Federico Rotolo

List of Publications by Year in descending order

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840776 580821 25 751 11 25 citations h-index g-index papers 30 30 30 1608 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Tumor Mutation Burden as a Biomarker in Resected Non–Small-Cell Lung Cancer. Journal of Clinical Oncology, 2018, 36, 2995-3006.	1.6	223
2	IPH4102, a first-in-class anti-KIR3DL2 monoclonal antibody, in patients with relapsed or refractory cutaneous T-cell lymphoma: an international, first-in-human, open-label, phase 1 trial. Lancet Oncology, The, 2019, 20, 1160-1170.	10.7	119
3	Empirical extensions of the lasso penalty to reduce the false discovery rate in highâ€dimensional Cox regression models. Statistics in Medicine, 2016, 35, 2561-2573.	1.6	74
4	parfm : Parametric Frailty Models in <i>R</i> . Journal of Statistical Software, 2012, 51, .	3.7	48
5	Surrogate End Points for Overall Survival in Loco-Regionally Advanced Nasopharyngeal Carcinoma: An Individual Patient Data Meta-analysis. Journal of the National Cancer Institute, 2017, 109, .	6.3	37
6	Statistical controversies in clinical research: prognostic gene signatures are not (yet) useful in clinical practice. Annals of Oncology, 2016, 27, 2160-2167.	1.2	33
7	surrosurv: An R package for the evaluation of failure time surrogate endpoints in individual patient data meta-analyses of randomized clinical trials. Computer Methods and Programs in Biomedicine, 2018, 155, 189-198.	4.7	29
8	Bias and precision of methods for estimating the difference in restricted mean survival time from an individual patient data meta-analysis. BMC Medical Research Methodology, 2016, 16, 37.	3.1	28
9	Adjuvant cisplatin-based chemotherapy in nonsmall-cell lung cancer: new insights into the effect on failure type via a multistate approach. Annals of Oncology, 2014, 25, 2162-2166.	1.2	26
10	Identification of biomarkerâ€byâ€treatment interactions in randomized clinical trials with survival outcomes and highâ€dimensional spaces. Biometrical Journal, 2017, 59, 685-701.	1.0	26
11	biospear: an R package for biomarker selection in penalized Cox regression. Bioinformatics, 2018, 34, 112-113.	4.1	16
12	A simulation procedure based on copulas to generate clustered multi-state survival data. Computer Methods and Programs in Biomedicine, 2013, 109, 305-312.	4.7	13
13	A Poisson approach to the validation of failure time surrogate endpoints in individual patient data meta-analyses. Statistical Methods in Medical Research, 2019, 28, 170-183.	1.5	12
14	Genome-wide copy number analyses of samples from LACE-Bio project identify novel prognostic and predictive markers in early stage non-small cell lung cancer. Translational Lung Cancer Research, 2018, 7, 416-427.	2.8	11
15	Predictive classifier for intensive treatment of head and neck cancer. Cancer, 2020, 126, 5263-5273.	4.1	11
16	Accounting for grouped predictor variables or pathways in high-dimensional penalized Cox regression models. BMC Bioinformatics, 2020, 21, 277.	2.6	11
17	Robust estimation of the expected survival probabilities from high-dimensional Cox models with biomarker-by-treatment interactions in randomized clinical trials. BMC Medical Research Methodology, 2017, 17, 83.	3.1	10
18	Evaluation of Treatment Effect with Paired Failure Times in a Single-Arm Phase II Trial in Oncology. Computational and Mathematical Methods in Medicine, 2018, 2018, 1-8.	1.3	8

#	Article	IF	Citations
19	Testing the treatment effect on competing causes of death in oncology clinical trials. BMC Medical Research Methodology, 2014, 14, 72.	3.1	7
20	First-line treatment in metastatic colorectal cancer: ImportantÂor crucial?. European Journal of Cancer, 2017, 84, 363-366.	2.8	4
21	Incorporation of nested frailties into semiparametric multiâ€state models. Statistics in Medicine, 2016, 35, 609-621.	1.6	1
22	Predictor of effectiveness of treatment intensification on overall survival in head and neck cancer (HNC). Annals of Oncology, 2018, 29, viii375-viii376.	1.2	1
23	An alternative trial-level measure for evaluating failure-time surrogate endpoints based on prediction error. Contemporary Clinical Trials Communications, 2019, 15, 100402.	1.1	1
24	Prediction of treatment benefit in high-dimensional cox models via gene signatures in randomized clinical trials. Trials, 2015, 16, .	1.6	0
25	Development and Validation of Genomic Signatures. , 2018, , .		0