down until you see “:RNA Fragment”.

Click table and then click the small dotted box to the right.



A new window will open. Click on the fragment name that needs to be corrected until it is highlighted. Rename appropriately (18S or 28S). Click OK.



Under the menu, select Electropherogram then select analyze. Select the results tab to view the corrected data.