

Surrogate CSF for the Bioanalysis of Oligonucleotide Therapeutics Through LBA

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INTRODUCTION

- Bioanalytical ligand binding assays (LBA) support pharmacokinetic (PK) drug and pharmacodynamic (PD) biomarker profiling for preclinical and clinical studies.
- Per the ICH M10 guidance, LBA for PK bioanalyses requires calibration standards prepared in the same biological matrix as the study samples.
- Additionally, per the ICH M10 guidance, the biological matrices are the recommended diluents for study samples.
- Therefore, each biological matrix lot must be qualified for specificity and selectivity against the analyte, prior to use.
- These requirements pose unique challenges for rare matrices such as cerebrospinal fluid:
 - Long lead times for procurement
 - Expensive
 - Limited volumes per collection lot
 - Performance variance between lots
 - Requiring robust criteria for qualification, making it further challenging.

CLINICAL APPLICATION

- Original bioanalytical method for an ASO validated using native human CSF and successfully used to support the Phase I clinical study.
- The above-mentioned challenges were observed during a longer-duration Phase II study, primarily through variance in lot-to-lot performance of native CSF.
- Therefore, the feasibility of a surrogate CSF method was explored through preliminary testing as depicted in **Figure 1**.
- Performance of 1% serum-supplemented PBS demonstrated in precision and accuracy evaluations with native human CSF (**Table 2**).

Table 2. Calibration standards in surrogate CSF. QC sample sets in surrogate and native CSF.

Precision and Accuracy	Prepared in surrogate CSF (1% serum supplemented PBS)					Prepared in native human CSF				
	LLOQ QC	LQC	MQC	HQC	ULOQ QC	LLOQ QC_CSf	LQC_CSf	MQC_CSf	HQC_CSf	ULOQ QC_CSf
	0.1500 ng/mL	0.4000 ng/mL	4.000 ng/mL	75.00 ng/mL	100.0 ng/mL	0.1500 ng/mL	0.4000 ng/mL	4.000 ng/mL	75.00 ng/mL	100.0 ng/mL
Replicate – 1	0.1782	0.3815	4.060	77.74	87.64	0.1542	0.4170	4.033	75.43	89.83
Replicate – 2	0.1723	0.3997	3.847	72.34	93.74	0.1628	0.4187	4.030	78.26	84.68
Replicate – 3	0.1733	0.3655	3.833	74.87	87.78	0.1560	0.4032	3.627	71.98	81.95
Intra-run Mean	0.1746	0.3822	3.913	74.98	89.72	0.1577	0.4130	3.897	75.22	85.49
Intra-run SD	0.003158	0.01711	0.1272	2.702	3.482	0.004536	0.008501	0.2335	3.145	4.001
Intra-run %CV	1.8	4.5	3.3	3.6	3.9	2.9	2.1	6.0	4.2	4.7
Intra-run %Bias	16.4	-4.4	-2.2	0.0	-10.3	5.1	3.2	-2.6	0.3	-14.5
Intra-run %Total Error	18.2	8.9	5.4	3.6	14.2	8.0	5.3	8.6	4.5	19.2
n	3	3	3	3	3	3	3	3	3	3

PRECLINICAL APPLICATION

- 1% plasma—PBS tested as surrogate CSF, for *de novo* development.
- Performance demonstrated in precision and accuracy evaluations with native monkey CSF (**Table 3**).
- Dilutional linearity established with surrogate CSF as diluent for reliable quantitation of preclinical study samples at C_{max} (**Table 4**).

Table 3. Calibration standards in surrogate CSF. QC samples sets in surrogate and native CSF.

Precision and Accuracy	Prepared in surrogate CSF (1% plasma supplemented PBS)					Prepared in native monkey CSF				
	LLOQ QC	LQC	MQC	HQC	ULOQ QC	LLOQ QC_CSf	LQC_CSf	MQC_CSf	HQC_CSf	ULOQ QC_CSf
	40 pg/mL	100 pg/mL	2000 pg/mL	150000 pg/mL	200000 pg/mL	40 pg/mL	100 pg/mL	2000 pg/mL	150000 pg/mL	200000 pg/mL
Replicate – 1	41.1	112	1900	164000	200000	34.4	102	2070	163000	200000
Replicate – 2	36.6	109	2090	175000	193000	34.7	99.1	1990	162000	192000
Replicate – 3	37.3	109	1960	165000	205000	38.7	106	2220	156000	203000
Intra-run Mean	38.3	110	1980	168000	199000	35.9	102	2090	160000	198000
Intra-run SD	2.42	1.73	97.1	6080	6030	2.40	3.46	117	3790	5690
Intra-run %CV	6.3	1.6	4.9	3.6	3.0	6.7	3.4	5.6	2.4	2.9
Intra-run %Bias	-4.2	10.0	-1.0	12.0	-0.5	-10.2	2.4	4.5	6.7	-1.0
Intra-run %Total Error	10.5	11.6	5.9	15.6	3.5	16.8	5.8	10.1	9.0	3.9
n	3	3	3	3	3	3	3	3	3	3

Table 4. Dilutional linearity of QC sample (C_{max}) in native CSF using surrogate CSF as diluent.

Dilution Linearity	Nominal concentration – 5,000,000,000 pg/mL						
	1X	10X	100X	1,000X	10,000X	100,000X	1,000,000X
	Measured concentration (pg/mL)						
Replicate – 1	>ULOQ	>ULOQ	>ULOQ	>ULOQ	>ULOQ	5150000000	4320000000
Replicate – 2	>ULOQ	>ULOQ	>ULOQ	>ULOQ	>ULOQ	5260000000	4450000000
Replicate – 3	>ULOQ	>ULOQ	>ULOQ	>ULOQ	>ULOQ	4640000000	4810000000
Mean	NA	NA	NA	NA	NA	5020000000	4530000000
S.D.	NA	NA	NA	NA	NA	331000000	254000000
n	1	3	3	3	3	3	3
% CV	NA	NA	NA	NA	NA	6.6	5.6
% Bias	NA	NA	NA	NA	NA	0.4	-9.4

SURROGATE CSF

- Compositional analyses indicate a 1000-fold lower protein content in CSF vs. blood-based matrices.
- Table 1** lists the identified composition from a review article.
- ¹Fogh JR et al. Anal Bioanal Chem 2020;412:1653-61*

To identify a suitable surrogate CSF:

- Dual-probe method for an ASO
- QC samples in native CSF
- Calibration standards in
 - 1% serum in PBS
 - 1% BSA in PBS
- Serum supplementation better mimics performance in native CSF

	CSF	Plasma
Proteins (µg/mL)	300–700	60,000–70,000
-Albumin	192	42,150
-Globulins	50	22,700
-Transferrin	14	2600
-Fibrinogen	0.6	2800
Lipids (µg/mL)	10–20	50,000
Na ⁺ (mmol/L)	140–145	135–147
Cl ⁻ (mmol/L)	115–120	95–105
Glucose (µg/mL)	600	900
pH	7.31	7.41

Table 1. Typical composition of CSF and plasma ¹

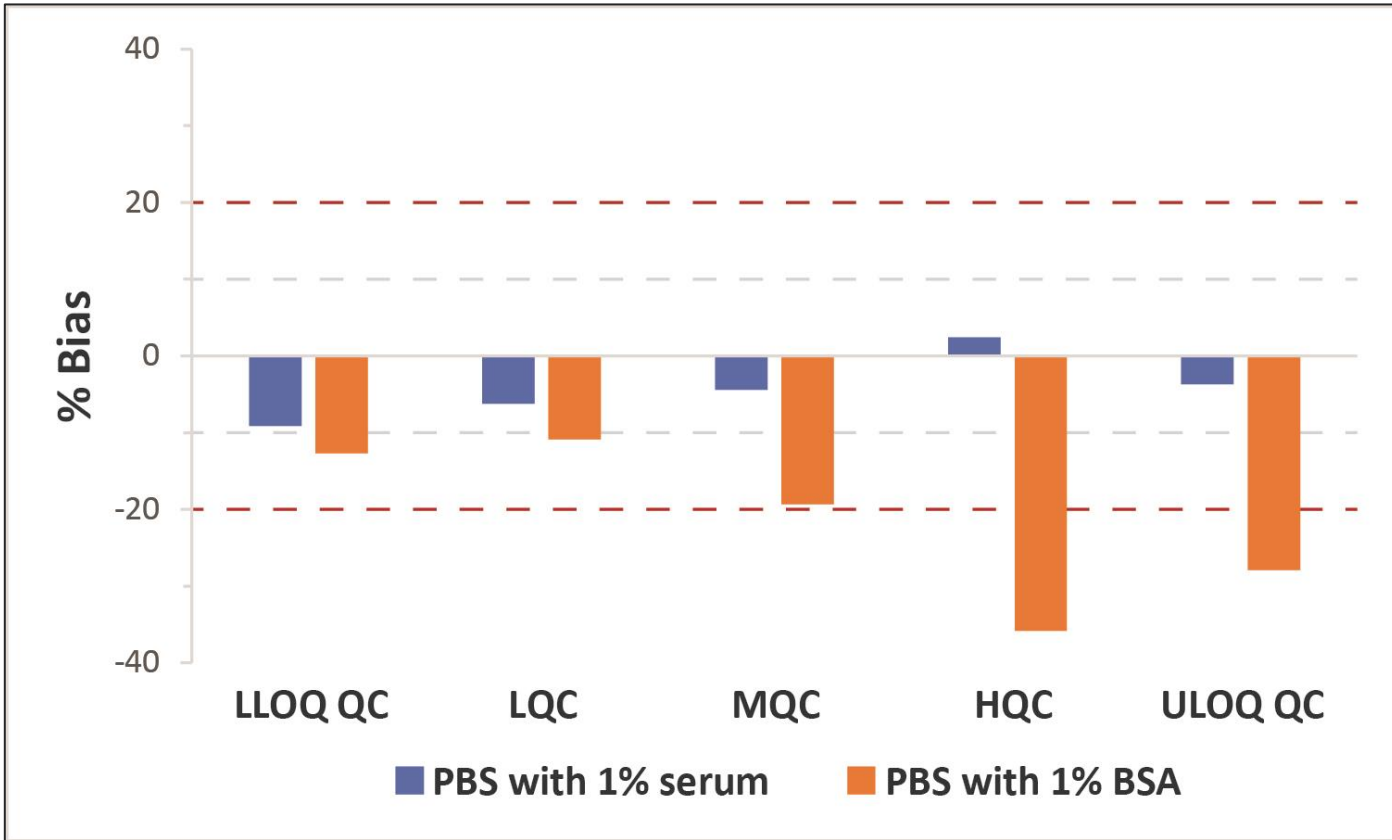


Figure 1. Serum vs. BSA-supplemented PBS as potential surrogate CSF candidates

Cross-validation of native CSF and surrogate CSF methods:

- Specific workflow used for method cross-validation:

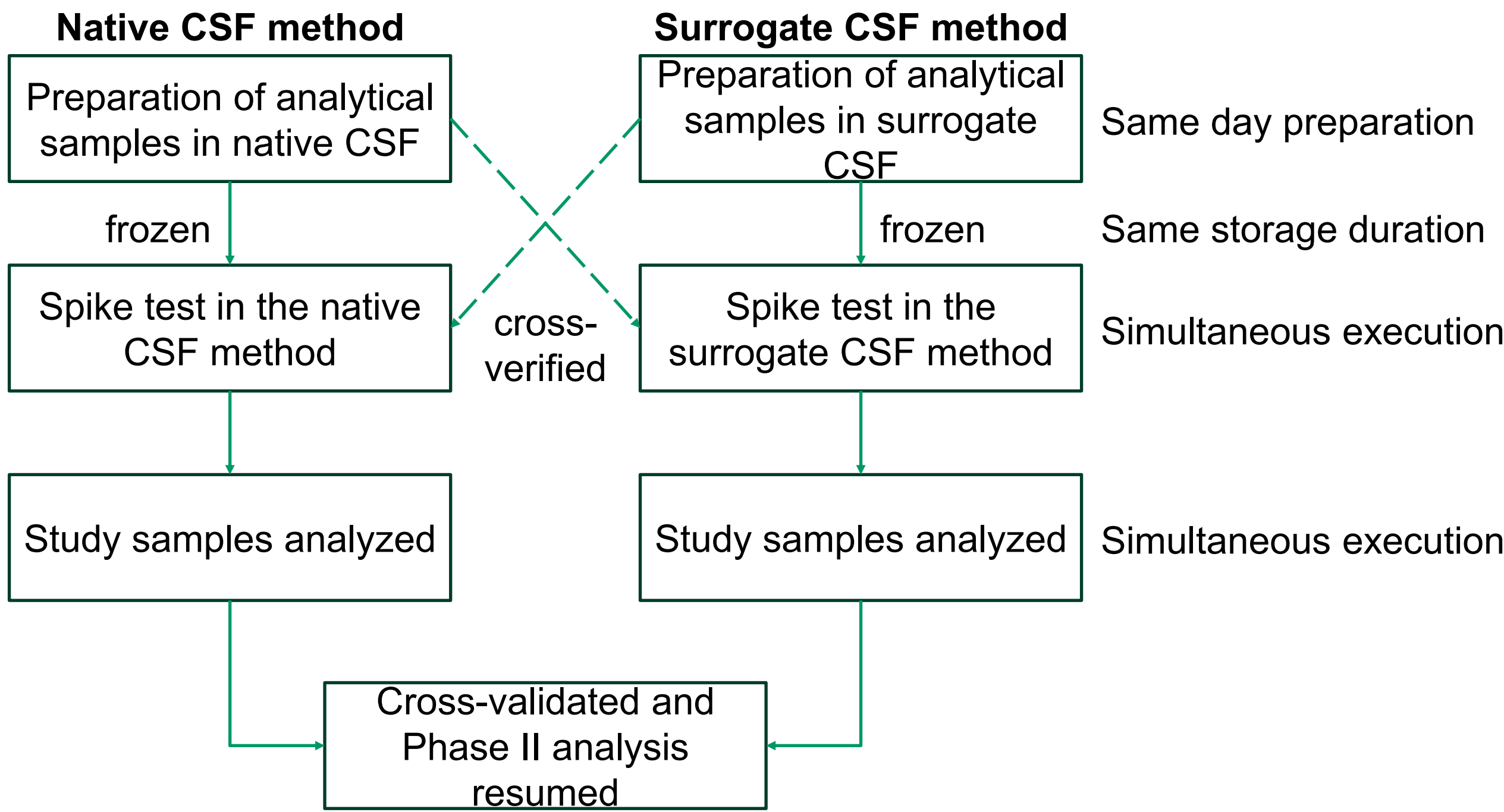


Figure 2. Cross-validation work-flow.

- Cross-validation first performed using QC samples.
- Cross-validation next performed using n≥30 study samples.
- All QC and study samples met acceptance criteria (100%).

CONCLUSIONS

- Serum or plasma (at 1% v/v) supplemented PBS mimics native CSF performance in bioanalytical methods for oligonucleotide therapeutics.
- This similarity in performance extends outside the analytical range, as demonstrated with dilution linearity up to 1,000,000-fold.
- The presented clinical study demonstrates the ease of transferring validated native CSF methods to surrogate CSF methods without extensive changes in the methodology.
- Surrogate CSF preparations perform consistently across different lots, enhancing method robustness and reliability.