

Volumetric Absorptive Micro Sampler (VAMS or Mitra) in Clinical Diagnostic:

-Therapeutic Drug Monitoring and Diagnostic Biomarkers

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Exagen Diagnostics Inc.

Acknowledgements

Neoteryx LLC (Provides All Mitra)

Dr. Stuart Kushon

Bobby Virasingh

Clinical Trial Patients Recruiting

Site 1: Dr. Smitha Reddy

Site 2: Dr. Puja Chitkara

Thermo Fisher Scientific

Dr. Anastasia Kalli

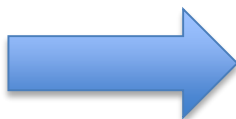
Dr. Raimund Fiedler (ThermoFisher)

Advanced Analytical Technology:

- LC-MS/MS and Immunological based Assay

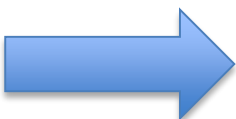


10 ml Venous
Whole Blood



Dried Blood Spot
(DBS), 30-50 μ l

The hematocrit (HCT) of blood affects its viscosity and so gives different-sized DBS spots.



VAMS (Mitra™)

HCT range: 20-65%
10.6 $\pm$ 0.4 μ l

- Spooner et al.,
Bioanalysis (2015), 7(6), 635-659

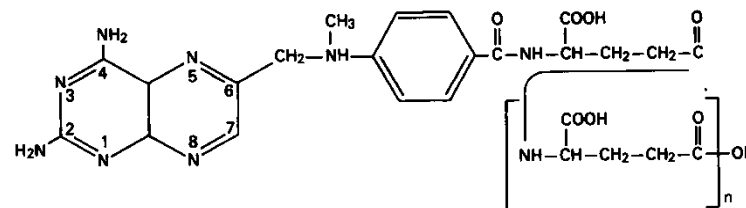
Comparison Study: DBS vs Mitra

- *Therapeutic Drug Monitoring: Methotrexate*
 - *LC-MS/MS method development for Methotrexate polyglutamates (MTXPG1-5);*
 - *Analytical Validation;*
 - *Clinical Validation;*
 - *clinical trial Study.*
- *Diagnostic Biomarker: Rheumatoid Factor (RF)*
 - *Immunological based assay for Anti-CCP IgG, Anti-Mrf IgM, Anti-Arf Ig A and Anti-MCV antibodies;*
 - *Analytical and Clinical Validation;*
 - *clinical trial Study (undergoing....)*

LC-MS/MS method for measuring MTXPGn in DBS and Mitra

Methotrexate (MTX) is a prodrug activated to MTX polyglutamates (MTXPGn)

MTXPG₁, MTXPG₂, MTXPG₃, MTXPG₄, MTXPG₅



DBS Sample Preparation:

- Add 30 µl whole blood (WB) on Whatman 903 DBS card;
- Dry it for at least 3 hours at RT;
- Punch DBS by using 6 mm puncher head into 96 well plate; One spot is equivalent to 14 µl whole blood;
- Add 70 µl water, vortex plate on speed 5 for 30 min;
- Remove 10 µl eluent for hemoglobin quantification;
- In 40 µl eluent, add 40 µl (50 nM) internal standard (IS): MTX-d3 and MTXPG₃-d3
- Add 10 µl 70% perchloric acid for protein precipitation;
- Centrifuge at 14,000 rpm for 5 min.
- Inject 20 µl supernatant to LC-MS/MS

Mitra Sample Preparation

Dip Mitra tip into WB for 5 sec. to absorb 10 µl WB;

- Dry it for at least 3 hours at RT;
- Put Mitra tip into a Eppendorf;
- Add 70 µl water, vortex plate on speed 1 for 30 min;
- Remove 10 µl eluent for hemoglobin quantification;
- In 40 µl eluent, add 40 µl (50 nM) internal standard (IS): MTX-d3 and MTXPG₃-d3
- Add 10 µl 70% perchloric acid for protein precipitation;
- Centrifuge at 14,000 rpm for 5 min.
- Inject 20 µl supernatant to LC-MS/MS

LC-MS/MS method for measuring MTXPGn in DBS and Mitra

LC Conditions

Column (Thermo column) :

Pentafluorophenyl (PFP):

2.1 x 50 mm 2.6 µm particle

Mobile Phase:

A: 0.1%formic acid, 0.01% Triethylamine in water,

B: 0.1%formic acid, 0.01% Triethylamine in ACN

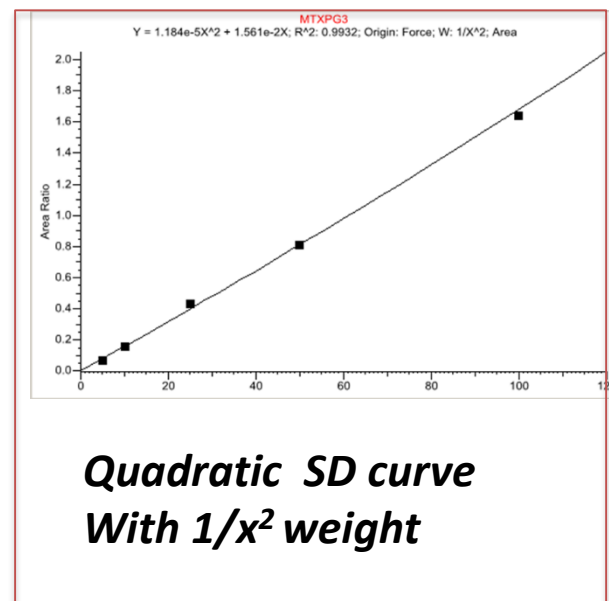
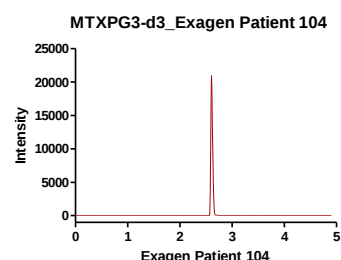
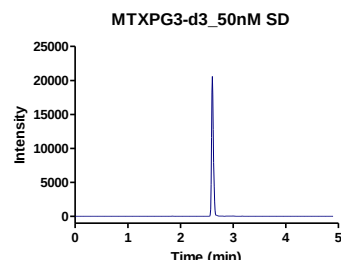
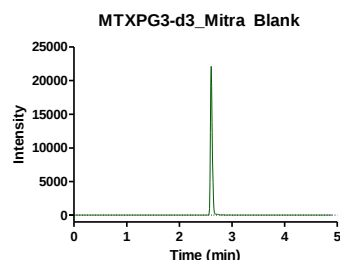
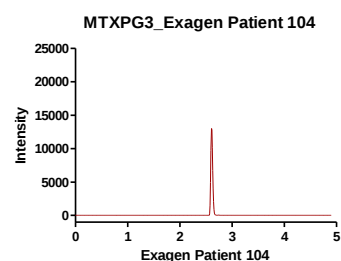
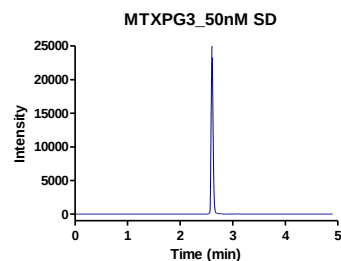
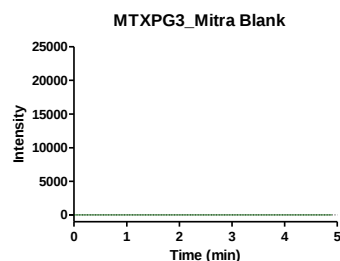
Time (min)	%B	Flow Rate (ml/min)
0.0	5	0.5
0.5	5	0.5
2.0	60	0.5
3.0	100	0.5
4.0	100	0.5
5.0	5	0.5
6.0	5	0.5

Mass Spectrometer Conditions

TSQ-Quantiva triple quadrupole MS
(ThermoScientific)

Parameters		Value
Spray Voltage (positive ion)		3700 (V)
Sheath Gas		45 (Arb)
Aux Gas		5 (Arb)
Sweep Gas		1 (Arb)
Ion Transfer Tube Temp		325 (°C)
Vaporizer Temp		317 (°C)
Compound	Precursor (m/z)	Product (m/z)
MTX	455.18	308.03
MTX-d3	458.22	311.12
MTXPG2	584.21	308.04
MTXPG3	713.23	308.09
MTXPG3-d3	716.36	311.13
MTXPG4	842.3	308.22
MTXPG5	971.33	308.12

LC-MS/MS method for measuring MTXPGn in DBS and Mitra



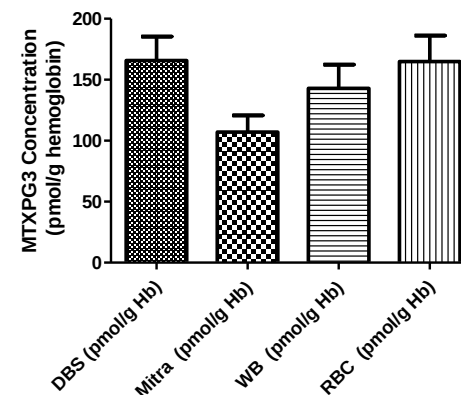
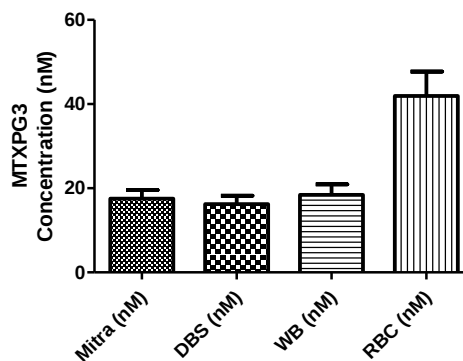
- No interfering peaks were observed in Mitra and DBS samples prepared from blank blood;
- No carryover;
- No Matrix effect for both Mitra and DBS extract (post column infusion experiment);
- Calibration range: 5-100 nM for DBS and Mitra, LOD: 1 nM and LOQ: 5 nM;
- Precision were within 15% for both DBS and Mitra, 20% at LOQ.

Clinical Validation by using Venous Whole Blood

28 Exagen patients which received MTX treatment

- Measured MTXPG₁₋₅ in Whole Blood, DBS, Mitra and RBC;
- Measured hemoglobin concentration in each sample;
- Measured hematocrit (HCT) value for each patient;

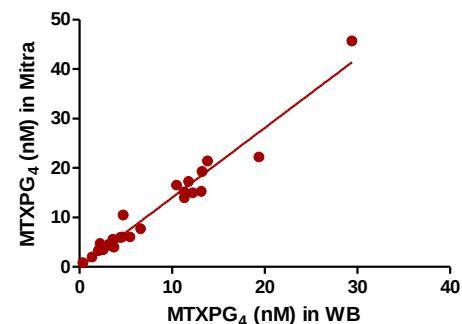
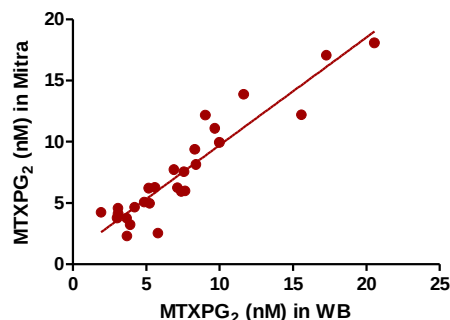
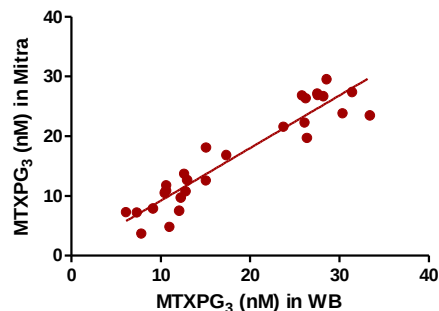
Hemoglobin Concntration				
Filename	DBS (g/L)	Mitra (g/L)	WB (g/L)	RBC (g/L)
16CE010099	95	154	134	228
16CE010100	105	189	156	252
16CE010101	80	154	102	224
16CE010102	115	189	147	216
16CE010103	85	168	125	268
16CE010104	85	147	124	244
16CE010105	90	147	115	292
16CE010106	100	182	127	270
16CE010107	95	182	134	280
16CE010109	100	175	134	210
16CE010110	85	154	129	272
16CE010111	85	161	132	270
16CE010113	110	175	132	274
16CE010114	145	182	137	270
16CE010115	115	147	129	226



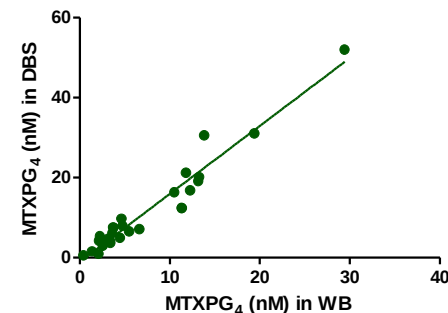
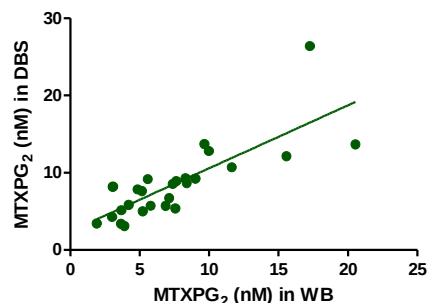
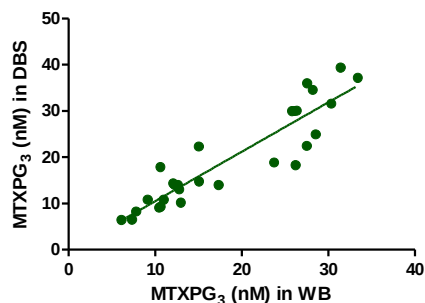
Elution recovery of hemoglobin from DBS and Mitra are different.

Results from Venous Blood Prepared Mitra and DBS (n=28)

Correlation of Mitra with Whole Blood



Correlation of DBS with Whole Blood



	Mitra (nM)	DBS (nM)
r^2	0.8839	0.8368
Slope	0.9344 ± 0.06641	1.184 ± 0.1026

	Mitra (nM)	DBS (nM)
r^2	0.8755	0.6166
Slope	0.9364 ± 0.07062	1.049 ± 0.1655

	Mitra (nM)	DBS (nM)
r^2	0.9596	0.9399
Slope	1.450 ± 0.05839	1.778 ± 0.08822

clinical trial: 44 Patients were Recruited in Two Clinical Sites

Clinical Study Case# 1

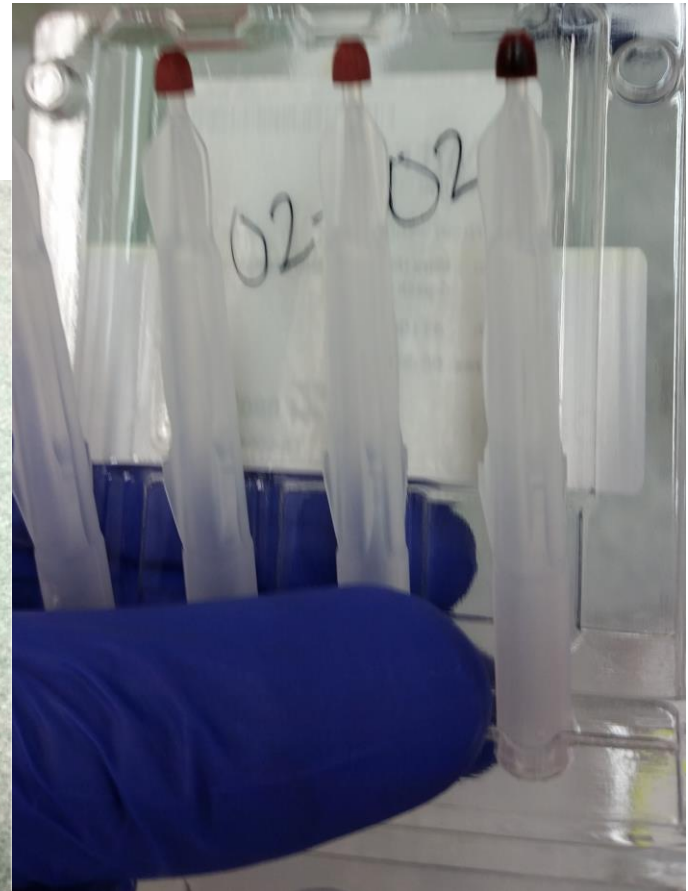
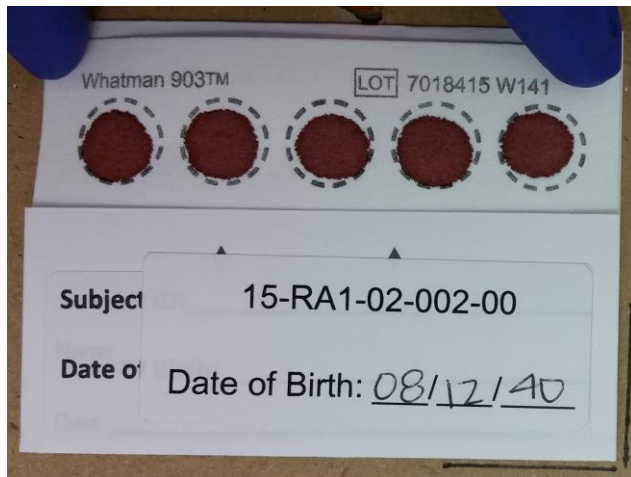
MTXPG₃ (nM)

Capillary Blood:

DBS: 14.6 nM;

Mitra: 13.9 nM;

Venous Whole Blood: 13.9 nM



clinical trial: 44 Patients were Recruited in Two Sites

Clinical Study Case# 2

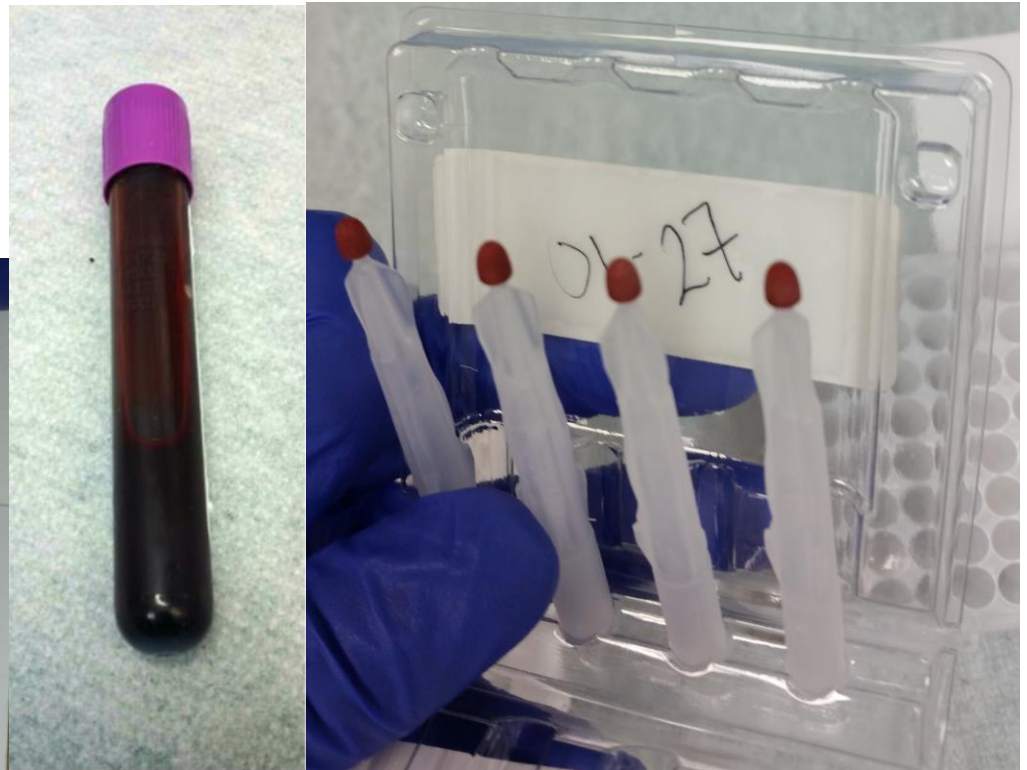
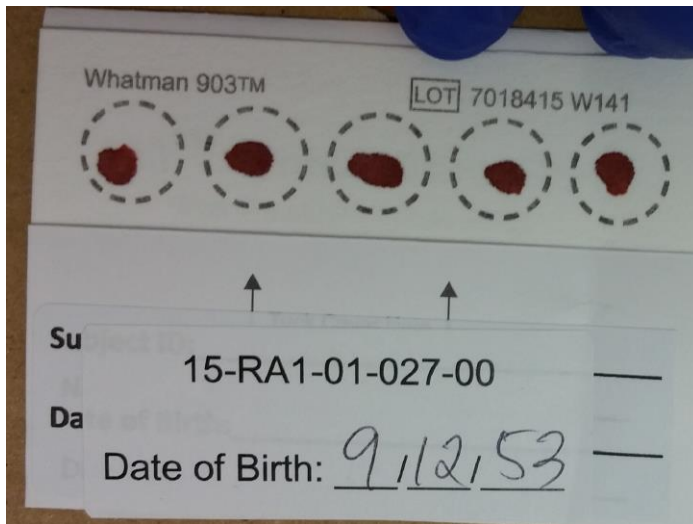
MTXPG₃ (nM)

Capillary Blood:

DBS: 19.4 nM;

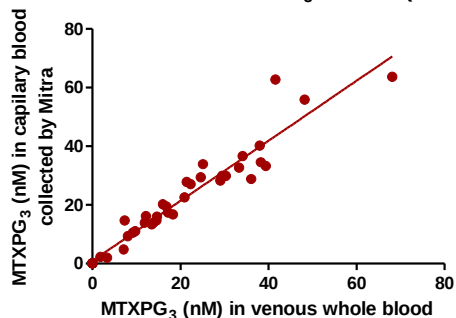
Mitra: 28.2 nM;

Venous Whole Blood: 29.0 nM

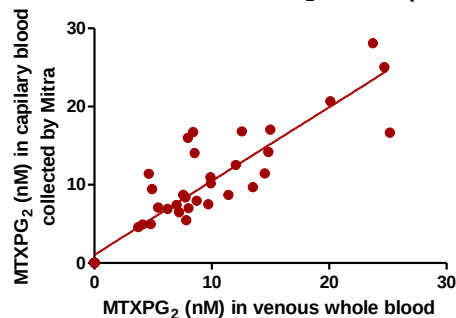


Results Comparison: Venous Blood and Capillary Blood (Mitra and DBS)

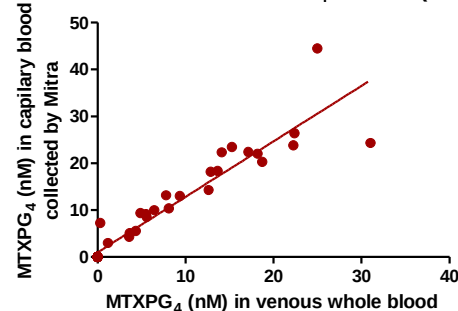
Correlation of MTXPG₃ in Mitra (n=44)



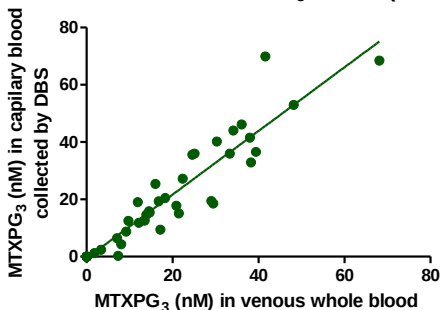
Correlation of MTXPG₂ in Mitra (n=44)



Correlation of MTXPG₄ in Mitra (n=44)

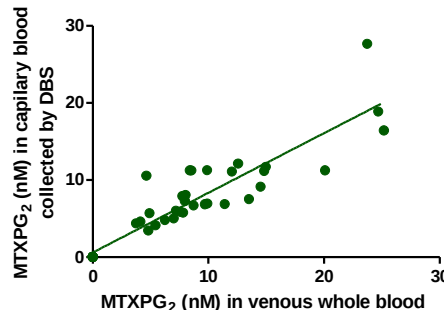


Correlation of MTXPG₃ in DBS (n=44)



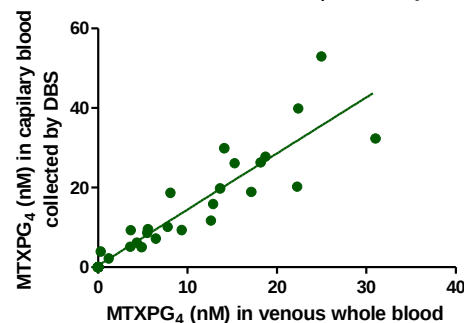
	Mitra (nM)	DBS (nM)
r ²	0.9318	0.8844
	Mitra (nM)	DBS (nM)
Slope	1.063 ± 0.04439	1.194 ± 0.06663

Correlation of MTXPG₂ in DBS (n=44)



	Mitra (nM)	DBS (nM)
r ²	0.8333	0.8349
	Mitra (nM)	DBS (nM)
Slope	1.038 ± 0.07165	0.8343 ± 0.05724

Correlation of MTXPG₄ in DBS (n=44)



	Mitra (nM)	DBS (nM)
r ²	0.8914	0.8652
	Mitra (nM)	DBS (nM)
Slope	1.272 ± 0.06848	1.561 ± 0.09624

Conclusions:

*Using Mitra to collect capillary blood
could represent venous whole blood
for monitoring MTX treatment in patients.*

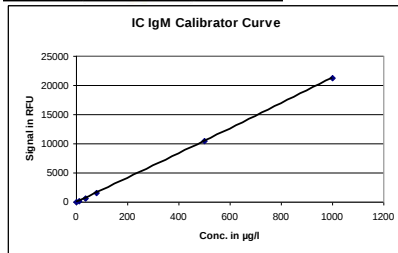
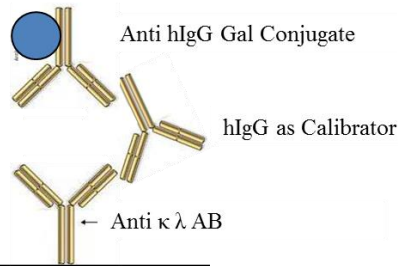
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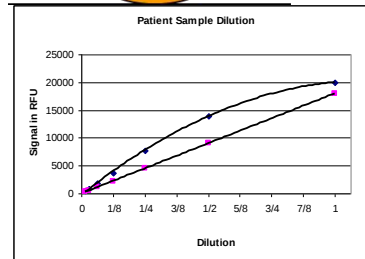
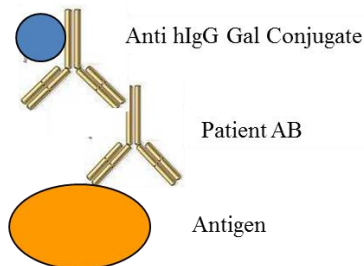
Rheumatoid Arthritis (RA) Diagnostic Panel

- Four serological markers measured in our clinical laboratory: anti-IgM RF, anti-IgA RF, anti-CCP and anti-MCV
- An indirect solid phase enzyme immunoassay (ELISA) was used for the quantitative measurement of Anti IgG class autoantibodies against mutated citrullinated vimentin (MCV);
- The method for determining anti-IgM RF, anti-IgA RF, anti-CCP is a quantitative Fluoroenzyme immunoassay on the Phadia 250 instrument (EliA system);
- All EliA test of one method (IgG, Ig M, Ig A) can be processed with same conjugate and calibration curve; Antigen: CCP, Mrf, Arf, 70, ro, la et. Al. 25 different markers

Substrate



Substrate



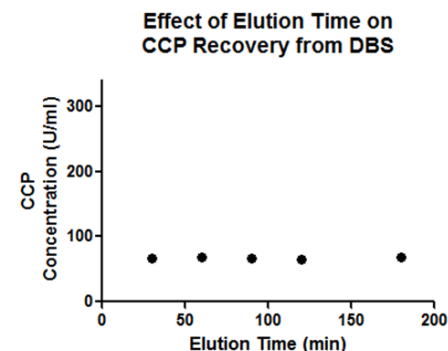
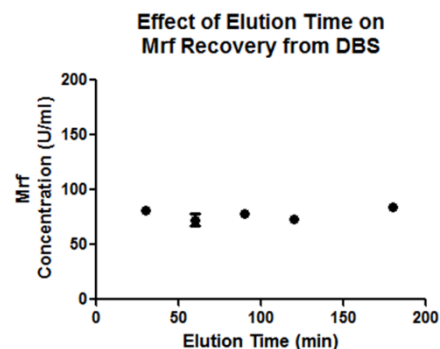
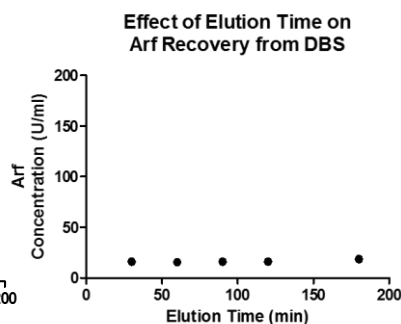
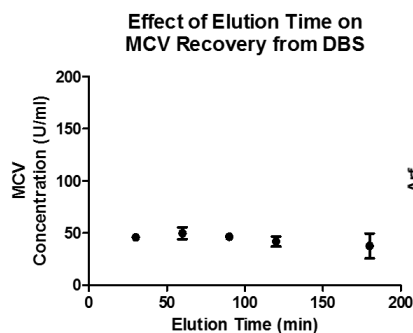
Sample Preparation for Immunological Based Assay

DBS Sample Preparation:

- Add 30 μ l venous whole blood (WB) on Whatman 903 card;
- Dry it for at least 3 hours at RT;
- Punch DBS by using 6 mm puncher head into 96 well plate; One spot is equivalent to 14 μ l whole blood;
- Add 1.0 ml EliA Diluent which are PBS buffer containing BSA, detergent and sodium azide (0.095 %). Shake on a test tube rocker for 30 min;
- Put tube on Phadia instrument and measure anti-IgM RF, anti-IgA RF, anti-CCP.

Mitra Sample Preparation:

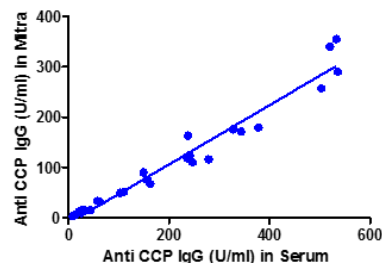
- Dip Mitra tip into venous WB for 5 sec. to absorb 10 μ l WB;
- Dry it for at least 3 hours at RT;
- Put Mitra tip into a test tube;
- Add 1.0 ml EliA Diluent which are PBS buffer containing BSA, detergent and sodium azide (0.095 %). Shake on a test tube rocker for 30 min;
- Remove Mitra tip;
- Put tube on Phadia instrument and measure anti-IgM RF, anti-IgA RF, anti-CCP.



Anti-CCP IgG in DBS and Mitra

Select Anti-CCP positive patients, prepare DBS and Mitra by using venous WB

Patients Anti-CCP Ig G comparison between serum and Mitra (n=32)



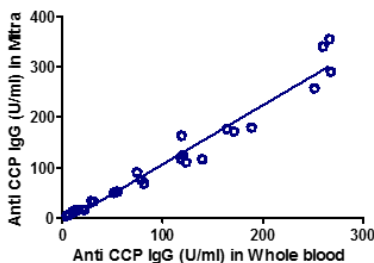
Deming regression line slope

Slope 0.5952 ± 0.02351

The correlation coefficient of linear regression:

r^2 0.9553

Patients Anti-CCP Ig G comparison between WB and Mitra (n=32)



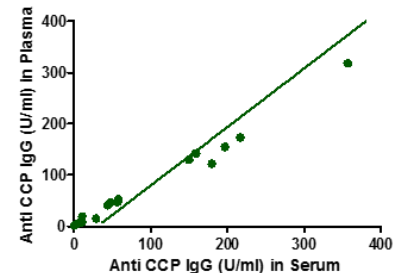
Deming regression line slope

Slope 1.209 ± 0.04774

The correlation coefficient of linear regression:

r^2 0.9553

Patients Anti-CCP Ig G comparison between serum and plasma (n=12)



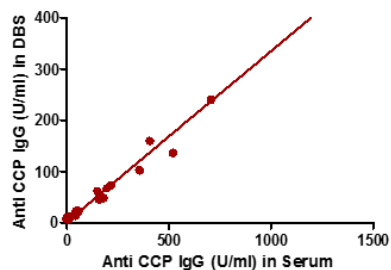
The correlation coefficient of linear regression:

Slope 1.157 ± 0.04120

Deming regression line slope

r^2 0.9777

Patients Anti-CCP Ig G comparison between serum and DBS (n=20)



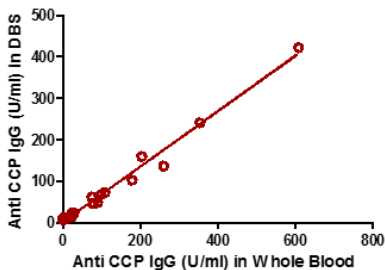
The correlation coefficient of linear regression:

Slope 0.3340 ± 0.009926

Deming regression line slope

r^2 0.9844

Patients Anti-CCP Ig G comparison between WB and DBS (n=20)



The correlation coefficient of linear regression:

Slope 0.6284 to 0.7121

Deming regression line slope

r^2 0.9844

*Original Method:
Measure anti-CCP
in patient serum.*

Anti-CCP IgG in DBS and Mitra

Select Anti-CCP negative patients, prepare DBS and Mitra by using venous WB

Patient ID	CCP Serum	CCP DBS
70916	1.5	8.3
70917	2.2	12
70881	1	7.4
70883	1.5	8.7
70826	1.6	12
70874	1.2	7.2
70914	1.1	7.5
70960	0.9	48
70909	1.6	9.9
70915	1.1	9.3
70923	2	12
70919	1.2	12
70829	1.5	18
70922	0.9	12
70875	0.9	7.9
70832	0.7	9
70913	1	7.8
70836	0.4	9.2
70820	1.8	9
70818	0.5	7

Patient ID	CCP serum	CCP Mitra
88630	0.8	1.3
88633	1.0	1.7
88802	1.4	2.2
88827	1.6	3.0
88819	1.3	2.0
88823	0.9	1.7
88845	1.4	3.4
88876	1.1	2.5
88872	2.4	2.3
88376	3.1	3.1
88414	1.0	2.7
88389	1.4	1.8
88384	1.5	1.9
88085	1.2	2.5
88350	1.6	2.1
88306	0.9	1.3
88308	1.0	1.0

**Non specific absorption
cause high background in DBS*

Continue to finish our clinical trial....