

TREATMENT OPTIONS FOR *BRAF*-MUTANT NSCLC


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April 20, 2024

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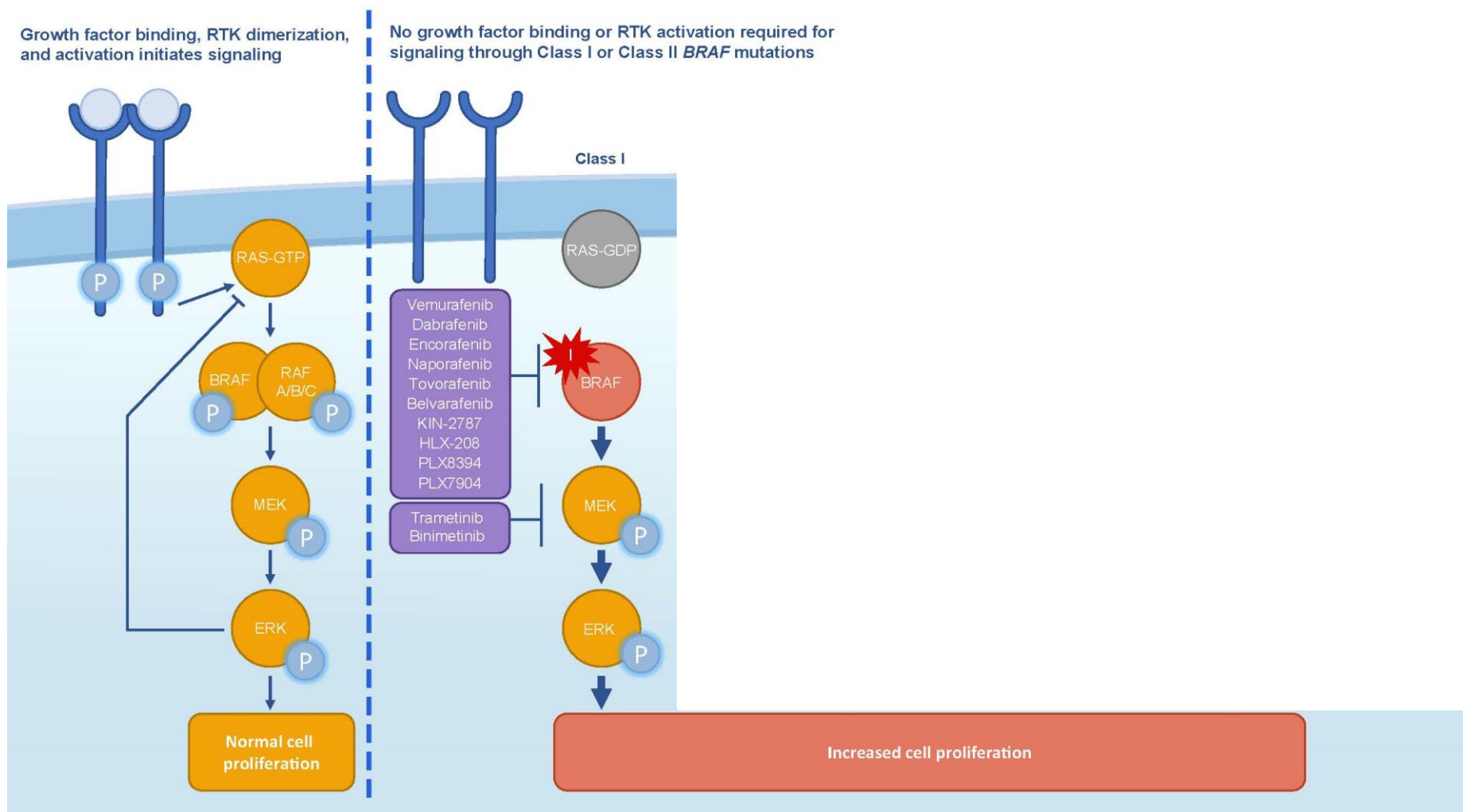
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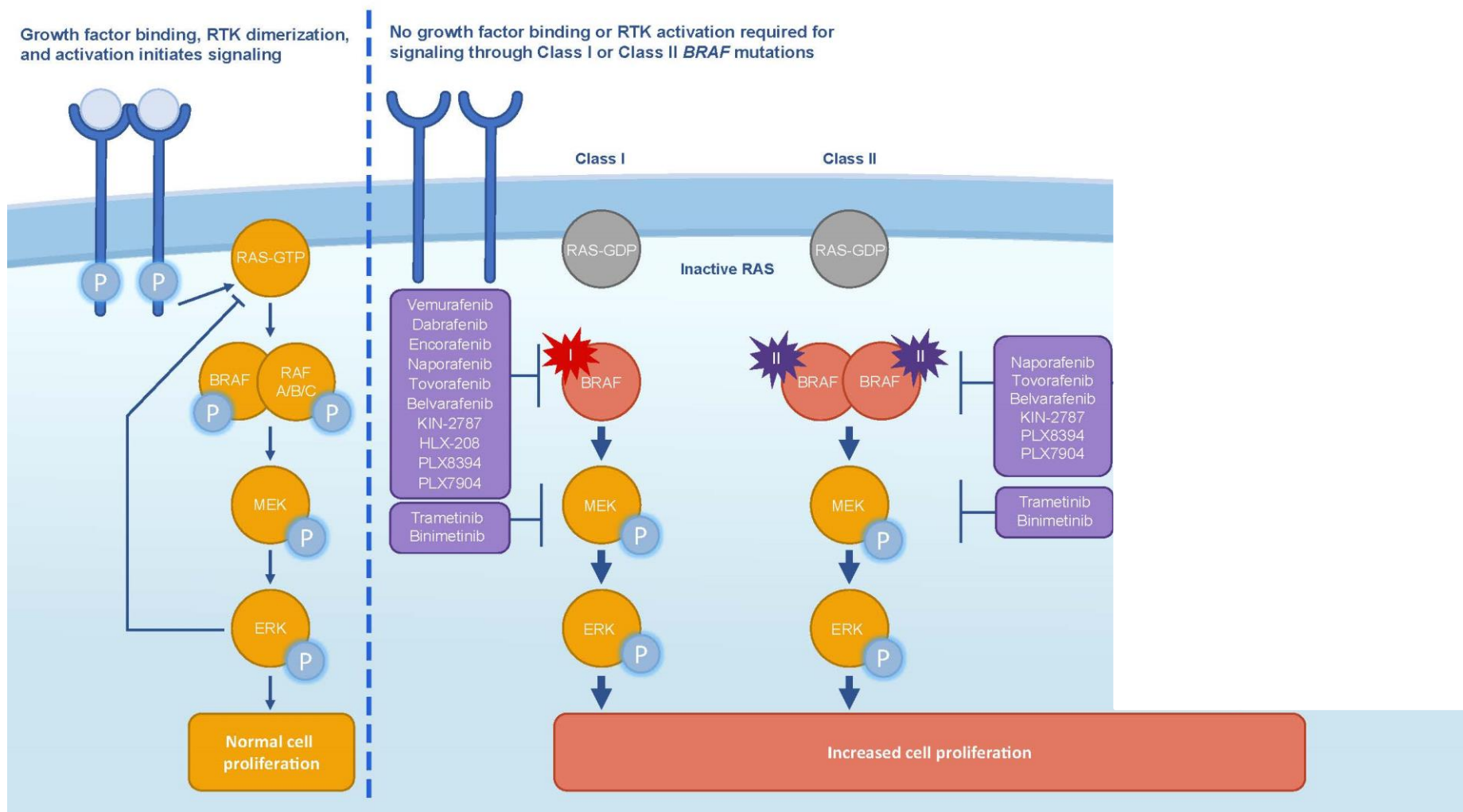


BRAF mutations can be classified into three groups



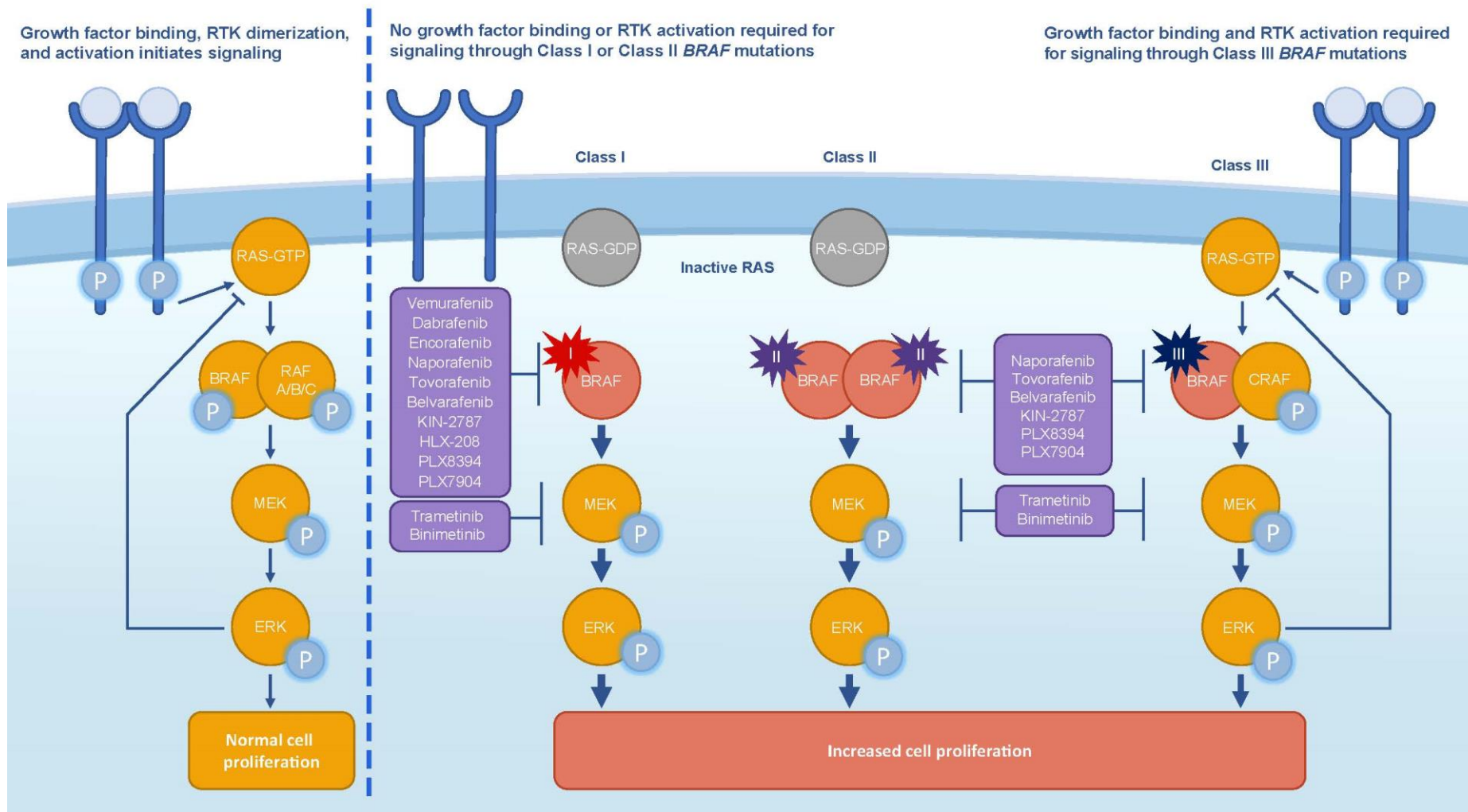
Adapted from Planchard, Sanborn, Negrao, Vaishnavi, Smit. Accepted for publication - *NPJ Precision Oncology*

BRAF mutations can be classified into three groups



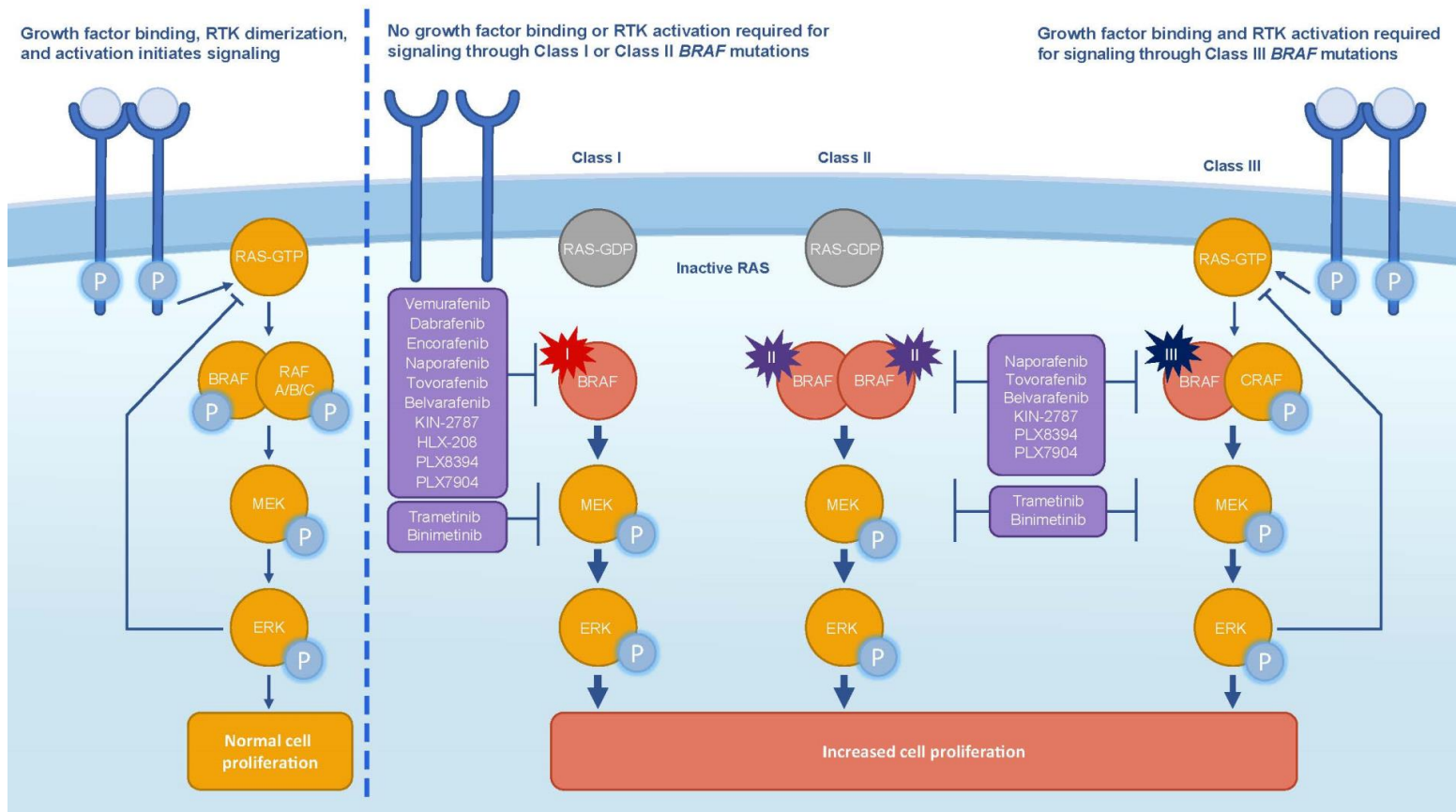
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BRAF mutations can be classified into three groups



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There are several topics for discussion



Class 1

- BRAF+MEKi
- Sequencing options
- Mechanisms of resistance
- Novel therapies
- Treatment of stage I-III disease

Class 2

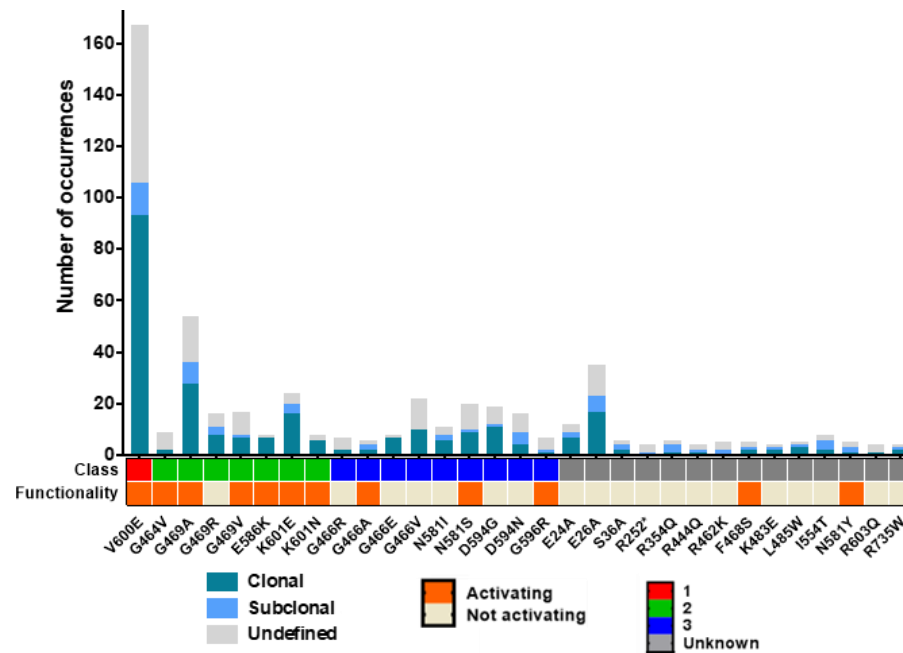
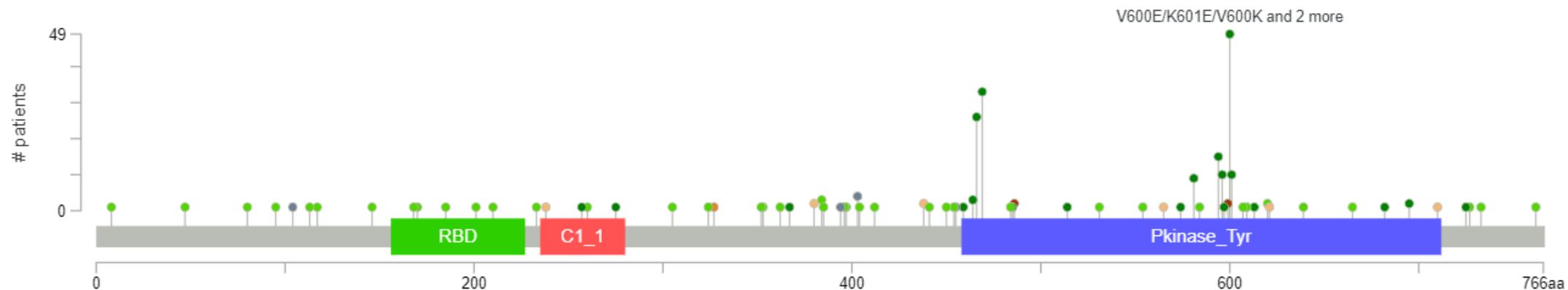
- BRAF+MEKi efficacy
- Novel therapies

Class 3

- Treatment options
- Novel therapies

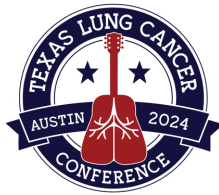
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V600X are the most prevalent *BRAF* mutations in NSCLC



cbiportal.org – March/2024 / Negrao et al. JTO 2020

BRAF+MEK inhibitor combos are effective for treatment of *BRAF* V600E NSCLC



| | Encorafenib+Binimetinib | | Dabrafenib+Trametinib | |
|-----|-------------------------|-----------------|-----------------------|-----------------|
| | Naïve (N=59) | Prior tx (N=39) | Naïve (N=36) | Prior tx (N=57) |
| ORR | 75% | 46% | 64% | 68% |
| DOR | NR | 16.7 mo | 10.2 mo | 9.8 mo |
| PFS | NR | 9.3 mo | 10.8 mo | 10.2 mo |
| OS | NA | NA | 17.3 mo | 18.2 mo |

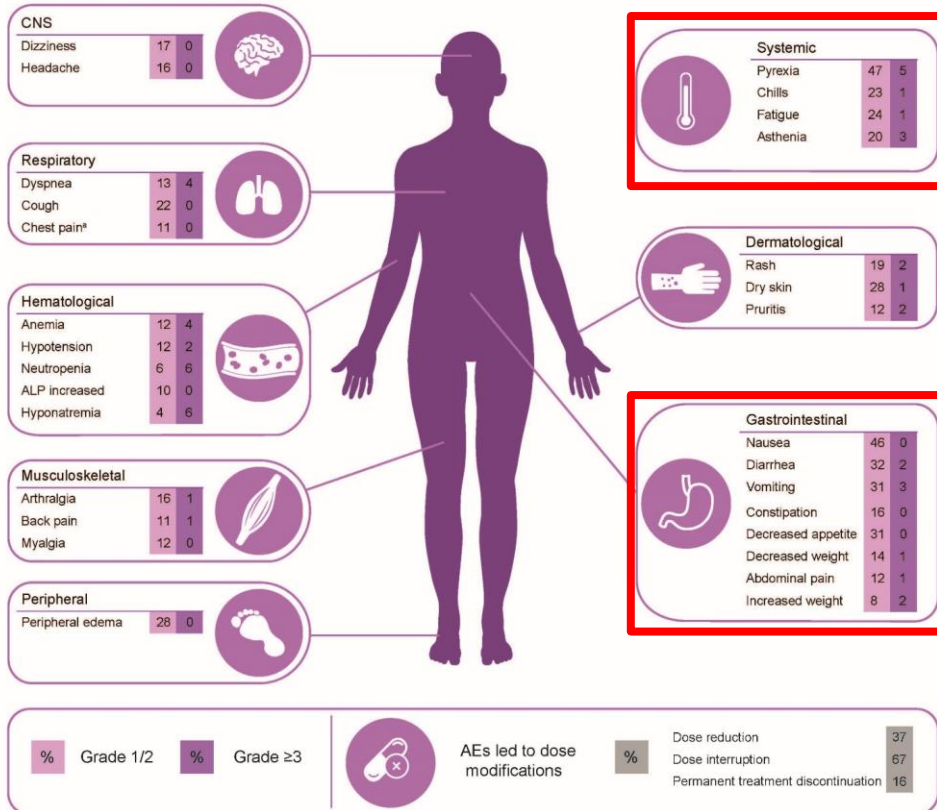
Brain mets underrepresented in both trials – PHAROS: 8/98; D+T - tx-naïve: 2/36; prior tx: 1/57
 PHAROS: prior ICI: 59% / ICI post-PD: 17% for tx naïve vs 26% for prior tx
 D+T trial: prior ICI – none / ICI post-PD: ~25%

Adapted from Riely et al. JCO 2023, Planchard et al. JTO 2021

Side effect profile differs for D+T and E+B

a

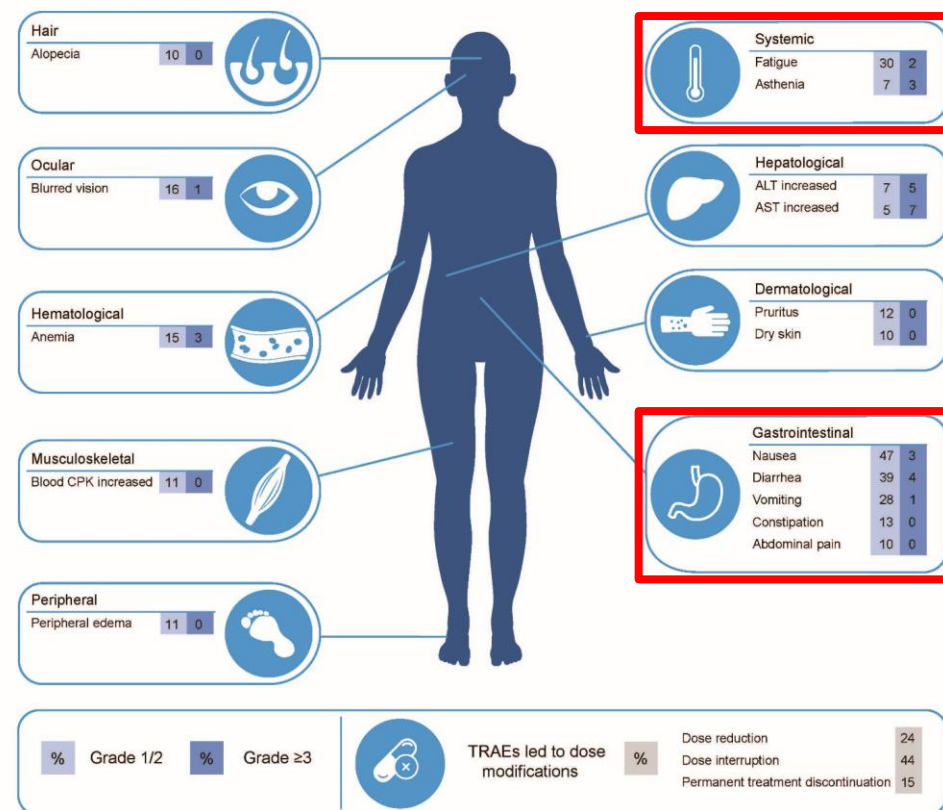
All-causality AEs for dabrafenib plus trametinib



Grade 3-4 TRAE: 66%

b

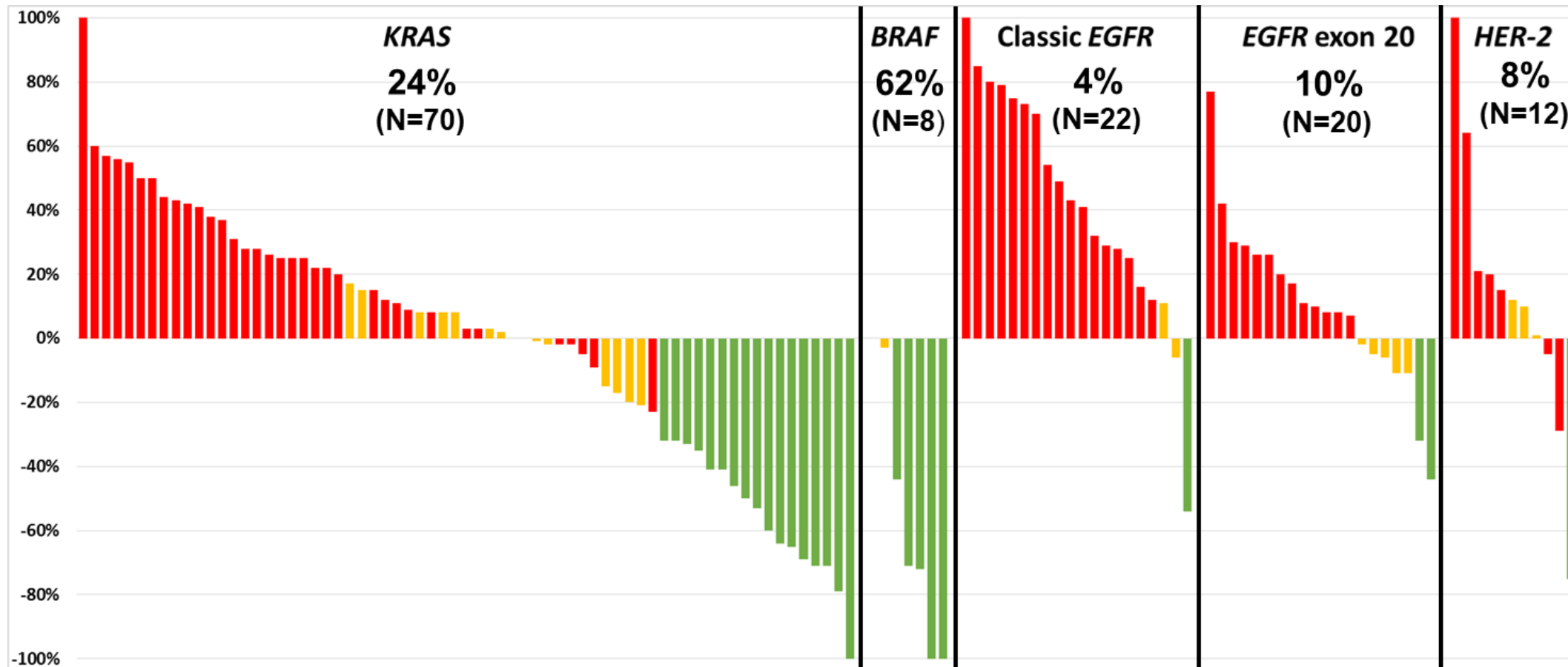
Treatment-related AEs for encorafenib plus binimetinib



Grade 3-4 TRAE: 41%

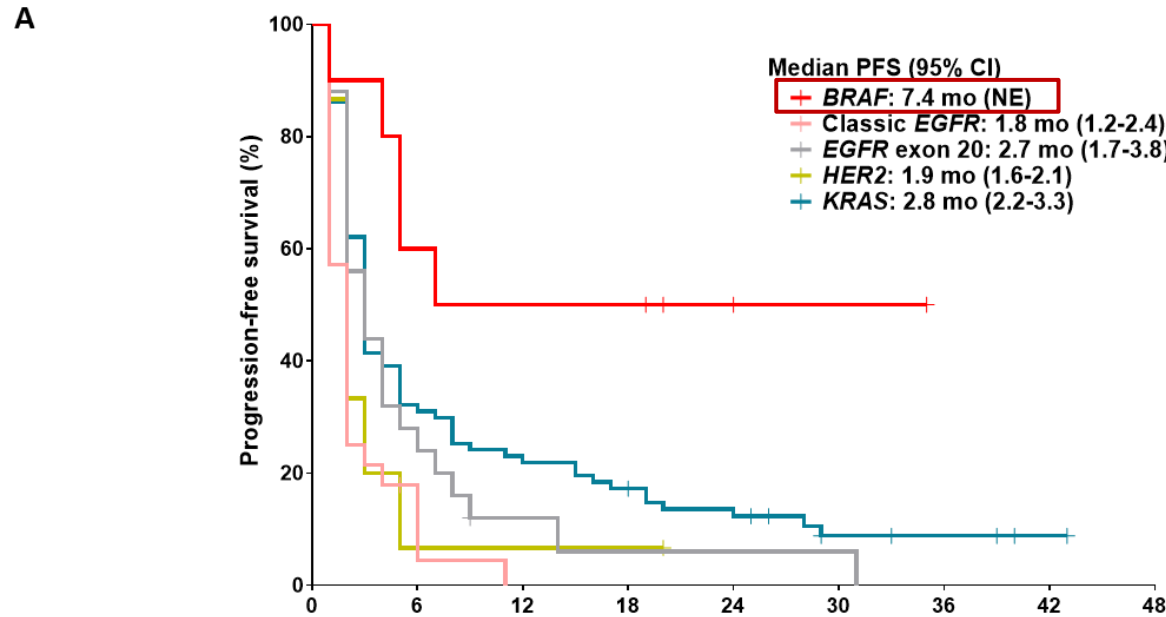
Planchard et al. JTO 2021; Riely et al. JCO 2023; Planchard, Sanborn, Negrao, Vaishnavi, Smit. Accepted for publication - NPJ Precision Oncology

***BRAF*-mutant NSCLC patients may have more favorable clinical outcomes to PD-1/PD-L1 inhibitor therapy...**



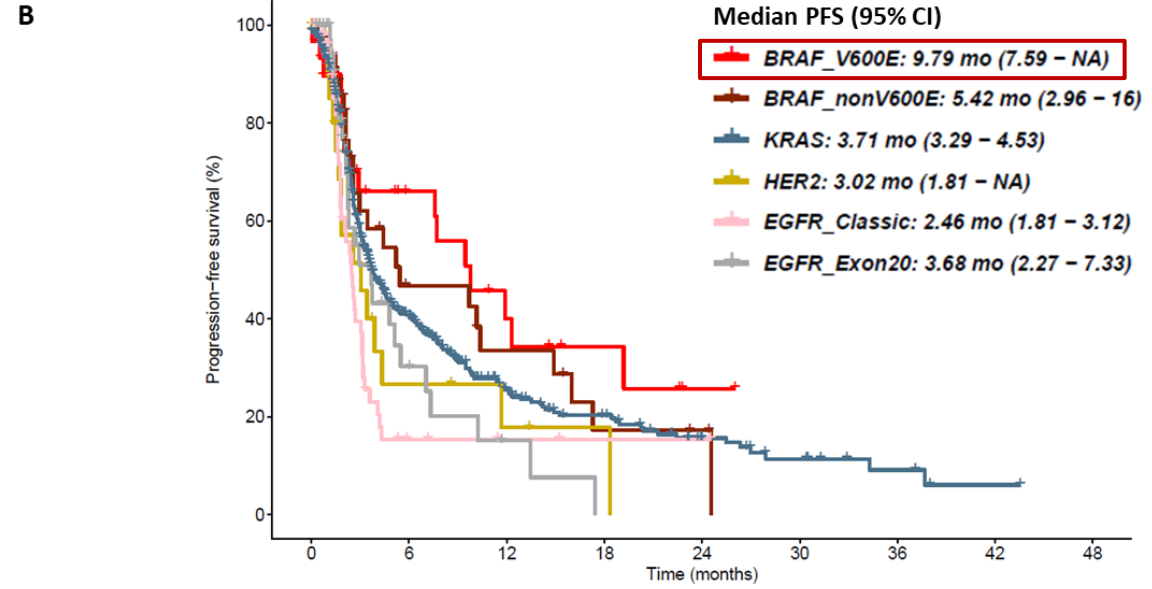
Negrao et al. WCLC 2019

BRAF-mutant NSCLC patients may have more favorable clinical outcomes to PD-1/PD-L1 inhibitor therapy...



Number at risk

| | 0 | 6 | 12 | 18 | 24 | 30 | 36 | 42 | 48 |
|---------------------|----|----|----|----|----|----|----|----|----|
| BRAF | 10 | 6 | 5 | 3 | 1 | 1 | 0 | 0 | 0 |
| Classic EGFR | 28 | 2 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| EGFR Exon 20 | 25 | 7 | 2 | 1 | 1 | 1 | 0 | 0 | 0 |
| HER2 | 15 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | 0 |
| KRAS | 87 | 27 | 20 | 15 | 9 | 4 | 2 | 1 | 0 |



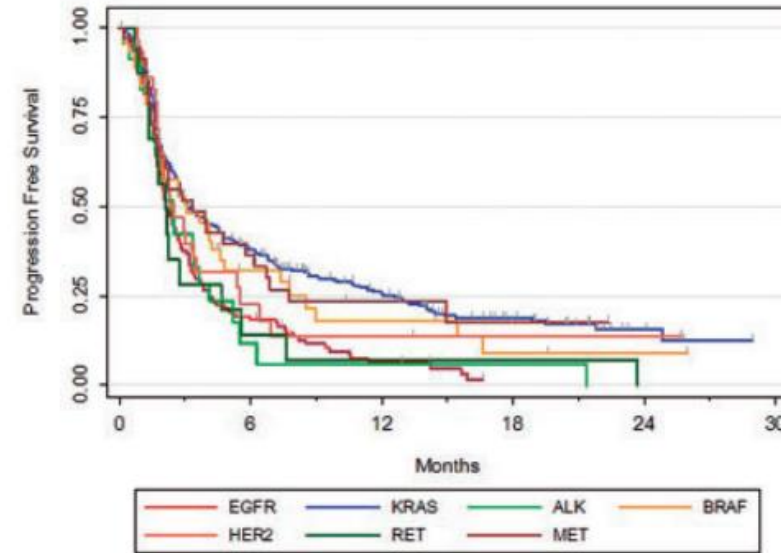
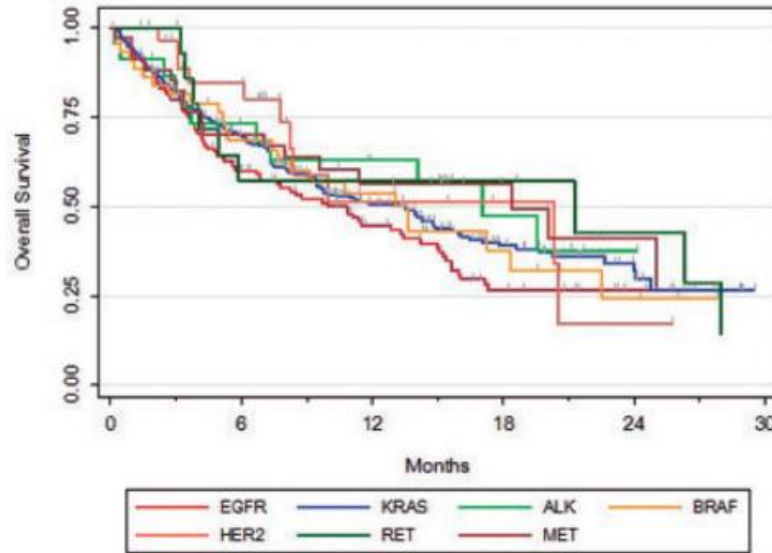
Number at risk

| | 0 | 6 | 12 | 18 | 24 | 30 | 36 | 42 | 48 |
|-----------------------|-----|-----|----|----|----|----|----|----|----|
| BRAFV600E | 30 | 13 | 7 | 4 | 1 | 0 | 0 | 0 | 0 |
| BRAF non-V600E | 36 | 11 | 7 | 3 | 2 | 0 | 0 | 0 | 0 |
| Classic EGFR | 54 | 4 | 2 | 1 | 1 | 0 | 0 | 0 | 0 |
| EGFR Exon 20 | 37 | 7 | 2 | 0 | 0 | 0 | 0 | 0 | 0 |
| HER2 | 21 | 4 | 2 | 1 | 0 | 0 | 0 | 0 | 0 |
| KRAS | 529 | 138 | 53 | 33 | 17 | 9 | 4 | 1 | 0 |

Negrao et al. JITC 2021

... or maybe not

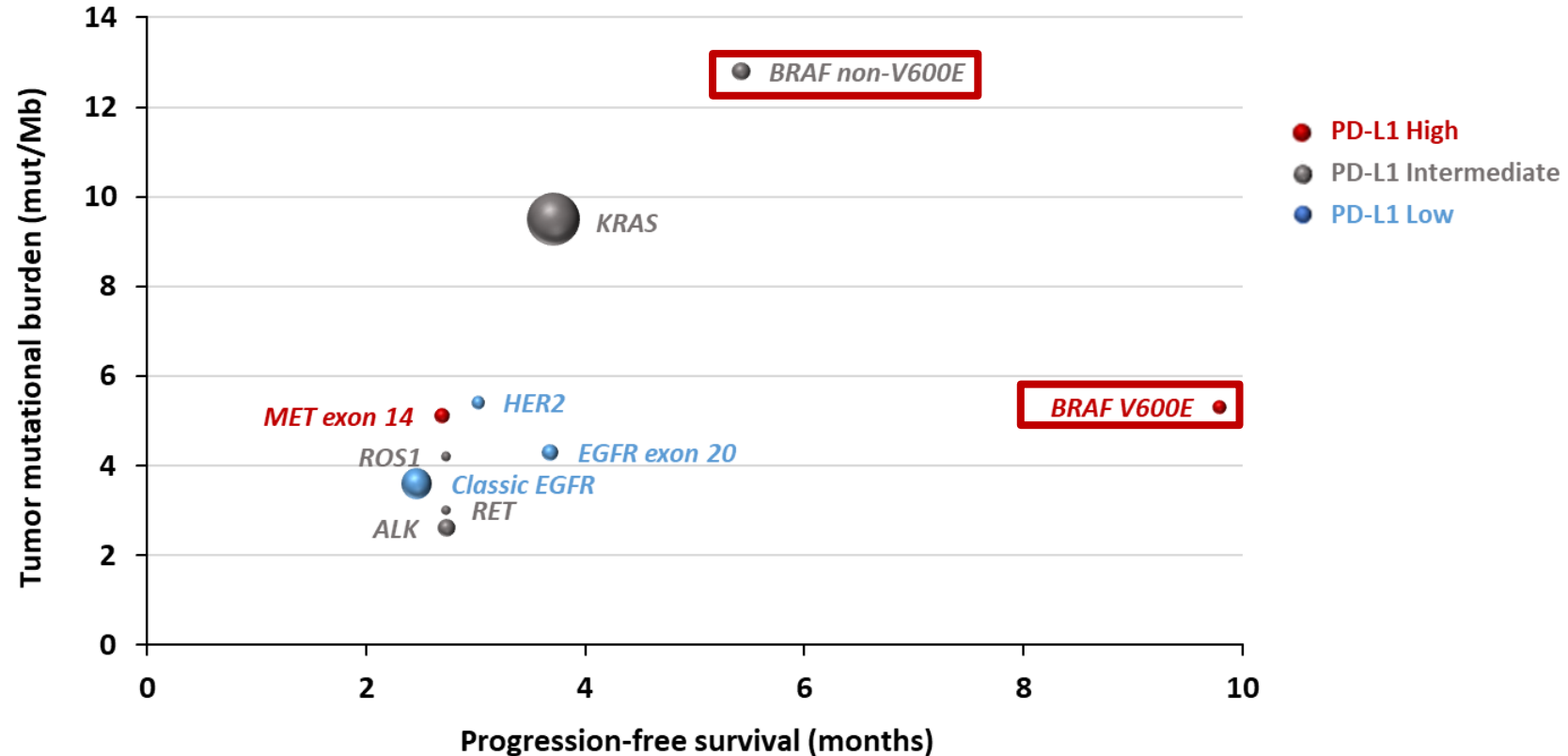
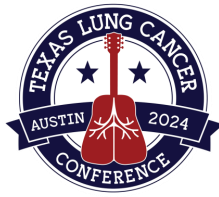
Oncogenic driver subgroups
n=543



| | EVT/N | Median PFS [95% CI] (months) | 6-month PFS [95% CI] | 12-month PFS [95% CI] |
|------|---------|------------------------------|----------------------|-----------------------|
| KRAS | 208/271 | 3.2 [2.7; 4.5] | 37.9 [32.1; 49.8] | 25.6 [20.2; 31.3] |
| EGFR | 117/125 | 2.1 [1.8; 2.7] | 18.4 [12.1; 25.6] | 6.4 [2.7; 12.1] |
| BRAF | 34/43 | 3.1 [1.8; 4.6] | 32.1 [18.3; 46.6] | 18.0 [7.2; 32.7] |
| HER2 | 23/29 | 2.5 [1.8; 3.5] | 22.7 [8.9; 40.2] | 13.6 [3.6; 30.1] |
| MET | 26/36 | 3.4 [1.7; 6.2] | 36.5 [20.7; 52.4] | 23.4 [10.6; 39.0] |
| ALK | 21/23 | 2.5 [1.5; 3.7] | 11.8 [2.2; 30.2] | 5.9 [0.4; 23.0] |
| ROS1 | - | - | - | - |
| RET | 15/16 | 2.1 [1.3; 4.7] | 14.1 [2.3; 35.9] | 7.0 [0.4; 27.1] |

Mazieres et al. Annals of Oncology 2019

BRAF-mutant NSCLC patients can have favorable TMB and PD-L1 expression profile for PD-1/PD-L1 inhibitor therapy

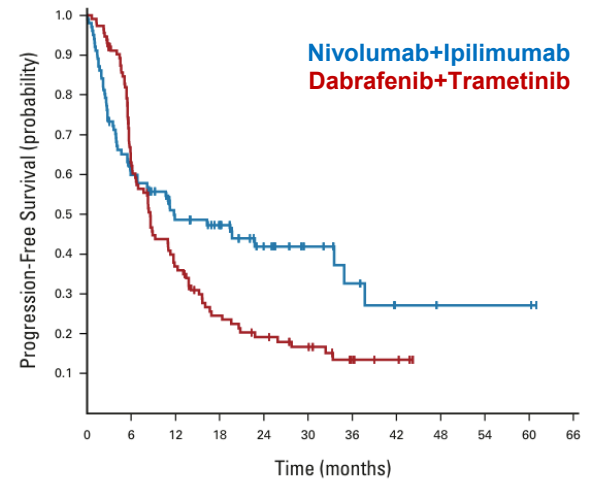
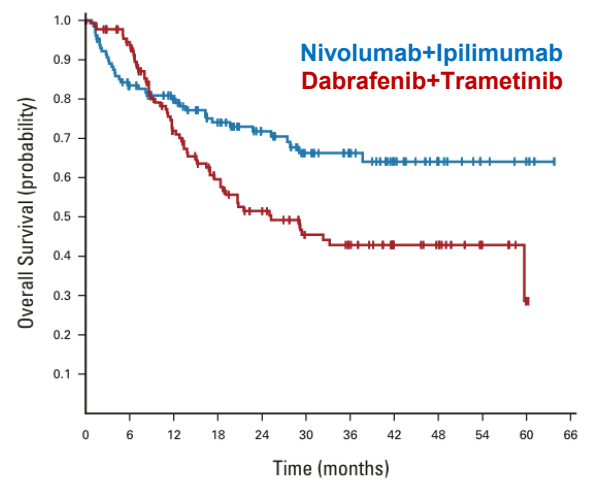
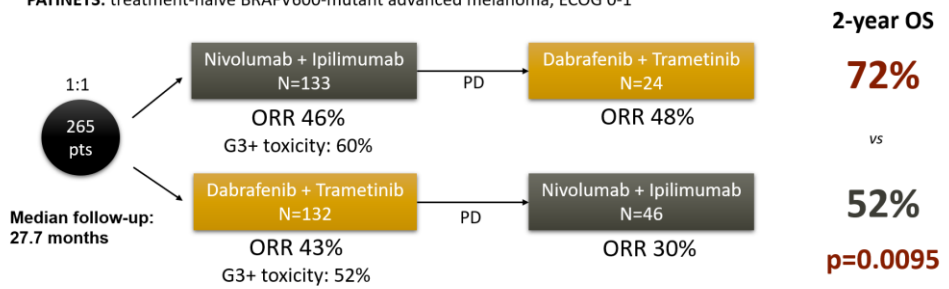


Negrao et al. JITC 2021

Melanoma trials suggest frontline anti-PD-1+CTLA-4 combo is superior to BRAF+MEK inhibitor combo

DREAMseq (Doublet, Randomized Evaluation in Advanced Melanoma Sequencing): A phase III trial—ECOG-ACRIN EA6134

AIM: to compare the efficacy and toxicity of the sequence of nivolumab/ipilimumab followed by dabrafenib/trametinib to the converse sequence
PATINETS: treatment-naïve BRAFV600-mutant advanced melanoma, ECOG 0-1



| No. at risk: | 0-6 | 6-12 | 12-18 | 18-24 | 24-30 | 30-36 | 36-42 | 42-48 | 48-54 | 54-60 | 60-66 |
|--------------|-----|------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| A/C | 133 | 99 | 87 | 71 | 55 | 42 | 33 | 23 | 15 | 6 | 3 |
| B/D | 132 | 115 | 78 | 60 | 47 | 35 | 30 | 18 | 15 | 6 | 1 |

| No. at risk: | 0-6 | 6-12 | 12-18 | 18-24 | 24-30 | 30-36 | 36-42 | 42-48 | 48-54 | 54-60 | 60-66 |
|--------------|-----|------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| A | 101 | 57 | 40 | 32 | 19 | 12 | 7 | 3 | 2 | 2 | 2 |
| B | 113 | 66 | 38 | 23 | 17 | 13 | 6 | 3 | 0 | 0 | 0 |

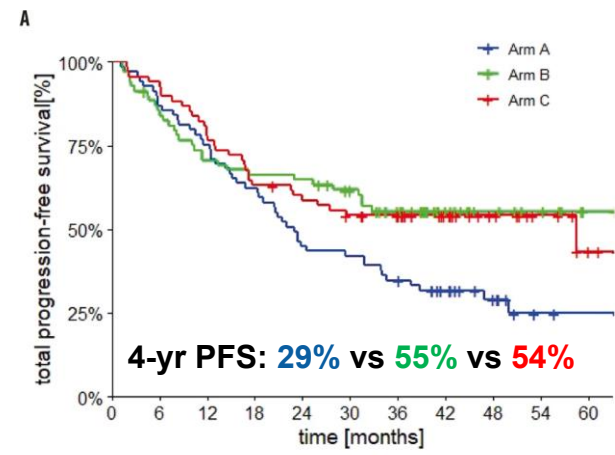
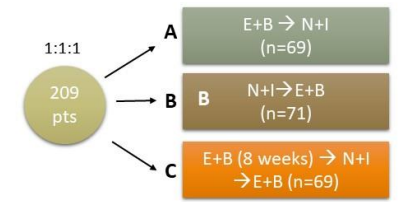
Atkins et al. ASCO Plenary Series 2021 and JCO 2022 / Ascierto et al. ESMO 2021 and Nature Communications 2024

2021 ESMO Congress

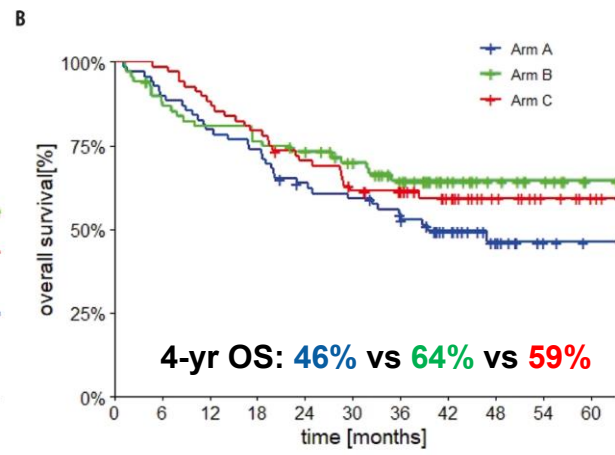
SECOMBIT: The best sequential approach with combo immunotherapy [ipilimumab (I) /nivolumab (N)] and combo target therapy [encorafenib (E)/binimetinib (B)] in patients with BRAF mutated metastatic melanoma: A phase II randomized study

Population:
 - metastatic BRAFV600-mutated melanoma
 - treatment naïve
 - ECOG PS 0 or 1

Primary endpoint:
 - overall survival

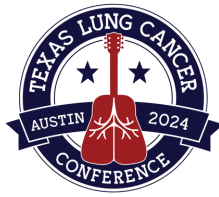


| | | | | | | | | | | | |
|---|----|----|----|----|----|----|----|----|----|---|---|
| ■ | 69 | 60 | 52 | 43 | 31 | 29 | 24 | 17 | 11 | 4 | 3 |
| ■ | 69 | 58 | 48 | 45 | 44 | 39 | 30 | 18 | 11 | 8 | 3 |
| ■ | 68 | 64 | 53 | 43 | 39 | 35 | 32 | 21 | 12 | 7 | 3 |



| | | | | | | | | | | | |
|---|----|----|----|----|----|----|----|----|----|---|---|
| ■ | 69 | 62 | 55 | 51 | 42 | 39 | 34 | 22 | 13 | 6 | 3 |
| ■ | 69 | 59 | 54 | 51 | 48 | 41 | 32 | 20 | 13 | 9 | 3 |
| ■ | 68 | 67 | 60 | 54 | 47 | 39 | 36 | 24 | 13 | 7 | 4 |

Launch of a multi-institution collaborative effort to determine the optimal treatment sequence for *BRAF* V600E NSCLC patients



Multi-institution project

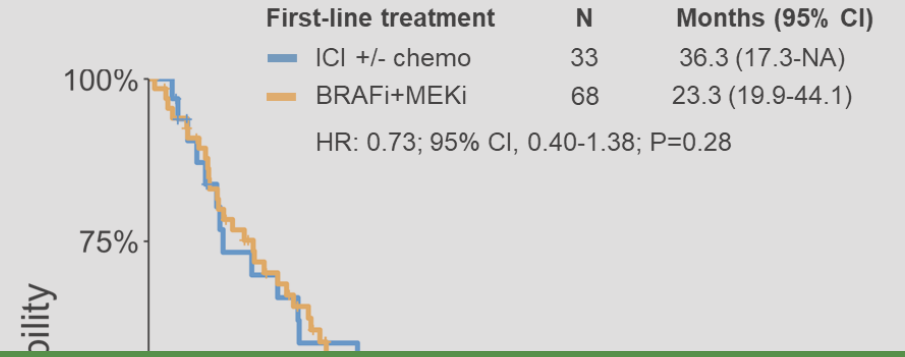
DFCI – Ricciuti / Di Federico
 MSKCC – Offin / Chen
 Gustave Roussy – Aldea
 Bologna – Ardizzoni
 Yale – Stockhammer
 Detroit – Rous
 UChicago – Garassino / Tawhani

Aim: determine best treatment sequence in *BRAF* V600E NSCLC

Primary endpoint: OS

Population:

- Advanced NSCLC
- *BRAF* V600X



Interest and support are welcome for this important effort

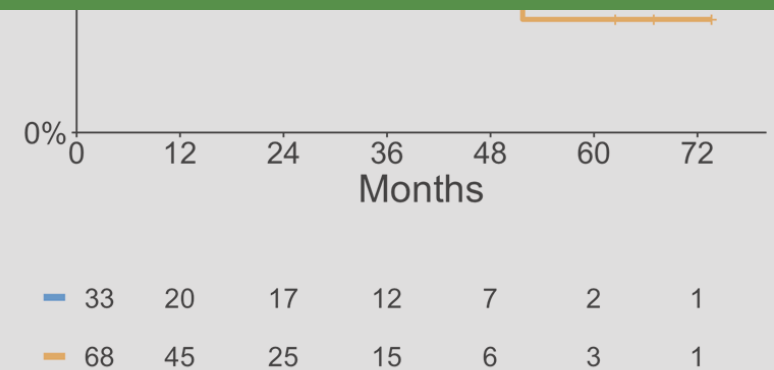
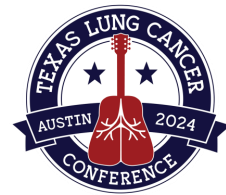


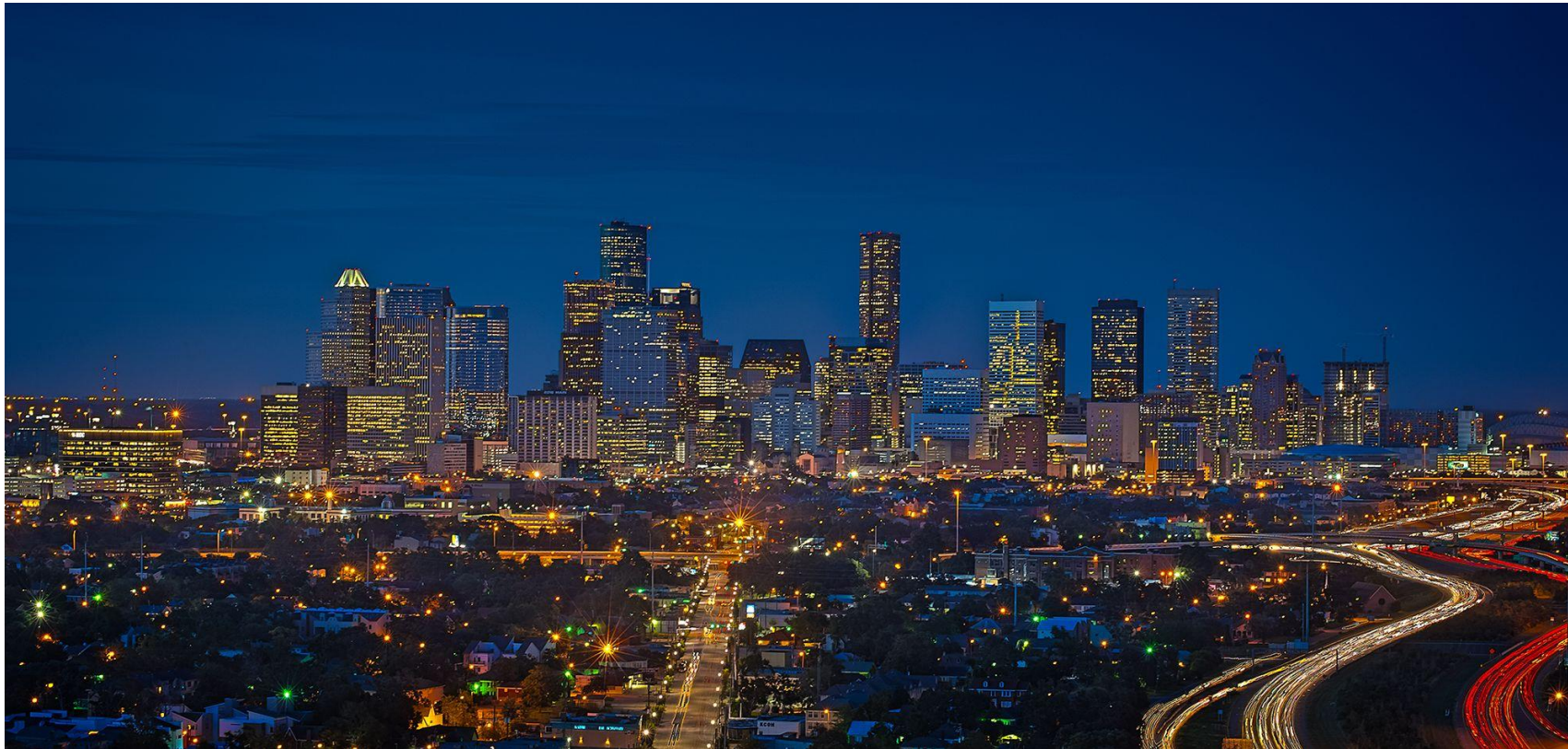
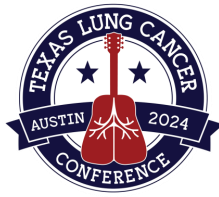
Figure courtesy of Biagio Ricciuti



Conclusions

- *BRAF* mutations are categorized into three classes based on their functional properties
- *BRAF* V600X are the most prevalent mutations
- BRAF+MEK inhibitors: effective for frontline and salvage therapy with distinctions in safety profile
- Role of ICI remains subject of further investigation – PD-L1 and/or TMB high status may favor ICI therapy over BRAF+MEK inhibitor combo
- Multi-institution efforts collaborations are key in this disease space
 - Necessary to answer relevant clinical questions since *BRAF* V600X mutations are uncommon drivers in NSCLC
 - May facilitate clinical trial development - particularly in the frontline setting

Thank you for your time and attention!



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