## Tommaso De Marchi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparative Proteome Analysis Revealing an 11-Protein Signature for Aggressive Triple-Negative Breast Cancer. Journal of the National Cancer Institute, 2014, 106, djt376.	6.3	77
2	Quantitative Proteomic Analysis of Microdissected Breast Cancer Tissues: Comparison of Label-Free and SILAC-based Quantification with Shotgun, Directed, and Targeted MS Approaches. Journal of Proteome Research, 2013, 12, 4627-4641.	3.7	54
3	Endocrine therapy resistance in estrogen receptor (ER)-positive breast cancer. Drug Discovery Today, 2016, 21, 1181-1188.	6.4	53
4	Ferritin Heavy Chain in Triple Negative Breast Cancer: A Favorable Prognostic Marker that Relates to a Cluster of Differentiation 8 Positive (CD8+) Effector T-cell Response. Molecular and Cellular Proteomics, 2014, 13, 1814-1827.	3.8	44
5	The advantage of laserâ€capture microdissection over whole tissue analysis in proteomic profiling studies. Proteomics, 2016, 16, 1474-1485.	2.2	38
6	Inhibition of Histone Demethylases LSD1 and UTX Regulates ERα Signaling in Breast Cancer. Cancers, 2019, 11, 2027.	3.7	34
7	Phosphoserine aminotransferase 1 is associated to poor outcome on tamoxifen therapy in recurrent breast cancer. Scientific Reports, 2017, 7, 2099.	3.3	33
8	4â€protein signature predicting tamoxifen treatment outcome in recurrent breast cancer. Molecular Oncology, 2016, 10, 24-39.	4.6	31
9	Annexin-A1 and caldesmon are associated with resistance to tamoxifen in estrogen receptor positive recurrent breast cancer. Oncotarget, 2016, 7, 3098-3110.	1.8	26
10	Targeted MS Assay Predicting Tamoxifen Resistance in Estrogen-Receptor-Positive Breast Cancer Tissues and Sera. Journal of Proteome Research, 2016, 15, 1230-1242.	3.7	21
11	Prognostic significance of nuclear expression of UMP-CMP kinase in triple negative breast cancer patients. Scientific Reports, 2016, 6, 32027.	3.3	19
12	Proteogenomic Workflow Reveals Molecular Phenotypes Related to Breast Cancer Mammographic Appearance. Journal of Proteome Research, 2021, 20, 2983-3001.	3.7	14
13	Regulatory Interplay between miR-181a-5p and Estrogen Receptor Signaling Cascade in Breast Cancer. Cancers, 2021, 13, 543.	3.7	10
14	Antibody-Based Capture of Target Peptides in Multiple Reaction Monitoring Experiments. Methods in Molecular Biology, 2015, 1293, 123-135.	0.9	6
15	A somatic mutation in moesin drives progression into acute myeloid leukemia. Science Advances, 2022, 8, eabm9987.	10.3	2
16	Global proteomic characterization of microdissected estrogen receptor positive breast tumors. Data in Brief, 2015, 5, 399-402.	1.0	1