





WELCOME TO

ESPN/ERKNet Educational Webinars on Pediatric Nephrology & Rare Kidney Diseases

• Date: 01 June 2021

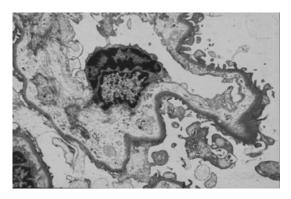
• **Topic:** Atypical Hemolytic Uremic Syndrome

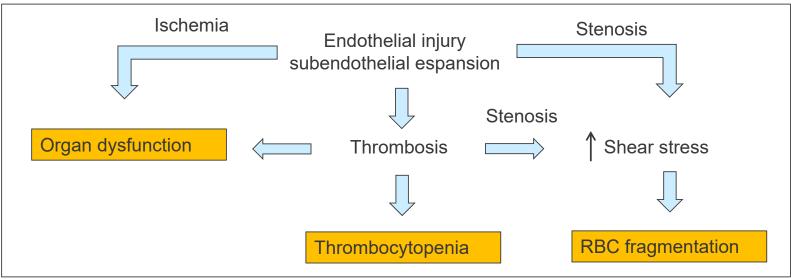
• Speaker: Marina Noris

• Moderator: Elena Levtchenko

HEMOLYTIC UREMIC SYNDROME

Histology lesions: Swelling and detachment of endothelial cells, accumulation of fluffy material in the subendothelium, thrombi and obstruction of the vessel lumina.





Ruggenenti and Remuzzi, Kidney Int 2001

Adhesion molecules and dHUS STEC P-selectin up-regulation

Comp Gene aHUS mutation

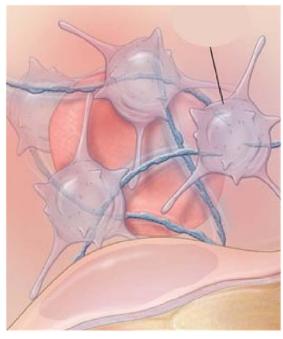
Defective complement regulation

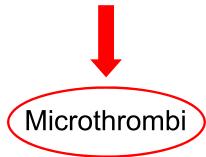
TTP ADAMTS13 **UL-VWF** multimers

sHUS

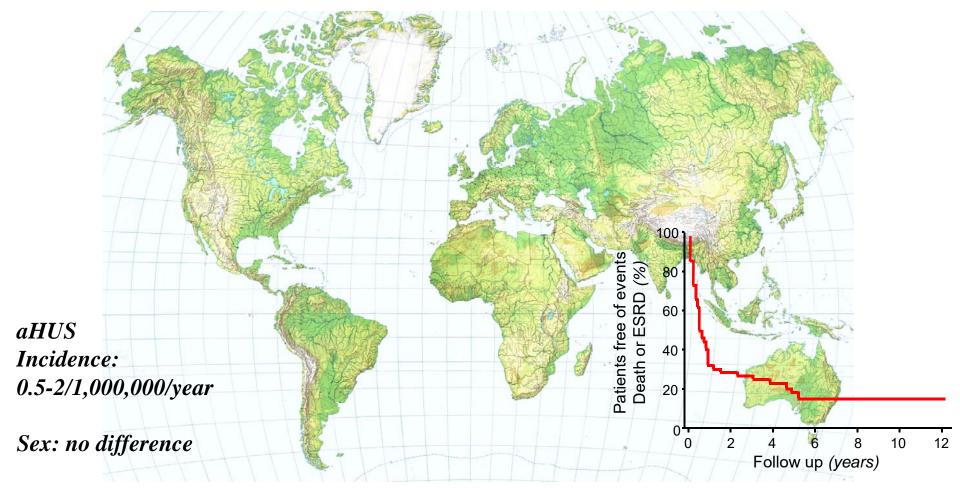
diseases, tx, infections glomerulopathies,

Autoimmune or systemic Complement activating conditions

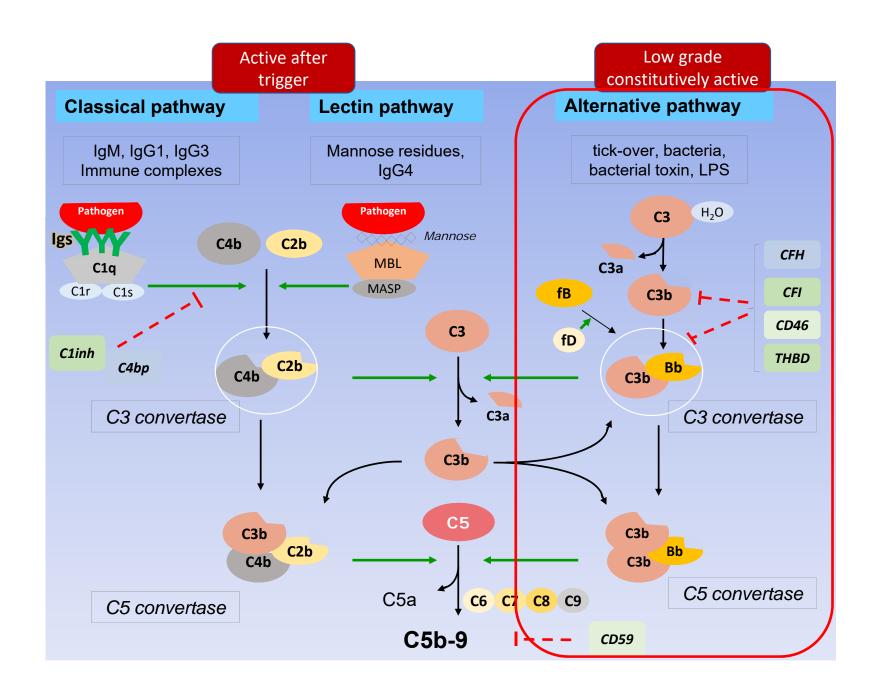




ATYPICAL HEMOLYTIC UREMIC SYNDROME

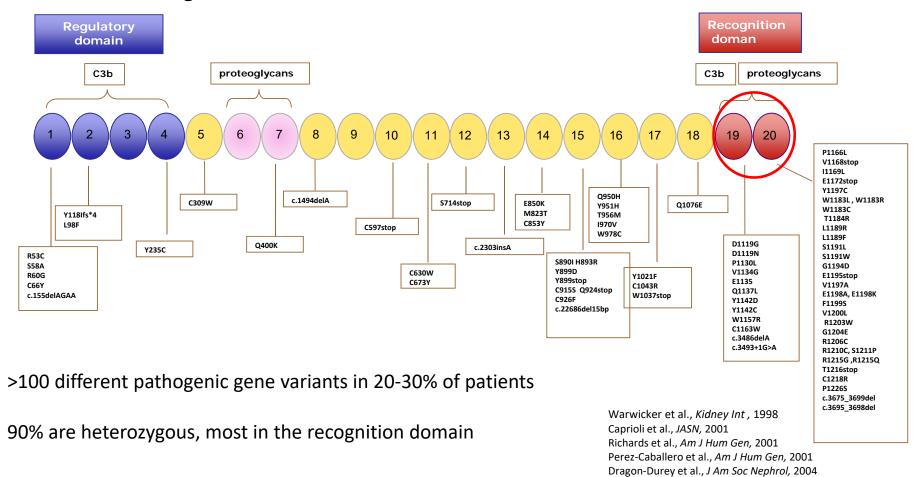


A life threatening multisystem disease of microangiopathic hemolytic anemia, thrombocytopenia, and acute renal failure with predominant but not exclusive renal involvement.



Factor H plays a pivotal role in the regulation of the alternative pathway of complement activation

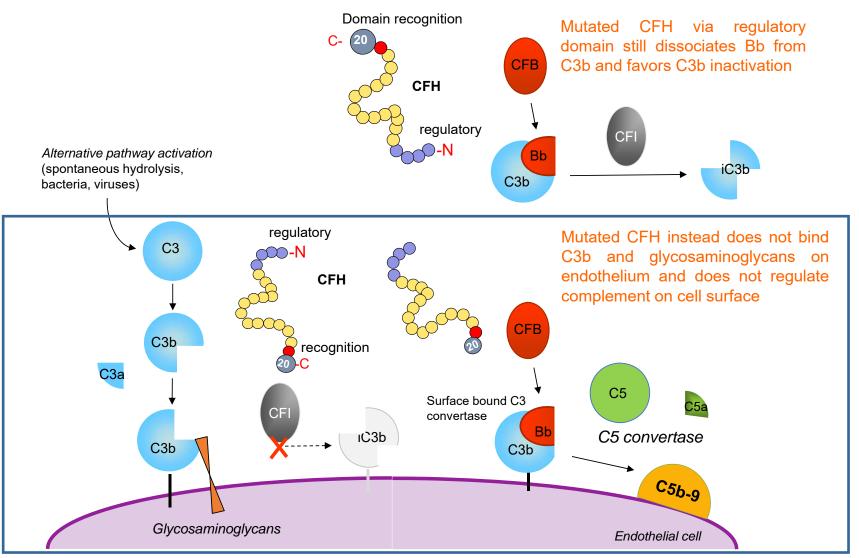
Produced mainly in the liver as a single peptide glycoprotein, factor H circulates in plasma at a concentration of 50 mg/dl



Caprioli et al., Blood, 2006

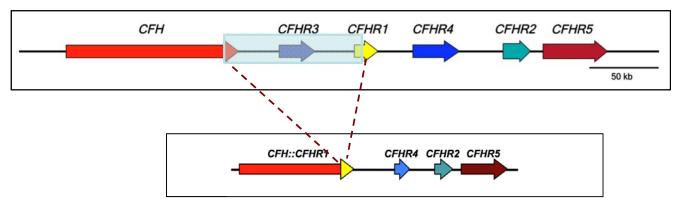
Noris et al, CJASN 2010

SOLID-PHASE RESTRICTED COMPLEMENT ACTIVATION IN aHUS



Manuelian et al., *JCI*, 2003 Sanchez-Corral et al., *Am J Hum Gen*, 2002 Heinen et al., *JASN*, 2007 Noris et al, Blood 2014 Merinero HM et al, Kidney Int 2018

HIGH HOMOLOGY IN THE REGULATORS OF COMPLEMENT ACTIVATION GENE CLUSTER

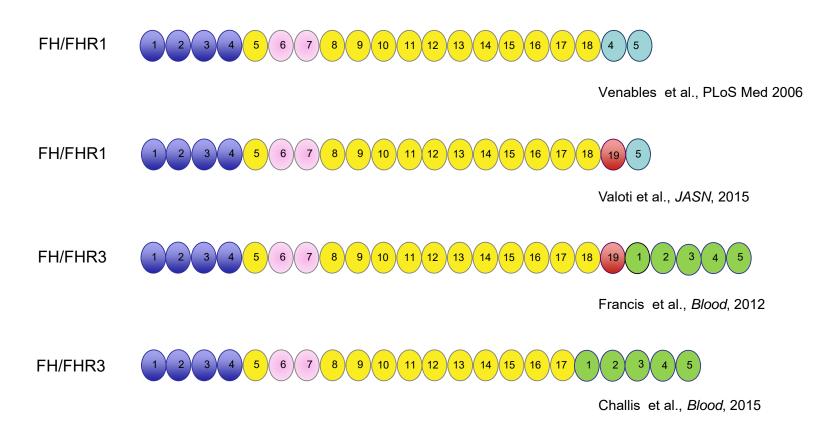


CFH/CFHR1 hybrid gene

Venables et al., Plos Medicine, 2006

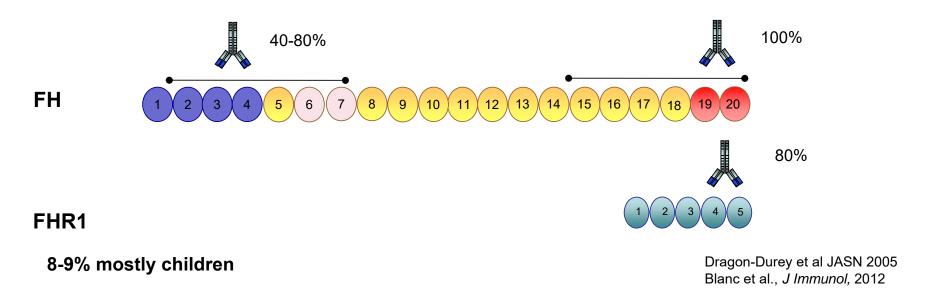
- High degree of sequence identity between the gene for factor H and the genes for the five factor H-related proteins (CFHR1 to 5) which favors non-allelic homologous recombinations giving rise to hybrid genes.
- Copy number variation assays (CGH arrays or MLPA) are required to detect hybrid genes
- Identified in 34/485 aHUS patients (7%).

FH/FHR HYBRID PROTEINS FOUND IN PATIENTS WITH AHUS



In the FH/FHR hybrid molecules the C-terminal SCRs of CFH are substituted with those of FHR1 or with the entire FHR3, resulting in decreased complement regulatory activity on endothelial cell surface

ANTI-FH AUTOANTIBODIES IN AHUS

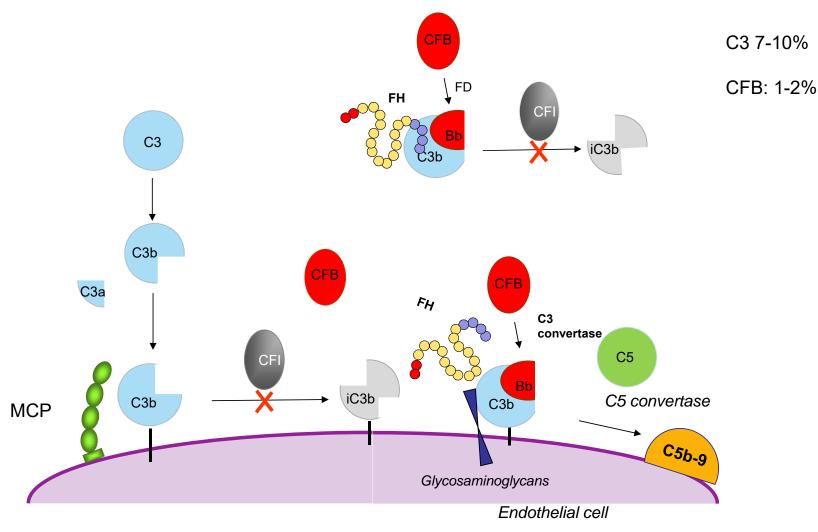


- Anti-FH abs mainly target the FH C-terminus
- Most patients with aHUS and FH autoantibodies are homozygous for a deletion of genes encoding FH related proteins 1 and 3.
- Failure of central and/or peripheral tolerance to FH related 1 and FH?

MCP VARIANTS IN aHUS spontaneous hydrolysis, bacteria, viruses C35X C35Y (n=3) E36X P50T C3 R59X (n=4) C64P K65D 192T>C+193-198del R103W SCRs G130V (96-129)del+G130I+Y132T+L133X Y189D G204R C3b T267fs270X (858-872)del+D277N+P278S F242C C3a STP A353V TM Y328X CT MCP 10% C5 CFI MCP C5 convertase iC3b C3b C5b-9 Endothelial cell

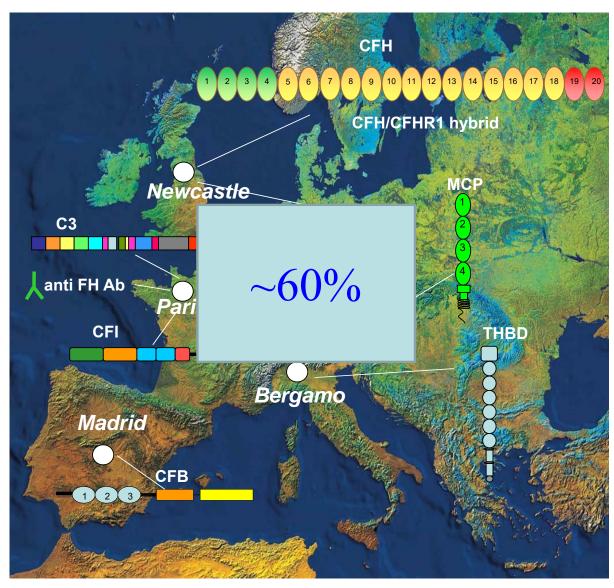
Loss of function heterozygous variants: low expression or reduced C3b binding and cofactor activity

C3 AND CFB GAIN OF FUNCTION VARIANTS IN aHUS

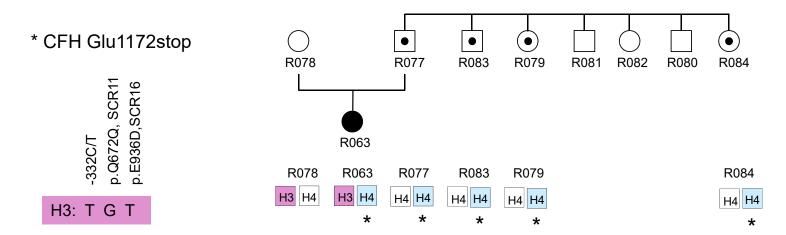


Goicoechea et al., *PNAS*, 2007 Roumenina et al., *Blood*, 2009 Fremeaux-Bacchi et al, Blood 2008

GENETICS OF AHUS: AN EUROPEAN DISCOVERY



THE CASE OF GENE POLYMORPHISMS IN CARRIERS OF PATHOGENETIC VARIANTS TO EXPLAIN INCOMPLETE DISEASE PENETRANCE



Caprioli et al., Hum Molec Genet, 2003

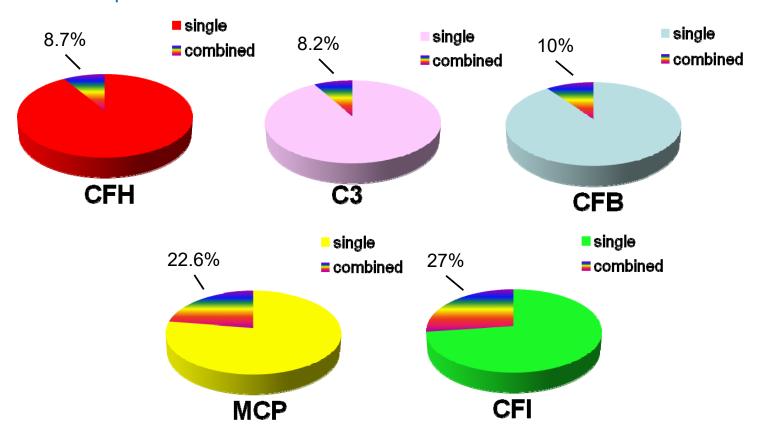
•Among 21 families with CFH pathogenetic variants (26 patients with aHUS and 47 unaffected carrier relatives) the H3 haplotype on wild-type allele was more prevalent in affected (69%) than in unaffected (17%) subjects.

Breno M et al., unpublished

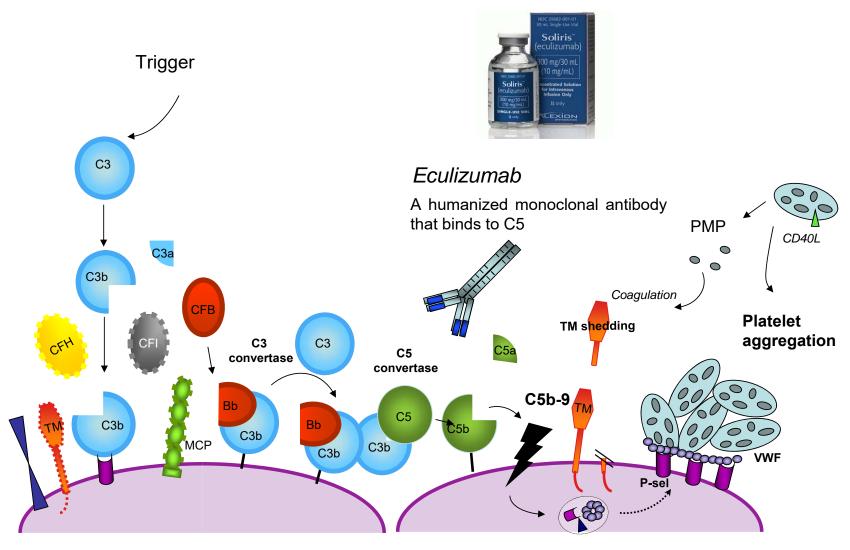
• In 80 pedigrees with a mutation in complement genes (CFH, or MCP or C3 or CFB) aHUS penetrance was 60% in subjects who also carried the CFH-H3 and the MCPggaac risk haplotypes, while only 18% of mutation carriers with neither risk haplotypes developed the disease.

COMBINED COMPLEMENT GENE ABNORMALITIES IN aHUS

795 patients from 4 European cohorts



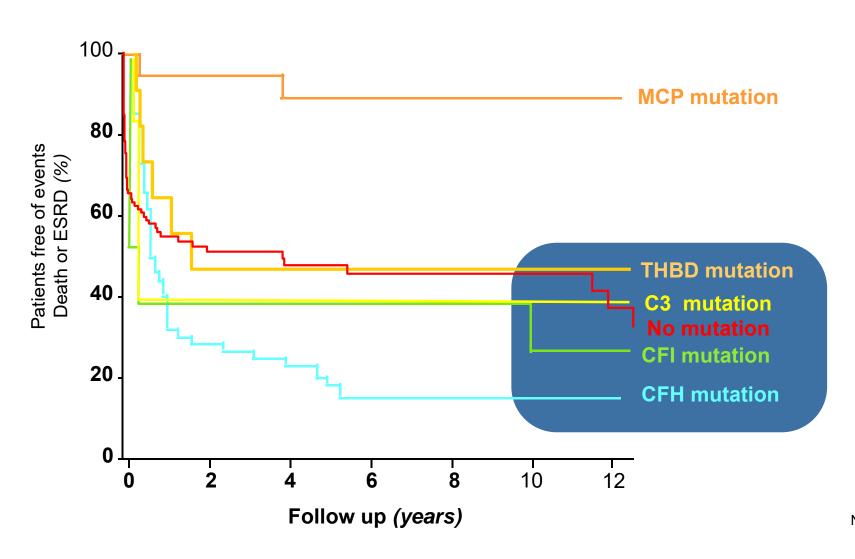
Combined complement gene abnormaties were found in around 10% of patients with CFH or C3 or CFB pathogenetic variants, and about 25% of patients with MCP or CFI pathogenetic variants.



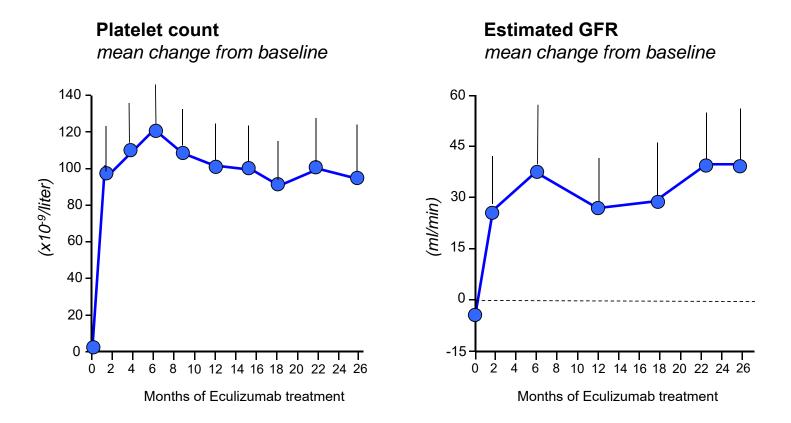
Endothelial cell

Endothelial cell

LONG TERM OUTCOME OF aHUS PATIENTS

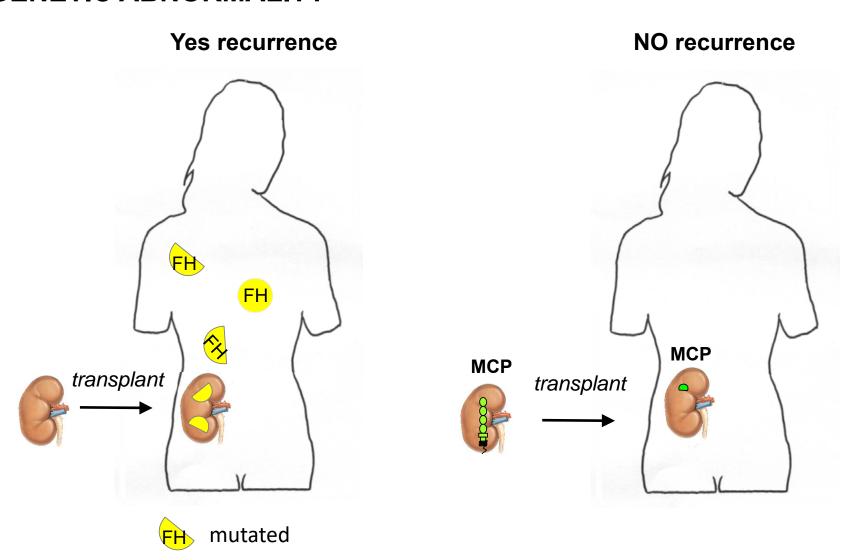


Noris et al CJASN 2010

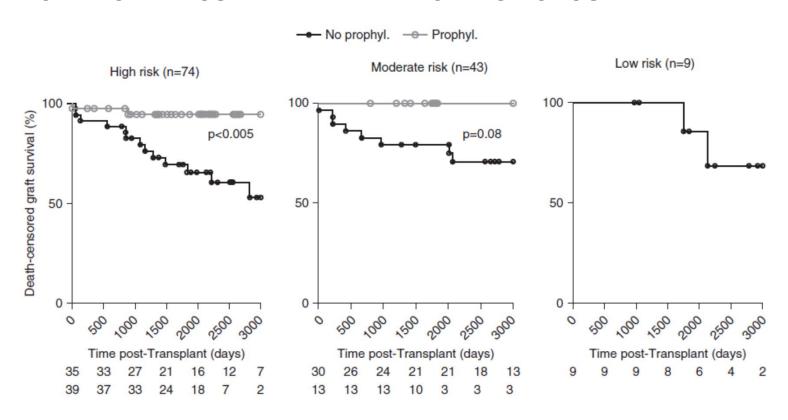


Treatment effect was sustained for up to 26 months

THE RISK OF POST-TRANSPLANT RECURRENCE DEPENDS ON THE GENETIC ABNORMALITY



ECULIZUMAB TREATMENT TAILORING BASED ON THE RISK STRATIFICATION IMPROVED GRAFT SURVIVAL IN THE HIGH RISK GROUP



- High risk: recurrence in a previous graft, and/or pathogenic variants of CFH/C3/CFB
- Moderate risk: negative complement screening results or a pathogenic variant in CFI
- Low risk: isolated variants in MCP or anti-CFH autoantibody no longer detected at the time of transplantation

•A 26-year old male presents with suspicion of TMA after 1 week of vomiting and progressive asthenia:

•Platelets: 39 x10³/µL

•LDH: 1900 IU/L

•Hemoglobin: 8.2 g/dL

Creatinine: 2.7 mg/dL

•ADAMTS13: >10%

•STEC tests: negative

Serum C3, and C4 levels: normal

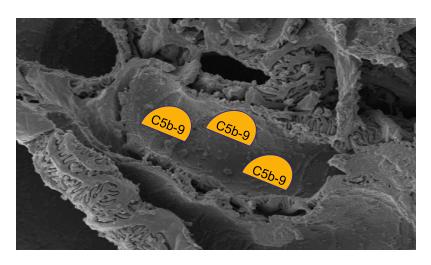
•Plasma C5b-9 levels: normal

•Can this be aHUS?

- •No, because circulating complement parameters are normal
- Yes, could be, but further tests are needed

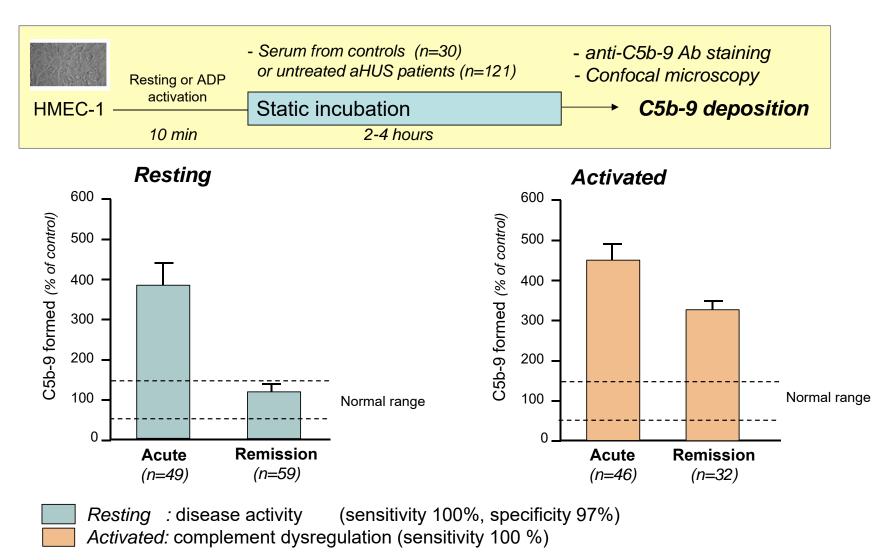
Dynamics of complement activation in aHUS and how to monitor eculizumab therapy

Marina Noris,¹ Miriam Galbusera,¹ Sara Gastoldi,¹ Paolo Macor,² Federica Banterla,¹ Elena Bresin,¹ Claudio Tripodo,³ Serena Bettoni,¹ Roberta Donadelli,¹ Elisabetta Valoti,¹ Francesco Tedesco,⁴ Alessandro Amore,⁵ Rosanna Coppo,⁵ Piero Ruggenenti,⁶ Eliana Gotti,⁶ and Giuseppe Remuzzi^{1,6} BLOOD, 11 SEPTEMBER 2014

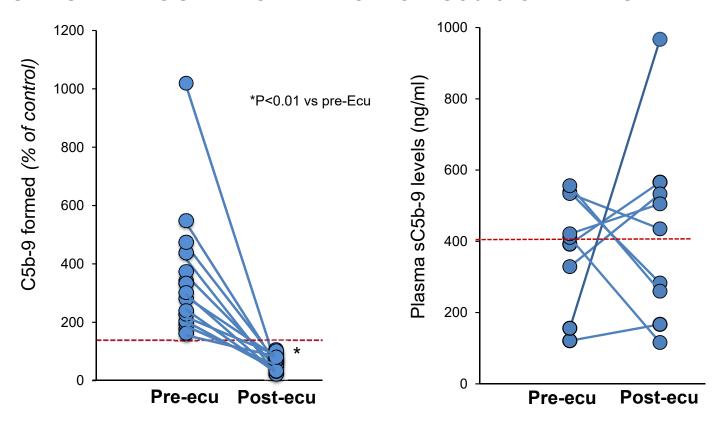


Both during the acute phase of the disease and at remission about half of aHUS patients had normal serum C3 and plasma sC5b-9 levels

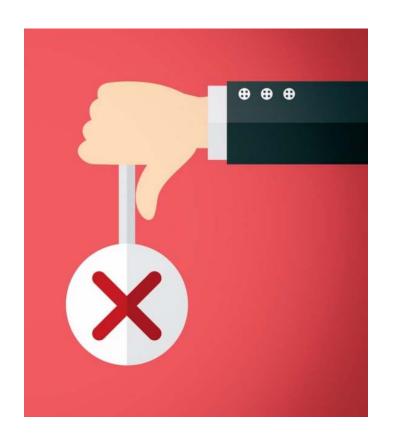
EX VIVO C5b-9 ENDOTHELIAL DEPOSITION IN A LARGE COHORT OF AHUS PATIENTS



ECULIZUMAB TREATMENT IN aHUS PATIENTS FULLY NORMALIZED EX VIVO SERUM-INDUCED FORMATION OF C5b-9 ON HMEC-1

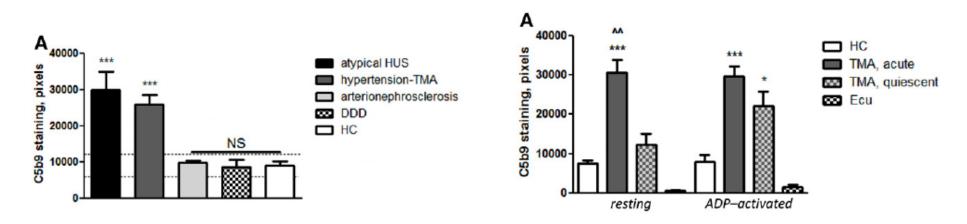


- In 20 aHUS cases treated with Eculizumab, serum-induced C5b-9 deposits on ADP-activated HMEC-1 ex-vivo normalized after treatment
- No significant change was observed in pre- and post-Eculizumab plasma sC5b-9 levels



The assay must be reproduced by other specialized laboratories so that clinicians could consider using it

SERUM FROM PATIENTS WITH HYPERTENSION-ASSOCIATED HUS INDUCED C5b-9 DEPOSITS ON HMEC-1



- Compared with serum from healthy controls (HC), serum from patients with aHUS or hypertensionassociated HUS induced abnormal C5b-9 formation on HMEC-1. C5b-9 deposits normalized after eculizumab treatment.
- Either serum or activated plasma from patients with acute aHUS induced more C5b-9 deposition on HMEC-1 than control sera.
- C5b-9 deposits were at control levels or lower with activated-plasma from aHUS patients treated with eculizumab



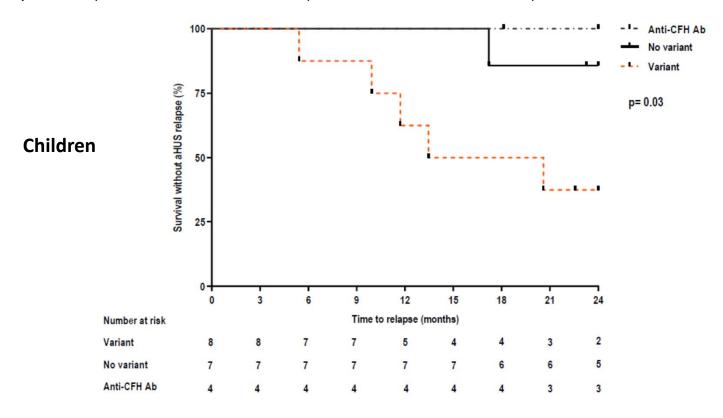
Every 15 days forever?

How to monitor and possibly tapering?



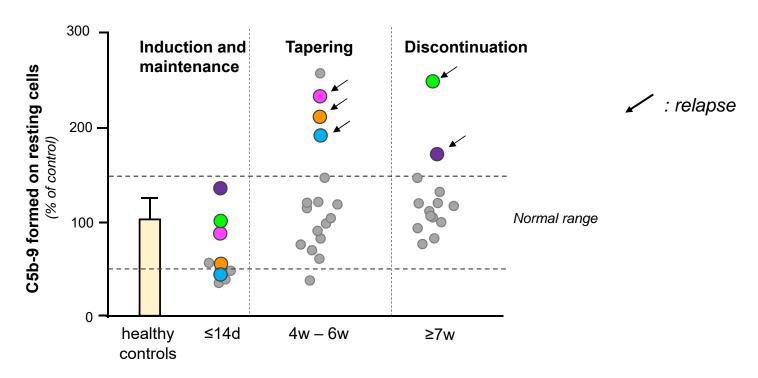
RISK OF AHUS RELAPSE AFTER ECULIZUMAB DISCONTINUATION: A PROSPECTIVE NATIONAL MULTICENTRIC OPEN-LABEL STUDY

55 patients (19 children and 36 adults) discontinued eculizumab (mean duration of treatment, 16,5 months)



During follow-up 13 (23%) patients experienced aHUS relapse. In multivariable analysis, the presence of a rare complement gene variant was associated with an increased risk of relapse (OR 16.20 [1.78-147.72]).

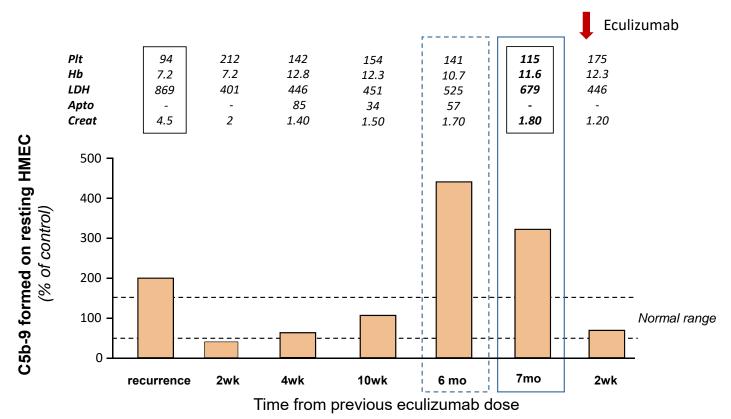
EX VIVO AHUS SERUM-INDUCED C5b-9 FORMATION ON RESTING HMEC-1 DURING ECULIZUMAB TAPERING/DISCONTINUATION



- The large majority of patients taking eculizumab at extended interdose intervals retained normal serum-induced C5b-9 deposits on unstimulated endothelium
- All five patients (colored dots) manifesting relapses showed elevated C5b-9 deposition on resting endothelial cells, in concomitance with or before worsening of clinical parameters (sensitivity for disease relapse 100%)

42 year old woman heterozygous CFI mutation (p.R187Q)

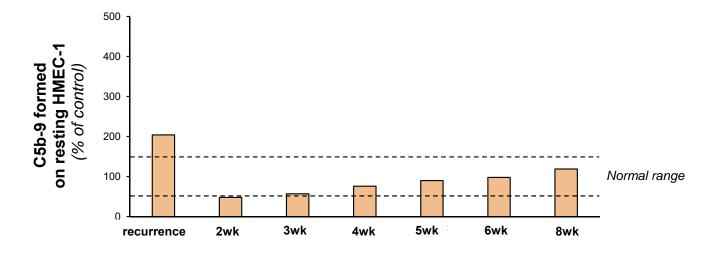
- During a recurrence the patient was treated with eculizumab resulting in disease remission
- After 7 months of eculizumab every 2 weeks, the interval between doses was increased till discontinuation
- Six months after eculizumab cessation, C5b-9 deposits on resting HMEC-1 rose above normal levels
- One month later the patient developed a disease relapse



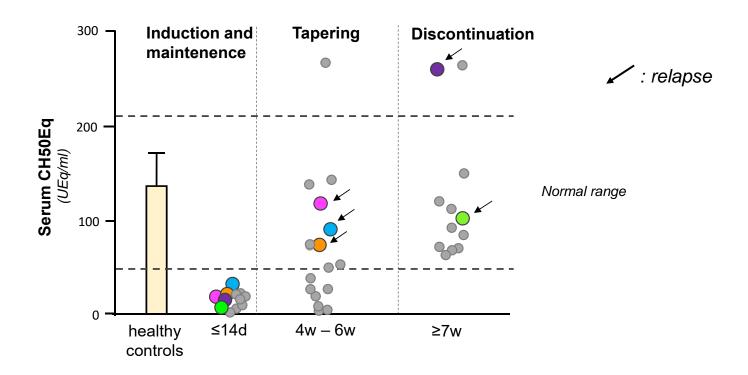
16 year old girl

- recurrent aHUS (onset at 14 years of age)
- no identified complement gene abnormalities
- During a recurrence the patient was treated with eculizumab resulting in disease remission
- After 2 year eculizumab every 2 weeks, the interval between doses was progressively increased
- Hematological and renal parameters and C5b-9 deposits on resting HMEC-1 remained stably normal

Plt	180	147	190	181	203	175	172
Hb	9	11	11.7	11.4	11.6	11.3	10.6
LDH	-	301	349	341	378	313	408
Apto	-	50	82	49	74	83	55
Creat	1.3	0.74	0.81	0.8	0.79	0.66	0.67



CH50 DID NOT EFFICIENTLY DETECT RELAPSE DURING ECULIZUMAB TAPERING/DISCONTINUATION



 At 5-week or longer intervals between eculizumab doses, total complement activity (CH50) levels did not differ between patients who had relapse (coloured dots,132±92 UEq/ml) and those who did not (grey dots,113±72 UEq/ml).

EX VIVO C5b-9 DEPOSITION HIGHLIGHTS COMPLEMENT DYSREGULATION AND DISEASE RELAPSES WHILE THE OTHER BIOMARKERS DO NOT

	Acute aHUS	Remission aHUS		Eculizumab discontinuation		
		No therapy	Eculizumab	No relapse	Relapse	
Plasma sC5b-9	N/†	N/†	N/†	N/†	N/†	
CH50	N	N	↓ ↓	N	N	
C5b-9 deposition on: - resting HMEC	† †	N	N	N	† †	

- Plasma sC5b-9 levels did not differ between acute aHUS and remission and did not consistently change during eculizumab treatment or after discontinuation
- CH50 was normal in aHUS, was suppressed by eculizumab and returned to normal after discontinuation, independently from disease relapses
- C5b-9 deposits on resting HMEC were elevated only in patients with acute aHUS and in those with signs of relapses during eculizumab discontinuation

Take Home Messages

- In Atypical HUS local complement activation on endothelial surface (rather than fluid phase) plays a pathogenic role.
- Genetic abnormalities (mainly heterozygous) causing loss of function of AP complement regulators or gain of function of the components of the AP C3 convertase predispose to aHUS.
- Both gene mutations and risk polymorphism concur to determine disease penetrance.
- C5-inhibition efficiently induces disease remission in aHUS and prevents disease recurrences
- The presence of a pathogenetic complement gene variant increases the risk of relapse upon anti-C5 therapy discontinuation, however relapses are still difficult to predict and early and reliable markers are required

ATYPICAL HEMOLYTIC UREMIC SYNDROME: A DISEASE IN A DISH

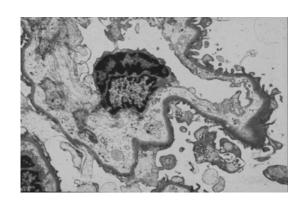


Diagnosis





Monitoring





Thanks!

Caterina Mele

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Matteo Breno

Miriam Galbusera

Sara Gastoldi

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Veronique-Fremeaux Bacchi

Peter Zipfel

Marina Botto

Matthew Pickering

John Atkinson

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Lubka Roumenina

Fadi Fakhouri

David Kavanagh

Josh Thurman

Augustin Tortajada

Next Webinars









- ESPN/ERKNet Educational Webinars on Pediatric Nephrology & Rare Kidney Diseases
- Date: 15 June 2021
- Speaker: Rosa Vargas Poussou
- Topic: Dent Disease
- ERKNet/ERA-EDTA Advanced Webinars on Rare Kidney Disorders
- Date: 29 June 2021
- Speaker: Jürgen Floege
- Topic: Update on KDIGO on Immune Glomerulopathies
- ESPN/ERKNet Educational Webinars on Pediatric Nephrology & Rare Kidney Diseases
- Date: 07 September 2021
- Speaker: Dieter Haffner
- Topic: Renal hypophosphatemia

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