

# Barbara Murphy

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4123512/publications.pdf>

Version: 2024-02-01

27  
papers

1,887  
citations

394421

19  
h-index

501196

28  
g-index

29  
all docs

29  
docs citations

29  
times ranked

3709  
citing authors

#	ARTICLE	IF	CITATIONS
1	Deep learning identified pathological abnormalities predictive of graft loss in kidney transplant biopsies. <i>Kidney International</i> , 2022, 101, 288-298.	5.2	28
2	Outcomes of Patients on Maintenance Dialysis Hospitalized with COVID-19. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2021, 16, 452-455.	4.5	25
3	AKI in Hospitalized Patients with COVID-19. <i>Journal of the American Society of Nephrology: JASN</i> , 2021, 32, 151-160.	6.1	500
4	Derivation and validation of a machine learning risk score using biomarker and electronic patient data to predict progression of diabetic kidney disease. <i>Diabetologia</i> , 2021, 64, 1504-1515.	6.3	61
5	DACH1 protects podocytes from experimental diabetic injury and modulates PTIP-H3K4Me3 activity. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	23
6	AMPK mediates regulation of glomerular volume and podocyte survival. <i>JCI Insight</i> , 2021, 6, .	5.0	16
7	APOL1 Long-term Kidney Transplantation Outcomes Network (APOLLO): Design and Rationale. <i>Kidney International Reports</i> , 2020, 5, 278-288.	0.8	62
8	Kidney Failure Associates With T Cell Exhaustion and Imbalanced Follicular Helper T Cells. <i>Frontiers in Immunology</i> , 2020, 11, 583702.	4.8	46
9	COVID-19 and kidney transplantation: Results from the TANGO International Transplant Consortium. <i>American Journal of Transplantation</i> , 2020, 20, 3140-3148.	4.7	305
10	Genome-wide non-HLA donor-recipient genetic differences influence renal allograft survival via early allograft fibrosis. <i>Kidney International</i> , 2020, 98, 758-768.	5.2	25
11	Key driver genes as potential therapeutic targets in renal allograft rejection. <i>JCI Insight</i> , 2020, 5, .	5.0	9
12	Machine Learning to Predict Mortality and Critical Events in a Cohort of Patients With COVID-19 in New York City: Model Development and Validation. <i>Journal of Medical Internet Research</i> , 2020, 22, e24018.	4.3	174
13	Initial Validation of a Machine Learning-Derived Prognostic Test (KidneyIntelX) Integrating Biomarkers and Electronic Health Record Data To Predict Longitudinal Kidney Outcomes. <i>Kidney360</i> , 2020, 1, 731-739.	2.1	15
14	A Peripheral Blood Gene Expression Signature to Diagnose Subclinical Acute Rejection. <i>Journal of the American Society of Nephrology: JASN</i> , 2019, 30, 1481-1494.	6.1	67
15	Disruption of podocyte cytoskeletal biomechanics by dasatinib leads to nephrotoxicity. <i>Nature Communications</i> , 2019, 10, 2061.	12.8	54
16	Pretransplant transcriptomic signature in peripheral blood predicts early acute rejection. <i>JCI Insight</i> , 2019, 4, .	5.0	26
17	NPHP1 (Nephrocystin-1) Gene Deletions Cause Adult-Onset ESRD. <i>Journal of the American Society of Nephrology: JASN</i> , 2018, 29, 1772-1779.	6.1	74
18	Analysis of OPTN/UNOS registry suggests the number of HLA matches and not mismatches is a stronger independent predictor of kidney transplant survival. <i>Kidney International</i> , 2018, 93, 482-490.	5.2	26

#	ARTICLE	IF	CITATIONS
19	Novel Therapeutics Identification for Fibrosis in Renal Allograft Using Integrative Informatics Approach. <i>Scientific Reports</i> , 2017, 7, 39487.	3.3	28
20	Moving Biomarkers toward Clinical Implementation in Kidney Transplantation. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 735-747.	6.1	46
21	Biopsy transcriptome expression profiling: proper validation is key – Authors' reply. <i>Lancet, The</i> , 2017, 389, 601.	13.7	2
22	APOL1: a case in point for replacing race with genetics. <i>Kidney International</i> , 2017, 91, 768-770.	5.2	9
23	APOL1 G2 risk allele – clarifying nomenclature. <i>Kidney International</i> , 2017, 92, 518-519.	5.2	4
24	Biopsy transcriptome expression profiling to identify kidney transplants at risk of chronic injury: a multicentre, prospective study. <i>Lancet, The</i> , 2016, 388, 983-993.	13.7	148
25	Concept and design of a genome-wide association genotyping array tailored for transplantation-specific studies. <i>Genome Medicine</i> , 2015, 7, 90.	8.2	49
26	A Novel Mechanism for the Immunomodulatory Functions of Class II MHC – Derived Peptides. <i>Journal of the American Society of Nephrology: JASN</i> , 2003, 14, 1053-1065.	6.1	8
27	Indirect Allorecognition of Donor Class I and II Major Histocompatibility Complex Peptides Promotes the Development of Transplant Vasculopathy. <i>Journal of the American Society of Nephrology: JASN</i> , 2001, 12, 2500-2506.	6.1	42