

Belen Peral

List of Publications by Year in descending order

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49
papers

6,835
citations

136950

32
h-index

197818

49
g-index

49
all docs

49
docs citations

49
times ranked

8026
citing authors

#	ARTICLE	IF	CITATIONS
1	Improved integrative analysis of the thiol redox proteome using filter-aided sample preparation. <i>Journal of Proteomics</i> , 2020, 214, 103624.	2.4	14
2	Cytoskeletal transgelin 2 contributes to gender-dependent adipose tissue expandability and immune function. <i>FASEB Journal</i> , 2019, 33, 9656-9671.	0.5	6
3	Mitoproteomics: Tackling Mitochondrial Dysfunction in Human Disease. <i>Oxidative Medicine and Cellular Longevity</i> , 2018, 2018, 1-26.	4.0	19
4	Differential proteomic and oxidative profiles unveil dysfunctional protein import to adipocyte mitochondria in obesity-associated aging and diabetes. <i>Redox Biology</i> , 2017, 11, 415-428.	9.0	40
5	N-acetylcysteine inhibits kinase phosphorylation during 3T3-L1 adipocyte differentiation. <i>Redox Report</i> , 2017, 22, 265-271.	4.5	5
6	Proteome-wide alterations on adipose tissue from obese patients as age-, diabetes- and gender-specific hallmarks. <i>Scientific Reports</i> , 2016, 6, 25756.	3.3	61
7	Enhanced fatty acid oxidation in adipocytes and macrophages reduces lipid-induced triglyceride accumulation and inflammation. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2015, 308, E756-E769.	3.5	143
8	Parathyroid Hormone-Related Protein, Human Adipose-Derived Stem Cells Adipogenic Capacity and Healthy Obesity. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, E826-E835.	3.6	11
9	Transducin-like enhancer of split 3 (TLE3) in adipose tissue is increased in situations characterized by decreased PPAR α gene expression. <i>Journal of Molecular Medicine</i> , 2015, 93, 83-92.	3.9	5
10	ITCH Deficiency Protects From Diet-Induced Obesity. <i>Diabetes</i> , 2014, 63, 550-561.	0.6	24
11	N-Acetylcysteine affects obesity-related protein expression in 3T3-L1 adipocytes. <i>Redox Report</i> , 2013, 18, 210-218.	4.5	23
12	The MRC1/CD68 Ratio Is Positively Associated with Adipose Tissue Lipogenesis and with Muscle Mitochondrial Gene Expression in Humans. <i>PLoS ONE</i> , 2013, 8, e70810.	2.5	17
13	Attenuated metabolism is a hallmark of obesity as revealed by comparative proteomic analysis of human omental adipose tissue. <i>Journal of Proteomics</i> , 2012, 75, 783-795.	2.4	39
14	Uncovering Suitable Reference Proteins for Expression Studies in Human Adipose Tissue with Relevance to Obesity. <i>PLoS ONE</i> , 2012, 7, e30326.	2.5	25
15	FABP4 Dynamics in Obesity: Discrepancies in Adipose Tissue and Liver Expression Regarding Circulating Plasma Levels. <i>PLoS ONE</i> , 2012, 7, e48605.	2.5	67
16	Breast Cancer 1 (BrCa1) May Be behind Decreased Lipogenesis in Adipose Tissue from Obese Subjects. <i>PLoS ONE</i> , 2012, 7, e33233.	2.5	18
17	Genetic variation near IRS1 associates with reduced adiposity and an impaired metabolic profile. <i>Nature Genetics</i> , 2011, 43, 753-760.	21.4	289
18	Study of the Potential Association of Adipose Tissue GLP-1 Receptor with Obesity and Insulin Resistance. <i>Endocrinology</i> , 2011, 152, 4072-4079.	2.8	121

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19	Decreased STAMP2 Expression in Association with Visceral Adipose Tissue Dysfunction. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011, 96, E1816-E1825.	3.6	34
20	The Gene Expression of the Main Lipogenic Enzymes is Downregulated in Visceral Adipose Tissue of Obese Subjects. <i>Obesity</i> , 2010, 18, 13-20.	3.0	99
21	MiRNA Expression Profile of Human Subcutaneous Adipose and during Adipocyte Differentiation. <i>PLoS ONE</i> , 2010, 5, e9022.	2.5	316
22	Study of the proinflammatory role of human differentiated omental adipocytes. <i>Journal of Cellular Biochemistry</i> , 2009, 107, 1107-1117.	2.6	64
23	Subcutaneous Fat Shows Higher Thyroid Hormone Receptor α 1 Gene Expression Than Omental Fat. <i>Obesity</i> , 2009, 17, 2134-2141.	3.0	39
24	Tackling the human adipose tissue proteome to gain insight into obesity and related pathologies. <i>Expert Review of Proteomics</i> , 2009, 6, 353-361.	3.0	24
25	Differential Proteomics of Omental and Subcutaneous Adipose Tissue Reflects Their Unalike Biochemical and Metabolic Properties. <i>Journal of Proteome Research</i> , 2009, 8, 1682-1693.	3.7	94
26	Proteomics and genomics: A hypothesis-free approach to the study of the role of visceral adiposity in the pathogenesis of the polycystic ovary syndrome. <i>Proteomics - Clinical Applications</i> , 2008, 2, 444-455.	1.6	12
27	Proteomic analysis of human omental adipose tissue in the polycystic ovary syndrome using two-dimensional difference gel electrophoresis and mass spectrometry. <i>Human Reproduction</i> , 2008, 23, 651-661.	0.9	108
28	Differential Gene Expression Profile in Omental Adipose Tissue in Women with Polycystic Ovary Syndrome. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2007, 92, 328-337.	3.6	155
29	Improved resolution of the human adipose tissue proteome at alkaline and wide range pH by the addition of hydroxyethyl disulfide. <i>Proteomics</i> , 2004, 4, 438-441.	2.2	55
30	Association of the Polycystic Ovary Syndrome with Genomic Variants Related to Insulin Resistance, Type 2 Diabetes Mellitus, and Obesity. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2004, 89, 2640-2646.	3.6	146
31	Characterization of a 6p21 translocation breakpoint in a family with idiopathic generalized epilepsy. <i>Epilepsy Research</i> , 2003, 56, 155-163.	1.6	8
32	The Methionine 196 Arginine Polymorphism in Exon 6 of the TNF Receptor 2 Gene (TNFRSF1B) Is Associated with the Polycystic Ovary Syndrome and Hyperandrogenism. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2002, 87, 3977-3983.	3.6	92
33	Common single nucleotide polymorphisms in intron 3 of the calpain-10 gene influence hirsutism. <i>Fertility and Sterility</i> , 2002, 77, 581-587.	1.0	33
34	The Methionine 196 Arginine Polymorphism in Exon 6 of the TNF Receptor 2 Gene (TNFRSF1B) Is Associated with the Polycystic Ovary Syndrome and Hyperandrogenism. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2002, 87, 3977-3983.	3.6	31
35	Mutation Analysis of the Entire PKD1 Gene: Genetic and Diagnostic Implications. <i>American Journal of Human Genetics</i> , 2001, 68, 46-63.	6.2	196
36	A Polymorphic Genomic Duplication on Human Chromosome 15 Is a Susceptibility Factor for Panic and Phobic Disorders. <i>Cell</i> , 2001, 106, 367-379.	28.9	219

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37	Additional Complexity on Human Chromosome 15q: Identification of a Set of Newly Recognized Duplicons (LCR15) on 15q11-q13, 15q24, and 15q26. <i>Genome Research</i> , 2001, 11, 98-111.	5.5	60
38	HMG20A and HMG20B map to human chromosomes 15q24 and 19p13.3 and constitute a distinct class of HMG-box genes with ubiquitous expression. <i>Cytogenetic and Genome Research</i> , 2000, 88, 62-67.	1.1	36
39	Recurrence of the PKD1 nonsense mutation Q4041X in Spanish, Italian, and British families. <i>Human Mutation</i> , 1998, 11, S117-S120.	2.5	16
40	Identification of Mutations in the Duplicated Region of the Polycystic Kidney Disease 1 Gene (PKD1) by a Novel Approach. <i>American Journal of Human Genetics</i> , 1997, 60, 1399-1410.	6.2	100
41	A stable, nonsense mutation associated with a case of infantile onset polycystic kidney disease 1 (PKD1). <i>Human Molecular Genetics</i> , 1996, 5, 539-542.	2.9	94
42	The polycystic kidney disease 1 (PKD1) gene encodes a novel protein with multiple cell recognition domains. <i>Nature Genetics</i> , 1995, 10, 151-160.	21.4	846
43	Splicing mutations of the polycystic kidney disease 1 (PKD1) gene induced by intronic deletion. <i>Human Molecular Genetics</i> , 1995, 4, 569-574.	2.9	90
44	Autosomal dominant polycystic kidney disease: molecular analysis. <i>Human Molecular Genetics</i> , 1995, 4, 1745-1749.	2.9	44
45	Deletion of the TSC2 and PKD1 genes associated with severe infantile polycystic kidney disease a contiguous gene syndrome. <i>Nature Genetics</i> , 1994, 8, 328-332.	21.4	466
46	The polycystic kidney disease 1 gene encodes a 14 kb transcript and lies within a duplicated region on chromosome 16. <i>Cell</i> , 1994, 77, 881-894.	28.9	784
47	Identification and characterization of the tuberous sclerosis gene on chromosome 16. <i>Cell</i> , 1993, 75, 1305-1315.	28.9	1,604
48	Estimating locus heterogeneity in autosomal dominant polycystic kidney disease (ADPKD) in the Spanish population. <i>Journal of Medical Genetics</i> , 1993, 30, 910-913.	3.2	6
49	Fine genetic localization of the gene for autosomal dominant polycystic kidney disease (PKD1) with respect to physically mapped markers. <i>Genomics</i> , 1992, 13, 152-158.	2.9	37