

# Pilar SÃ¡nchez-Corral

## List of Publications by Year in descending order

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Version: 2024-02-01

30  
papers

2,925  
citations

304743

22  
h-index

501196

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docs citations

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times ranked

1864  
citing authors

#	ARTICLE	IF	CITATIONS
1	Contribución de variantes funcionales y cuantitativas del Factor H y las proteínas FHRs (Factor) Tj ETQq1 1 0.784314 rgBT 0 Overlo	0.4	0
2	Low factor H-related 5 levels contribute to infection-triggered haemolytic uraemic syndrome and membranoproliferative glomerulonephritis. CKJ: Clinical Kidney Journal, 2021, 14, 707-709.	2.9	3
3	Complement Genetic Variants and FH Desialylation in S. pneumoniae-Haemolytic Uraemic Syndrome. Frontiers in Immunology, 2021, 12, 641656.	4.8	14
4	Case Report: Combined Liver-Kidney Transplantation to Correct a Mutation in Complement Factor B in an Atypical Hemolytic Uremic Syndrome Patient. Frontiers in Immunology, 2021, 12, 751093.	4.8	3
5	Xenoantibodies and Complement Activity Determinations by Flow Cytometry in Pig-to-Primate Xenotransplantation. Methods in Molecular Biology, 2020, 2110, 73-81.	0.9	0
6	Nephritic Factors: An Overview of Classification, Diagnostic Tools and Clinical Associations. Frontiers in Immunology, 2019, 10, 886.	4.8	52
7	Potential of complement regulator factor H protects human endothelial cells from complement attack in aHUS sera. Blood Advances, 2019, 3, 621-632.	5.2	18
8	Complement as a diagnostic tool in immunopathology. Seminars in Cell and Developmental Biology, 2019, 85, 86-97.	5.0	33
9	Self-Damage Caused by Dysregulation of the Complement Alternative Pathway: Relevance of the Factor H Protein Family. Frontiers in Immunology, 2018, 9, 1607.	4.8	39
10	High Complement Factor H-Related (FHR)-3 Levels Are Associated With the Atypical Hemolytic-Uremic Syndrome-Risk Allele CFHR3*B. Frontiers in Immunology, 2018, 9, 848.	4.8	26
11	Complement factor H, FHR-3 and FHR-1 variants associate in an extended haplotype conferring increased risk of atypical hemolytic uremic syndrome. Molecular Immunology, 2015, 67, 276-286.	2.2	49
12	Atypical aHUS: State of the art. Molecular Immunology, 2015, 67, 31-42.	2.2	236
13	The molecular and structural bases for the association of complement C3 mutations with atypical hemolytic uremic syndrome. Molecular Immunology, 2015, 66, 263-273.	2.2	47
14	Complement Mutations in Diacylglycerol Kinase-Associated Atypical Hemolytic Uremic Syndrome. Clinical Journal of the American Society of Nephrology: CJASN, 2014, 9, 1611-1619.	4.5	61
15	Eculizumab long-term therapy for pediatric renal transplant in aHUS with CFH/CFHR1 hybrid gene. Pediatric Nephrology, 2014, 29, 149-153.	1.7	34
16	Autoantibodies to complement components in C3 glomerulopathy and atypical hemolytic uremic syndrome. Immunology Letters, 2014, 160, 163-171.	2.5	50
17	An ELISA assay with two monoclonal antibodies allows the estimation of free factor H and identifies patients with acquired deficiency of this complement regulator. Molecular Immunology, 2014, 58, 194-200.	2.2	20
18	An Engineered Construct Combining Complement Regulatory and Surface-Recognition Domains Represents a Minimal-Size Functional Factor H. Journal of Immunology, 2013, 191, 912-921.	0.8	70

#	ARTICLE	IF	CITATIONS
19	Combined Complement Gene Mutations in Atypical Hemolytic Uremic Syndrome Influence Clinical Phenotype. <i>Journal of the American Society of Nephrology: JASN</i> , 2013, 24, 475-486.	6.1	308
20	C3 glomerulopathy-associated CFHR1 mutation alters FHR oligomerization and complement regulation. <i>Journal of Clinical Investigation</i> , 2013, 123, 2434-2446.	8.2	176
21	Complement factor H variants I890 and L1007 while commonly associated with atypical hemolytic uremic syndrome are polymorphisms with no functional significance. <i>Kidney International</i> , 2012, 81, 56-63.	5.2	34
22	Anti-factor H antibody affecting factor H cofactor activity in a patient with dense deposit disease. <i>CKJ: Clinical Kidney Journal</i> , 2012, 5, 133-136.	2.9	20
23	Factor H-related protein 1 neutralizes anti-factor H autoantibodies in autoimmune hemolytic uremic syndrome. <i>Kidney International</i> , 2011, 80, 397-404.	5.2	70
24	Advances in understanding the aetiology of atypical Haemolytic Uraemic Syndrome. <i>British Journal of Haematology</i> , 2010, 150, 529-542.	2.5	40
25	Characterization of complement factor H-related (CFHR) proteins in plasma reveals novel genetic variations of CFHR1 associated with atypical hemolytic uremic syndrome. <i>Blood</i> , 2009, 114, 4261-4271.	1.4	190
26	Gain-of-function mutations in complement factor B are associated with atypical hemolytic uremic syndrome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 240-245.	7.1	429
27	De novo gene conversion in the RCA gene cluster (1q32) causes mutations in complement factor H associated with atypical hemolytic uremic syndrome. <i>Human Mutation</i> , 2006, 27, 292-293.	2.5	143
28	Predisposition to atypical hemolytic uremic syndrome involves the concurrence of different susceptibility alleles in the regulators of complement activation gene cluster in 1q32. <i>Human Molecular Genetics</i> , 2005, 14, 703-712.	2.9	272
29	Structural and Functional Characterization of Factor H Mutations Associated with Atypical Hemolytic Uremic Syndrome. <i>American Journal of Human Genetics</i> , 2002, 71, 1285-1295.	6.2	208
30	Clustering of Missense Mutations in the C-Terminal Region of Factor H in Atypical Hemolytic Uremic Syndrome. <i>American Journal of Human Genetics</i> , 2001, 68, 478-484.	6.2	280