

Method Development for the Detection of Exicure's AST-008 in Human Plasma

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Introduction

AST-008 is a novel spherical nucleic acid (SNA) configuration of a toll-like receptor 9 (TLR9) agonist oligonucleotide, designed to trigger innate and adaptive immune responses that are useful in oncology applications. AST-008 activated key immune cells and cytokines predictive for an anti-tumor effect in a Phase 1 healthy-volunteer study.

A sensitive bioanalytical method was required to determine the concentrations of AST-008 in human plasma with minimal detection of metabolites to support pharmacokinetic analysis in clinical studies.

Objective

The concentrations of AST-008 in clinical samples were initially determined by hybridizing a complementary, fluorescently-labeled peptide nucleic acid (PNA) probe and using liquid chromatography with fluorescence detection (LC-FD). All samples, when analyzed with this method, were below the limit of quantification (BLQ) of 10 ng/mL; therefore, a more sensitive method was required. The objective of this study was to develop a bioanalytical method with greater sensitivity, as determined by the lower limit of quantitation (LLOQ). Two hybridization methods (Dual Hybridization and Hybridization-Ligation) and two platforms (Fluorescence and ECL) were compared in order to select the one with greatest sensitivity and selectivity for further validation.

Figure 1: SNA Structure

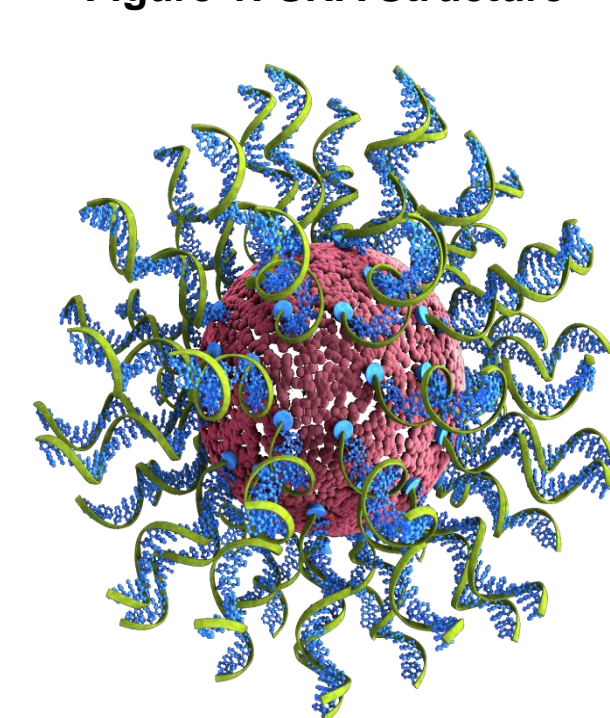
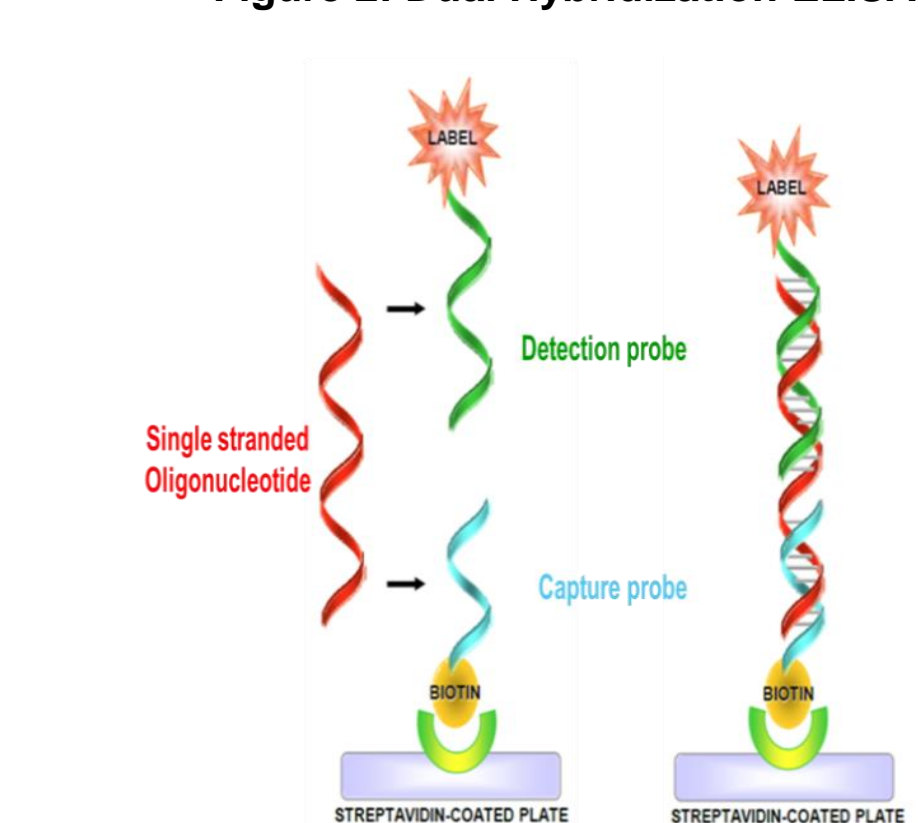


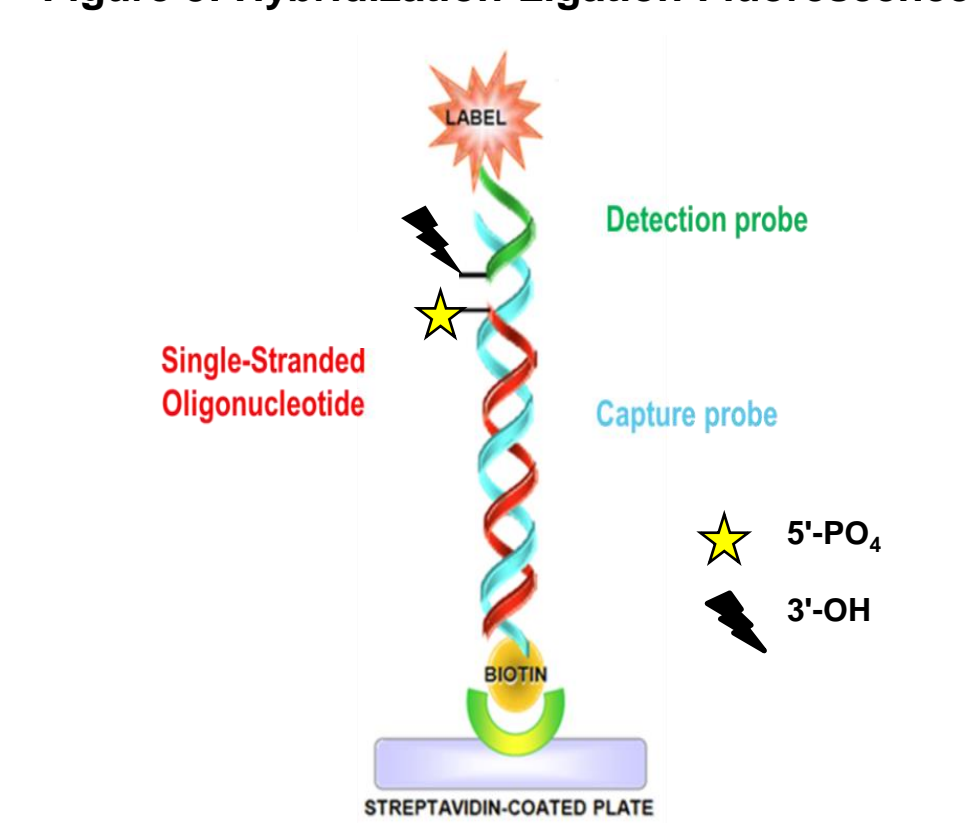
Figure 2: Dual Hybridization ELISA



- Hybridization of oligonucleotide (ON) with capture and detection probes
- Non-Hybridized probe will be washed away
- Advantage: Enhanced sensitivity and wider dynamic range
- Disadvantage: Cross-Reactivity with 3' and 5' metabolites

Efler, SM et al.(2005) Oligonucleotides..15 (2) 119-131

Figure 3: Hybridization-Ligation Fluorescence



- Ligation of ON with T4 polynucleotide kinase (PNK) and DNA ligase/ATP
- Non-Ligated probe will be washed away
- Advantage: Minimal 5'-metabolite cross-reactivity
- Disadvantage: Reduced sensitivity

Yu RZ et al.(2002) Anal. Biochem. 304, 19-25

Results

Table 2: Sensitivity of LC-FD versus Dual Hybridization Methods

	Dual Hybridization	LC-FD
LLOQ (Plasma)	1.000 ng/mL	10.00 ng/mL

- The Dual Hybridization method was evaluated first using the fluorescence platform and although the LLOQ was lower than the LC-FD method, higher sensitivity was pursued utilizing additional hybridization methods and platforms.

Table 3: Metabolite Cross-Reactivity with Hybridization-Ligation and Dual Hybridization

Nominal Metabolite Concentration (ng/mL)	Hybridization-Ligation ECL Method				Dual Hybridization ECL Method					
	105.0									
	Run ID	Metabolite Identity		Metabolite Identity		Run ID	Metabolite Identity			
	N-1 AST-008	N-2 AST-008	N-1 AST-008	N-2 AST-008	N-1 AST-008	N-2 AST-008	N-1 AST-008	N-2 AST-008		
Observed Concentration	EXI557.33	7.401	6.573	118.4	104.3	EXI557.10	7.470	6.493	123.8	95.86
N		2	2	2	2					
Mean		7.436	6.533	121.1	100.1					
Cross-Reactivity (%)		7.1	6.2	115.3	95.3					

$$\%Cross - Reactivity = (Mean\ observed\ concentration) \div (Nominal\ metabolite\ concentration) \times 100$$

- The hybridization-ligation ECL method had a minimal cross-reactivity with N-1 or N-2 metabolites, whereas the dual hybridization ECL method detected 100% of both metabolites.

Table 4: Hybridization-Ligation ECL Method Precision and Accuracy

Nominal concentration (ng/mL)	LQCA	LQCB	QC1A	QC1B	QC2	QC3	ULQ
0.5110	1.009	1.587	3.186	29.08	111.8	158.4	
0.5066	1.020	1.527	3.174	33.23	113.0	164.3	
0.5293	1.018	1.450	3.039	30.97	114.5	172.4	
0.5343	1.048	1.477	2.926	30.60	115.8	175.4	
0.5519	1.016	1.471	3.075	34.46	125.4	181.0	
N	5	5	5	5	5	5	5
Mean	0.5266	1.022	1.502	3.080	31.67	116.1	170.3
SD	0.0183	0.0151	0.0551	0.1065	2.153	5.399	8.979
%CV	3.5	1.5	3.7	3.5	6.8	4.7	5.3
%RE	-5.3	2.2	0.2	2.7	5.6	10.6	13.5

- The hybridization-ligation ECL method exhibited acceptable precision and accuracy (%CV ≤ 20% and %RE ± 20%) indicating a sensitivity of 0.5000 ng/mL.

Table 5: Hybridization-Ligation ECL Method Matrix Effect Selectivity

Nominal Concentration (ng/mL)	Blank	LQCA	LQCB	QC3				
0.0000	0.5000	1.000	105.0					
Run ID	Lot #	AST-008	AST-008	% RE	AST-008	% RE	AST-008	% RE
EXI557.33	1	BLQ	0.4553	-8.9	0.8991	-10.1	101.2	-3.6
EXI557.41	2	BLQ	0.5232	4.6	1.0490	4.9	114.6	9.2
	3	BLQ	0.3786	-24.3	0.8248	-17.5	93.77	-10.7
	4	BLQ	0.3275	-34.5	0.6893	-31.1	82.11	-21.8
	5	BLQ	0.4091	-18.2	0.8099	-19.0	90.14	-14.2
	6	BLQ	0.4202	-16.0	0.8099	-19.0	96.62	-8.0
	7	BLQ	0.5327	6.5	0.9983	-0.2	116.9	11.3
	8	BLQ	0.4987	-0.3	0.9983	-0.2	115.9	10.4
	9	BLQ	0.4710	-5.8	0.8746	-12.5	102.7	-2.2
	10	BLQ	0.4631	-7.4	0.9237	-7.6	105.1	0.1
Mean	BLQ	0.4479	-10.4	0.8877	-11.2	101.9	-2.9	

- The hybridization-ligation ECL method exhibited acceptable selectivity of 10 lots with %RE ± 20% at the 0.5000 ng/mL and 105.0 ng/mL levels.

Table 6: Hybridization-Ligation ECL Method Dilution Linearity

Nominal concentration (ng/mL)	20000			
Dilution factor	1:200	1:500	1:5000	
Concentration after dilution (ng/mL)	100.0	40.00	4.000	
Run ID	Observed concentration (ng/mL)	22163	24146	21408
EXI557.38		20399	18413	18245
		20175	18966	18305
		20616	19912	18494
N		4	4	4
Mean		20838	20359	19113
SD		901.4	2599	1533
%CV		4.3	12.8	8.0
%RE		4.2	1.8	-4.4

- Sample dilution was found to be acceptable (%QV ≤ 20% and %Nominal ± 20%) up to 5000 fold using the hybridization-ligation ECL method.

Table 7: Validation Summary for the Determination of AST-008 in Human Plasma Using the Hybridization-Ligation ECL Method

Short description of method	Hybridization-Ligation ECL
Biological matrix	Human Plasma (K ₂ EDTA)
Analyte	AST-008
Calibration concentrations	0.5000 to 150.0 ng/mL
Sensitivity	0.5000 ng/mL (LOQ QC)
Lower limit of quantification	LLOQ (ng/mL) 0.5000
	Between-run accuracy 99%
	Between-run precision 12%
	Within-run accuracy 95%
	Within-run precision 2%
Between-run accuracy	91% to 99%
Between-run precision	4% to 12%
Within-run accuracy	90% to 95%
Within-run precision	1% to 5%
Matrix effect	No significant interference was observed in 9 out of 10 individual human plasma lots: acceptance criteria were met at all tested levels (Blank [un-spiked], LOQ QC and QC3). Acceptance criteria were met at all tested levels (Blank [un-spiked], LOQ QC and QC3) in hemolyzed (up to 2%) human plasma and lipemic (>300 mg/dL triglycerides) human plasma. Minimal cross-reactivity observed for N-1_AST-008 (4%) and N-2_AST-008 (1%) at QC3 level (105.0 ng/mL).
Metabolite cross-reactivity	
Metabolite interference	No effect on the determination of AST-008 in human plasma tested at QC1 and QC3 levels in the presence of either metabolite N-1_AST-008 or N-2_AST-008.
Hook effect	No hook effect observed up to 20000 ng/mL.
Dilution linearity	DQC1 at 20000 ng/mL was used for dilution factors of 1/200 and 1/5000, whereas DQC2 at 200.0 ng/mL was used for dilution factor of 1/5.
	%Nominal %CV
	Diluted 5-fold 86% 3%
	Diluted 200-fold 87% 4%
	Diluted 5000-fold 95% 3%
Whole blood stability	Reported up to 2 hours on ice/water bath.
Stock stability at -20 °C Nominal	Reported up to 19 days at -20 °C nominal. Extended period will be conducted on a later date.
Short-term matrix stability at 22 °C Nominal	Reported up to 26.2 hours at ambient room temperature.
Freeze-thaw matrix stability	Reported up to 5 cycles.
Long-term matrix stability at -20 °C Nominal	Reported up to 11 days. Extended period will be conducted on a later date.

- The hybridization-ligation ECL method is fully validated with respect to accuracy, precision, selectivity (matrix effects, metabolite cross-reactivity and interference), hook effect, dilution linearity and stability.

Conclusions

- The hybridization-ligation ECL method to measure AST-008 concentrations in human plasma was successfully validated with the range of 0.5000 ng/mL to 150.0 ng/mL.
- This method is 20-fold more sensitive than the PNA probe, LC-FD method that was formerly validated for the same analyte.
- The level of sensitivity obtained using the hybridization-ligation ECL method enabled the detection of low concentrations of AST-008 in clinical study samples that had previously been undetectable using the PNA probe LC-FD method.