Louise Nordfors

List of Publications by Year in descending order

43 papers	2,736 citations	279701 23 h-index	265120 42 g-index
44 all docs	44 docs citations	44 times ranked	3952 citing authors

#	Article	IF	Citations
1	Overexpression of the obese (ob) gene in adipose tissue of human obese subjects. Nature Medicine, 1995, 1, 950-953.	15.2	680
2	Low fetuin-A levels are associated with cardiovascular death: Impact of variations in the gene encoding fetuin. Kidney International, 2005, 67, 2383-2392.	2.6	274
3	Genetic loci influencing kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 373-375.	9.4	246
4	Adiponectin in renal disease: Relationship to phenotype and genetic variation in the gene encoding adiponectin. Kidney International, 2004, 65, 274-281.	2.6	160
5	Low leptin gene expression and hyperleptinemia in chronic renal failure. Kidney International, 1998, 54, 1267-1275.	2.6	148
6	Circulating Levels of Visfatin/Preâ€“B-Cell Colonyâ€“Enhancing Factor 1 in Relation to Genotype, GFR, Body Composition, and Survival in Patients With CKD. American Journal of Kidney Diseases, 2007, 49, 237-244.	2.1	109
7	Genetic and clinical factors influence the baseline permeability of the peritoneal membrane. Kidney International, 2005, 67, 2477-2487.	2.6	108
8	A functional variant of the myeloperoxidase gene is associated with cardiovascular disease in end-stage renal disease patients. Kidney International, 2003, 63, S172-S176.	2.6	105
9	Increased circulating sclerostin levels in end-stage renal disease predict biopsy-verified vascular medial calcification and coronary artery calcification. Kidney International, 2015, 88, 1356-1364.	2.6	102
10	Fat tissue accumulation during peritoneal dialysis is associated with a polymorphism in uncoupling protein 2. Kidney International, 2000, 57, 1713-1719.	2.6	68
11	Large-scale genotyping of single nucleotide polymorphisms by Pyrosequencing? and validation against the 5?nuclease (Taqmanï¿½) assay. Human Mutation, 2002, 19, 395-401.	1.1	66
12	CCR5 Deletion Protects Against Inflammation-Associated Mortality in Dialysis Patients. Journal of the American Society of Nephrology: JASN, 2009, 20, 1641-1649.	3.0	66
13	Reduced gene expression of adiponectin in fat tissue from patients with end-stage renal disease. Kidney International, 2004, 66, 46-50.	2.6	57
14	CDKN2A/p16INK4a expression is associated with vascular progeria in chronic kidney disease. Aging, 2017, 9, 494-507.	1.4	52
15	Visfatin is increased in chronic kidney disease patients with poor appetite and correlates negatively with fasting serum amino acids and triglyceride levels. Nephrology Dialysis Transplantation, 2010, 25, 901-906.	0.4	50
16	AHSG gene variant is associated with leanness among Swedish men. Human Genetics, 2005, 117, 54-60.	1.8	47
17	Changes in fat mass after initiation of maintenance dialysis is influenced by the uncoupling protein 2 exon 8 insertion/deletion polymorphism. Nephrology Dialysis Transplantation, 2006, 22, 196-202.	0.4	30
18	Is Fetuin-A/Î±2-Heremans-Schmid Glycoprotein Associated with the Metabolic Syndrome in Patients with Chronic Kidney Disease?. American Journal of Nephrology, 2008, 28, 669-676.	1.4	30

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19	Expression of Inflammatory and Insulin Signaling Genes in Adipose Tissue in Response to Elective Surgery. Journal of Clinical Endocrinology and Metabolism, 2010, 95, 3460-3469.	1.8	27
20	Genetic approaches in the clinical investigation of complex disorders: Malnutrition, inflammation, and atherosclerosis (MIA) as a prototype. Kidney International, 2003, 63, S162-S167.	2.6	26
21	Increased expression of inflammatory pathway genes in skeletal muscle during surgery. Clinical Nutrition, 2009, 28, 291-298.	2.3	25
22	Progress in Uremic Toxin Research: Genetics/Genomics in Chronic Kidney Diseaseâ€"Towards Personalized Medicine?. Seminars in Dialysis, 2009, 22, 417-422.	0.7	25
23	Is fetuin-A a mortality risk factor in dialysis patients or a mere risk marker? A Mendelian randomization approach. Nephrology Dialysis Transplantation, 2011, 26, 239-245.	0.4	25
24	Interleukin-1 Gene Cluster Polymorphisms Are Associated with Nutritional Status and Inflammation in Patients with End-Stage Renal Disease. Blood Purification, 2005, 23, 384-393.	0.9	21
25	Associations between the CYBA 242C/T and the MPO â€"463G/A Polymorphisms, Oxidative Stress and Cardiovascular Disease in Chronic Kidney Disease Patients. Blood Purification, 2007, 25, 210-218.	0.9	20
26	Relationship Between the ?374T/A Receptor of Advanced Glycation End Products Gene Polymorphism and Peritoneal Solute Transport Status at the Initiation of Peritoneal Dialysis. Therapeutic Apheresis and Dialysis, 2007, 11, 301-305.	0.4	18
27	Expression of osteoprotegerin in human fat tissue; implications for chronic kidney disease. European Journal of Clinical Investigation, 2011, 41, 498-506.	1.7	15
28	How can genetics and epigenetics help the nephrologist improve the diagnosis and treatment of chronic kidney disease patients?. Nephrology Dialysis Transplantation, 2014, 29, 972-980.	0.4	13
29	Influence of cytokine gene polymorphisms on erythropoetin dose requirements in chronic haemodialysis patients. Nephrology Dialysis Transplantation, 2007, 22, 3586-3592.	0.4	12
30	Understanding the role of genetic polymorphisms in chronic kidney disease. Pediatric Nephrology, 2008, 23, 1941-1949.	0.9	12
31	Influence of the CYP2D6 polymorphism and hemodialysis on codeine disposition in patients with end-stage renal disease. European Journal of Clinical Pharmacology, 2010, 66, 269-273.	0.8	12
32	Telomere Attrition and Elongation after Chronic Dialysis Initiation in Patients with End-Stage Renal Disease. Blood Purification, 2016, 41, 25-33.	0.9	11
33	Current epigenetic aspects the clinical kidney researcher should embrace. Clinical Science, 2017, 131, 1649-1667.	1.8	11
34	Longitudinal genome-wide DNA methylation changes in response to kidney failure replacement therapy. Scientific Reports, 2022, 12, 470.	1.6	11
35	Novel insights from genetic and epigenetic studies in understanding the complex uraemic phenotype. Nephrology Dialysis Transplantation, 2014, 29, 964-971.	0.4	9
36	Genotypic and phenotypic predictors of inflammation in patients with chronic kidney disease. Nephrology Dialysis Transplantation, 2016, 31, 2033-2040.	0.4	8

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37	Is UCP2 Gene Polymorphism Associated With Decreased Resting Energy Expenditure in Nondialyzed Chronic Kidney Disease Patients?. , 2008, 18, 489-494.		7
38	Genetic studies in chronic kidney disease: interpretation and clinical applicability. Journal of Nephrology, 2012, 25, 851-864.	0.9	7
39	Use of Single-Nucleotide Polymorphisms in the Search for Genetic Modifiers of the Uremic Phenotype. , 2007, 17, 17-22.		6
40	Association between oestrogen receptor Â gene polymorphism and mortality in female end-stage renal disease patients. Nephrology Dialysis Transplantation, 2007, 22, 2571-2577.	0.4	5
41	Genetic studies in chronic kidney disease: basic concepts. Journal of Nephrology, 2012, 25, 141-149.	0.9	5
42	Selection of Genetic and Phenotypic Features Associated with Inflammatory Status of Patients on Dialysis Using Relaxed Linear Separability Method. PLoS ONE, 2014, 9, e86630.	1.1	4
43	FC 123RENAL TRANSPLANTATION MITIGATES INCREASED BIOLOGICAL (EPIGENETIC) AGE IN CHRONIC KIDNEY DISEASE. Nephrology Dialysis Transplantation, 2021, 36, .	0.4	1