

Alignment of Fc-binding domains									
E	---	-----	AQQ	NAFYQVLNMP	NLNADQRNGF	IQSLKDDPSQ	SANVLGEAQK	LNSQAPK	(SEQ ID NO: 1)
D	ADA	QONKFNKDQQ	SAFYEILNMP	NLNEEQRNGF	IQSLKDDPSQ	IQSLKDDPSQ	STNVLGEAKK	LNESQAPK	(SEQ ID NO: 2)
A	--A	DNN-FNKEQQ	NAFYEILNMP	NLNEEQRNGF	IQSLKDDPSQ	IQSLKDDPSQ	SANLLAEAKK	LNESQAPK	(SEQ ID NO: 3)
B	---	ADNKFNKEQQ	NAFYEILHLP	NLNEEQRNGF	IQSLKDDPSQ	IQSLKDDPSQ	SANLLAEAKK	LNDAAQAPK	(SEQ ID NO: 4)
C	---	ADNKFNKEQQ	NAFYEILHLP	NLTEEQRNGF	IQSLKDDPSV	IQSLKDDPSV	SKEILAEAKK	LNDAAQAPK	(SEQ ID NO: 5)
Z	---	VDNKFNKEQQ	NAFYEILHLP	NLTEEQRNAF	IQSLKDDPSQ	IQSLKDDPSQ	SANLLAEAKK	LNDAAQAPK	(SEQ ID NO: 6)
Zvar	---	VDAKFDKEQQ	NAFYEILHLP	NLTEEQRNAF	IQSLKDDPSQ	IQSLKDDPSQ	SANLLAEAKK	LNDAAQAPK	(SEQ ID NO: 7)
	---	-----	KEQQ	NAFYEILHLP	NLTEEQRNAF	IQSLKDDPSQ	SANLLAEAKK	LNDAAQAPK	(SEQ ID NO: 51)
	---	-----	KEQQ	NAFYEILHLP	NLTEEQRNGF	IQSLKDDPSV	SKEILAEAKK	LNDAAQAPK	(SEQ ID NO: 52)
Pos	1	10	20	30	40	50	58		

Fig. 1

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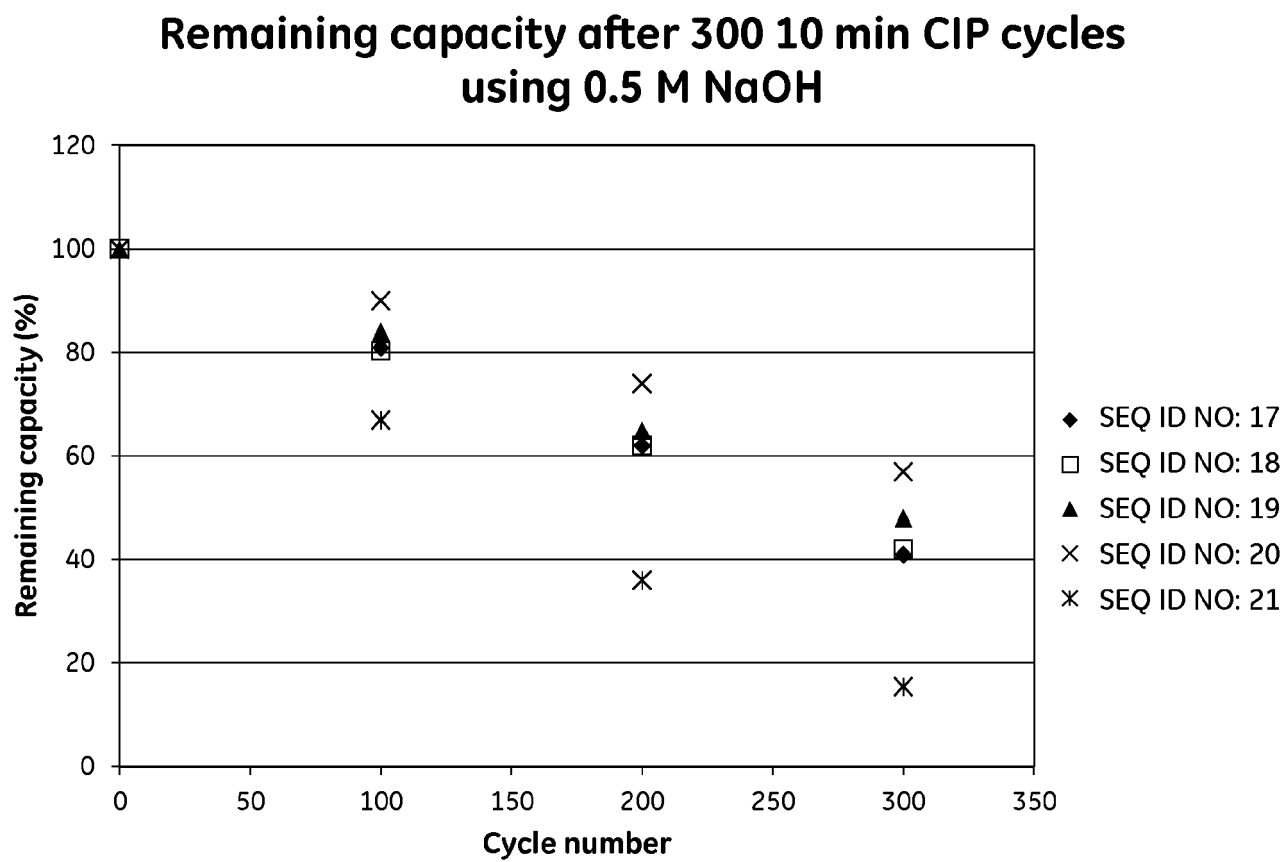


Fig. 2.

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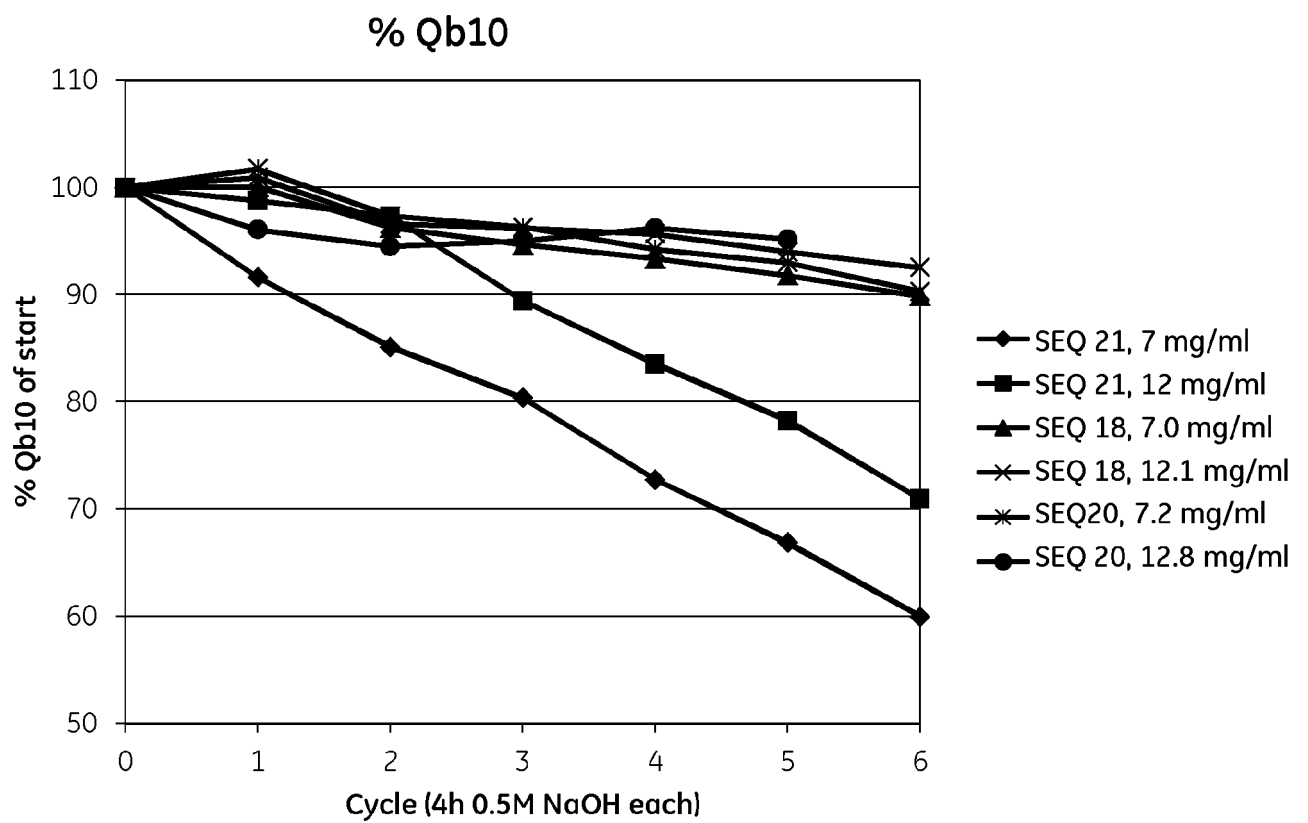


Fig. 3.

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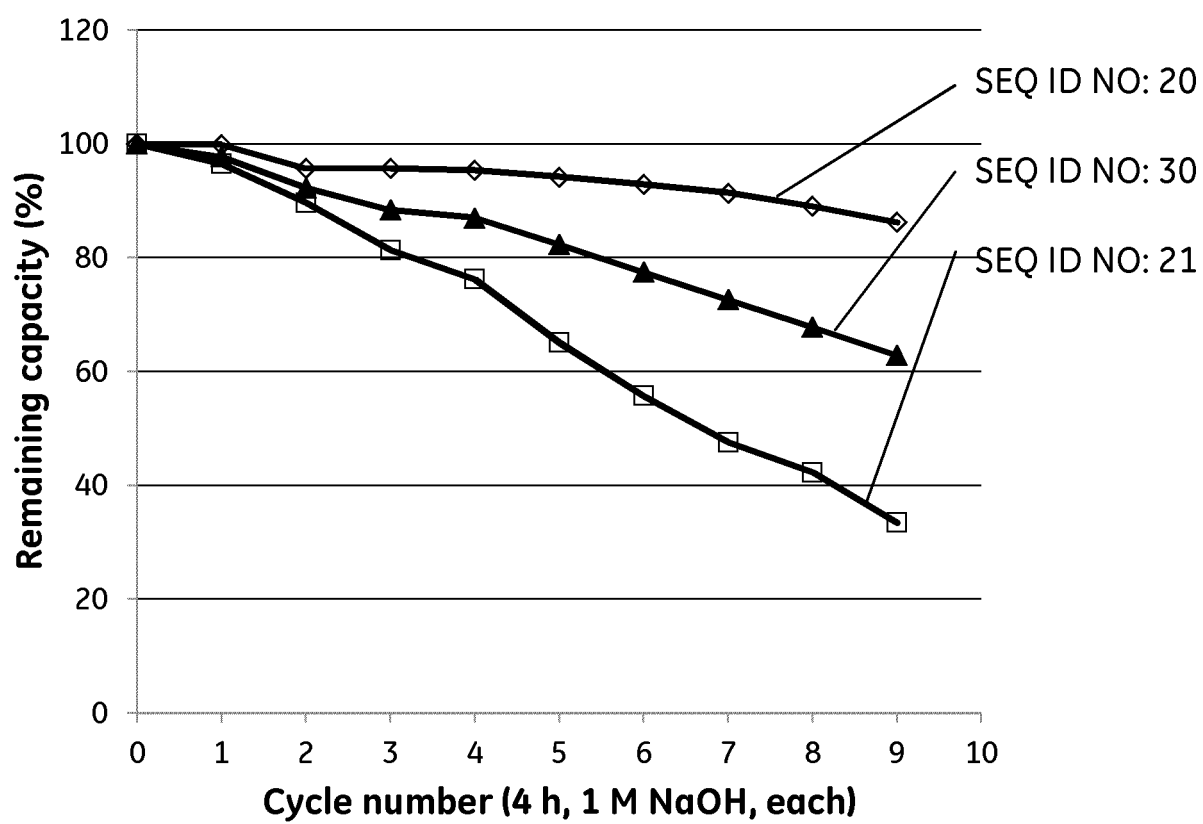


Fig. 4.