

## Summary results

### BLOT-LINE ANA

					Summary results BLOT-LINE ANA									
Lot: 23290666 Performed by: n.vijaya Evaluation time: 9/27/2023 12:59:48 PM					Borderline area from 8.00 to 10.00									
Scan image(1 - 10):														
	Sample no. 1	Sample no. 2	Sample no. 3	Sample no. 4	Sample no. 5	Sample no. 6	Sample no. 7	Sample no. 8 (+)	Sample no. 9	Sample no. 10				
														
Results(1 - 10):														
	Sample no. 1	Sample no. 2	Sample no. 3	Sample no. 4	Sample no. 5	Sample no. 6	Sample no. 7	Sample no. 8 (+)	Sample no. 9	Sample no. 10				
Description				Mrs. SHRADDAH BIDKAR	Dr. MADHAVI RUDRA									
<b>IgG Control</b>	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓				
<b>Jo-1</b>	<b>44.82[P]</b>	0.51[N]	1.98[N]	0.00[N]	0.98[N]	0.49[N]	0.97[N]	0.00[N]	0.00[N]	0.47[N]				
<b>PL-7</b>	1.00[N]	0.51[N]	0.50[N]	0.00[N]	0.00[N]	0.49[N]	0.49[N]	0.00[N]	0.47[N]	0.00[N]				
<b>PL-12</b>	2.00[N]	0.51[N]	0.50[N]	0.49[N]	2.44[N]	0.49[N]	1.45[N]	1.43[N]	0.47[N]	1.42[N]				
<b>PmScl</b>	1.01[N]	0.00[N]	0.00[N]	1.48[N]	0.99[N]	0.98[N]	0.00[N]	2.86[N]	0.47[N]	0.00[N]				
<b>CENP-A</b>	2.01[N]	<b>18.14[P]</b>	0.50[N]	0.00[N]	1.96[N]	0.00[N]	0.00[N]	1.91[N]	0.00[N]	0.00[N]				
<b>CENP-B</b>	0.00[N]	0.51[N]	0.00[N]	0.00[N]	0.49[N]	0.49[N]	0.00[N]	0.00[N]	0.48[N]	0.00[N]				
<b>Scl-70</b>	2.01[N]	<b>35.06[P]</b>	0.00[N]	0.00[N]	0.00[N]	0.49[N]	0.00[N]	3.82[N]	0.00[N]	0.00[N]				
<b>Ro52</b>	5.92[N]	<b>19.51[P]</b>	0.00[N]	1.48[N]	0.99[N]	0.00[N]	3.87[N]	<b>29.51[P]</b>	0.00[N]	0.00[N]				
<b>Ro60</b>	<b>28.86[P]</b>	0.00[N]	0.00[N]	1.96[N]	0.00[N]	0.00[N]	<b>13.20[P]</b>	<b>22.94[P]</b>	1.43[N]	0.95[N]				
<b>La</b>	1.51[N]	0.00[N]	0.00[N]	1.46[N]	0.99[N]	0.00[N]	0.00[N]	8.98[B]	0.00[N]	0.00[N]				
<b>RNP-A</b>	0.00[N]	0.00[N]	1.00[N]	0.49[N]	1.48[N]	0.00[N]	<b>43.47[P]</b>	6.19[N]	<b>24.10[P]</b>	0.00[N]				
<b>RNP-C</b>	0.00[N]	0.00[N]	0.50[N]	0.49[N]	1.96[N]	0.00[N]	<b>41.77[P]</b>	5.26[N]	8.42[B]	0.00[N]				
<b>RNP-68</b>	1.01[N]	2.51[N]	1.01[N]	0.00[N]	0.99[N]	0.49[N]	2.91[N]	1.93[N]	<b>28.93[P]</b>	0.48[N]				
<b>SmB</b>	0.50[N]	4.46[N]	1.02[N]	2.44[N]	0.49[N]	0.00[N]	<b>30.15[P]</b>	2.41[N]	5.68[N]	0.48[N]				
<b>SmD</b>	4.46[N]	8.79[B]!	1.02[N]	0.49[N]	0.49[N]	0.00[N]	<b>44.81[P]</b>	9.93[B]	0.48[N]	0.00[N]				
<b>P0</b>	0.00[N]	0.00[N]	1.02[N]	0.49[N]	0.99[N]	0.00[N]	3.41[N]	<b>10.85[P]</b>	0.00[N]	0.00[N]				
<b>PCNA</b>	0.00[N]	0.98[N]	1.02[N]	1.47[N]	0.49[N]	0.49[N]	0.49[N]	0.00[N]	0.48[N]	0.00[N]				
<b>dsDNA</b>	2.50[N]	0.00[N]	1.52[N]	2.93[N]	1.48[N]	1.48[N]	3.89[N]	<b>32.21[P]</b>	0.96[N]	1.43[N]				
<b>Histone</b>	2.50[N]	2.96[N]	1.02[N]	1.47[N]	0.49[N]	0.49[N]	1.47[N]	<b>19.73[P]</b>	0.48[N]	0.96[N]				
<b>Nucleosom</b>	2.50[N]	1.98[N]	1.02[N]	7.65[N]	2.94[N]	0.00[N]	1.96[N]	<b>27.42[P]</b>	1.45[N]	1.43[N]				
<b>DFS70</b>	1.01[N]	0.00[N]	0.00[N]	2.94[N]	0.00[N]	0.00[N]	0.49[N]	<b>10.88[P]</b>	0.00[N]	0.48[N]				
<b>M2</b>	1.51[N]	1.49[N]	0.00[N]	0.50[N]	0.49[N]	0.00[N]	0.00[N]	2.43[N]	0.00[N]	0.00[N]				
<b>ANA</b>	<b>Positive</b>	<b>Positive</b>	Negative	Negative	Negative	Negative	Negative	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	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

				Summary results BLOT-LINE ANA					
Lot: 23290666 Performed by: n.vijaya Evaluation time: 9/27/2023 12:59:48 PM				Borderline area from 8.00 to 10.00					
Scan image(11 - 19):									
	Sample no. 11	Sample no. 12	Sample no. 14	Sample no. 15	Sample no. 16	Sample no. 17	Sample no. 18	Sample no. 19	
									
Results(11 - 19):									
	Sample no. 11	Sample no. 12	Sample no. 14	Sample no. 15	Sample no. 16	Sample no. 17	Sample no. 18	Sample no. 19	
<b>Description</b>									
<b>IgG Control</b>	√	√	√	√	X	√	√	√	
<b>Jo-1</b>	0.00[N]	0.00[N]	1.96[N]	0.00[N]	0.50[N]	0.00[N]	0.00[N]	1.01[N]	
<b>PL-7</b>	0.48[N]	0.98[N]	0.50[N]	1.48[N]	1.01[N]	1.01[N]	0.51[N]	0.51[N]	
<b>PL-12</b>	0.48[N]	1.47[N]	0.99[N]	0.00[N]	0.00[N]	1.01[N]	1.02[N]	0.00[N]	
<b>PmScl</b>	0.48[N]	4.85[N]	0.99[N]	0.00[N]	1.01[N]	0.00[N]	0.00[N]	0.51[N]	
<b>CENP-A</b>	0.00[N]	0.98[N]	0.99[N]	0.49[N]	0.00[N]	0.00[N]	0.00[N]	0.00[N]	
<b>CENP-B</b>	0.00[N]	0.49[N]	0.00[N]	1.48[N]	1.51[N]	0.00[N]	0.00[N]	1.02[N]	
<b>Scl-70</b>	0.00[N]	0.00[N]	4.86[N]	0.50[N]	0.00[N]	0.00[N]	1.53[N]	2.52[N]	
<b>Ro52</b>	2.39[N]	<b>23.16[P]</b>	1.48[N]	0.00[N]	1.49[N]	2.01[N]	<b>29.06[P]</b>	0.00[N]	
<b>Ro60</b>	0.96[N]	<b>23.16[P]</b>	0.99[N]	<b>19.72[P]</b>	0.00[N]	0.51[N]	3.53[N]	0.00[N]	
<b>La</b>	1.91[N]	<b>23.16[P]</b>	0.00[N]	0.50[N]	3.47[N]	0.00[N]	0.51[N]	0.00[N]	
<b>RNP-A</b>	0.48[N]	0.98[N]	2.45[N]	3.42[N]	0.00[N]	1.02[N]	3.51[N]	0.00[N]	
<b>RNP-C</b>	0.48[N]	4.83[N]	4.86[N]	6.77[N]	0.00[N]	0.00[N]	<b>21.38[P]</b>	1.52[N]	
<b>RNP-68</b>	0.48[N]	1.47[N]	4.38[N]	0.49[N]	0.50[N]	0.00[N]	0.51[N]	0.51[N]	
<b>SmB</b>	2.39[N]	4.36[N]	2.94[N]	0.00[N]	0.00[N]	2.01[N]	1.52[N]	0.00[N]	
<b>SmD</b>	0.00[N]	0.00[N]	<b>15.59[P]</b>	0.00[N]	0.50[N]	0.00[N]	0.00[N]	0.51[N]	
<b>P0</b>	0.48[N]	0.00[N]	0.99[N]	0.00[N]	0.00[N]	0.51[N]	0.00[N]	1.02[N]	
<b>PCNA</b>	0.96[N]	1.47[N]	0.49[N]	0.49[N]	0.00[N]	1.01[N]	0.00[N]	0.00[N]	
<b>dsDNA</b>	1.44[N]	0.98[N]	<b>10.50[P]</b>	0.49[N]	1.99[N]	1.51[N]	0.51[N]	1.52[N]	
<b>Histone</b>	0.48[N]	2.92[N]	<b>18.28[P]</b>	0.98[N]	1.00[N]	1.01[N]	1.02[N]	1.02[N]	
<b>Nucleosom</b>	2.87[N]	2.92[N]	6.75[N]	2.44[N]	1.51[N]	1.51[N]	1.52[N]	3.02[N]	
<b>DFS70</b>	0.00[N]	0.00[N]	1.47[N]	0.00[N]	0.50[N]	0.00[N]	0.00[N]	1.01[N]	
<b>M2</b>	0.00[N]	2.43[N]	0.49[N]	0.00[N]	0.00[N]	0.00[N]	0.51[N]	1.01[N]	
<b>ANA</b>	Negative	<b>Positive</b>	<b>Positive</b>	<b>Positive</b>	Negative	Negative	<b>Positive</b>	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.									
X - Conjugate control line was not found. Please check the possibility of substitution									