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Tumor cell sensitivity to vemurafenib can be predicted from protein expression in a BRAF-V600E basket trial setting

DOI: 10.1186/s12885-019-6175-2 BMC Cancer, 2019, 19, 1025.

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#	Paper	IF	Citations
7	The Evolution of Master Protocol Clinical Trial Designs: A Systematic Literature Review. <i>Clinical Therapeutics</i> , 2020 , 42, 1330-1360	3.5	31
6	METTL3 induces PLX4032 resistance in melanoma by promoting mA-dependent EGFR translation. <i>Cancer Letters</i> , 2021 , 522, 44-56	9.9	3
5	Analysis of alternative mRNA splicing in vemurafenib-resistant melanoma cells.		1
4	Analysis of Alternative mRNA Splicing in Vemurafenib-Resistant Melanoma Cells. <i>Biomolecules</i> , 2022 , 12, 993	5.9	
3	A Subset of Secreted Proteins in Ascites Can Predict Platinum-Free Interval in Ovarian Cancer. 2022 , 14, 4291		O
2	AI + precision medicine. 2023 , 69-98		О
1	MUC1-C is necessary for SHP2 activation and BRAF inhibitor resistance in BRAF(V600E) mutant colorectal cancer. 2023 , 559, 216116		Ο