

AFFORDABLE ANALYTICAL SERVICE FOR CAPPING EFFICIENCY



Capping efficiency and poly(A) length are considered critical quality attributes (CQA) for mRNA. Areterna uses the LC-MS method to identify and quantify mRNA. Simply provide us with the 5'- and 3'- sequences of your mRNA construct, and we will design and run the LC-MS assays for you, delivering accurate and reproducible results.



Our Advantages



EXPERIENCED TEAM

>400 capping efficiency and poly(A) length assays have been performed



AFFORDABLE PRICE

\$950 per sample, per assay*



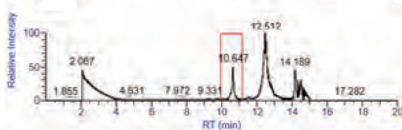
QUICK RESULTS

2 weeks turn-around time upon receipt of the sample

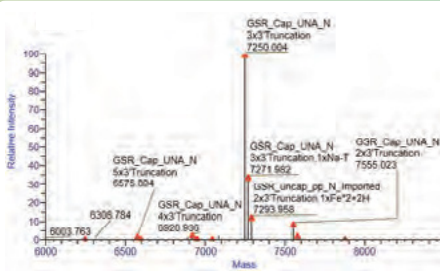
TESTING CASE 1

The capping efficiency is 99.3%

Total Ion Chromatography



Deconvolution Mass Spectrum



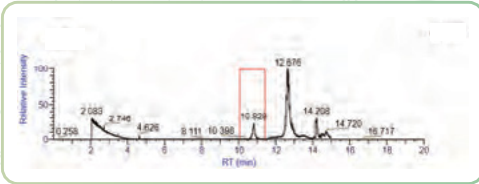
Sequence Name	Modification	Monoisotopic Mass	Theoretical Mass (Da)	Matched Mass Error (ppm)	Average Mass	Sum Intensity
GSR_Cap_UNA_N	3x3 Truncation	7250.004	7250.012	1.2	7253.39	1.61E+07
GSR_Cap_UNA_N	3x3 Truncation, 1xNa-T	7271.982	7271.993	1.6	7275.38	5.45E+06
GSR_Cap_UNA_N	3x3 Truncation, 2xNa-T	7293.958	7293.843	15.7	7297.37	2.01E+06
GSR_Cap_UNA_N	2x3 Truncation	7555.023	7555.054	4.0	7558.55	1.45E+06
GSR_Cap_UNA_N	4x3 Truncation	6920.936	6920.960	3.5	6924.17	4.80E+05
GSR_Cap_UNA_N	2x3 Truncation, 1xNa-T	7577.001	7577.035	4.4	7580.54	4.06E+05
GSR_Cap_UNA_N	5x3 Truncation	6575.894	6575.912	2.8	6578.96	3.04E+05
GSR_Cap_UNA_N	4x3 Truncation, 1xNa-T	6942.911	6942.941	4.3	6946.16	1.18E+05
GSR_uncap_N	2x3 Truncation, 1xNa	7042.910	7043.017	15.2	7046.20	1.17E+05
GSR_Cap_UNA_N	6x3 Truncation	6246.849	6246.860	1.8	6249.77	1.14E+05
GSR_Cap_UNA_N	1x3 Truncation	7875.057	7875.095	4.7	7878.74	1.00E+05
GSR_uncap_ppp_N	3x3 Truncation	6954.851	6954.884	4.7	6958.10	8.23E+04
GSR_Cap_UNA_N	5x3 Truncation, 1xNa-T	6597.881	6597.893	1.8	6600.96	7.89E+04

*Price is subject to change without notice

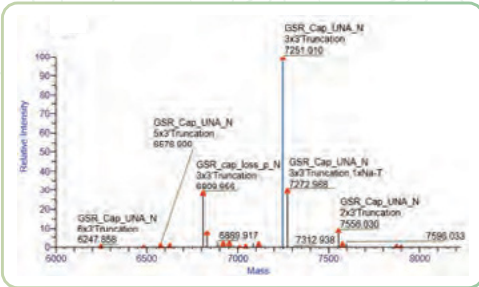
TESTING CASE 2

The capping efficiency is 77.3%

Total Ion Chromatography



Deconvolution Mass Spectrum



Sequence Name	Modification	Monoisotopic Mass	Theoretical Mass (Da)	Matched Mass Error (ppm)	Average Mass	Sum Intensity
GSR_Cap_UNA_N	3x3 Truncation	7251.010	7251.020	1.4	7254.40	8.46E+06
GSR_Cap_UNA_N	3x3 Truncation, 1xNa-T	7272.988	7273.001	1.8	7276.39	2.53E+06
GSR_cap_loss_p_N	3x3 Truncation	6809.956	6809.975	2.7	6813.14	2.44E+06
GSR_Cap_UNA_N	2x3 Truncation	7556.030	7556.061	4.1	7559.56	7.60E+05
GSR_cap_loss_p_N	3x3 Truncation, 1xNa-T	6831.928	6831.956	4.0	6835.12	6.58E+05
GSR_uncap_ppp_N_Imported	3x3 Truncation	6955.878	6955.892	2.0	6959.13	1.82E+05
GSR_Cap_UNA_N	4x3 Truncation	6921.928	6921.967	5.7	6925.16	1.79E+05
GSR_cap_loss_p_N	2x3 Truncation	7114.979	7115.016	5.1	7118.31	1.50E+05
GSR_Cap_UNA_N	2x3 Truncation, 1xNa-T	7577.999	7578.042	5.7	7581.54	1.50E+05
GSR_Cap_UNA_N	5x3 Truncation	6576.900	6576.920	3.0	6579.97	8.66E+04
GSR_uncap_ppp_N_Imported	4x3 Truncation	6626.811	6626.839	4.2	6629.90	7.97E+04
GSR_uncap_N_Imported	2x3 Truncation, 1xNa	7043.910	7044.024	16.1	7047.20	6.88E+04
GSR_Cap_UNA_N	6x3 Truncation	6247.858	6247.867	1.4	6250.78	6.09E+04
GSR_Cap_UNA_N	1x3 Truncation	7876.070	7876.102	4.1	7879.75	5.98E+04
GSR_cap_loss_p_N	4x3 Truncation	6480.903	6480.922	2.9	6483.92	2.36E+04
GSR_Cap_UNA_N	1x3 Truncation, 1xNa	7899.063	7899.092	3.7	7902.75	1.94E+04
GSR_uncap_ppp_N_Imported	3x3 Truncation, 1xFe-T	7012.788	7012.802	1.9	7016.07	6.38E+03

For customers who wish to run their own capping and tailing assays, Areterna offers analytical method development services.

- Design construct-specific cleavage probes.
- Develop construct-specific analytical methods and tech transfer to customers



Other Analytical Services for Analytical Method Development and Validation

