

Hernando Escobar

List of Publications by Year in descending order

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14
papers

476
citations

1040056

9
h-index

1125743

13
g-index

14
all docs

14
docs citations

14
times ranked

945
citing authors

#	ARTICLE	IF	CITATIONS
1	Development of UHPLC/Q-TOF Analysis Method to Screen Glycerin for Direct Detection of Process Contaminants 3-Monochloropropane-1,2-diol Esters (3-MCPDEs) and Glycidyl Esters (GEs). <i>Molecules</i> , 2021, 26, 2449.	3.8	5
2	Peptidomic analysis of type 1 diabetes associated HLA-DQ molecules and the impact of HLA-DM on peptide repertoire editing. <i>European Journal of Immunology</i> , 2017, 47, 314-326.	2.9	22
3	Type 1 diabetes associated HLA-DQ2 and DQ8 molecules are relatively resistant to HLA-DM mediated release of invariant chain-derived CLIP peptides. <i>European Journal of Immunology</i> , 2016, 46, 834-845.	2.9	21
4	High Sensitivity Measurement of Pancreatic Polypeptide and Its Variant in Serum and Plasma by LC-MS/MS. <i>Methods in Molecular Biology</i> , 2016, 1378, 199-210.	0.9	5
5	Measurement of Pancreatic Polypeptide and Its Peptide Variant in Human Serum and Plasma by Immunocapture-Liquid-Chromatography-Tandem Mass Spectrometry. Reference Intervals and Practical Assay Considerations. <i>Biochemistry & Physiology</i> , 2014, 03, .	0.2	4
6	Expression of the Mouse MHC Class Ib H2-T11 Gene Product, a Paralog of H2-T23 (Qa-1) with Shared Peptide-Binding Specificity. <i>Journal of Immunology</i> , 2014, 193, 1427-1439.	0.8	10
7	ERAAP and Tapasin Independently Edit the Amino and Carboxyl Termini of MHC Class I Peptides. <i>Journal of Immunology</i> , 2013, 191, 1547-1555.	0.8	29
8	Regulation of HLA-DR peptide occupancy by histone deacetylase inhibitors. <i>Human Vaccines and Immunotherapeutics</i> , 2013, 9, 784-789.	3.3	13
9	Mice completely lacking immunoproteasomes show major changes in antigen presentation. <i>Nature Immunology</i> , 2012, 13, 129-135.	14.5	222
10	Utility of Characteristic QTOF MS/MS Fragmentation for MHC Class I Peptides. <i>Journal of Proteome Research</i> , 2011, 10, 2494-2507.	3.7	18
11	Endoplasmic Reticulum Aminopeptidase Associated with Antigen Processing Defines the Composition and Structure of MHC Class I Peptide Repertoire in Normal and Virus-Infected Cells. <i>Journal of Immunology</i> , 2010, 184, 3033-3042.	0.8	79
12	Identification of naturally processed ligands in the C57BL/6 mouse using large-scale mass spectrometric peptide sequencing and bioinformatics prediction. <i>Immunogenetics</i> , 2009, 61, 241-246.	2.4	12
13	Large Scale Mass Spectrometric Profiling of Peptides Eluted from HLA Molecules Reveals N-Terminal-Extended Peptide Motifs. <i>Journal of Immunology</i> , 2008, 181, 4874-4882.	0.8	36
14	Large Scale Mass Spectrometric Sequencing of Peptides Eluted from HLA-B35 Molecules Revealed Broad and N-Terminal Extended Peptide Motifs. <i>FASEB Journal</i> , 2008, 22, 423-423.	0.5	0