

Kelly A Avery-Kiejda

List of Publications by Citations

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37
papers

1,092
citations

17
h-index

33
g-index

43
ext. papers

1,319
ext. citations

5.4
avg, IF

4.16
L-index

#	Paper	IF	Citations
37	Methylome sequencing in triple-negative breast cancer reveals distinct methylation clusters with prognostic value. <i>Nature Communications</i> , 2015 , 6, 5899	17.4	118
36	Up-regulation of Mcl-1 is critical for survival of human melanoma cells upon endoplasmic reticulum stress. <i>Cancer Research</i> , 2008 , 68, 6708-17	10.1	118
35	Small molecular weight variants of p53 are expressed in human melanoma cells and are induced by the DNA-damaging agent cisplatin. <i>Clinical Cancer Research</i> , 2008 , 14, 1659-68	12.9	92
34	BRIP1, PALB2, and RAD51C mutation analysis reveals their relative importance as genetic susceptibility factors for breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011 , 127, 853-9	4.4	87
33	Genetic insights into breast cancer risk. <i>Hereditary Cancer in Clinical Practice</i> , 2015 , 13,	2.3	78
32	P53 in human melanoma fails to regulate target genes associated with apoptosis and the cell cycle and may contribute to proliferation. <i>BMC Cancer</i> , 2011 , 11, 203	4.8	73
31	Glucose-regulated protein 78 antagonizes cisplatin and adriamycin in human melanoma cells. <i>Carcinogenesis</i> , 2009 , 30, 197-204	4.6	64
30	Decreased expression of key tumour suppressor microRNAs is associated with lymph node metastases in triple negative breast cancer. <i>BMC Cancer</i> , 2014 , 14, 51	4.8	62
29	Proteotranscriptomic Profiling of 231-BR Breast Cancer Cells: Identification of Potential Biomarkers and Therapeutic Targets for Brain Metastasis. <i>Molecular and Cellular Proteomics</i> , 2015 , 14, 2316-30	7.6	50
28	MiRNAs and Other Epigenetic Changes as Biomarkers in Triple Negative Breast Cancer. <i>International Journal of Molecular Sciences</i> , 2015 , 16, 28347-76	6.3	46
27	The relative mRNA expression of p53 isoforms in breast cancer is associated with clinical features and outcome. <i>Carcinogenesis</i> , 2014 , 35, 586-96	4.6	45
26	Novel genes associated with lymph node metastasis in triple negative breast cancer. <i>Scientific Reports</i> , 2015 , 5, 15832	4.9	31
25	The expression of Dicer and Drosha in matched normal tissues, tumours and lymph node metastases in triple negative breast cancer. <i>BMC Cancer</i> , 2014 , 14, 253	4.8	28
24	DNA methylation profile of triple negative breast cancer-specific genes comparing lymph node positive patients to lymph node negative patients. <i>Scientific Reports</i> , 2016 , 6, 33435	4.9	27
23	STaRRRT: a table of short tandem repeats in regulatory regions of the human genome. <i>BMC Genomics</i> , 2013 , 14, 795	4.5	24
22	Activation of Jun N-terminal kinase is a mediator of vincristine-induced apoptosis of melanoma cells. <i>Anti-Cancer Drugs</i> , 2008 , 19, 189-200	2.4	24
21	Nucleotide excision repair gene expression after Cisplatin treatment in melanoma. <i>Cancer Research</i> , 2010 , 70, 7918-26	10.1	20

20	Low prevalence of germline PALB2 mutations in Australian triple-negative breast cancer. <i>International Journal of Cancer</i> , 2014 , 134, 301-5	7.5	13
19	Regulation of the human placental (pro)renin receptor-prorenin-angiotensin system by microRNAs. <i>Molecular Human Reproduction</i> , 2018 , 24, 453-464	4.4	12
18	The Roles of Non-Coding RNAs in Tumor-Associated Lymphangiogenesis. <i>Cancers</i> , 2020 , 12,	6.6	11
17	Regulation of the interferon-gamma (IFN- γ) pathway by p63 and $\Delta 33p53$ isoform in different breast cancer subtypes. <i>Oncotarget</i> , 2018 , 9, 29146-29161	3.3	11
16	Molecular patterns of cancer colonisation in lymph nodes of breast cancer patients. <i>Breast Cancer Research</i> , 2018 , 20, 143	8.3	10
15	The presence of the intron 3 16 bp duplication polymorphism of p53 (rs17878362) in breast cancer is associated with a low $\Delta 0p53:p53$ ratio and better outcome. <i>Carcinogenesis</i> , 2016 , 37, 81-86	4.6	7
14	Regulators of global genome repair do not respond to DNA damaging therapy but correlate with survival in melanoma. <i>PLoS ONE</i> , 2013 , 8, e70424	3.7	7
13	Genome-wide miRNA, gene and methylation analysis of triple negative breast cancer to identify changes associated with lymph node metastases. <i>Genomics Data</i> , 2017 , 14, 1-4		6
12	Good Cop, Bad Cop: Defining the Roles of $\Delta 0p53$ in Cancer and Aging. <i>Cancers</i> , 2020 , 12,	6.6	5
11	Comparison of Three Different Methods for Determining Cell Proliferation in Breast Cancer Cell Lines. <i>Journal of Visualized Experiments</i> , 2016 ,	1.6	4
10	Intronic Polymorphisms Are Associated with Increased Transcript, Immune Infiltration and Cancer Risk. <i>Cancers</i> , 2020 , 12,	6.6	4
9	Copy number variation in triple-negative breast cancer samples associated with lymph node metastasis. <i>Neoplasia</i> , 2021 , 23, 743-753	6.4	4
8	Comparison of the QuantiGene 2.0 Assay and Real-Time RT-PCR in the Detection of p53 Isoform mRNA Expression in Formalin-Fixed Paraffin-Embedded Tissues- A Preliminary Study. <i>PLoS ONE</i> , 2016 , 11, e0165930	3.7	3
7	Crosstalk Between microRNAs and the Pathological Features of Secondary Lymphedema. <i>Frontiers in Cell and Developmental Biology</i> , 2021 , 9, 732415	5.7	1
6	Cross-Cultural Adaptation of the Functional Assessment of Cancer Therapy-Breast (FACT-B) in Malaysian Breast Cancer Survivors. <i>Asian Pacific Journal of Cancer Prevention</i> , 2021 , 22, 1055-1061	1.7	1
5	A novel polymorphic repeat in the upstream regulatory region of the estrogen-induced gene EIG121 is not associated with the risk of developing breast or endometrial cancer. <i>BMC Research Notes</i> , 2016 , 9, 287	2.3	1
4	A polymorphic repeat in the IGF1 promoter influences the risk of endometrial cancer. <i>Endocrine Connections</i> , 2016 , 5, 115-22	3.5	1
3	The intron 3 16bp duplication polymorphism of p53 (rs17878362) is not associated with increased risk of developing triple-negative breast cancer. <i>Breast Cancer Research and Treatment</i> , 2019 , 173, 727-733	4.4	1

2	Assessment of Potential Risk Factors and Skin Ultrasound Presentation Associated with Breast Cancer-Related Lymphedema in Long-Term Breast Cancer Survivors. <i>Diagnostics</i> , 2021 , 11,	3.8	1
1	Verification and Validation of a Four-Gene Panel as a Prognostic Indicator in Triple Negative Breast Cancer.. <i>Frontiers in Oncology</i> , 2022 , 12, 821334	5.3	