

PharmOptima LLC

Value-Driven Research Services

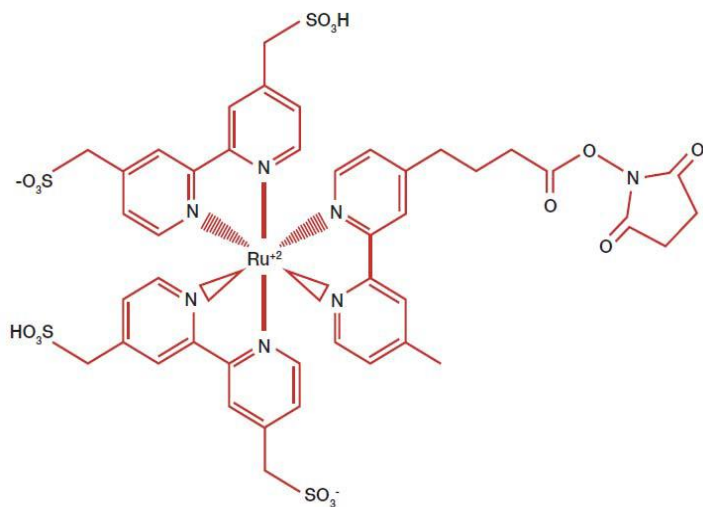


SMA
FOUNDATION

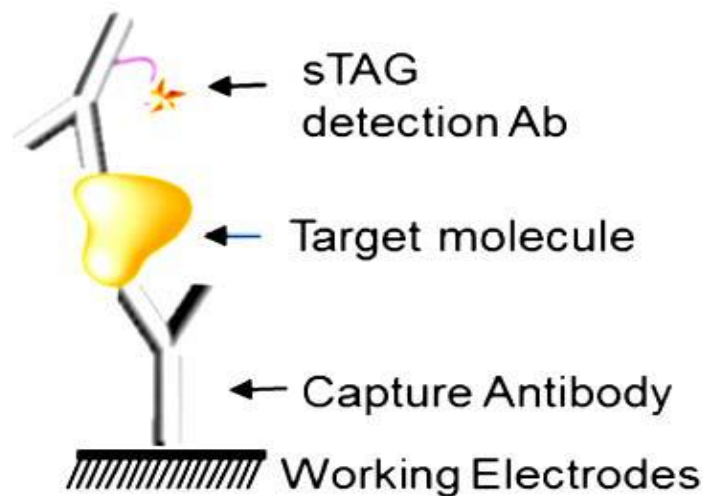
Measurement
of Survival
Motor Neuron
in Whole Blood

Meso Scale Discovery Platform Electrochemiluminescence (ECL)

Signal: Caged Ruthenium tag
emits light in response to
an electric current

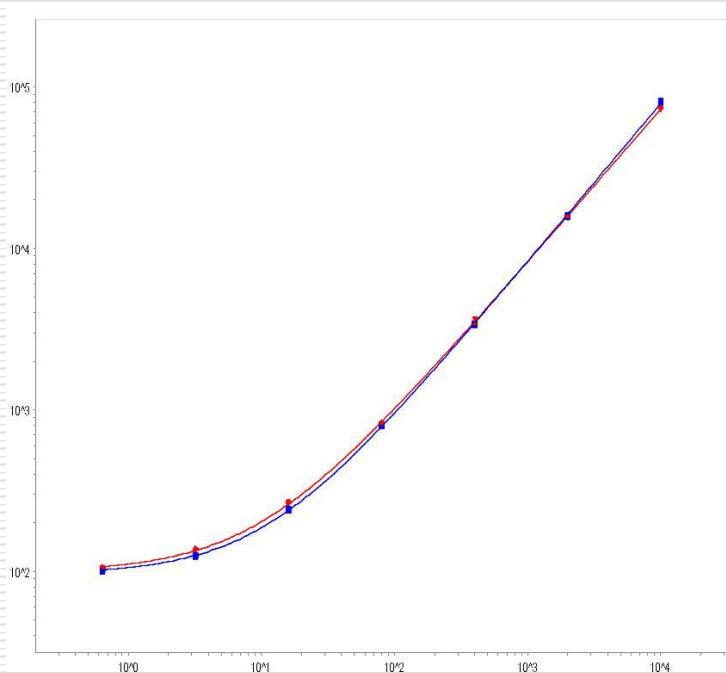


Assay Format: Sandwich Immunoassay



SMN-ECL Assay Characteristics

Comparison of replicate SMN Standard curves



Capture antibody: mouse monoclonal (2B1) anti-human SMN

Detection antibody: rabbit polyclonal anti-human SMN

Sensitivity: lower limits of detection
2-3 pg/ml

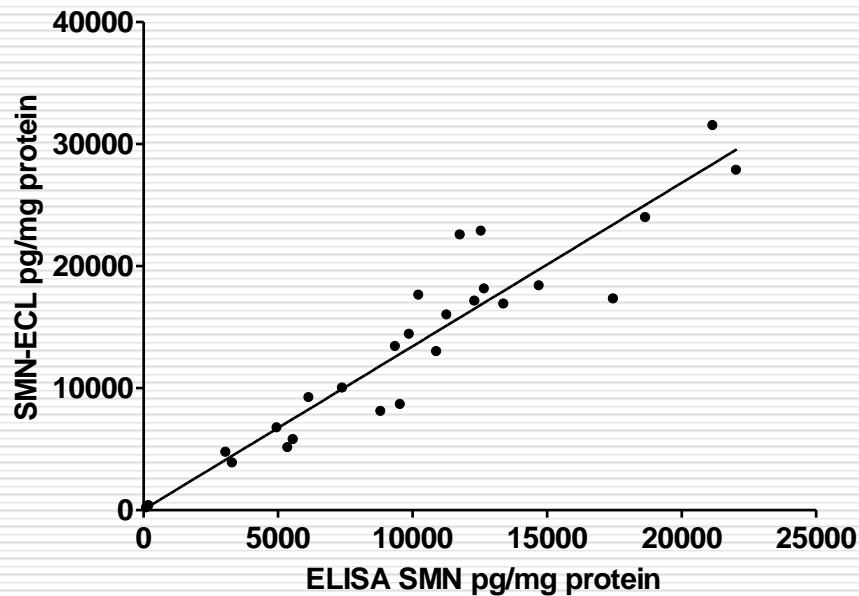
Dynamic range: 3-10,000 pg/ml

Assay time: 3-4 h

Format: 96 well plate

SMN-ECL Assay Correlates With SMN ELISA

SMN-ECL compared to SMN-ELISA in human PBMC lysates



SMAF/PharmOptima Study Report 2011-096: Collection, processing, and ELISA analysis of SMN from PBMC, peripheral blood mononuclear cells, from SMA, spinal muscular atrophy, patients enrolled in a pilot study at Jasper Clinic

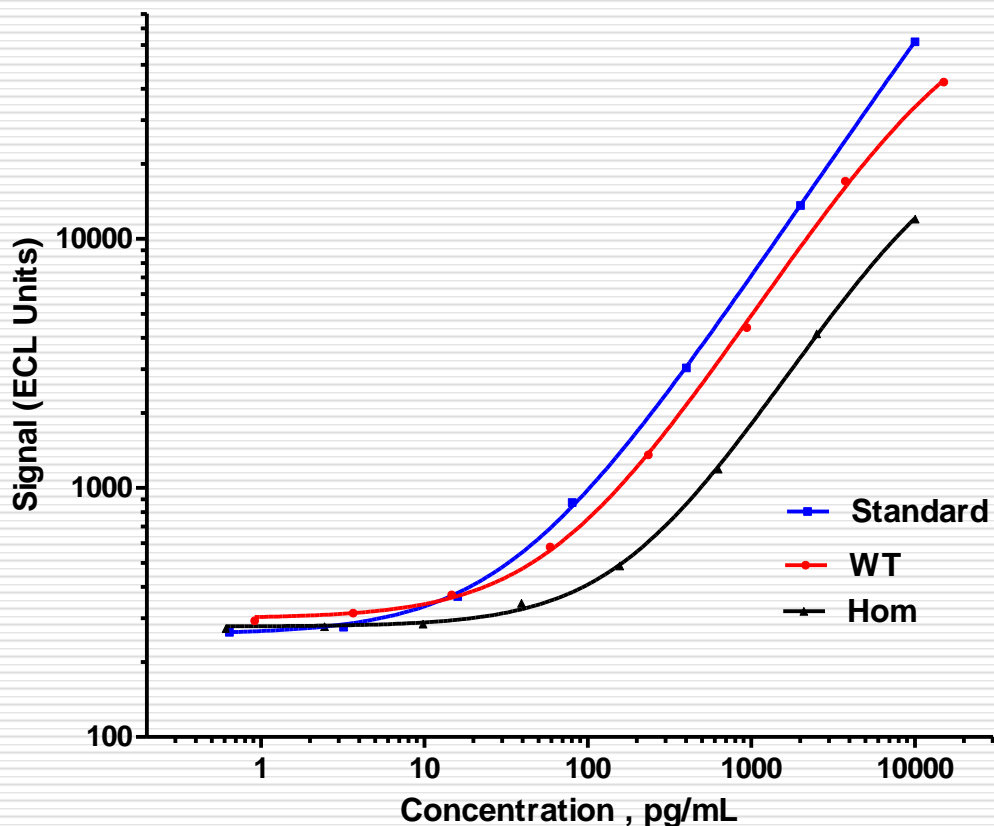
Effect of whole blood contamination on SMN levels in C/C mouse plasma

Sample	Visual Appearance	Mean [SMN] pg/ml	Standard Deviation
Plasma only	amber	14.5	7.1
Plasma, 2% WB lysate spike	pink	154.3	11.4
Plasma, 4% WB lysate spike	pink	412.5	18.3

Effect of whole blood contamination on SMN levels in C/C mouse cerebral spinal fluid (CSF)

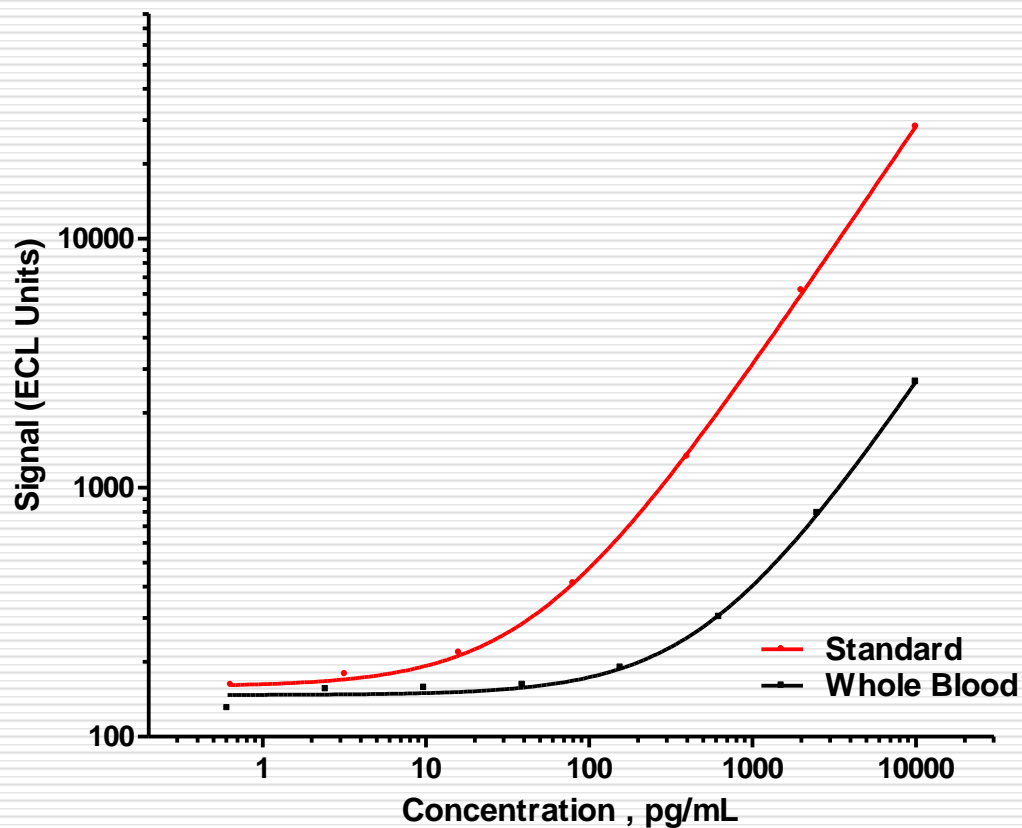
Animal ID	Visual Appearance	Mean [SMN] pg/mL	Standard Deviation
HOM-A18	pink	330.7	5.3
HOM-A26	clear colorless	69.1	7.9
HOM-A48	clear colorless	62.9	0.0

Parallelism between a SMN standard curve and a mouse whole blood dilution curve



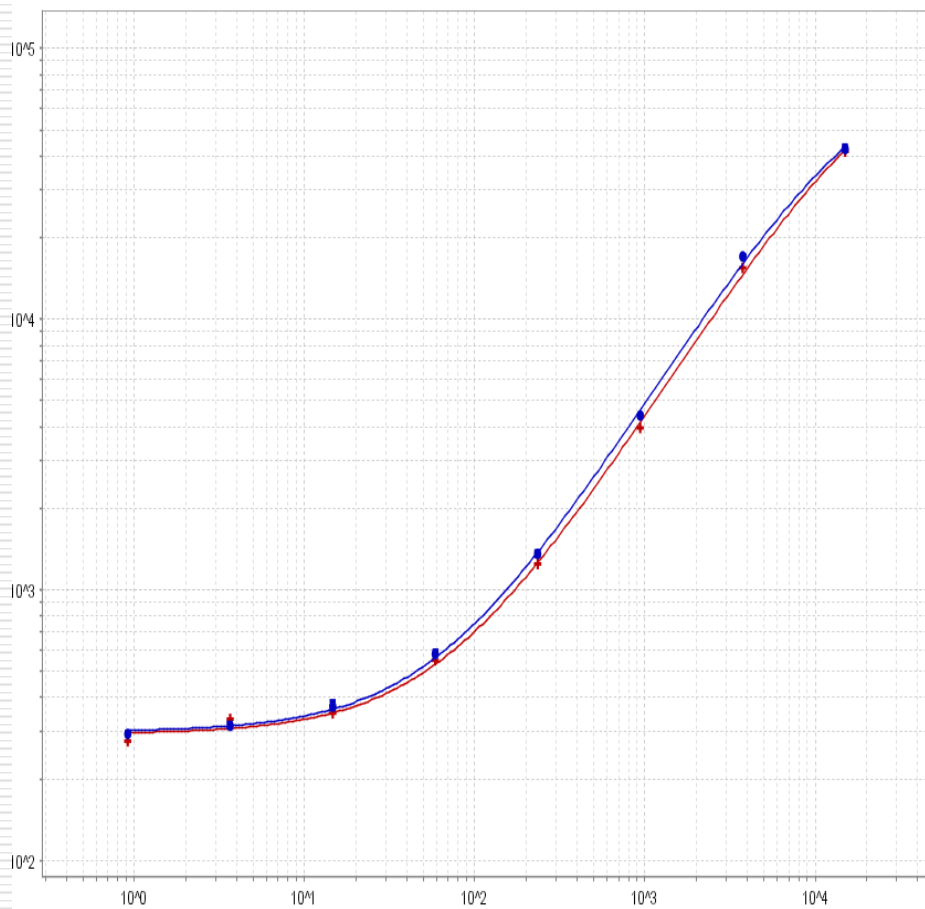
SMN spike (pg/ml) final concentration	% spike recovery
500	90.7
250	83.7
100	90.8

Parallelism between a SMN standard curve and a human whole blood dilution curve



SMN spike (pg/ml) final concentration	% spike recovery
5000	95.9
1000	97.2
200	102.5

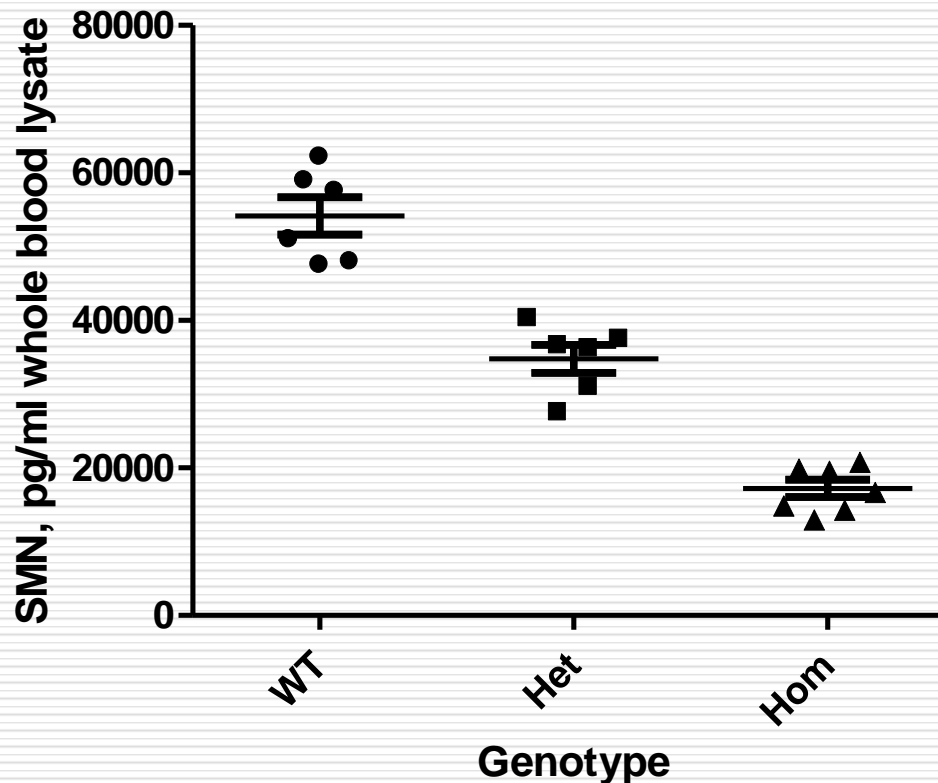
Comparison of mouse whole blood dilution curves following a single freeze/thaw



- **No Freeze Thaw**
- **Freeze Thaw**

SMN in mouse whole blood – SMN changes in C/C mouse genotypes

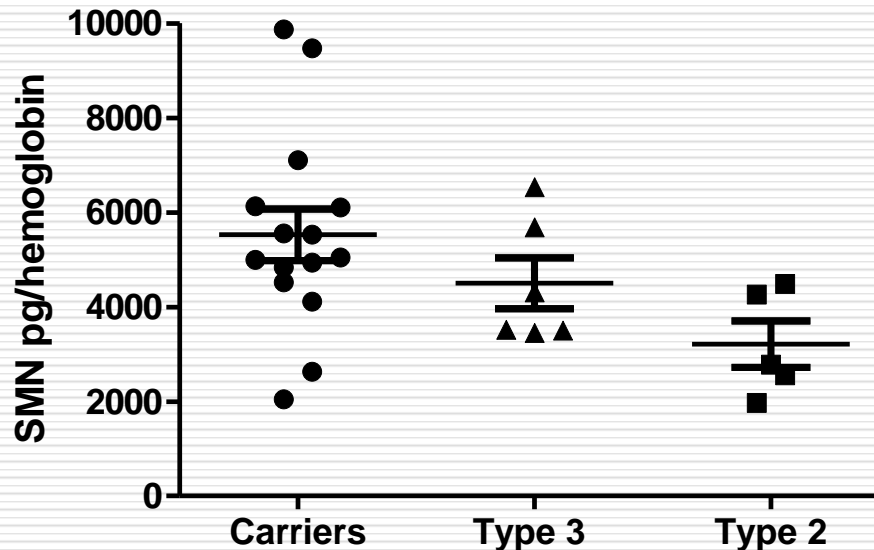
- Comparison of SMN levels in wild type, heterozygous and homozygous C/C mice results in **statistically significant differences between the genotypes**



SMN in human whole blood

SMN changes in human carriers & patients

- Frozen human whole blood from SMA carriers, Type 2 and Type 3 SMA patients demonstrated **statistically significant differences between carriers and Type 2 patients**
- This difference was not seen using PBMCs from the same samples



References

- SMAF/PharmOptima Study Report 2011-096: Collection, processing, and ELISA analysis of SMN from peripheral blood mononuclear cells (PBMC) from SMA (spinal muscular atrophy) patients enrolled in a pilot study at Jasper Clinic
- Evaluation of Peripheral Blood Mononuclear Cell Processing and Analysis for Survival Motor Neuron Protein (Kobayashi et al 2012) PLOS ONE November 2012 | Volume 7 | Issue 11 | e50763

Whole blood fractionation study

Methods:

- Platelet isolation: Low speed centrifugation and harvest of platelet rich plasma
- PBMC isolation:
 - Ficol gradient centrifugation
 - Buffy coat harvest from low speed centrifugation
- Reticulocyte isolation: Anti-CD71 coated magnetic bead (Myltenii)
- Granulocyte isolation: Treatment of red blood cell pellet from low speed centrifugation with erythrocyte lysis buffer.
- RBC isolation:
 - Ficol interference with the SMN-ECL assay.
 - RBC fraction contribution to total SMN signal determined by subtraction

Blood fractionation by centrifugation (Preliminary Results)

Cell Type	Number of cells per mL	Circulation Lifetime	% Total Whole Blood SMN per mL
RBC	$4 - 6 \times 10^9$	100 - 120 days	20 - 40
Granulocyte	$2 - 8 \times 10^6$	hours - days	
Reticulocyte	$4 - 6 \times 10^7$	1 day	2
Platelet*	$1.5 - 4 \times 10^8$	5 - 9 days	40
PBMC	$0.2 - 3 \times 10^6$	Varies greatly	20

*Denis et al 2005 Cell vol 122, 379-391: Escaping the nuclear confines: Signal-dependent pre-mRNA splicing in anucleate platelets.

Summary

- SMN can be measured in whole blood
 - As little as 5 μ l whole blood sufficient for assay
 - Low sample requirements allows for longitudinal studies in mice
 - Statistically significant differences noted between genotypes of SMA C/C mice
 - Statistically significant differences noted between Type 2 patients and carriers
 - SMN appears to be distributed throughout all whole blood cell types examined
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Thank You Contributors



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Why PharmOptima?

- Project team of scientists find optimal experimental approaches
- Experts who understand the data required to move IP up the value chain
- Consultation on best approaches to solve difficult challenges
- Scientists with broad expertise to execute your studies
- Years of experience developing, validating and executing GLP studies
- Full attention of a small company allowing rapid study start and completion and competitive pricing

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