## Elena LÃ<sup>3</sup>pez-Knowles

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Analysis of <i>ESR1</i> mutation in circulating tumor DNA demonstrates evolution during therapy for metastatic breast cancer. Science Translational Medicine, 2015, 7, 313ra182.	5.8	460
2	Prediction of late distant recurrence in patients with oestrogen-receptor-positive breast cancer: a prospective comparison of the breast-cancer index (BCI) assay, 21-gene recurrence score, and IHC4 in the TransATAC study population. Lancet Oncology, The, 2013, 14, 1067-1076.	5.1	332
3	Prediction of Late Distant Recurrence After 5 Years of Endocrine Treatment: A Combined Analysis of Patients From the Austrian Breast and Colorectal Cancer Study Group 8 and Arimidex, Tamoxifen Alone or in Combination Randomized Trials Using the PAM50 Risk of Recurrence Score. Journal of Clinical Oncology. 2015. 33. 916-922.	0.8	189
4	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386.	9.4	125
5	Tumor <i>PIK3CA</i> Genotype and Prognosis in Early-Stage Breast Cancer: A Pooled Analysis of Individual Patient Data. Journal of Clinical Oncology, 2018, 36, 981-990.	0.8	95
6	Inactivating <i>NF1</i> Mutations Are Enriched in Advanced Breast Cancer and Contribute to Endocrine Therapy Resistance. Clinical Cancer Research, 2020, 26, 608-622.	3.2	71
7	Impact of a Panel of 88 Single Nucleotide Polymorphisms on the Risk of Breast Cancer in High-Risk Women: Results From Two Randomized Tamoxifen Prevention Trials. Journal of Clinical Oncology, 2017, 35, 743-750.	0.8	58
8	Estrogen Receptor Expression in 21-Gene Recurrence Score Predicts Increased Late Recurrence for Estrogen-Positive/HER2-Negative Breast Cancer. Clinical Cancer Research, 2015, 21, 2763-2770.	3.2	36
9	Impact of mutational profiles on response of primary oestrogen receptor-positive breast cancers to oestrogen deprivation. Nature Communications, 2016, 7, 13294.	5.8	34
10	Heterogeneity in global gene expression profiles between biopsy specimens taken peri-surgically from primary ER-positive breast carcinomas. Breast Cancer Research, 2016, 18, 39.	2.2	24
11	Impact of aromatase inhibitor treatment on global gene expression and its association with antiproliferative response in ER+ breast cancer in postmenopausal patients. Breast Cancer Research, 2020, 22, 2.	2.2	15
12	Molecular characterisation of aromatase inhibitor-resistant advanced breast cancer: the phenotypic effect of ESR1 mutations. British Journal of Cancer, 2019, 120, 247-255.	2.9	13
13	Major Impact of Sampling Methodology on Gene Expression in Estrogen Receptor–Positive Breast Cancer. JNCI Cancer Spectrum, 2018, 2, pky005.	1.4	11
14	Integrative analyses identify modulators of response to neoadjuvant aromatase inhibitors in patients with early breast cancer. Breast Cancer Research, 2015, 17, 35.	2.2	8
15	Impact of Duration of Neoadjuvant Aromatase Inhibitors on Molecular Expression Profiles in Estrogen Receptor–positive Breast Cancers. Clinical Cancer Research, 2022, 28, 1217-1228.	3.2	6