

# Amaia Jauregi-Miguel

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7630742/publications.pdf>

Version: 2024-02-01

18  
papers

355  
citations

933447

10  
h-index

888059

17  
g-index

20  
all docs

20  
docs citations

20  
times ranked

579  
citing authors

#	ARTICLE	IF	CITATIONS
1	The tight junction and the epithelial barrier in coeliac disease. <i>International Review of Cell and Molecular Biology</i> , 2021, 358, 105-132.	3.2	21
2	Seminal Plasma Triggers the Differential Expression of the Glucocorticoid Receptor (NR3C1/GR) in the Rabbit Reproductive Tract. <i>Animals</i> , 2020, 10, 2158.	2.3	3
3	Semen Modulates Inflammation and Angiogenesis in the Reproductive Tract of Female Rabbits. <i>Animals</i> , 2020, 10, 2207.	2.3	3
4	Allogeneic Embryos Disregulate Leukemia Inhibitory Factor (LIF) and Its Receptor in the Porcine Endometrium During Implantation. <i>Frontiers in Veterinary Science</i> , 2020, 7, 611598.	2.2	6
5	Semen Modulates the Expression of NGF, ABHD2, VCAN, and CTEN in the Reproductive Tract of Female Rabbits. <i>Genes</i> , 2020, 11, 758.	2.4	7
6	TBX3 acts as tissue-specific component of the Wnt/ $\beta$ -catenin transcriptional complex. <i>ELife</i> , 2020, 9, .	6.0	33
7	A novel RT-QPCR-based assay for the relative quantification of residue specific m6A RNA methylation. <i>Scientific Reports</i> , 2019, 9, 4220.	3.3	33
8	The methylome of the celiac intestinal epithelium harbours genotype-independent alterations in the HLA region. <i>Scientific Reports</i> , 2019, 9, 1298.	3.3	23
9	MAGI2 Gene Region and Celiac Disease. <i>Frontiers in Nutrition</i> , 2019, 6, 187.	3.7	8
10	DEXI, a candidate gene for type 1 diabetes, modulates rat and human pancreatic beta cell inflammation via regulation of the type I IFN/STAT signalling pathway. <i>Diabetologia</i> , 2019, 62, 459-472.	6.3	32
11	Celiac Disease-associated lncRNA Named <i>HCG14</i> Regulates <i>NOD1</i> Expression in Intestinal Cells. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2018, 67, 225-231.	1.8	13
12	Transcription Factor Binding Site Enrichment Analysis in Co-Expression Modules in Celiac Disease. <i>Genes</i> , 2018, 9, 245.	2.4	5
13	Subcellular Fractionation from Fresh and Frozen Gastrointestinal Specimens. <i>Journal of Visualized Experiments</i> , 2018, , .	0.3	0
14	Ancestry-based stratified analysis of Immunochip data identifies novel associations with celiac disease. <i>European Journal of Human Genetics</i> , 2016, 24, 1831-1834.	2.8	15
15	Expression analysis in intestinal mucosa reveals complex relations among genes under the association peaks in celiac disease. <i>European Journal of Human Genetics</i> , 2015, 23, 1100-1105.	2.8	38
16	Coregulation and modulation of NF $\kappa$ B-related genes in celiac disease: uncovered aspects of gut mucosal inflammation. <i>Human Molecular Genetics</i> , 2014, 23, 1298-1310.	2.9	74
17	Alteration of Tight Junction Gene Expression in Celiac Disease. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2014, 58, 762-767.	1.8	33
18	Cubic regression-based degree of correction predicts the performance of whole bisulfite amplified DNA methylation analysis. <i>Epigenetics</i> , 2012, 7, 1349-1354.	2.7	5