

Tesa M Severson

List of Publications by Year in descending order

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31
papers

5,235
citations

331670

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docs citations

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citing authors

#	ARTICLE	IF	CITATIONS
1	Somatic mutations altering EZH2 (Tyr641) in follicular and diffuse large B-cell lymphomas of germinal-center origin. <i>Nature Genetics</i> , 2010, 42, 181-185.	21.4	1,504
2	Frequent mutation of histone-modifying genes in non-Hodgkin lymphoma. <i>Nature</i> , 2011, 476, 298-303.	27.8	1,428
3	Mutational evolution in a lobular breast tumour profiled at single nucleotide resolution. <i>Nature</i> , 2009, 461, 809-813.	27.8	984
4	Prostate cancer reactivates developmental epigenomic programs during metastatic progression. <i>Nature Genetics</i> , 2020, 52, 790-799.	21.4	174
5	Acquired <i>TNFRSF14</i> Mutations in Follicular Lymphoma Are Associated with Worse Prognosis. <i>Cancer Research</i> , 2010, 70, 9166-9174.	0.9	160
6	Integration of genomic, transcriptomic and proteomic data identifies two biologically distinct subtypes of invasive lobular breast cancer. <i>Scientific Reports</i> , 2016, 6, 18517.	3.3	143
7	Evidence for Archaic Asian Ancestry on the Human X Chromosome. <i>Molecular Biology and Evolution</i> , 2005, 22, 189-192.	8.9	81
8	Assessment of PD-L1 expression across breast cancer molecular subtypes, in relation to mutation rate, <i>BRCA1</i> -like status, tumor-infiltrating immune cells and survival. <i>Oncolmmunology</i> , 2018, 7, e1509820.	4.6	80
9	A novel DNA sequence database for analyzing human demographic history. <i>Genome Research</i> , 2008, 18, 1354-1361.	5.5	74
10	ARID1A mutation sensitizes most ovarian clear cell carcinomas to BET inhibitors. <i>Oncogene</i> , 2018, 37, 4611-4625.	5.9	72
11	Hypomorphic Temperature-Sensitive Alleles of NSDHL Cause CK Syndrome. <i>American Journal of Human Genetics</i> , 2010, 87, 905-914.	6.2	64
12	DNA repair deficiency biomarkers and the 70-gene ultra-high risk signature as predictors of veliparib/carboplatin response in the I-SPY 2 breast cancer trial. <i>Npj Breast Cancer</i> , 2017, 3, 31.	5.2	64
13	The <i>BRCA1</i> ness signature is associated significantly with response to PARP inhibitor treatment versus control in the I-SPY 2 randomized neoadjuvant setting. <i>Breast Cancer Research</i> , 2017, 19, 99.	5.0	58
14	<i>BRCA1</i> -like signature in triple negative breast cancer: Molecular and clinical characterization reveals subgroups with therapeutic potential. <i>Molecular Oncology</i> , 2015, 9, 1528-1538.	4.6	54
15	Lysine specific demethylase 1 inactivation enhances differentiation and promotes cytotoxic response when combined with all- <i>trans</i> retinoic acid in acute myeloid leukemia across subtypes. <i>Haematologica</i> , 2019, 104, 1156-1167.	3.5	50
16	Fanconi anemia and homologous recombination gene variants are associated with functional DNA repair defects <i>in vitro</i> and poor outcome in patients with advanced head and neck squamous cell carcinoma. <i>Oncotarget</i> , 2018, 9, 18198-18213.	1.8	37
17	EZH2 Is Overexpressed in <i>BRCA1</i> -like Breast Tumors and Predictive for Sensitivity to High-Dose Platinum-Based Chemotherapy. <i>Clinical Cancer Research</i> , 2019, 25, 4351-4362.	7.0	33
18	Comparative Cistromics Reveals Genomic Cross-talk between FOXA1 and ER α in Tamoxifen-Associated Endometrial Carcinomas. <i>Cancer Research</i> , 2016, 76, 3773-3784.	0.9	30

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19	Robust BRCA1-like classification of copy number profiles of samples repeated across different datasets and platforms. <i>Molecular Oncology</i> , 2015, 9, 1274-1286.	4.6	29
20	A review of estrogen receptor/androgen receptor genomics in male breast cancer. <i>Endocrine-Related Cancer</i> , 2017, 24, R27-R34.	3.1	23
21	Expression profile and genome location of cDNA clones from an infant human trabecular meshwork cell library. <i>Investigative Ophthalmology and Visual Science</i> , 2002, 43, 3698-704.	3.3	22
22	Drug-Induced Epigenomic Plasticity Reprograms Circadian Rhythm Regulation to Drive Prostate Cancer toward Androgen Independence. <i>Cancer Discovery</i> , 2022, 12, 2074-2097.	9.4	22
23	The MSP-ERON axis stimulates cancer cell growth in models of triple negative breast cancer. <i>Molecular Oncology</i> , 2020, 14, 1868-1880.	4.6	15
24	Neoadjuvant tamoxifen synchronizes ER \pm binding and gene expression profiles related to outcome and proliferation. <i>Oncotarget</i> , 2016, 7, 33901-33918.	1.8	13
25	Epigenetic and transcriptional analysis reveals a core transcriptional program conserved in clonal prostate cancer metastases. <i>Molecular Oncology</i> , 2021, 15, 1942-1955.	4.6	10
26	Mutations In MLL2 and MEF2B Genes In Follicular Lymphoma and Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2010, 116, 473-473.	1.4	6
27	Tyrosine 641 of the EZH2 Oncogene Is Frequently Mutated in Follicular and Diffuse Large B-Cell Lymphomas of Germinal Center Origin.. <i>Blood</i> , 2009, 114, 139-139.	1.4	3
28	Identification of Genes Frequently Mutated In FL and DLBCL with Transcriptome, Genome and Exome Sequencing. <i>Blood</i> , 2010, 116, 804-804.	1.4	1
29	TNFRSF14 Is Mutated in a Subset of Follicular Lymphoma and Correlated with Inferior Prognosis.. <i>Blood</i> , 2009, 114, 1919-1919.	1.4	1
30	Base-Pair Resolution of Somatic and Germline-Derived Genome Rearrangement Breakpoints in Follicular Lymphoma.. <i>Blood</i> , 2009, 114, 439-439.	1.4	0
31	FAS Mutations in Follicular Lymphoma Are Rare but Associated with Aggressive Clinical Behavior.. <i>Blood</i> , 2009, 114, 3967-3967.	1.4	0