

Tracey C Fleischer

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

1,957
citations

840776

11
h-index

940533

16
g-index

18
all docs

18
docs citations

18
times ranked

2591
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone Deacetylase Activity Is Required for Full Transcriptional Repression by mSin3A. <i>Cell</i> , 1997, 89, 341-347.	28.9	705
2	A role for histone deacetylase activity in HDAC1-mediated transcriptional repression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 3519-3524.	7.1	350
3	Systematic identification and functional screens of uncharacterized proteins associated with eukaryotic ribosomal complexes. <i>Genes and Development</i> , 2006, 20, 1294-1307.	5.9	248
4	Identification of a <i>Drosophila</i> Myb-E2F2/RBF transcriptional repressor complex. <i>Genes and Development</i> , 2004, 18, 2929-2940.	5.9	233
5	Identification and Characterization of Three New Components of the mSin3A Corepressor Complex. <i>Molecular and Cellular Biology</i> , 2003, 23, 3456-3467.	2.3	164
6	Development and validation of a spontaneous preterm delivery predictor in asymptomatic women. <i>American Journal of Obstetrics and Gynecology</i> , 2016, 214, 633.e1-633.e24.	1.3	88
7	Chemical Proteomics Identifies Nampt as the Target of CB30865, An Orphan Cytotoxic Compound. <i>Chemistry and Biology</i> , 2010, 17, 659-664.	6.0	52
8	Purifying protein complexes for mass spectrometry: applications to protein translation. <i>Methods</i> , 2005, 35, 274-290.	3.8	30
9	Performance of a proteomic preterm delivery predictor in a large independent prospective cohort. <i>American Journal of Obstetrics & Gynecology MFM</i> , 2020, 2, 100140.	2.6	27
10	Analogues of 4-[(7-Bromo-2-methyl-4-oxo-3 <i>H</i> -quinazolin-6-yl)methylprop-2-ynylamino]- <i>N</i> -(3-pyridylmethyl)benzamide (CB-30865) as Potent Inhibitors of Nicotinamide Phosphoribosyltransferase (Nampt). <i>Journal of Medicinal Chemistry</i> , 2010, 53, 8734-8746.	6.4	25
11	Analytical validation of protein biomarkers for risk of spontaneous preterm birth. <i>Clinical Mass Spectrometry</i> , 2017, 3, 25-38.	1.9	17
12	Clinical Validation of a Proteomic Biomarker Threshold for Increased Risk of Spontaneous Preterm Birth and Associated Clinical Outcomes: A Replication Study. <i>Journal of Clinical Medicine</i> , 2021, 10, 5088.	2.4	6
13	337: Use of a second trimester serum-based proteomic risk classifier for prediction of spontaneous and medically-indicated preterm birth. <i>American Journal of Obstetrics and Gynecology</i> , 2017, 216, S204-S205.	1.3	3
14	Better Estimation of Spontaneous Preterm Birth Prediction Performance through Improved Gestational Age Dating. <i>Journal of Clinical Medicine</i> , 2022, 11, 2885.	2.4	3
15	Targeted Identification of Protein Interactions in Eukaryotic mRNA Translation. <i>Proteomics</i> , 2020, 20, 1900177.	2.2	2
16	Performance of a validated spontaneous preterm delivery predictor in South Asian and Sub-Saharan African women: a nested case control study. <i>Journal of Maternal-Fetal and Neonatal Medicine</i> , 2022, 35, 8878-8886.	1.5	2