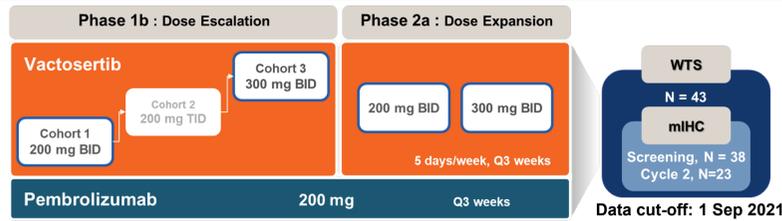


BACKGROUND

- Immune checkpoint inhibitor is an established treatment option for microsatellite instability-high (MSI-H) colorectal cancer^{1,2}, however, only 3.5-6.5% of metastatic colorectal cancer (mCRC) patients show MSI-H^{3,4,5}
- MP-VAC-204, a phase 1b/2a study evaluating the combination of vactosertib (TGF-β receptor type 1 inhibitor) with pembrolizumab in previously treated microsatellite stable (MSS) mCRC (Clinical trial: NCT03724851) demonstrated promising anti-tumor efficacy and manageable safety at ASCO 2021⁶
 - Overall Response Rate (ORR) were 16.0% (RECIST) / 20.0% (iRECIST) and median Overall Survival (mOS) was 15.8 months without cardiac valvulopathy
- Meanwhile, expression of PD-L1 and tumor-infiltrating CD8 T cells were reported to have a decisive effect on the immunotherapy response⁷
- This study has been conducted to understand the basis of the clinical responses of anti-PD-1 and TGF-β inhibitor combination therapy in MSS mCRC, conducting a comprehensive analysis of survival outcome, whole transcriptome sequencing (WTS) data, and multiplex immunohistochemistry (mIHC) data

METHODS



Clinical Analysis

- Response assessed by RECIST and iRECIST
- Responder was defined as best overall response (BOR) complete remission (CR), partial response (PR), and iPR
- Non-responder was defined as BOR progression disease (PD)

Whole Transcriptome Sequencing (WTS)

- Screening and Cycle 2 post-treatment biopsies were used
- WTS data was generated by Illumina platform
- CD274 and CD8A expression cut-off were calculated as median value in the merged data set of TCGA Pan-cancer and MP-VAC-204 study
- Tumor microenvironment immune status with a combination of CD274 and CD8A gene expression (high or low) was classified into four subtypes

Subtype	CD8A high	CD8A low
CD274 high	Subtype 1	Subtype 3
CD274 low	Subtype 4	Subtype 2

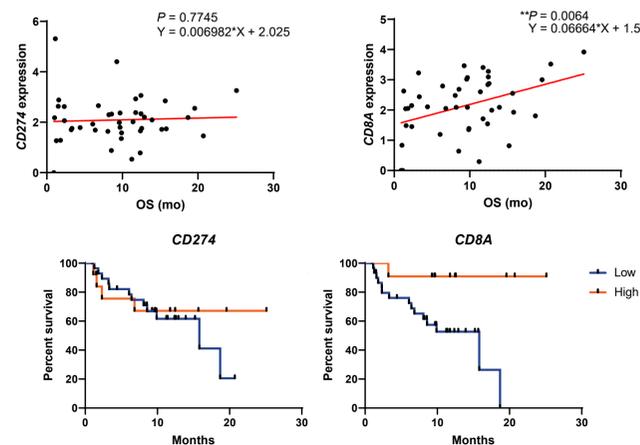
Multiplex Immunohistochemistry (mIHC)

- Screening and Cycle 2 post-treatment biopsies were used
- As part of the multiplex panel, expression of PD-L1 (clone 22C3), CD8 (clone SP16), and granzyme B (GZB, clone D6E9W) were measured in tumor nest and stroma

Table 1. Baseline Characteristics

	Subtype 1 (N=6)	Subtype 2 (N=25)	Subtype 3 (N=7)	Subtype 4 (N=5)	Overall (N=43)
Age, years, median (range)	62 (39-72)	60 (39-83)	56 (40-67)	63 (46-70)	60 (39-83)
Sex, n (%)					
Male	5 (83)	12 (48)	3 (43)	5 (100)	25 (58)
Female	1 (17)	13 (52)	4 (57)	0 (0)	18 (42)
Race, n (%)					
Asian	6 (100)	25 (100)	7 (100)	5 (100)	43 (100)
ECOG Performance Status, n (%)					
0	3 (50)	9 (36)	1 (14)	1 (20)	14 (33)
1	3 (50)	16 (64)	6 (86)	4 (80)	29 (67)
KRAS mutation, n (%)	2 (33)	13 (52)	2 (29)	0 (0)	17 (40)
BRAF mutation, n (%)	0 (0)	1 (4)	0 (0)	0 (0)	1 (2)
Microsatellite Instability, n (%)					
MSS	6 (100)	25 (100)	7 (100)	5 (100)	43 (100)
TMB (var No./MBs), n (%)					
< 10	4 (67)	18 (72)	4 (57)	3 (60)	29 (67)
≥ 10	1 (17)	7 (28)	3 (43)	2 (40)	13 (30)
Unknown	1 (17)	0 (0)	0 (0)	0 (0)	1 (2)
No. of prior anticancer therapies, n (%)					
1-2	4 (67)	9 (36)	1 (14)	1 (20)	15 (35)
3	1 (17)	6 (24)	2 (29)	2 (40)	11 (26)
≥ 4	1 (17)	10 (40)	4 (57)	2 (40)	17 (40)

Figure 3. Correlation and Survival Analysis by expression of CD274 and CD8A

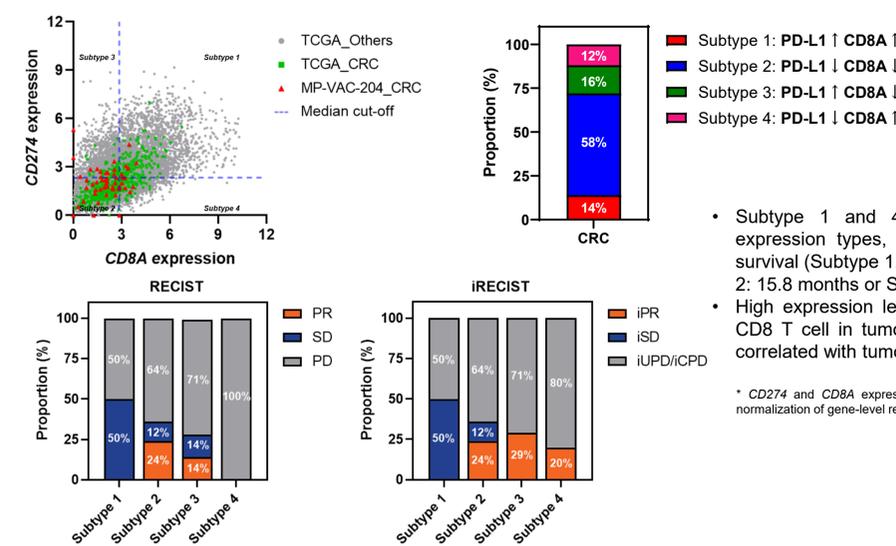


Log-rank test	CD274		CD8A	
	Low	High	Low	High
mOS (months)	15.8	Not reached	15.8	Not reached
P value	0.5968 (ns)		0.0091 (**)	
Hazard Ratio	1.35		8.65	
95% CI	0.47 - 3.90		3.13 - 23.93	

- Overall survival and CD8A gene expression showed positive correlation (**P=0.0064)
- Survival analysis by expression of CD274 (PD-L1) or CD8A suggested that high expression of CD8A was associated with prolonged mOS than CD274 (15.8 months vs Not reached, **P=0.0091, hazard ratio 8.65 [95% CI 3.13-23.93])

RESULTS

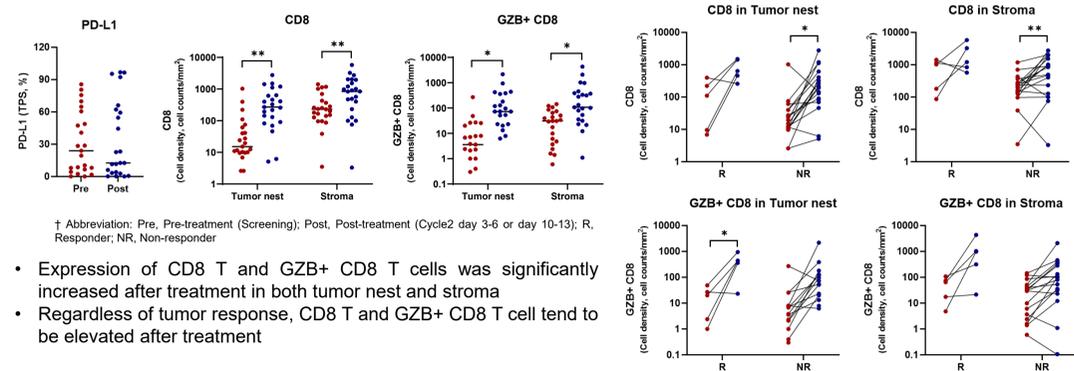
Figure 1. Tumor Microenvironment Immune Subtypes



- Subtype 1 and 4, the CD8A gene high expression types, showed increased overall survival (Subtype 1,4: Not reached vs Subtype 2: 15.8 months or Subtype 3: 6.8 months)
- High expression levels of CD8 T and GZB+ CD8 T cell in tumor nest at Screening were correlated with tumor response

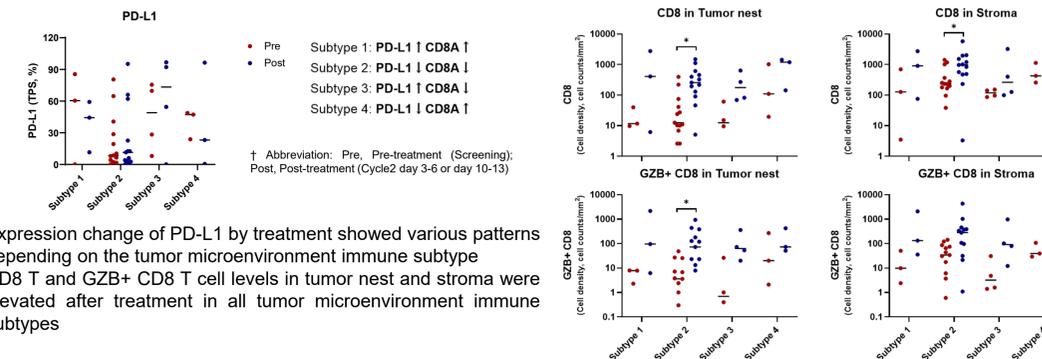
* CD274 and CD8A expression level was calculated based on normalization of gene-level read counts from RNA sequencing data

Figure 4. Changes in the expression of PD-L1, CD8 T cell, and GZB+ CD8 T cell after treatment



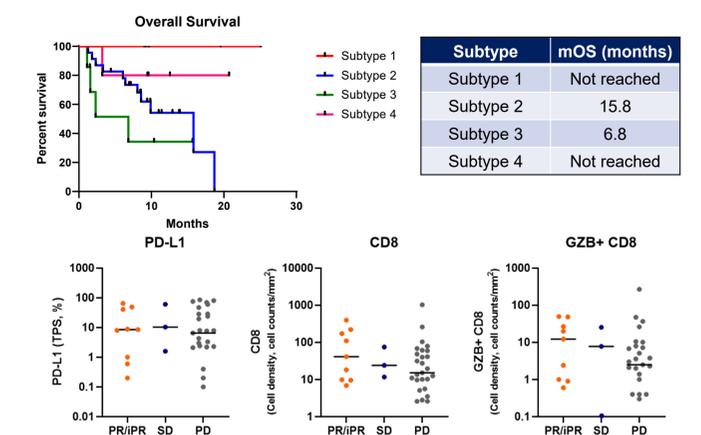
- Expression of CD8 T and GZB+ CD8 T cells was significantly increased after treatment in both tumor nest and stroma
- Regardless of tumor response, CD8 T and GZB+ CD8 T cell tend to be elevated after treatment

Figure 5. Expression of PD-L1, CD8 T cell, and GZB+ CD8 T cell after treatment by Tumor Microenvironment Immune Subtypes



- Expression change of PD-L1 by treatment showed various patterns depending on the tumor microenvironment immune subtype
- CD8 T and GZB+ CD8 T cell levels in tumor nest and stroma were elevated after treatment in all tumor microenvironment immune subtypes

Figure 2. Survival Analysis by Tumor Microenvironment Immune Subtypes



- 58% of MP-VAC-204 CRC patients were categorized as Subtype 2, the "immune-cold" subtype
- 24% of patients with "immune-cold" tumor microenvironment showed PR (RECIST 1.1 and iRECIST)

SUMMARY and CONCLUSIONS

- Vactosertib and pembrolizumab combination therapy was well tolerated and demonstrated promising anti-tumor efficacy in this phase 1b/2a study with previously treated MSS mCRC patients
- With respect to overall survival, better clinical outcome with this dual inhibition regimen of TGF-β and PD-1 was associated with CD8 T cell infiltration status in tumor microenvironment
- The increase in CD8 T cell infiltration driven by vactosertib and pembrolizumab treatment was observed regardless of tumor response based on RECIST 1.1
- It is possible that this increase in TIL is one of the main contributing factors of better survival in these previously treated MSS mCRC patients
- Further investigation is warranted

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