

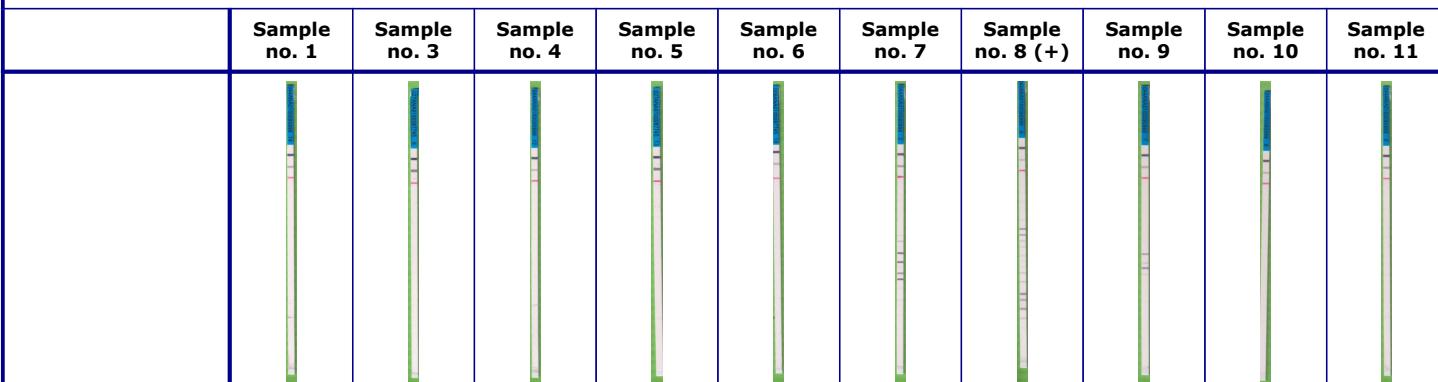
## Summary results

### BLOT-LINE ANA

Lot: 23904752  
 Performed by: N.VIJAYA  
 Evaluation time: 11/14/2023 11:50:36 AM

Borderline area from 8.00 to 10.00

Scan image(1 - 11):



Results(1 - 11):

	Sample no. 1	Sample no. 3	Sample no. 4	Sample no. 5	Sample no. 6	Sample no. 7	Sample no. 8 (+)	Sample no. 9	Sample no. 10	Sample no. 11
<b>Description</b>					Mr. KUKATLA NARSING RAO					
<b>IgG Control</b>	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓
<b>Jo-1</b>	1.01[N]	0.50[N]	0.00[N]	0.49[N]	0.97[N]	0.96[N]	0.00[N]	0.47[N]	0.00[N]	0.00[N]
<b>PL-7</b>	0.00[N]	0.99[N]	0.50[N]	0.00[N]	0.00[N]	0.00[N]	0.48[N]	0.47[N]	0.00[N]	0.00[N]
<b>PL-12</b>	0.00[N]	1.49[N]	1.50[N]	2.93[N]	0.00[N]	1.45[N]	1.91[N]	0.00[N]	0.00[N]	0.00[N]
<b>PmScl</b>	0.51[N]	0.00[N]	1.00[N]	3.41[N]	0.49[N]	0.00[N]	2.85[N]	0.47[N]	0.00[N]	0.48[N]
<b>CENP-A</b>	0.51[N]	0.00[N]	0.00[N]	1.47[N]	0.00[N]	0.49[N]	1.91[N]	0.00[N]	0.00[N]	0.00[N]
<b>CENP-B</b>	0.00[N]	0.50[N]	0.50[N]	0.00[N]	0.00[N]	0.00[N]	0.00[N]	0.47[N]	0.00[N]	0.00[N]
<b>Scl-70</b>	1.02[N]	0.00[N]	0.00[N]	2.93[N]	0.00[N]	0.00[N]	3.81[N]	0.47[N]	0.47[N]	0.96[N]
<b>Ro52</b>	1.01[N]	0.00[N]	1.50[N]	0.49[N]	0.00[N]	3.85[N]	<b>28.59[P]</b>	0.00[N]	0.00[N]	2.37[N]
<b>Ro60</b>	0.00[N]	0.00[N]	0.50[N]	0.99[N]	0.00[N]	<b>12.72[P]</b>	<b>21.57[P]</b>	0.47[N]	0.95[N]	0.95[N]
<b>La</b>	0.00[N]	0.50[N]	0.00[N]	0.99[N]	0.00[N]	0.00[N]	8.96[B]	0.00[N]	0.00[N]	0.00[N]
<b>RNP-A</b>	0.51[N]	0.50[N]	1.01[N]	0.49[N]	0.00[N]	<b>42.98[P]</b>	6.17[N]	<b>24.10[P]</b>	1.42[N]	0.00[N]
<b>RNP-C</b>	1.02[N]	0.50[N]	1.00[N]	1.96[N]	0.00[N]	<b>41.28[P]</b>	4.77[N]	8.88[B]	0.00[N]	0.48[N]
<b>RNP-68</b>	0.51[N]	1.50[N]	0.00[N]	1.48[N]	0.49[N]	2.42[N]	1.93[N]	<b>28.50[P]</b>	0.00[N]	0.48[N]
<b>SmB</b>	0.00[N]	1.52[N]	1.50[N]	0.00[N]	0.00[N]	<b>29.22[P]</b>	2.88[N]	5.66[N]	0.95[N]	2.39[N]
<b>SmD</b>	0.00[N]	1.02[N]	0.00[N]	0.00[N]	0.00[N]	<b>44.32[P]</b>	9.91[B]	0.48[N]	0.00[N]	0.48[N]
<b>P0</b>	0.00[N]	1.52[N]	1.00[N]	0.49[N]	0.00[N]	3.40[N]	<b>10.83[P]</b>	0.48[N]	0.48[N]	0.48[N]
<b>PCNA</b>	0.00[N]	1.02[N]	1.00[N]	0.00[N]	0.00[N]	0.00[N]	0.00[N]	0.48[N]	0.48[N]	0.00[N]
<b>dsDNA</b>	1.51[N]	1.51[N]	8.73[B]	0.98[N]	0.98[N]	3.88[N]	<b>31.29[P]</b>	1.44[N]	1.90[N]	0.96[N]
<b>Histone</b>	4.48[N]	1.01[N]	0.50[N]	0.49[N]	0.49[N]	0.98[N]	<b>19.25[P]</b>	0.00[N]	0.95[N]	0.48[N]
<b>Nucleosom</b>	<b>19.97[P]!</b>	1.01[N]	1.49[N]	2.44[N]	0.98[N]	1.47[N]	<b>26.99[P]</b>	0.96[N]	0.95[N]	2.87[N]
<b>DFS70</b>	1.00[N]	1.02[N]	1.49[N]	0.49[N]	0.00[N]	0.00[N]	<b>10.88[P]</b>	0.48[N]	0.00[N]	0.00[N]
<b>M2</b>	0.50[N]	1.02[N]	0.00[N]	0.49[N]	0.00[N]	0.00[N]	2.42[N]	0.00[N]	0.00[N]	0.00[N]
<b>ANA</b>	<b>Positive</b>	Negative	Borderline	Negative	Negative	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	Negative	Negative

Strips assigned (+) - In case of negative result of ANA and positive result of DFS70: low probability of SARD. In case of positive result of M2: potential risk of primary biliary cirrhosis progression.

Check the assignment of bands marked [!] because of a possible interference with the strip background or unclear band interpretation

## Summary results

### BLOT-LINE ANA

					Borderline area from 8.00 to 10.00					
					Borderline area from 8.00 to 10.00					
Scan image(12 - 20):										
	Sample no. 12	Sample no. 13	Sample no. 14	Sample no. 15	Sample no. 16	Sample no. 17	Sample no. 18 (#)	Sample no. 19	Sample no. 20	
										
Results(12 - 20):										
	Sample no. 12	Sample no. 13	Sample no. 14	Sample no. 15	Sample no. 16	Sample no. 17	Sample no. 18 (#)	Sample no. 19	Sample no. 20	
Description										
<b>IgG Control</b>	✓	✓	✓	✓	X	✓	✓	✓	✓	
<b>Jo-1</b>	0.00[N]	0.00[N]	2.46[N]	0.00[N]	0.00[N]	0.00[N]	<b>42.76[P]</b>	0.00[N]	1.53[N]	
<b>PL-7</b>	0.97[N]	0.00[N]	0.50[N]	0.98[N]	2.00[N]	0.00[N]	0.00[N]	0.00[N]	0.00[N]	
<b>PL-12</b>	0.98[N]	0.50[N]	0.50[N]	0.00[N]	0.00[N]	0.00[N]	0.00[N]	3.96[N]	0.00[N]	
<b>PmScl</b>	4.82[N]	0.49[N]	0.50[N]	0.00[N]	1.51[N]	0.50[N]	0.00[N]	1.99[N]	0.51[N]	
<b>CENP-A</b>	0.49[N]	0.00[N]	0.99[N]	0.99[N]	0.00[N]	0.50[N]	0.00[N]	1.50[N]	1.02[N]	
<b>CENP-B</b>	0.00[N]	0.00[N]	0.00[N]	0.00[N]	1.51[N]	0.50[N]	0.51[N]	1.99[N]	0.51[N]	
<b>Scl-70</b>	0.49[N]	0.00[N]	5.82[N]	0.00[N]	0.50[N]	0.00[N]	0.00[N]	0.00[N]	0.51[N]	
<b>Ro52</b>	<b>21.74[P]</b>	<b>24.14[P]</b>	1.48[N]	0.00[N]	0.00[N]	<b>36.59[P]</b>	1.01[N]	2.98[N]	9.40[B]	
<b>Ro60</b>	<b>22.18[P]</b>	<b>45.23[P]</b>	1.48[N]	<b>18.77[P]</b>	1.49[N]	<b>43.89[P]</b>	0.00[N]	1.00[N]	0.00[N]	
<b>La</b>	<b>22.67[P]</b>	<b>46.08[P]</b>	0.50[N]	0.00[N]	3.44[N]	<b>29.61[P]</b>	<b>34.56[P]</b>	1.00[N]	0.51[N]	
<b>RNP-A</b>	1.46[N]	3.89[N]	2.45[N]	2.94[N]	0.00[N]	1.00[N]	<b>26.71[P]</b>	0.00[N]	0.00[N]	
<b>RNP-C</b>	3.38[N]	0.49[N]	4.38[N]	6.27[N]	0.00[N]	0.00[N]	<b>45.83[P]</b>	0.00[N]	0.00[N]	
<b>RNP-68</b>	1.46[N]	0.99[N]	4.38[N]	0.49[N]	0.00[N]	0.50[N]	0.00[N]	0.50[N]	0.51[N]	
<b>SmB</b>	3.85[N]	0.00[N]	2.45[N]	5.80[N]	0.00[N]	0.50[N]	<b>43.54[P]</b>	2.98[N]	1.02[N]	
<b>SmD</b>	0.00[N]	0.49[N]	<b>14.68[P]</b>	0.00[N]	1.00[N]	0.00[N]	0.51[N]	0.50[N]	0.00[N]	
<b>P0</b>	0.00[N]	0.49[N]	0.49[N]	0.00[N]	0.00[N]	0.00[N]	0.51[N]	1.51[N]	1.02[N]	
<b>PCNA</b>	1.94[N]	0.00[N]	0.49[N]	0.00[N]	0.00[N]	0.50[N]	0.00[N]	1.01[N]	0.00[N]	
<b>dsDNA</b>	1.46[N]	1.48[N]	<b>10.06[P]</b>	0.98[N]	1.99[N]	1.00[N]	0.00[N]	1.50[N]	1.53[N]	
<b>Histone</b>	2.91[N]	0.49[N]	<b>17.37[P]</b>	0.98[N]	0.50[N]	0.50[N]	2.52[N]	1.00[N]	0.00[N]	
<b>Nucleosom</b>	1.94[N]	1.96[N]	6.27[N]	1.95[N]	1.00[N]	1.49[N]	1.52[N]	<b>13.87[P]</b>	1.02[N]	
<b>DFS70</b>	0.00[N]	0.49[N]	0.00[N]	0.00[N]	0.50[N]	0.00[N]	1.51[N]	0.50[N]	0.00[N]	
<b>M2</b>	1.94[N]	0.00[N]	0.00[N]	0.00[N]	0.50[N]	0.00[N]	0.00[N]	0.00[N]	0.00[N]	
<b>ANA</b>	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	Negative	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	Borderline
Strips assigned (+) - In case of negative result of ANA and positive result of DFS70: low probability of SARD. In case of positive result of M2: potential risk of primary biliary cirrhosis progression.										
Strips assigned (#) - Control line located incorrectly, SW evaluation result is not valid										
X - Conjugate control line was not found. Please check the possibility of substitution										