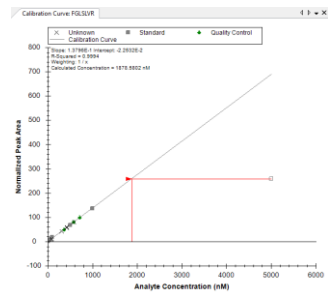


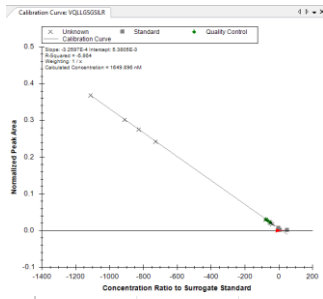
1)

Protein	Protein Note	Peptide	Replicate Name	Sample Dilution Factor	Average Measured Retention Time	Standard Type	Total Area	Calculated Concentration	Calibration Curve	Conc Multi	Normalization Method
	standard	FGLSLVR	01	1	3.1	Surrog...	129	1.906...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	05	1	3.1	Surrog...	741	3.071...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	1	1	3.1	Surrog...	2123	5.732...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	5	1	3.1	Surrog...	22365	4.493...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	10	1	3.1	Surrog...	43958	8.480...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	50	1	3.1	Surrog...	270825	4.927...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	100	1	3.1	Surrog...	643964	1.199...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	500	1	3.1	Surrog...	2799750	4.909...	Slope: 1.3796E...		Ratio to Global S...
	standard	FGLSLVR	1000	1	3.1	Surrog...	5632577	9.923...	Slope: 1.3796E...		Ratio to Global S...

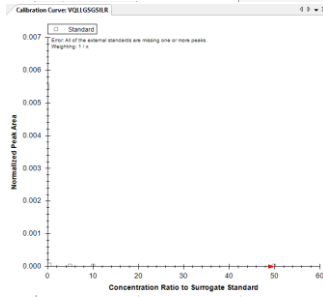


2)

3)



Protein	Protein Note	Peptide	Replicat Name	Sample Dilution Factor	Average Measured Retention Time	Standard Type	Total Area	Calculat Concent	Calibration Curve	Conc Multi	Normalization Method
		VQLLGSGSILR	1	1	3.22		12	-3.301...	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	5	1	3.22		#N/A	#N/A	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	10	1	3.22		#N/A	#N/A	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	50	1	3.22		12	1.636...	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	100	1	3.22		#N/A	#N/A	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	500	1	3.22		31	1.647...	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	1000	1	3.22		8	1.650...	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	5000	1	3.22		26	1.649...	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	GW1	6.52	3.22		63072	-4.736...	Slope: -3.2597E...		Ratio to surrogat...
		VQLLGSGSILR	GW2	6.52	3.22		77664	-7.241...	Slope: -3.2597E...		Ratio to surrogat...



Protein	Protein Note	Peptide	Replicat Name	Sample Dilution Factor	Average Measured Retention Time	Standard Type	Total Area	Calculat Concent	Calibration Curve	Conc Multi	Normalization Method
		VQLLGSGSILR	1	1	3.22		12	#N/A	Error: All of the e...		Ratio to surro...
		VQLLGSGSILR	5	1	3.22		#N/A	#N/A	Error: All of the e...		Ratio to surrogat...
		VQLLGSGSILR	10	1	3.22		#N/A	#N/A	Error: All of the e...		Ratio to surrogat...
		VQLLGSGSILR	50	1	3.22		12	#N/A	Error: All of the e...		Ratio to surrogat...
		VQLLGSGSILR	100	1	3.22		#N/A	#N/A	Error: All of the e...		Ratio to surrogat...
		VQLLGSGSILR	500	1	3.22		31	#N/A	Error: All of the e...		Ratio to surrogat...