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When to test for BRAF and what are the consequences?

by Dr. Chiara Cremolini, Dr. Armin Gerger and Dr. Guillem Argilés

WHEN TO TEST FOR BRAF AND WHAT ARE THE CONSEQUENCES?

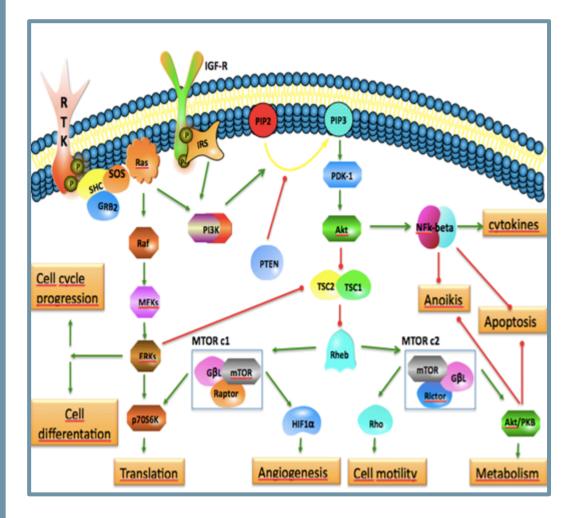
BY

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BRAF MUTATION, GENERAL CONSIDERATIONS



- BRAF V600E is found in approx. 5-10% of mCRC
 - 4% of non-hypermutat CRC
 - 46% of hypermutant CRC
- Driver mutation
- Considered mutually exclusive with RAS mutations

(though ultra-sensitive NGS platform unveiled concomitancy with minor RAS mut. allele fractions in certain tumors)

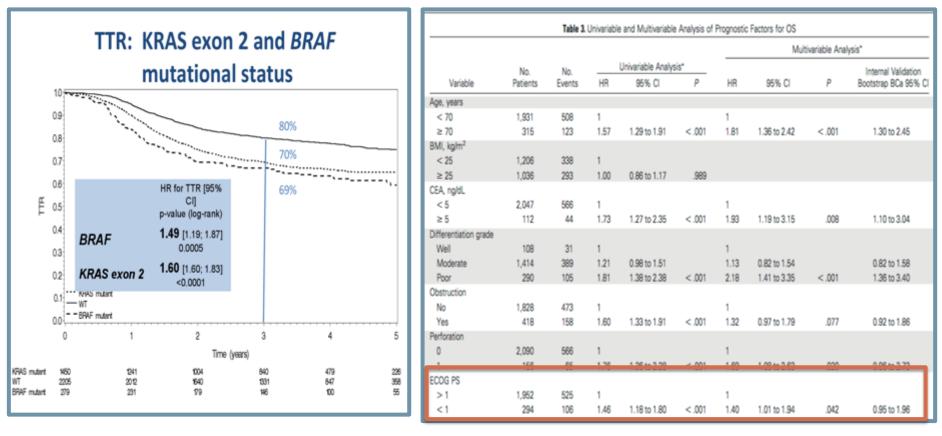


ROLE OF BRAF V600E IN LOCALIZED CRC

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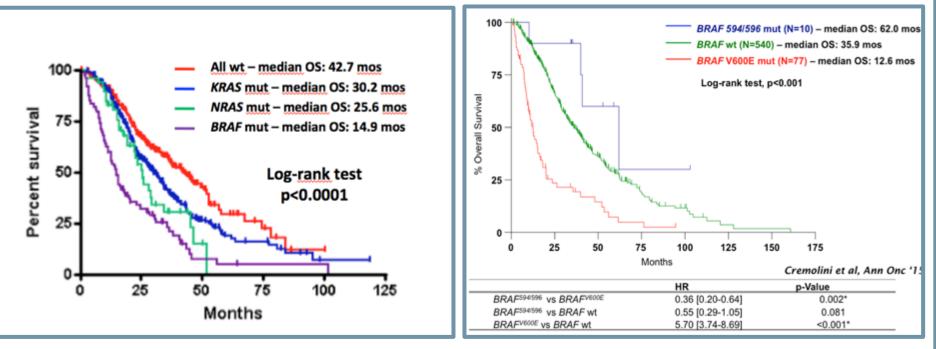


BRAF testing on limited disease not recommended, prognostic and predictive data are pending to be clarified (disparate results in different trials)

PROGNOSTIC ROLE OF BRAF MUTATION IN MCRC

BRAF mutations confer bad prognosis

Different biology of BRAF mutations

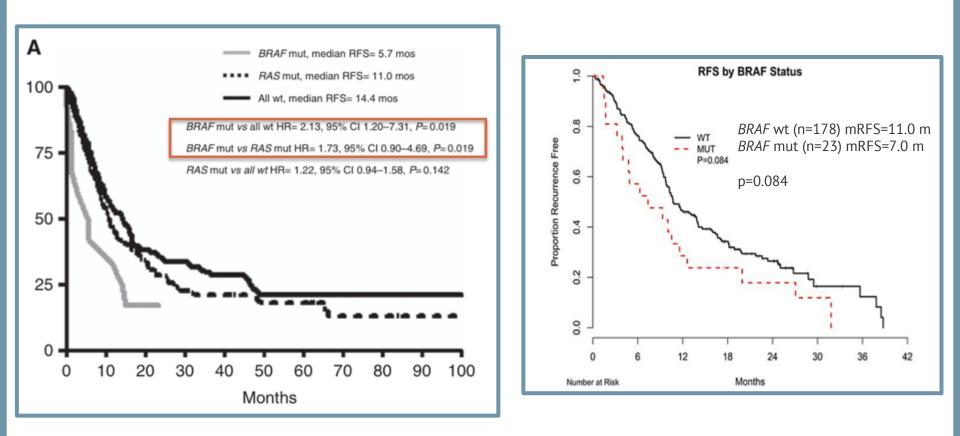


BRAF V600E mutation confers bad prognosis in the metastatic setting



Schirripa et al, Int J Canc 2005, Cremolini et al, Ann Onc 2015

PROGNOSTIC ROLE OF BRAF MUT. IN mCRC (2)



The trend towards bad prognosis still persist in BRAF V600E metastatic patients undergoing resection



Schirripa et al, BJC 2015 Yaeger et al, Cancer 2014

PREDICTIVE ROLE OF BRAF MUTATION IN MCRC

Role of BRAF Mut anti-EGFR predictive factor

Subgroup Sample size of Tx groups PFS hazard ratio [95% CI] study Cmab/pmab Comparator RAS WT / BRAF WT PRIME 228 218 0.68 [0.54 . 0.87] CRYSTAL and OPUS 0.64 [0.52 , 0.79] 349 381 CO.17 101 97 0.41 [0.30 , 0.55] 20020408 63 52 0.37 [0.24 , 0.55] PICCOLO 183 188 0.71 [0.57 . 0.88] 20050181 186 190 0.68 [0.51 , 0.90] COIN 292 289 0.93 [0.78 , 1.10] 1402 1415 0.62 [0.50 , 0.77] Summary: Test for effect: P < 0.001 Heterogeneity: /2 = 82%, P < 0.001 RAS WT / BRAF WT PRIME 24 29 0.58 [0.29 . 1.15] CRYSTAL and OPUS 32 38 0.67 [0.34 , 1.29] 0.76 [0.19 . 3.08] CO.17 4 6 0.34 [0.09 , 1.24] 20020408 9 6 PICCOLO 37 1.40 [0.82 , 2.39] 31 0.69 [0.32 , 1.49] 20050181 22 23 40 50 1.25 [0.81 , 1.94] COIN 0.86[0.61.1.21] Summary: 168 183 Test for effect: P = 0.38 Heterogeneity: /2 = 39%, P = 0.13 0.2 0.5 1 2 HR (log scale) Favours Cmab/Pmab Favours compara

				Hazard Ratio	Hazard Ratio
Study or Subgroup	log[Hazard Ratio]	SE	Weight	IV, Random, 95% Cl	I Year IV, Random, 95% CI
Bokemeyer 2012	-0.4	0.34	12.6%	0.67 [0.34, 1.31]	2012
Peeters 2013	-1.079	0.669	3.8%	0.34 [0.09, 1.26]	2013
Seymour 2013	0.336	0.273	17.5%	1.40 [0.82, 2.39]	2013
Douillard 2013	-0.545	0.351	12.0%	0.58 [0.29, 1.15]	2013
Smith 2013	0.131	0.207	25.1%	1.14 [0.76, 1.71]	2013
Karapetis 2013	-0.274	0.711	3.4%	0.76 [0.19, 3.06]	2013
Peeters 2014	-0.371	0.392	10.0%	0.69 [0.32, 1.49]	2014
Stintzing 2014	-0.139	0.297	15.5%	0.87 [0.49, 1.56]	2014
Total (95% CI)			100.0%	0.88 [0.67, 1.14]	•
Heterogeneity: Tau ² = 0.03; Chi ² = 8.88, df = 7 (P = 0.26); l ² = 21%					
Test for overall effect: Z = 0.98 (P = 0.33)				Favours anti-EGFR MoAbs Favours control	

TRIBE: subgroup analysis according to RAS/BRAF status

	N	FOLFIRI + bev Median OS	FOLFOXIRI + bev Median OS	HR [95% CI]	р
ITT population	508	25.8	29.8	0.80 [0.65-0.98]	0.030
RAS and BRAF evaluable	357	24.9	28.6	0.84 [0.66-1.07]	0.159
RAS and BRAF wt	93	33.5	41.7	0.77 [0.46-1.27]	
RAS mutated	236	23.9	27.3	0.88 [0.65-1.18]	0.522*
BRAF mutated	28	10.7	19.0	0.54 [0.24-1.20]	

* P for interaction

BRAF V600E diminish benefit derived from anti-EGFR MoAbs

However intensive strategies using extended cytostatic combinations seem to improve patient outcomes



FIRST GENERATION OF BRAF THERAPEUTICS

Dabrafenib (D) + Trametinib (T): Vemurafenib Monotherapy: Limited Activity in **BRAFm** CRC Not Effective in BRAFm CRC 5% Response Rate 3.1 m. PFS TRAME Confirmed Response Rate (CR + PR): 7% Median Progression-Free Survival: 3.5 months Kopetz et al, J Clin Oncol, 2010. BRF113220 - Corcoran et al, J Clin Oncol, 2014

Contrary to melanoma, initial trials with BRAF V600E inhibitors s/a failed to demonstrate clinical activity in mCRC



BRAF INHIBITORS + EGFR INHIBITORS HAVE IN VIVO ACTIVITY IN BRAF V600E MUTATED CRC XENOGRAFTS

LETTER

Unresponsiveness of colon cancer to BRAF(V600E) inhibition through feedback activation of EGFR

Anirudh Prahallad1+, Chong Sun1+, Sidong Huang1+, Federica Di Nicolantonio2.1+, Ramon Salazar4, Davide Zecchin2, Roderick L. Beijersbergen¹, Alberto Bardelli^{2,3} & René Bernards¹

drug PLX4032 (vemurafenib) is highly effective in the treatment of melanoma¹. However, colon cancer patients harbouring the same BRAF(V600E) oncogenic lesion have poor prognosis and show only a very limited response to this drug²⁻⁴. To investigate the cause of the limited therapeutic effect of PLX4032 in BRAF(V600E) mutant colon tumours, here we performed an RNA-interferencebased genetic screen in human cells to search for kinases whose knockdown synergizes with BRAF(V600E) inhibition. We report that blockade of the epidermal growth factor receptor (EGFR) shows strong synergy with BRAF(V600E) inhibition. We find in multiple BRAF(V800E) mutant colon cancers that inhibition of EGFR by the antibody drug cetuximab or the small-molecule drugs gefitinib or erlotinib is strongly synergistic with BRAF(V600E) inhibition, both in vitro and in vivo. Mechanistically, we find that BRAF(V600E) inhibition causes a rapid feedback activation of EGFR and are therefore not subject to this feedback activation. Consistent with this, we find that ectopic expression of EGFR in melanoma cells is sufficient to cause resistance to PLX4032. Our data suggest that BRAF(V600E) mutant colon cancers (approximately 8-10% of all colon cancers123), for which there are currently no targeted treatment options available, might benefit from combination therapy consisting of BRAF and EGFR inhibitors.

Activating mutations in the BRAF oncogene (BRAF(V600E)) are seen in some 70% of primary melanomas*, some 10% of colorectal cancers' and some 30-70% of papillary thyroid carcinoma*-10. However, clinical responses to the highly selective small-molecule inhibitor of the BRAF(V600E) oncoprotein, PLX4032, differ widely, ranging from a response rate of approximately 80% in melanoma to DOM: N IN

Inhibition of the BRAF(V600E) oncoprotein by the small-molecule determined by next generation sequencing of the barcode identifiers present in each shRNA vector (Fig. 1c; see Methods). We arbitrarily considered only shRNA vectors that had been sequenced at least 300 times and which were depleted at least fivefold by the drug treatment. Figure 1d shows that only very few of the 3,388 shRNA vectors in the library met this stringent selection criterion, among which were three independent shRNA vectors targeting the EGFR (see Supplementary Table 2 for all selected shRNAs). This suggested that suppression of EGFR synergizes with BRAF inhibition in these CRC cells. To validate this finding, we infected WiDr cells with each of these three EGFR shRNA vectors (all of which reduced EGFR levels; Fig. 1f) and cultured these cells with or without PLX4032 for 2 weeks. Figure 1e shows that inhibition of EGFR does not significantly affect proliferation of EGFR in WiDr cells, consistent with the clinical observations that KRAS or BRAF mutant CRC cells do not respond to EGFR-targeted monoclonal antibodies73534. In contrast, suppression of EGFR in combination with EGFR, which supports continued proliferation in the presence of PLX4032 caused a marked inhibition of proliferation in WiDr cells BRAF(V600E) inhibition. Melanoma cells express low levels of (Fig. 1e). This suggested that BRAF(V600E) mutant CRC cells are responsive to treatment with a combination of BRAF inhibitor plus an EGFR inhibitor.

doi:10.1038/nature10868

At present, two classes of anti-EGFR drugs are clinically available; these include the monoclonal antibodies cetuximab and panitumumab, and the small-molecule kinase inhibitors gefitinib and erlotinib. We found that three BRAF mutant CRC cell lines (WiDr, VACO432 and KM20) all lack a significant response to monotherapy with PLX4032, cetuximab or gefitinib. However, strong synergy was seen when PLX4032 was combined with either cetuximab or gelitinib (Fig. 2a and Supplementary Fig. 1A, C) or erlotinib (data not shown), consistent with the notion derived from the shRNA screen that EGFR inhibition is required to elicit a response to BRAF inhibition in CRC cells.

To address the molecular mechanism underlying the synergy BOAR of BOARD LAND

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000 K 20000000000 EGFR-mediated re-activation of MAPK signaling contributes to insensitivity of BRAF mutant colorectal cancers to RAF inhibition with vemurafenib

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Abstract

NIH-PA AL

BRAF mutations occur in 10-15% of colorectal cancers (CRCs) and confer adverse outcome While RAF inhibitors such as vemurafenib (PLX4032) have proven effective in BRAF mutant. melanoma, they are surprisingly ineffective in BRAF mutant CRCs, and the reason for this disparity remains unclear. Compared to BRAF mutant melanoma cells, BRAF mutant CRC cells were less sensitive to vemurafenib, and P-ERK suppression was not sustained in response to treatment. Although transient inhibition of phospho-ERK by vemurafenib was observed in CRC, rapid ERK re-activation occurred through EGFR-mediated activation of RAS and CRAF. BRAF mutant CRCs expressed higher levels of phospho-EGFR than BRAF mutant melanomas,

Ouput signail

MEKs

ERKs

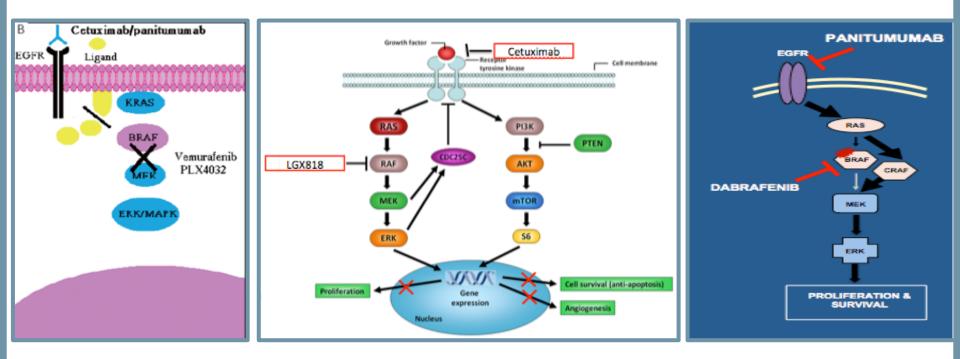
Ras

GRA:

The genetic context is different in CRC. EGFR constitutive expression lead to a feedback crosstalk with BRAF downstream effectors that functionally rescue BRAF inhibition

connect Corcoran RB et al, Cancer Discov 2012; 2:227-35, Prahallad A et al, Nature 2012; 483:100-3 POWERED BY CORRE

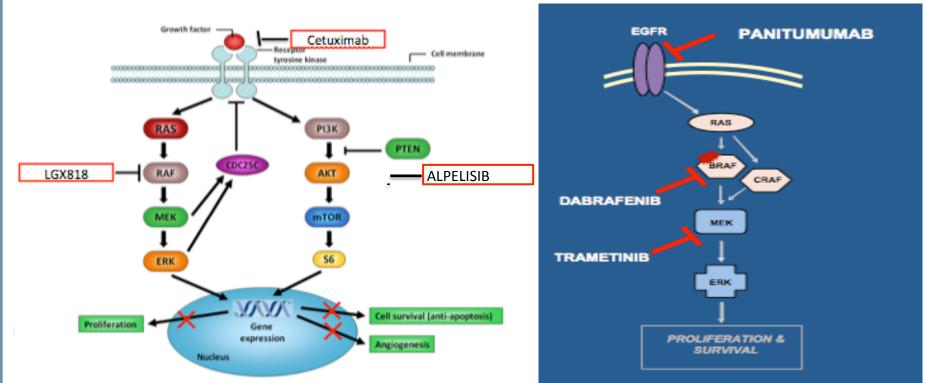
SECOND GENERATION BRAF THERAPEUTICS



Regimen	Ν	PR/CR (%)	SD (%)	mPFS (m)
Dabrafenib + Panitumumab	20	10%	80	3.4 Van Cutsem WGIC 2015
Encorafenib + Cetuximab (ph II)		11%	54 (53)	3.7 Elez WGIC 2015
Vemurafenib + Cetuximab	26	4%	16(40)	3.7 Hyman NEJM 2015
GI 📿				

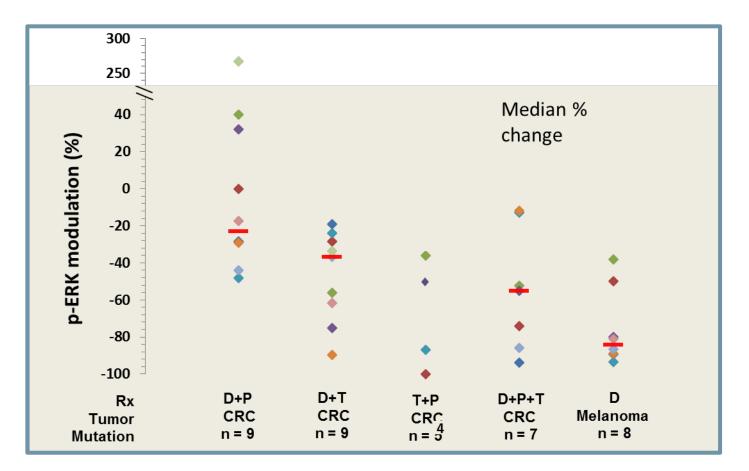
CONNECT

THIRD GENERATION BRAF INHIBITORS COMBOS



Regimen		PR/CR (%)	SD (%)	mPFS (m)	
Dabrafenib + Trabetinib + Panitumumab	35	26%	50	4.1 Van Cutsem WGIC 2015	
Encorafenib + Cetuximab + Alpelisib (ph II)		32%	44	4.3 Elez WGIC 2015	
Vemurafenib + Cetuximab + CPT 11		ONGOING			
Connect Policies by concept					

DIFFERENTIAL DEGREE OF MODULATION OF PERK BY VARIOUS TREATMENTS IN BRAF V600MUT CRC AND MELANOMA



Even so... numbers are still far distant fom those seen in melanoma. We have a long and fascinating way to walk



CONCLUSIONS

- BRAF testing can not be recommended in localized setting
- BRAF V600E testing should be perform at the debut of metastatic disease, based on:
 - Bad prognostic implications
 - Need from intensive chemotherapy combos to overcome bad outcome (FOLFOXIRI-bevacizumab)
 - Less benefit from anti-EGFR monoclonal antibodies
 - Refer patients to trials including BRAF inhibitor combos with anti-EGFR monoclonal antibodies





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