Performance Characteristics of *iLite*[®] **Cell-based Reporter Gene Assays:** Case Study of *iLite*[®] IL-23 Assay Ready Cells

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Introduction

Biological activity is an essential quality attribute for the assessment of the potency and stability of biological drugs, and the successful development of such biologics is dependent upon the establishment of validated and standardized assays that allow direct comparisons of the relative potency and stability of different batches.

We have developed a high throughput assay platform, *iLite*[®], based on the use of engineered reporter-gene cell lines that is applicable to most biologics and that allows the direct comparison of drug potency and stability in the same assay. Furthermore, since biological activity is measured, the same assay system can be used to determine the presence of neutralizing antibodies.

Here, using our *iLite[®]* IL-23 Assay Ready Cells as an example, we show the robustness of our assay system, describing parameters such as accuracy, precision, and plate homogeneity, as well as assay characteristics such as EC50 and LLOQ.

Cell Line Development

The IL-23 reporter gene assay was established using the avian B-cell line DT-40 that does not require IL-23 or other human cytokines in order to proliferate and that is unresponsive to the growth factors present in human serum. Thus, DT-40 cells were co-transfected with the IL-23R and IL-12Rb1 receptor chains together with a STAT5 expression vector, a STAT5 responsive Firefly luciferase (FL) reporter gene construct, and the Renilla luciferase (RL) normalization gene under the control of a constitutive promoter.

Methods

Precision

CV%

7%

2%

1%

5%

0%

1%

91%

105%

94%

7%

5%

5%

22

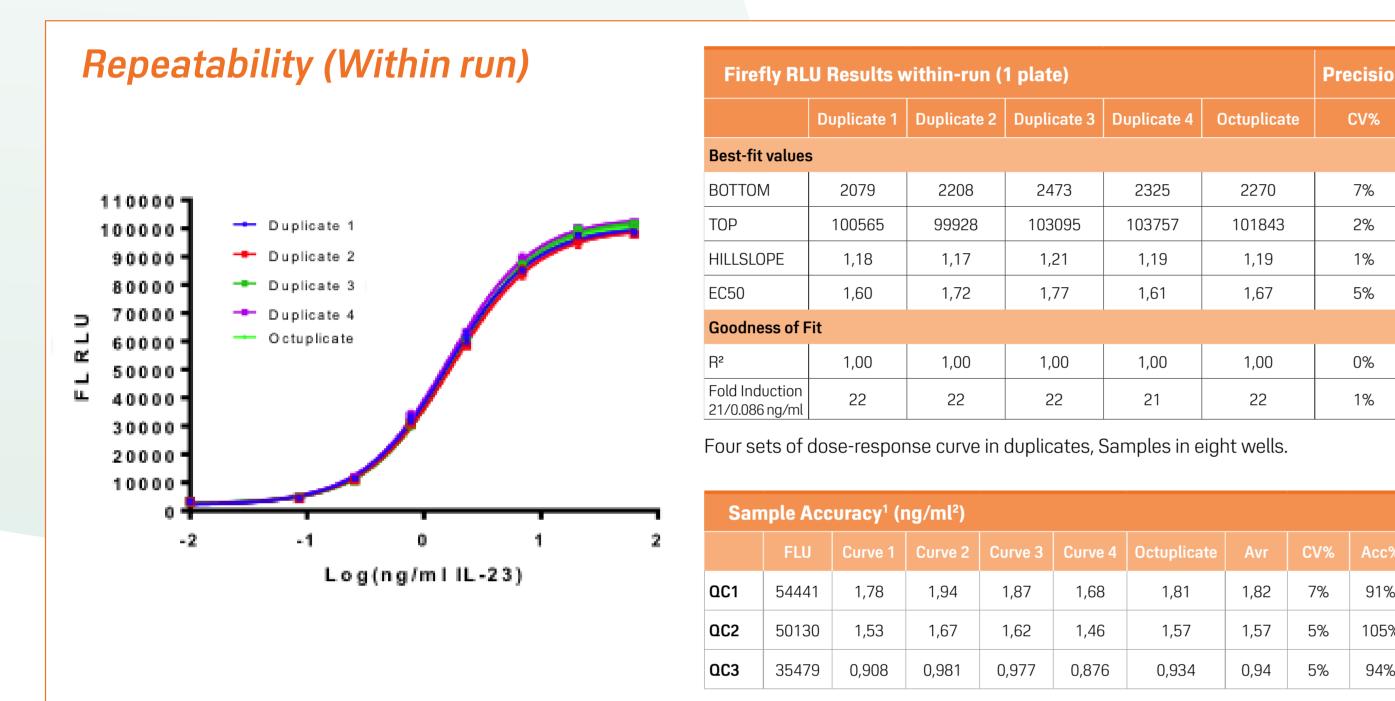
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Assays ready IL-23 cells were thawed and diluted in RPMI + 9% HI-FBS +1% P/S. IL-23 protein (R&D Systems) was diluted in same buffer and both incubated for 5h at 37°C/5% CO2. Detection of Firefly and Renilla Luciferase activity was determined with Dual-Glo[®] Luciferase Assay System (Promega) and read sequentially. Standard dose response curves were run with IL-23 protein 0-63 ng/ml, 3-fold dilution steps (63, 21, 6.9, 2.3, 0.77, 0.26, 0.086 and 0 ng/ml). Samples of IL-23 protein, Q1, Q2 and Q3 were diluted to 2.0 ng/ml, 1.5 ng/ml and 1.0 ng/ml. Assay plate homogeneity (FL RLU) was explored using a solution of 0.5 ng/ml IL-23 protein in all 96 wells of a plate.

Results

The robustness of the *iLite[®]* IL-23 Assay Ready Cells was explored by testing a number of different characteristics and parameters including repeatability, intermediate precision, accuracy and plate homogeneity. The results show that the assay has high repeatability and precision as well as good accuracy. In addition, the assay gives a high fold induction (19x) as well as a lower limit of quantification (LLOQ) in the low nanoliter range (0.1ng/mL). Taken together, the results show that the *iLite[®]* Technology and the Assay Ready format is capable of generating a high performing bioassay, suitable for many applications within the drug continuum.



¹Firefly RLU Results between-runs (4 plates) ²IL-23 protein diluted to 2.0 ng/ml, 1.5 ng/ml and 1.0 ng/ml

Assay characteris	tics
Dose response curve	0-63 ng/ml, 3-fold dilution
EC50†	1,40 (1,10-1,72)
Fold induction†	19 (15-22)
LLOQ*	0.1 ng/ml
ULOQ*	5.1 ng/ml

* LLOQ, ULOQ – defined as lowest and highest concentration of analyte which can be quantified in this assay set-up with 75-125% accuracy and CV<25% precision.

† EC50 and fold-induction - average from 20 dose response curves, tests at separate days and with different cell batches (90% CI)

Repeata	ability (Between runs)	Fir	efly RLl	J Result	s betwe	en-runs	(
				Plat	te 1	Plate 2	
		Best-	fit values				
110000 –		BOTT	ОМ	25	73	2822	
100000	Set 1	TOP		706	61	68969	
90000 -	- Set 2	HILLS	LOPE	1,3	31	1,37	
80000-	- Set 3	EC50		1,2	27	1,20	
70000	Set 4 All together	Good	ness of Fi	t			
60000 50000		R ²		1,0	00	1,00	
40000			nduction)86 ng/ml	1	7	16	
30000 - 20000 - 10000 -			set of d plates. A			urve in d	h
o 芉			ample A	ccuracy	¹ (ng/ml	²)	
-2	-1 0 1 Log(ng/mIIL-23)	2	Plate 1	Plate 2	Plate 3	Plate 4	
	209(iig/iii i i 20)	QC1	45938	45010	45138	44365	
		002	43563	42783	43415	43945	

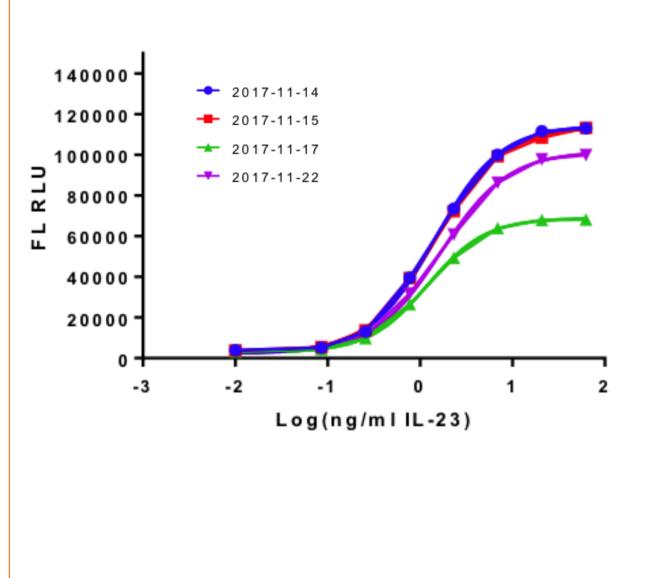
Firefly RLU Results between-runs (4 plates)											sion	
		Plat	e 1	Plate 2	2	Plate 3		Plate 4		CV	%	
Best-	fit values											
BOTT	ОМ	257	73	2822		2626		2748		49	6	
TOP		706	61	68969		70921		68494		29	6	
HILLS	HILLSLOPE 1,31 1,37 1,30 1,36 3%										6	
EC50 1,27 1,20 1,28 1,21 3%												
Goodness of Fit												
R²	R ² 1,00 1,00 1,00 1,00 0%									6		
	nduction)86 ng/ml	17	7	16		16		16		29	6	
One set of dose response curve in duplicates, and each sample in quadruplicate. Four plates. All same day.												
Sa	mple A	ccuracy	(ng/ml	²)								
	Plate 1	Plate 2	Plate 3	Plate 4	Plate 1	Plate 2	Plate 3	Plate 4	Avr	CV%	Acc%	
001	15038	45010	45138	11365	1 95	1 82	1 88	1 81	1 86	3%	03%	

QC1	45938	45010	45138	44365	1,95	1,82	1,88	1,81	1,86	3%	93%
QC2	43563	42783	43415	43945	1,74	1,64	1,73	1,77	1,72	3%	115%
QC3	33498	32278	31475	31745	1,10	1,02	1,01	1,02	1,04	4%	104%

¹Firefly RLU Results between-runs (4 plates) ²IL-23 protein diluted to 2.0 ng/ml, 1.5 ng/ml and 1.0 ng/ml

Repeatability (Retween rune)

Intermediate precision (Between Days)



	Day 1	Day 2	Day 3	Day 4	CV%
Best-fit values					
BOTTOM	2503	2418	2822	2270	7%
ТОР	114790	114007	68969	101843	22%
HILLSLOPE	1,23	1,19	1,37	1,19	6%
EC50	1,45	1,46	1,20	1,67	9%
Goodness of Fit					
R ²	1,00	1,00	1,00	1,00	0%
Fold Induction 21/0.086 ng/ml	21	20	16	22	5%

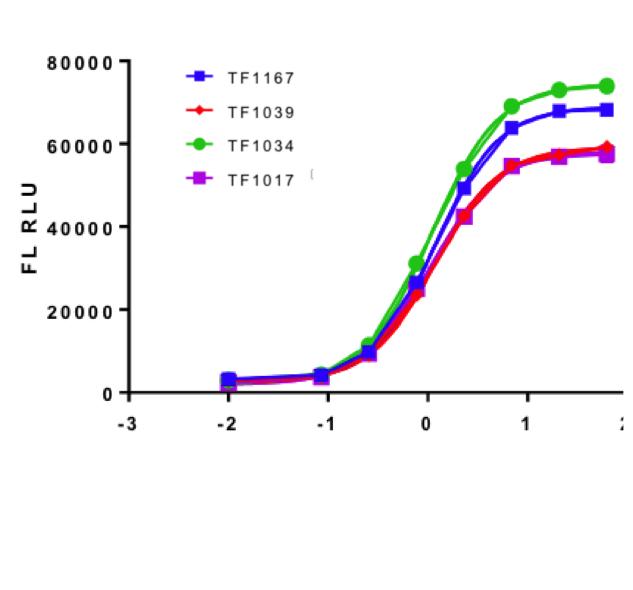
90 45010	54441	1,90	1,78	1,82	1,81	1,86	3%	93%
42783	50130	1,43	1,47	1,64	1,57	1,54	6%	102%
350 32278	35479	0,884	0,922	1,02	0,934	0,97	6%	97%
	s between	350 32278 35479 as between-runs (4 p	350 32278 35479 0,884 as between-runs (4 plates)	350 32278 35479 0,884 0,922 is between-runs (4 plates)	350 32278 35479 0,884 0,922 1,02	350 32278 35479 0,884 0,922 1,02 0,934 is between-runs (4 plates)	350 32278 35479 0,884 0,922 1,02 0,934 0,97 is between-runs (4 plates)	350 32278 35479 0,884 0,922 1,02 0,934 0,97 6% is between-runs (4 plates)

Plate Homogeneity

Assay plate homogeneity (FL RLU): 0,5 ng/ml in all 96 wells of a plate.

FL signal homogenity 32500

Between Assay Ready Cell batches



Fire		Preci	sion												
	TF1017 TF1034 TF 1039 TF 1167										%				
Best-fi	t values														
вотто	DTTOM 2822 2488 2267 1999 15%										%				
ТОР	OP 68969 59181 74523 58170 12%														
HILLSLOPE 1,37 1,33 1,30 1,31 2%															
EC50 1,20 1,17 1,09 1,06 6%															
Goodness of Fit															
R²	1,000 0,999 0,999 0,999					0%									
Fold Induction 16 15 17 15 5% 21/0.086 ng/ml 16 15 17 15 5%															
our cel	ne set of dose response curve in duplicates, and each sample in quadruplicate our cell batches. Sample Accuracy ¹ (ng/ml ²)														
	TF1017	TF1034	TF1039	TF1167	TF1017	TF1034	TF1039	TF1167	Avr	CV%	Acc%				
QC1	45010	39763	50893	37688	1,82	1,90	1,90	1,61	1,81	8%	90%				
QC2	42783	39100	48408	39783	1,64	1,83	1,69	1,83	1,75	6%	117%				

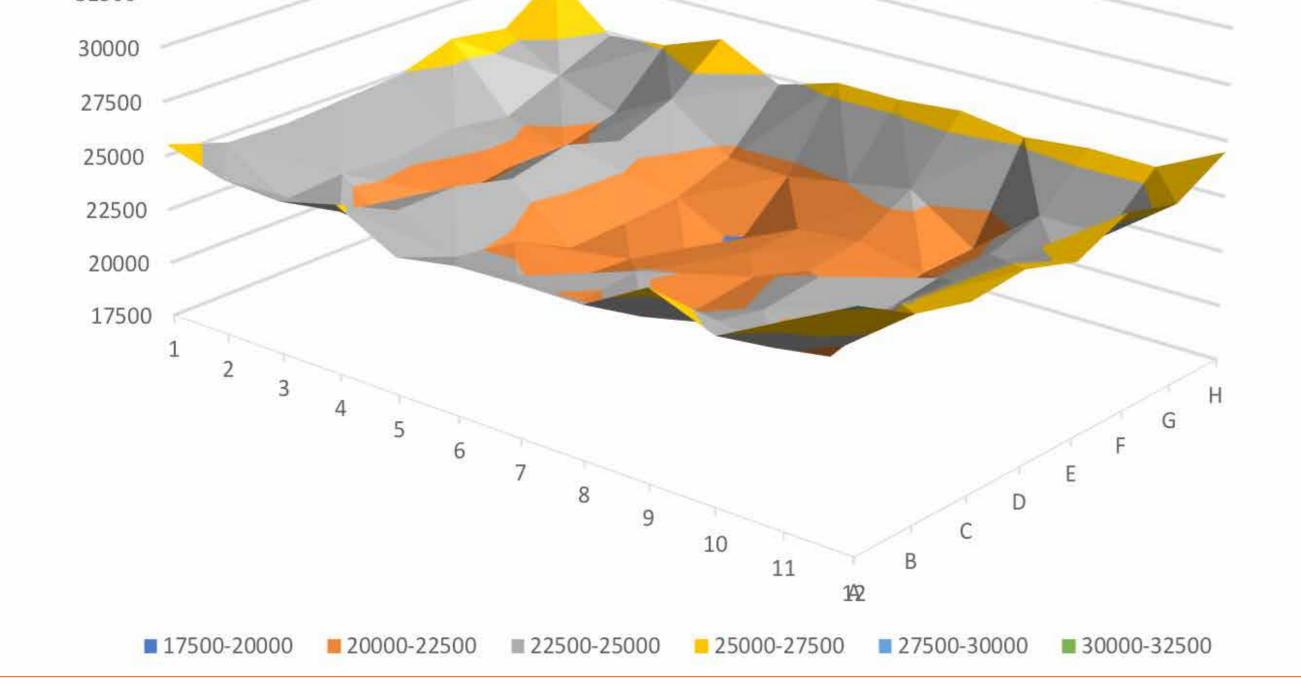
¹Firefly RLU Results between-runs (4 plates) ²IL-23 protein diluted to 2.0 ng/ml, 1.5 ng/ml and 1.0 ng/ml

Summary and Conclusions

By using the *iLite*[®] IL-23 Assay Ready Cells as an example, we have shown the robustness of our assay system, describing parameters such as accuracy, precision, and plate homogeneity, as well as assay characteristics such as EC50 and LLOQ. The results demonstrate that using this system, assay parameters normally associated with ligand binding assays can be achieved also with a cell-based setup.

Accuracy





	Expected value	Measured value	Accuracy
Sample 1	2,0	1,83 (1,68-1,95)	92% (84-98%)
Sample 2	1,5	1,61 (1,46-1,83)	107% (97-122%)
Sample 3	1,0	0,98 (0,88-1,07)	98% (88-107%)

FL RLU homogeneity (CV%)							
Row	<5%						
Column	<9%						
Plate	8%						

Average from 20 duplicates, tests at separate days and with different cell batches (90% CI)

Precision (CV%)

		EC50	Fold induction	Sample 1	Sample 2	Sample 3
Denestability	Within run (Octuplicates)	5%	1%	7%	5%	5%
Repeatability	Between Runs (4 plates)	3%	2%	3%	3%	4%
Intermediate	Between Days (4 days)	9%	12%	3%	6%	6%
precision	Between IARC batches (4 batches)	6%	5%	8%	6%	11%

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