

**Cell Systems, Volume 11**

**Supplemental Information**

**Comparing the Efficacy of Cancer Therapies  
between Subgroups in Basket Trials**

**Adam C. Palmer, Deborah Plana, and Peter K. Sorger**

**Supplemental Information**

**Comparing the efficacy of cancer therapies between subgroups in basket trials**

**Adam C. Palmer\*, Deborah Plana\*, Peter K. Sorger**

**\*These authors contributed equally**

**Inventory of Supplemental Information**

- 1. Supplementary Table S1. Related to Figure 2.**
- 2. Supplementary Table S2. Related to Figure 3.**
- 3. Supplementary Figure S1. Related to Figure 4.**
- 4. Supplementary Figure S2. Related to Figure 5.**

A.

Mutation Type (n= number of patients per subgroup)	Test for larger benefit in tumor volume		Conclusion based on 25% False Discovery Rate
	p-value	Benjamini-Hochberg critical value for 1-sided test	
L755 Hotspot (n =13)	0.031	0.025	N.S.
Exon20 Insertion Hotspot (n= 28)	0.162	0.050	N.S.
S310 Hotspot (n= 30)	0.267	0.075	N.S.
V777 Hotspot (n= 15)	0.276	0.100	N.S.
ERBB3 Nonhotspot (n= 4)	0.421	0.125	N.S.
Other Hotspot (n= 9)	0.529	0.150	N.S.
PKD Nonhotspot (n= 12)	0.651	0.175	N.S.
ERBB3 Hotspot (n= 12)	0.931	0.200	N.S.
PKD Hotspot (n= 14)	0.950	0.225	N.S.
Other Nonhotspot (n= 4)	0.954	0.250	N.S.

B.

Mutation Type (n= number of patients per subgroup)	Test for larger benefit in hazard ratio for progression		Conclusion based on 25% False Discovery Rate
	p-value	Benjamini-Hochberg critical value for 1-sided test	
<b>Exon20 Insertion Hotspot (n= 26)</b>	<b>0.011</b>	<b>0.025</b>	<b>Larger Benefit</b>
S310 Hotspot (n= 30)	0.094	0.050	N.S.
Other Hotspot (n= 8)	0.274	0.075	N.S.
L755 Hotspot (n= 13)	0.456	0.100	N.S.
PKD Hotspot (n= 14)	0.499	0.125	N.S.
PKD Nonhotspot (n= 11)	0.759	0.150	N.S.
V777 Hotspot (n= 15)	0.944	0.175	N.S.
ERBB3 Nonhotspot (n= 4)	0.962	0.200	N.S.
ERBB3 Hotspot (n= 12)	0.970	0.225	N.S.
Other Nonhotspot (n= 4)	0.985	0.250	N.S.

Supplementary Table S1. Benjamini-Hochberg critical values for analysis of neratinib tumor volume responses (A) and hazard ratios for progression (B) by specific mutation type. Related to Figure 2.

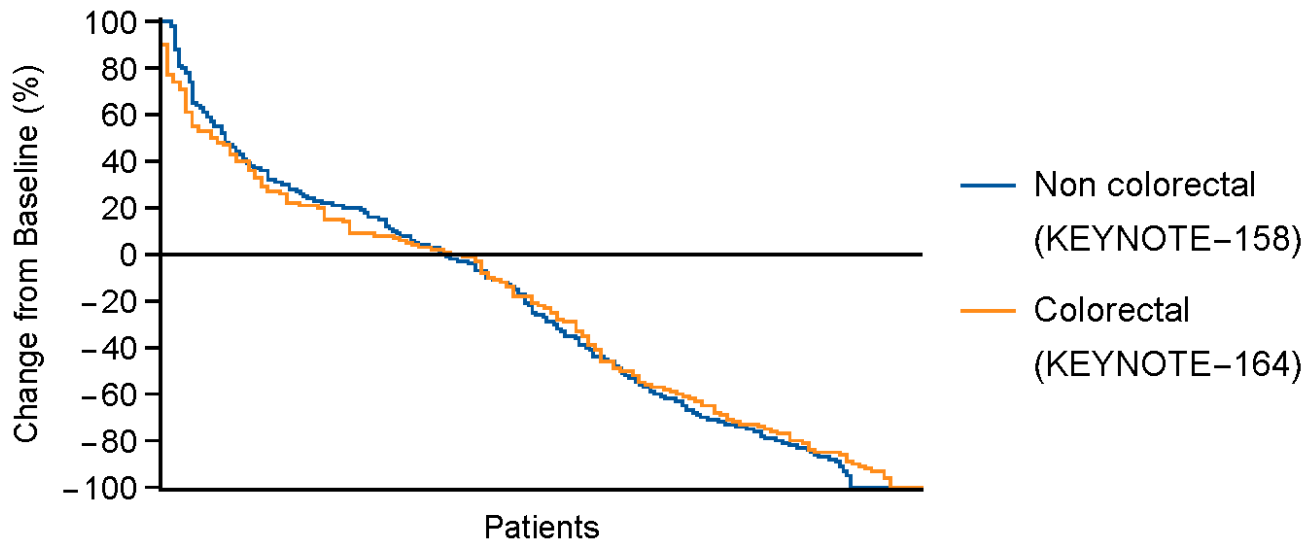
A.

NTRK paralog (n= number of patients per subgroup)	Test for larger benefit in tumor volume		Test for smaller benefit in tumor volume		Conclusion based on 25% False Discovery Rate
	p-value	Benjamini- Hochberg critical value for 2-sided test	p-value	Benjamini- Hochberg critical value for 2-sided test	
NTRK3 (n= 28)	0.147	0.063	0.854	0.125	N.S.
NTRK1 (n= 24)	0.804	0.125	0.197	0.063	N.S.

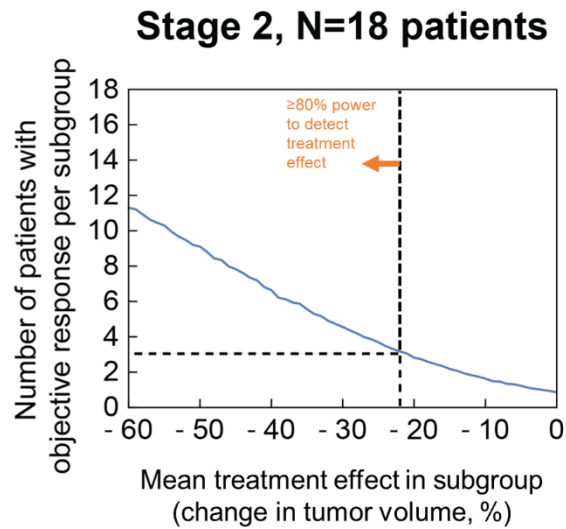
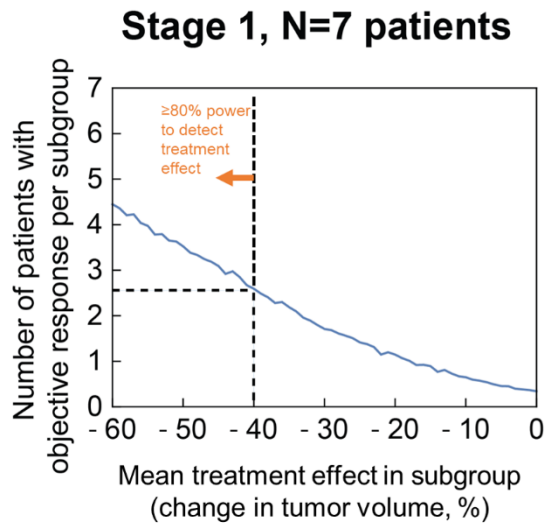
B.

NTRK fusion partner (n= number of patients per subgroup)	Test for larger benefit in tumor volume		Test for smaller benefit in tumor volume		Conclusion based on 25% False Discovery Rate
	p-value	Benjamini- Hochberg critical value for 2-sided test	p-value	Benjamini- Hochberg critical value for 2-sided test	
ETV6-NTRK3 (n= 27)	0.160	0.042	0.840	0.125	N.S.
TPM3-NTRK1 (n= 9)	0.757	0.083	0.246	0.083	N.S.
LMNA-NTRK1 (n= 5)	0.941	0.125	0.059	0.042	N.S.

**Supplementary Table S2. Benjamini-Hochberg critical values for analysis of larotrectinib tumor volume responses by genomic status. Related to Figure 3.**



**Supplementary Figure S1. Comparison of tumor volume changes in response to pembrolizumab in colorectal and non-colorectal cancers. Related to Figure 4.** Patient tumor volume change data was extracted from the KEYNOTE-158 (n=233) and KEYNOTE-164 (n=124) trials. Both trials tested the PD-L1 inhibitor, pembrolizumab, in colorectal and noncolorectal cancers.



**Supplementary Figure S2. Number of responsive patients needed to achieve 80% power using a permutation test. Related to Figure 5.** Computational simulation identifies number of patients with objective responses (greater than 30% decrease in tumor volume) in a subgroup, across different average treatment effects. Dotted lines indicate the number of responsive patients needed for 80% power in a permutation test (based on the simulation and tumor volume change values reported in Figure 5, row 2).