## Iryna Lobach

## List of Publications by Year in descending order

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567281 345221 1,389 39 15 36 citations h-index g-index papers 43 43 43 2826 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Diagnostic Accuracy of Amyloid versus <sup>18</sup> Fâ€Fluorodeoxyglucose Positron Emission Tomography in <scp>Autopsyâ€Confirmed</scp> Dementia. Annals of Neurology, 2021, 89, 389-401.	5.3	34
2	Reach, engagement and effectiveness of in-person and online lifestyle change programs to prevent diabetes. BMC Public Health, 2021, 21, 1314.	2.9	11
3	Pericardial Fat Enhancement. Journal of Thoracic Imaging, 2020, 35, 270-275.	1.5	7
4	Caseâ€control versus caseâ€only estimates of geneâ€environment interactions with common and misclassified clinical diagnosis. Genetic Epidemiology, 2020, 44, 4-15.	1.3	0
5	A second X chromosome contributes to resilience in a mouse model of Alzheimer's disease. Science Translational Medicine, 2020, 12, .	12.4	107
6	Performance of screening MRI in high risk patients at initial versus subsequent screen. Clinical Imaging, 2020, 66, 87-92.	1.5	2
7	Pathogenic DDX3X Mutations Impair RNA Metabolism and Neurogenesis during Fetal Cortical Development. Neuron, 2020, 106, 404-420.e8.	8.1	121
8	Genetic effect estimates in caseâ€control studies when a continuous variable is omitted from the model. Genetic Epidemiology, 2020, 44, 261-271.	1.3	0
9	A systematic review and meta-analysis examining the effects of cannabis and its derivatives in adults with malignant CNS tumors. Neuro-Oncology Practice, 2020, 7, 376-383.	1.6	6
10	Factors Predicting Metastatic Disease in <sup>68</sup> Ga-PSMA-11 PET–Positive Osseous Lesions in Prostate Cancer. Journal of Nuclear Medicine, 2020, 61, 1779-1785.	5.0	15
11	A simple approximation to bias in the genetic effect estimates when multiple disease states share a clinical diagnosis. Genetic Epidemiology, 2019, 43, 522-531.	1.3	1
12	The impact of technical parameters on ablation volume during MR-guided focused ultrasound of desmoid tumors. International Journal of Hyperthermia, 2019, 36, 472-475.	2.5	3
13	A Simple Approximation to Bias in Gene–Environment Interaction Estimates When a Case Might Not Be the Case. Frontiers in Genetics, 2019, 10, 886.	2.3	1
14	Female XX sex chromosomes increase survival and extend lifespan in aging mice. Aging Cell, 2019, 18, e12871.	6.7	46
15	The Gambler's Fallacy in Screening Mammography. Journal of the American College of Radiology, 2019, 16, 830-833.	1.8	1
16	A simple approximation to the bias of gene–environment interactions in case–control studies with silent disease. Genetic Epidemiology, 2019, 43, 292-299.	1.3	0
17	Hyperpolarized $\langle i \rangle$ in vivo $\langle i \rangle$ pH imaging reveals grade-dependent acidification in prostate cancer. Oncotarget, 2019, 10, 6096-6110.	1.8	16
18	Synovial Fluid Profile at the Time of Anterior Cruciate Ligament Reconstruction and Its Association With Cartilage Matrix Composition 3 Years After Surgery. American Journal of Sports Medicine, 2018, 46, 890-899.	4.2	64

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19	Rates of Amyloid Imaging Positivity in Patients With Primary Progressive Aphasia. JAMA Neurology, 2018, 75, 342.	9.0	76
20	Case-control studies of gene-environment interactions. When a case might not be the case. PLoS ONE, 2018, 13, e0201140.	2.5	5
21	Bias in parameter estimates due to omitting gene–environment interaction terms in caseâ€control studies. Genetic Epidemiology, 2018, 42, 838-845.	1.3	4
22	Gene–environment interactions in case–control studies with silent disease. Genetic Epidemiology, 2018, 42, 551-558.	1.3	2
23	A multicenter study to define sarcopenia in patients with endâ€stage liver disease. Liver Transplantation, 2017, 23, 625-633.	2.4	343
24	Tau prions from Alzheimer's disease and chronic traumatic encephalopathy patients propagate in cultured cells. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E8187-E8196.	7.1	141
25	Serum miRNA Signatures Are Indicative of Skeletal Fractures in Postmenopausal Women With and Without Type 2 Diabetes and Influence Osteogenic and Adipogenic Differentiation of Adipose Tissueâe"Derived Mesenchymal Stem Cells In Vitro. Journal of Bone and Mineral Research, 2016, 31, 2173-2192.	2.8	115
26	Intravenous heparin dosing strategy in hospitalized patients with atrial dysrhythmias. Journal of Thrombosis and Thrombolysis, 2016, 42, 179-185.	2.1	6
27	Clinical correlates of longitudinal brain atrophy in progressive supranuclear palsy. Parkinsonism and Related Disorders, 2016, 28, 29-35.	2.2	18
28	Gene Level Meta-Analysis of Quantitative Traits by Functional Linear Models. Genetics, 2015, 200, 1089-1104.	2.9	25
29	Divergent CSF Â alterations in two common tauopathies: Alzheimer's disease and progressive supranuclear palsy. Journal of Neurology, Neurosurgery and Psychiatry, 2015, 86, 244-250.	1.9	101
30	Generalized Functional Linear Models for Geneâ€Based Caseâ€Control Association Studies. Genetic Epidemiology, 2014, 38, 622-637.	1.3	22
31	Periprocedural Glycemic Control in Patients With Diabetes Mellitus Undergoing Coronary Angiography With Possible Percutaneous Coronary Intervention. American Journal of Cardiology, 2014, 113, 1474-1480.	1.6	7
32	A Risk Score for PredictionÂof Hyperglycemia at Coronary Angiography. JACC: Cardiovascular Interventions, 2014, 7, 1199-1201.	2.9	2
33	Genotype-based association models of complex diseases to detect gene-gene and gene-environment interactions. Statistics and Its Interface, 2014, 7, 51-60.	0.3	1
34	Genotype-Based Bayesian Analysis of Gene-Environment Interactions with Multiple Genetic Markers and Misclassification in Environmental Factors. Journal of Probability and Statistics, 2012, 2012, 1-15.	0.7	1
35	Inverse relationship between cirrhosis and massive tumours in hepatocellular carcinoma. Hpb, 2012, 14, 741-745.	0.3	10
36	Risk Factors for Early-Onset and Late-Onset Hepatocellular Carcinoma in Asian Immigrants With Hepatitis B in the United States. American Journal of Gastroenterology, 2011, 106, 1994-2000.	0.4	28

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37	Semiparametric Bayesian analysis of gene-environment interactions with error in measurement of environmental covariates and missing genetic data. Statistics and Its Interface, 2011, 4, 305-315.	0.3	9
38	Genotypeâ€based association mapping of complex diseases: geneâ€environment interactions with multiple genetic markers and measurement error in environmental exposures. Genetic Epidemiology, 2010, 34, 792-802.	1.3	14
39	Haplotypeâ€Based Regression Analysis and Inference of Case–Control Studies with Unphased Genotypes and Measurement Errors in Environmental Exposures. Biometrics, 2008, 64, 673-684.	1.4	21