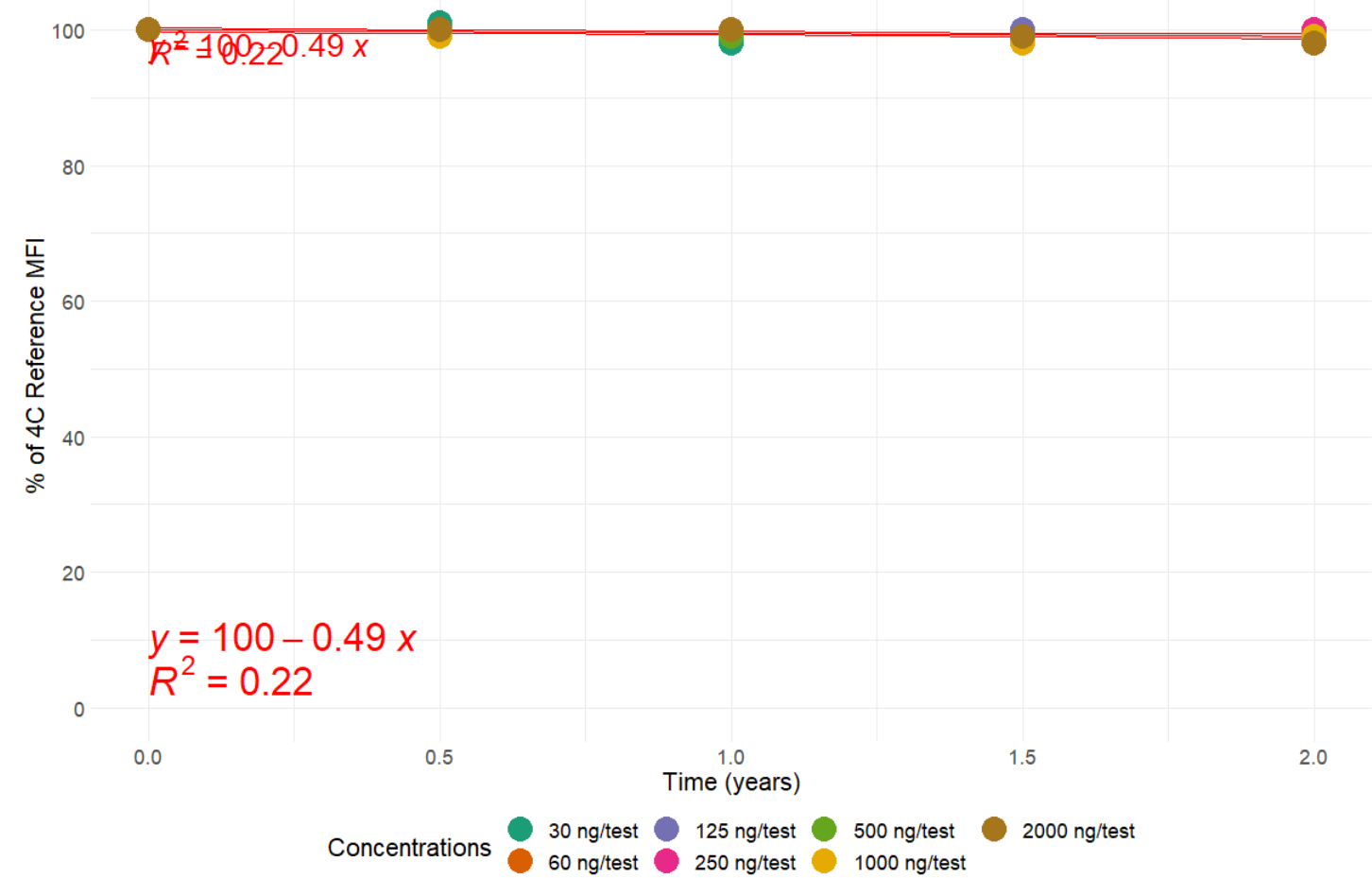


Gran

Hu CD45 X40 BV421

optimal = 0.2mg/ml

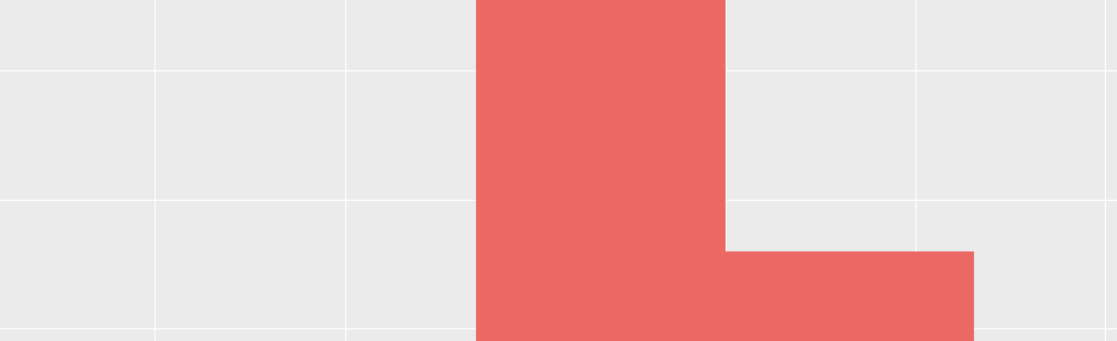


% of 0 Reference MFI							
Time	30 ng/test	60 ng/test	125 ng/test	250 ng/test	500 ng/test	1000 ng/test	2000 ng/test
0	100	100	100	100	100	100	100
0.5	101	100	100	100	100	99	100
1	98	100	100	99	99	100	100
1.5	100	99	100	99	98	98	99
2	100	100	99	100	99	99	98

Shelf-Life Estimates			
Raw Shelf-Life	Rounded Shelf-Life	R-squared	Model p-value
50.4 yrs (18406 days)	5 yrs (1825 days)	0.22	0.004

A scatter plot comparing predicted versus actual values for the percentage of 4C MFI. The x-axis is labeled "Predicted % of 4C MFI" and ranges from approximately 99.1 to 100.1, with major ticks at 99.25, 99.50, 99.75, and 100.00. The y-axis is unlabeled but has numerical labels at -1, 0, and 1. A horizontal black line is drawn at y = 0. There are 10 red circular data points plotted. Most points are clustered around the y=0 line, indicating good prediction accuracy, though there is some spread, particularly at higher predicted values.

Predicted % of 4C MFI	Actual % of 4C MFI (approx.)
99.15	-1.0
99.15	0.0
99.15	1.0
99.35	-1.3
99.35	-0.3
99.35	0.7
99.55	-1.6
99.55	-0.5
99.55	0.5
99.80	-0.8
99.80	0.2
99.80	1.2
100.05	0.0



A histogram showing the distribution of residuals. The x-axis is labeled 'Residuals' and ranges from -2 to 1. The y-axis is labeled '# of Residuals' and ranges from 0 to 15. The distribution is skewed to the right, with a peak at 0 (15 residuals) and a long tail extending to the right.

Residuals Range	# of Residuals
-2.0 to -1.5	1
-1.5 to -1.0	1
-1.0 to -0.5	4
-0.5 to 0.0	15
0.0 to 0.5	9
0.5 to 1.0	6
1.0 to 1.5	6

Anderson-Darling Normality Test p-value
0.059