Determination of the Target of Monoclonal Immunoglobulins using the MIAA Assay: A Novel Tool for the Diagnosis of Monoclonal Gammopathies

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Monoclonal Gammopathies

- Three stages: 1. Monoclonal Gammopathy of Undetermined Significance = MGUS
 - 2. Smoldering Multiple Myeloma = SMM
 - 3. Multiple Myeloma = MM



Disease may progress from the asymptomatic MGUS stage toward SMM, then toward overt MM.

MGUS, SMM and MYELOMA



Prevalence



MGUS / SMM

≈ 3-5% population after age 50
Monoclonal Immunoglobulin (Mc Ig) < 30 g/L
Plasma cells < 10% in bone marrow
No symptoms, No treatment – Monitoring of Mc Ig



Evolution MGUS -> SMM -> Myeloma

Slow: > 10 ans Markers of progression: few, and weakly predictive +1%/year/patient (= 15% in 10 years, 30% in 20 years)



Myeloma

Mc Ig > 30 g/L, Plasma cells > 10% in bone marrow **Symptoms: "CRAB"** Calcemia Renal insufficiency

- Anemia
- Bone lesions



Myeloma Treatment: Intensive therapy aimed at suppressing the plasmacytic clone Median survival: 6 years

France ≈ 900 000 MGUS

≈ 900 000 MGUS (40 000 new cases/year) ≈ 100 000 Myeloma Europe

≈ 5.5 millions MGUS

≈ 600 000 Myeloma





Rationale for a New Approach to MGUS and MM

✓ Treaments are rarely curative as long as the cause(s) of a disease is/are not known

Try to find (a) cause(s) for MGUS and MM

 The patient's Mc Ig is a major marker of MGUS and MM disease: yet Mc Ig were assumed not to have antibody function, and were not studied

Our Rationale

- ✓ Ig are typically produced to fight infections
- ✓ Chronic infection at the MGUS stage would explain the inflammation observed in MGUS

For subsets of patients, Mc Ig may target antigens from infectious pathogens

New assay to determine whether a Mc Ig targets an infectious pathogen or protein

Infectious Origin of Monoclonal Gammopathies?

- ✓ Latent infection could explain inflammation at the MGUS stage
- ✓ An abnormal Ig response to infectious pathogen may lead to MGUS (poly-, oligo-, then monoclonal Ig)
- ✓ Accumulation of genetic alterations in clonal plasma cells eventually lead to SMM then Myeloma



✓ Knowing the target of the Mc Ig would indicate the initial event of the clonal gammopathy

Novel therapeutic approach: Treat the target of the Monoclonal Ig to reduce or suppress antigen-induced stimulation of clonal plasma cells and Mc Ig production, hereby preventing evolution toward myeloma

Example: Mc Ig from HCV-infected Patients Typically Recognize a HCV protein

Hermouet et al. New Engl J Med 2003 (1 case)

Hepatitis C Virus, Human Herpesvirus 8, and the Development of Plasma-Cell Leukemia

and human herpesvirus 8 (HHV-8), two B-cell- primary-effusion lymphoma, a condition associattropic viruses, in B-cell proliferation¹ is illustrated ed with HHV-8.4 by the following unusual case of plasma-cell leukemia. In 1995, a 32-year-old man with a history of HCV-driven process leading to plasma-cell leukehepatitis A virus and HCV infection but who tested mia. Immunoblotting showed that the monoclonal negative for hepatitis B virus and human immuno- IgG kappa was directed against HCV core protein deficiency virus was admitted to the hospital with (Fig. 1B). The patient had HCV viremia (type 1a) septic shock, bilateral pneumonia, and hepatosple- and HHV-8 viremia (subtype C'). Reverse-transcripnomegaly. The hemoglobin level was 8.9 g per dec- tase polymerase-chain-reaction (PCR) and immuiliter; the white-cell count was 26.6×109 per liter nofluorescence studies revealed that his plasmawith 50 percent plasmablasts (Fig. 1A), 41 percent blasts were infected by HCV and produced HCV neutrophils, 7 percent lymphocytes, and 2 percent core protein (Fig. 1C). Immunofluorescence and monocytes; the platelet count was 99×109 per liter. real-time (TaqMan) quantitative PCR studies indi-

TO THE EDITOR: The role of hepatitis C virus (HCV) ed in tumor cells of an HCV-positive patient with

Retrospective studies were consistent with an

Bigot-Corbel et al. Blood 2008

For 9/11 HCV+ patients, the Mc Ig targets HCV

2008 112: 4357-4358 doi:10.1182/blood-2008-07-167569

Hepatitis C virus (HCV) infection, monoclonal immunoglobulin specific for HCV core protein, and plasma-cell malignancy

Edith Bigot-Corbel, Michelle Gassin, Isabelle Corre, Didier Le Carrer, Odile Delaroche and Sylvie Hermouet



Figure 1. Studies in a Patient with Plasma-Cell Leukemia.

Panel A shows May-Grunwald-Giemsa staining of blood plasmablasts (×100). Abnormal cells exhibited a plasmablastic morphology: a large nucleus with an irregular shape; fine, homogeneous chromatin and prominent nuclei; reduced cytoplasm with little or no perinuclear hof. Mitotic figures were frequent. Panel B shows the specificity of monoclonal IgG kappa for the hepatitis C virus (HCV) core protein. The patient's serum monoclonal IgG kappa was separated by charge from polyclonal immunoglobulin (Ig) and beta globulins on an agarose gel (Paragon SPE-II, Beckman Coulter). Parts of the gel corresponding to each protein fraction were carefully cut, and proteins were eluted. The purity of each protein fraction was confirmed by immunofixation analysis (Titan Immunogel, Helena Biosciences), and then purified immunoglobulin was subjected to a recombinant immunoblot assay (RIBA III, Ortho Diagnostic Systems), which detects immunoglobulin directed against fragments of the HCV nonstructural proteins NS-3, NS-4, and NS-5 and fragment C22-3 of the core protein. Two purified polyclonal immunoglobulin fractions recognized HCV NS-3 and NS-4 proteins. The monoclonal IgG kappa fraction recognized HCV core protein; it also weakly recognized NS-4, most likely because of overlap with one polyclonal immunoglobulin fraction. The beta globulin fraction, used as a negative control, did not recognize HCV peptides. Panels C and D show immunofluorescence studies. Plasmablasts were incubated with a primary antibody - a control antibody or an antibody specific for HCV core protein (Affinity Bioreagents) or specific for the human herpesvirus 8 (HHV-8) productive phase ORF 59 or ORF K8.1 (Advanced Biotechnologies) - and then with a secondary antibody labeled with fluorescein isothiocyanate. Images show the perinuclear, globular fluorescence characteristic of HCV. Magnification was ×100 for HCV and ×40 for HHV-8.

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Chronic Inflammation in MGUS and in Myeloma

Analysis of inflammation in the serum of MGUS and Myeloma patients:

- ✓ Cytokine profiles of 148 patients (68 MGUS, 6 SMM, 74 MM)
- ✓ 40 cytokines quantified using the Luminex technology (Bio-Plex 200, Bio-Rad)
- ✓ Serum levels of 36/40 cytokines were similarly elevated in MGUS and in MM

Only HGF, IL-11, SDF-1 α and RANTES were significantly higher in MM than in MGUS



Determination of the Target of Monoclonal Igs: Technical Approach (I)

Separation of the monoclonal Ig from other Igs in the serum of patients



Determination of the Target of Monoclonal Igs: Technical Approach (II)

The Multiplexed Infectious Antigen microArray (MIAA) assay

Choice of pathogens: those causing latent infection in humans



The MIAA Assay



16 pad slide used for the MIAA assay, to test simultaneously 7 patients (serum + mc lg) with positive & negative controls.

Feron et al. Anal Biochemistry 2013 2018: US patent N° 9915662 2022: EU patent N° 2877852

Infectious Targets of Monoclonal Igs already Identified

Infectious pathogens identified and confirmed as targets of monoclonal Igs (G/A) in MGUS, SMM and MM:

- 1 bacteria: Helicobacter pylori (H. pylori)
- 6 viruses: EBV, HSV-1, CMV, VZV, HCV, HBV

+ VP1 protein of Human Enteroviruses (2 peptide sequences)



Examples of Virus-Specific Monoclonal IgGs



Infectious pathogen-specific Mc IgG



EBV

HSV-1

Typical Results Obtained with the MIAA assay for EBV (EBNA-1), *H. pylori* and HCV



(A-B) and (C-D) Fluorescent signals obtained (triplicates) for the serum and monoclonal (mc) IgG of two patients whose mc IgG targets EBV: in serum, polyclonal Igs recognize several pathogens, while the mc Ig recognizes a single protein, EBV EBNA-1.

(E) Fluorescent signals obtained (triplicates) for the serum and mc IgG of a patient whose mc IgG targets *H. pylori*: in serum, polyclonal Igs recognize multiple pathogens, while the mc IgG recognizes *H. pylori* only.

(F) Fluorescent signals obtained (triplicates) for the serum and mc IgG of 3 patients whose mc IgG targets HCV: for patient 207, serum polyclonal Igs recognize HCV core and NS-4 proteins, while the mc IgG recognizes HCV NS-4 only.

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Results Obtained for 399 Patients

	Total (n=399)	MGUS (n=155)	MM (n=147)
Identified Targets of mc Ig	243 (60.9%)	120 (77.4%)	69 (46.9%)

Not shown: data from 97 SMM patients

(article in preparation)

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- EBV	115 (28.8%)	53 (34.2%)	38 (25.8%)
- CMV	18 (4.5%	4 (2.6%)	0
- HSV-1	18 (4.5%)	15 (9.7%)	2 (1.4%)
- H. pylori	10 (2.5%)	3 (1.9%)	2 (1.4%)
- VZV	8 (2.0%)	5 (3.2%)	3 (2.0%)
- HCV	3 (0.7%)	2 (1.3%)	1 (0.6%)
Mc Ig of Unknown Target	156 (39.1%)	35 (22.6%)	78 (53.1%)

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- Infection-initiated MGUS and Myeloma are frequent --- the majority in MGUS ! \succ
- Mc Igs from MGUS patients target infectious pathogens twice more frequently than Mc Igs from MM patients >
- CMV and HSV-1 are targets of Mc Igs in MGUS but not in MM \geq

→ Interest for Prognosis: CMV- and HSV-1-associated MGUS, low risk of transformation toward MM ?

MM with EBV EBNA-1-specific Mc IgG: More Severe Disease?

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JCI insight
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2017

RESEARCH ARTICLE

Table 4. Characteristics of SM/MM patients with an mc IgG specific for an infectious pathogen of the MIAA assay

	SM and MM						
	MIAA	MIAA-	Pvalue ^A	EBNA-1*	EBNA-1	P value [®]	Pvalue ^c
Patients, n	28	88		20	96		
Male sex, n (%)	23 (82.1%)	52 (59.1%)	0.04	18 (90.0%)	57 (59.4%)	0.009	0.009
Age at diagnosis (yr)							
Patients, n	28	84		20	92		
Median	65	63	NS	61	64	NS	NS
Range, min-max	37-84	40-90		37-84	40-90		
Leukocytes (10º/l)							
Patients, n	25	84		18	91		
Median	6.40	5.50	NS	6.60	5.50	NS	NS
Range, min-max	0.60-9.70	1.70-18.95		0.60-9.65	1.70-18.95		
Hemoglobin (g/dl)							
Patients, n	27	88		19	96		
Median	11.10	10.90	NS	11.50	10.90	NS	NS
Range, min-max	8.0-15.50	7.0-15.20		9.0-15.50	7.0-15.20		
Platelets (10º/l)							
Patients, n	26	88		18	96		
Median	212.0	211.0	NS	228.5	210.5	NS	NS
Range, min-max	15-428	78-736		91-428	15-736		
Bone Marrow Plasma Cells (%)							
Patients, n	24	69		18	75		
Median	33.5	16.0	0.007	33.5	17.0	0.039	0.023
Range, min-max	5 ⁰ -98	1 ^r -75		5 ⁰ -98	1 ^r -78		
Calcemia (mmol/l)							
Patients, n	24	86		17	93		
Median	2.33	2.33	NS	2.33	2.33	NS	NS
Range, min-max	1.98-3.00	1.71-3.66		1.99-3.00	1.71-3.66		
Creatinine (μmol/l)							
Patients, n	20	75		14	81		
Median	88.5	79.0	NS	93.0	81.0	0.046	0.044
Range, min-max	41-605	35-401		72-605	35-401		
β ₂ -Microgobulin (mg/l)							
Patients, n	22	54		16	61		
Median	4.7	2.9	0.003	4.5	3.1	0.041	0.015
>3.5 mg/l, n (%)	16 (72.3%)	17 (31.5%)	0.0004	11 (68.7%)	22 (36.1%)	0.024	0.010
Range min-max	2.4-16.0	1.3-16.4		2.4-16.0	1.3-16.4		

Myeloma with EBNA-1-specific mc lgG:

- Mostly men
- Greater invasion of bone marrow by clonal plasma cells
- Higher creatinin level
- Higher β2-microbulin level





Article



Characteristics of MGUS and Multiple Myeloma According to the Target of Monoclonal Immunoglobulins, Glucosylsphingosine, or Epstein-Barr Virus EBNA-1

Adrien Bosseboeuf ¹, Nicolas Mennesson ¹, Sophie Allain-Maillet ¹, Anne Tallet ², Eric Piver ^{2,3}, Olivier Decaux ⁴, Caroline Moreau ⁶, Philippe Moreau ⁶, Philippe Lehours ^{7,8}, Francis Mégraud ^{7,8}, Valéry Salle ⁹, Edith Bigot-Corbel ^{1,10}, Jean Harb ^{1,10,11} and Sylvie Hermouet ^{1,12,*}

ENTEROVIRUS VP1 PROTEINS

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Not shown: data from 97 SMM patients

(article in preparation)

- > Mc Igs target Enterovirus VP1 protein four times more frequently in MGUS than in MM
- > Few Enterovirus VP1 protein-associated MGUS seem to evolve toward MM

ENTEROVIRUS VP1 PROTEINS

TAVETGA

8

/GSQFGPYQLLRLI **MLRKRKRLPVFF** DVLLAGNAEADR KKAEEAKKINDKI VKKGNSYPKLSKS

ULQQLAVKTFGQF

SASVPALTAVET

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ADDDDDDETYRG(

HVSPTHYVPI VRDPROVPGR

Detection: PEPperCHIP® infectious disease epitope microarray



VKPRKISVD

Tested:

3760 linear B-cell epitopes, from 196 infectious pathogens

Result: Monoclonal IgGs bind to peptide sequences PALTAVETG and PALTAAETG of VP1 coat proteins of human poliovirus 1, 3 and coxsackievirus B1, B3.



cells

2021

MDPI

Comparison of Monoclonal Gammopathies Linked to Poliovirus or Coxsackievirus vs. Other Infectious Pathogens

Jean Harb 1,2,3,†, Nicolas Mennesson 1, Cassandra Lepetit 1, Maeva Fourny 1, Margaux Louvois 1 Adrien Bosseboeuf¹, Sophie Allain-Maillet¹, Olivier Decaux⁴, Caroline Moreau⁵, Anne Tallet⁶, Eric Piver^{7,8}, Philippe Moreau 9, Valéry Salle 10, Edith Bigot-Corbel 1,30 and Sylvie Hermouet 1,11,+,+0

ENTEROVIRUS VP1 PROTEINS

Confirmation by dot blot assays







Result: Monoclonal IgGs bind to peptide sequences PALTAVETG and PALTAAETG of VP1 coat proteins of human poliovirus 1, 3 and coxsackievirus B1, B3.

Figure 3. Results of the "PV/CVB" Dot Blotting Assays obtained for the 6 Monoclonal Igs found to be Specific for the PALTAV/AETG Epitopes with the PEPperCHIP[®] Infectious Epitope Arrays.



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Self-Antigen: Glucosylsphingosine (GlcSph)

BRIEF REPORT

Clonal Immunoglobulin against Lysolipids in the Origin of Myeloma

Shiny Nair, Ph.D., Andrew R. Branagan, M.D., Jun Liu, Ph.D., Chandra Sekhar Boddupalli, Ph.D., Pramod K. Mistry, M.D., and Madhav V. Dhodapkar, M.B., B.S.

Nair et al. New Engl J Med 2016





cancers

2020

Article

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Glucosylsphingosine-associated myeloma seems to be a mild form of myeloma \geq

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> Mc Igs target glucosylphingosine about as frequently in MGUS as in MM (less frequently in SMM)

Interest for the Prognosis of MGUS and Myeloma



PROGNOSIS

- MYELOMA EBV EBNA-1-associated Myeloma : More severe myeloma disease?
 - Glucosylsphingosine-associated Myeloma: Mild form of myeloma disease?

MGUS - Enterovirus VP1 protein-, CMV- and HSV-1-associated MGUS: Rarely evolve toward myeloma?

Interest for the Therapy of MGUS and Myeloma



Target-reducing therapy works!

2020: Complete remission obtained with Glucosylsphingosine-reducing therapy in two patients (MGUS, SMM) with a GlcSph-specific Mc Ig. Nair S et al. Mol Genet Metab (2020) 129:286

80% of Mc Ig from HCV-infected Patients **Recognize a HCV protein**

Hermouet et al. New Engl J Med 2003 (1 case)

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Efficacy of Target Antigen Reducing Therapy: HCV



60% of Mc Ig from HBV-infected Patients Recognize a HBV protein

Targets of the Mc Igs of 18 HBV-infected patients with MGUS (n=6) or MM (n=12)

- MIAA assay + confirmation by dot blots
- GlcSph assay

Identified Targets

- **HBV (n=11)** 4 MGUS, 7 MM
- EBV (n=3) 2 MGUS, 1 MM
- HSV-1 (n=2) 2 MM
- *H. pylori* (n=1) 1 MM
- GlcSph (n=1) 1 MM

HBV

Target of 61% HBV-infected MGUS/MM patients

Target proteins:

HBV X protein (n=7), HBeAg (n=2), HBc (n=2)



Efficacy of Target Antigen Reducing Therapy: HBV

Efficacy of anti-HBV and anti-HCV treatments in MM patients - Collaboration







		Patients in cohort	Patients with outcome	Survival probability at the end of time window
Ľ	Global MM post HBV infection with AVT	175	36	77.91 %
V coh	Global MM post HBV infection no AVT	1192	329	68.41 %
HB	Log-Rank Test	7.2 5.786	df 1	P 0.016
Ľ	Global MM post HCV infection with AVT	179	33	80.46 %
V coh	Global MM post HCV infection no AVT	1041	253	70.78 %
H	Log-Rank Test	<u>7,2</u>	df	р
		8.026	1	0.005

Anti-viral therapy significantly improves the overall survival at 3 years of MM patients infected with HBV or HCV.

Rodriguez-Garcia et al. Haematologica (2023)

CONCLUSION (I)

• Thanks to the MIAA assay (+ the GlcSph assay), it is now possible to determine the target of Mc Igs from the majority of patients diagnosed with MGUS, SMM or MM

• Knowing the target of a patient's Mc Ig is useful in terms of prognosis and therapy

 Target-reducing therapy is beneficial to MGUS patients (prevention of MM) and to MM patients (improved response to MM treatments)

CONCLUSION (II)



The efficacy of target-reducing therapy is demonstrated for:

✓	GlcSph	(MGUS, SMM)
✓	HCV	(MGUS, MM)
\checkmark	HBV	(MGUS, MM)

 ✓ *H. pylori* (SMM): on-going studies

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