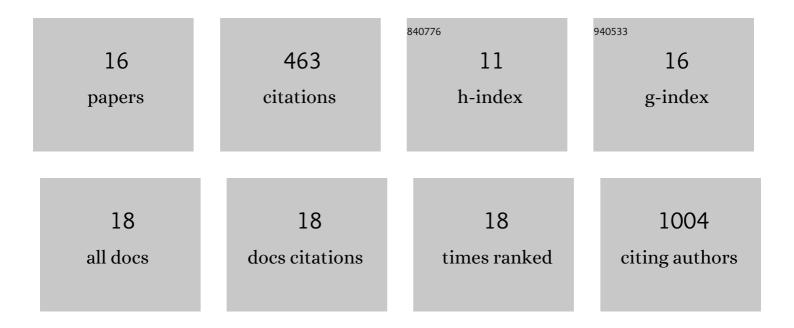
Tommaso De Marchi

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | A somatic mutation in moesin drives progression into acute myeloid leukemia. Science Advances, 2022, 8, eabm9987. | 10.3 | 2 |
| 2 | Regulatory Interplay between miR-181a-5p and Estrogen Receptor Signaling Cascade in Breast Cancer. Cancers, 2021, 13, 543. | 3.7 | 10 |
| 3 | Proteogenomic Workflow Reveals Molecular Phenotypes Related to Breast Cancer Mammographic Appearance. Journal of Proteome Research, 2021, 20, 2983-3001. | 3.7 | 14 |
| 4 | Inhibition of Histone Demethylases LSD1 and UTX Regulates ERα Signaling in Breast Cancer. Cancers, 2019, 11, 2027. | 3.7 | 34 |
| 5 | Phosphoserine aminotransferase 1 is associated to poor outcome on tamoxifen therapy in recurrent breast cancer. Scientific Reports, 2017, 7, 2099. | 3.3 | 33 |
| 6 | The advantage of laserâ€capture microdissection over whole tissue analysis in proteomic profiling studies. Proteomics, 2016, 16, 1474-1485. | 2.2 | 38 |
| 7 | Endocrine therapy resistance in estrogen receptor (ER)-positive breast cancer. Drug Discovery Today, 2016, 21, 1181-1188. | 6.4 | 53 |
| 8 | Prognostic significance of nuclear expression of UMP-CMP kinase in triple negative breast cancer patients. Scientific Reports, 2016, 6, 32027. | 3.3 | 19 |
| 9 | 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 |
| 10 | 4â€protein signature predicting tamoxifen treatment outcome in recurrent breast cancer. Molecular Oncology, 2016, 10, 24-39. | 4.6 | 31 |
| 11 | 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 |
| 12 | Global proteomic characterization of microdissected estrogen receptor positive breast tumors. Data in Brief, 2015, 5, 399-402. | 1.0 | 1 |
| 13 | Antibody-Based Capture of Target Peptides in Multiple Reaction Monitoring Experiments. Methods in Molecular Biology, 2015, 1293, 123-135. | 0.9 | 6 |
| 14 | 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 |
| 15 | 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 |
| 16 | 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 |