

Supplemental Tables for: A Randomized Phase II Trial of mFOLFOX6 + Bevacizumab Alone or with AdCEA Vaccine + Avelumab Immunotherapy for Untreated Metastatic Colorectal Cancer Jason Redman et al.

Supplemental Table 1. 138 peripheral immune cell subsets analyzed by flow cytometry. Ten parental phenotypes are identified as well as refined subsets of each relating to maturation and function. Expected function based on expression of specific markers within each subset is indicated in italics.

1. Total CD4+ T cells

- PD-L1+ CD4 activation/inhibition
- PD-1⁺CD4 activation/inhibition
- CTLA-4⁺ CD4 inhibition
- Tim-3⁺CD4 inhibition
- 41bb⁺ CD4 co-stimulation
- Ki67⁺ CD4 proliferation
- ICOS⁺ CD4 activation o ICOS+ PD-L1+ CD4 -
- activation/inhibition ICOS⁺ PD-1⁺ CD4 - activation/inhibition
- Total naïve (CCR7⁺CD45RA⁺) CD4
 - PD-L1⁺ naïve CD4 activation/inhibition PD-1⁺ naïve CD4 - activation/inhibition
 - CTLA-4⁺ naïve CD4 inhibition
 - Tim-3⁺ naïve CD4 inhibition
- Total central memory (CCR7⁺CD45RA⁻) 3. <u>Total Tregs</u> CD4
- PD-L1⁺ CM CD4 activation/inhibition
- PD-1⁺ CM CD4 activation/inhibition
- CTLA-4⁺ CM CD4 inhibition
- Tim-3⁺ CM CD4 inhibition
- o Ki67⁺ CM CD4 proliferation
- Total effector memory (CCR7⁻ CD45RA⁻) CD4
 - o PD-L1+ EM CD4 activation/inhibition
 - PD-1⁺EM CD4 activation/inhibition
 - CTLA-4⁺ EM CD4 inhibition Tim-3⁺ EM CD4 - inhibition
- o Ki67⁺ EM CD4 proliferation Total EMRA (CCR7-CD45RA⁺) CD4
- o PD-L1⁺ EMRA CD4 activation/inhibition
- o PD-1+EMRA CD4 activation/inhibition
- CTLA-4⁺ EMRA CD4 inhibition
- Tim-3⁺ EMRA CD4 inhibition
- o Ki67+ EMRA CD4 proliferation
- 2. Total CD8+ T cells

regulatory T cells.

- PD-L1⁺ CD8 activation/inhibition
- PD-1+CD8 activation/inhibition
- CTLA-4⁺ CD8 inhibition
- Tim-3⁺ CD8 inhibition
- 41bb⁺ CD8 co-stimulation
- Ki67⁺ CD8 proliferation
- Total naïve (CCR7+CD45RA+) CD8 o PD-L1+ naïve CD8 - activation/inhibition o PD-1+ naïve CD8 - activation/inhibition
- CTLA-4⁺ naïve CD8 inhibition
- o Tim-3+ naïve CD8 inhibition Total central memory (CCR7⁺CD45RA⁻)
- CD8
 - o PD-L1+ CM CD8 activation/inhibition
 - PD-1⁺ CM CD8 activation/inhibition
 - o CTLA-4+ CM CD8 inhibition

- Tim-3⁺ CM CD8 inhibition
- Ki67⁺ CM CD8 proliferation
- Total effector memory (CCR7⁻ CD45RA⁻) CD8
 - PD-L1⁺ EM CD8 activation/inhibition
- PD-1⁺EM CD8 activation/inhibition
- CTLA-4⁺ EM CD8 inhibition
- Tim-3⁺ EM CD8 inhibition
- o Ki67⁺ EM CD8 proliferation
- Total EMRA (CCR7-CD45RA+) CD8 PD-L1⁺ EMRA CD8
 - activation/inhibition
- PD-1⁺ EMRA CD8 activation/inhibition
- CTLA-4⁺ EMRA CD8 inhibition
- Tim-3⁺ EMRA CD8 inhibition
- o Ki67+ EMRA CD8 proliferation

- PD-L1⁺ Tregs activation/inhibition
- PD-1⁺ Tregs suppression
- CTLA-4⁺ Tregs suppression
- ICOS⁺ Tregs suppression
- CD45RA+ Tregs highly expandable in vitro
- CD49d Tregs suppression
- Ki67⁺ Tregs proliferation
- CD38⁺ Tregs suppression
- · HLA-DR+ Tregs suppression

4. Total B cells

- · PD-L1+ B cells activation/inhibition
- PD-1+B cells activation/inhibition

5. Total NK cells

- PD-L1⁺ NK inhibition
- PD-1⁺ NK activation/inhibition
- Tim-3⁺ NK activation/inhibition
- Ki67⁺NK proliferation
- NKp30⁺NK activation
- NKp46⁺ NK activation
- NKG2D⁺ NK activation
- CD226⁺ NK adhesion/activation
- Total mature (CD16⁺ CD56^{dim}) NK lytic PD-L1⁺ mature NK - inhibition
- PD-1⁺ mature NK activation/inhibition
- Tim-3⁺ mature NK activation/inhibition
- Total functional intermediate (CD16⁺
- CD56^{br}) NK lytic, cytokine production
- PD-L1⁺ functional intermediate NK inhibition
- PD-1⁺ functional intermediate NK activation/inhibition
- o Tim-3+ functional intermediate NK activation/inhibition
- · Total immature (CD16- CD56br) NK -
- cytokine production PD-L1⁺ immature NK - inhibition

cDC, conventional dendritic cells; CM, central memory; CTLA-4, cytotoxic T lymphocyteassociated protein-4; EM, effector memory; EMRA, terminally differentiated effector memory; FoxP3, forkhead box P3; gMDSCs, granulocytic myeloid derived suppressor cells; ICOS, inducible T cell co-stimulator; lin neg MDSCs, lineage negative MDSCs; mMDSCs, monocytic MDSCs; NK, natural killer; pDC, plasmacytoid DC; PD-1, programmed cell death-1; PD-L1, programmed cell death ligand-1; Tim-3, T cell immunoglobulin and mucin domain-3; Tregs,

- PD-1⁺ immature NK activation/inhibition
- Tim-3⁺ immature NK activation/inhibition
- Total unconventional (CD16⁻ CD56^{dim}) NK non-lytic, non-cytokine production
- PD-L1⁺ unconventional NK inhibition PD-1⁺ unconventional NK -
- activation/inhibition
- o Tim-3+ unconventional NK activation/inhibition

6. Total NK-T cells

7. Total cDC

8. Total pDC

9. Total MDSC

MDSC

10.Total Monocytes

 PD-L1⁺ NK-T - inhibition Tim-3⁺ NK-T - inhibition

· Ki67+ NK-T - proliferation

PD-L1⁺ cDC - inhibition

· Tim-3+ cDC - inhibition

Ki67⁺ cDC - proliferation

· PD-L1+ pDC - inhibition

Tim-3⁺ pDC - inhibition

Ki67⁺ pDC - proliferation

• PD-L1⁺ MDSC - inhibition

· PD-1+ NK-T - activation/inhibition

PD-1⁺ cDC - activation/inhibition

· PD-1+ pDC - activation/inhibition

PD-1⁺ MDSC - activation/inhibition

PD-L1⁺ mMDSC - inhibition

o PD-L1+ gMDSC - inhibition

CD16⁺ MDSC - immature/suppression

Total monocytic (CD14⁺ CD15⁻) MDSC

o PD-1+ mMDSC - activation/inhibition

Total granulocytic (CD14⁻ CD15⁺) MDSC

o PD-1+ gMDSC - activation/inhibition

Total lineage negative (CD14⁻ CD15⁻)

○ PD-L1⁺ lin neg MDSC - inhibition

PD-1⁺ lin neg MDSC -

activation/inhibition

o CD16+ lin neg MDSC -

· Intermediate Monocytes

immature/suppression

· Classical Monocytes - phagocytic

Non-Classical Monocytes - proinflammatory

· PD-1+ Monocytes - activation/inhibition

phagocvtic/proinflammatorv

PD-L1⁺ Monocytes - inhibition

CD16⁺ gMDSC - immature/suppression

CD16⁺ mMDSC - immature/suppression

Supplemental Table 2. Effect of treatment on classic and refined PBMC subsets

Classic Subset	Direction of Change	Pre (median)	D29 (median)	P value	# Increasing >25%	# Decreasing >25%
CD4	=	34.24	36.99	0.95	3/8	1/8
CD8	=	11.34	12.74	0.95	1/8	1/8
Treg	=	0.98	0.65	0.55	3/8	3/8
NK	=	6.18	8.11	0.08	4/8	1/8
NK-T	=	0.92	1.29	0.08	2/8	1/8
B cells	=	10.92	9.48	1.00	3/8	2/8
cDC	=	0.29	0.32	0.38	4/8	2/8
pDC	=	0.16	0.22	0.31	4/8	2/8
MDSC	=	7.76	8.05	0.64	2/8	4/8
Monocytes	=	14.43	16.4	0.15	1/8	4/8
Refined Subsets	Direction	Pre	D29	P value	# Increasing	# Decreasing
	of Change	(median)	(median)		>25%	>25%
CD4 ki67	1	0.59	0.97	0.02	6/8	0/8
CD4 CM ki67	1	0.17	0.31	0.04	5/8	0/8
NK Ki67	1	0.30	0.61	0.04	7/8	1/8
NK NKp30	\uparrow	0.70	1.32	0.01	8/8	0/8
NK Mature NKp30	1	0.60	1.11	0.01	8/8	0/8

A. SOC

B. SOC + IO

Classic Subset	Direction of Change	Pre (median)	D29 (median)	P value	# Increasing >25%	# Decreasing >25%
CD4	=	32.57	27.71	0.03*	0/11	1/11
CD8	=	13.31	14.17	0.04*	4/11	0/11
Treg	=	0.62	0.48	0.05	2/11	4/11
NK	=	9.87	9.65	0.64	3/11	1/11
NK-T	=	2.94	4.67	0.05	6/11	1/11
B cells	=	7.37	6.31	0.03*	1/11	5/11
cDC	=	0.48	0.46	0.70	5/11	3/11
pDC	=	0.1	0.21	0.07	8/11	1/11
MDSC	=	7.21	8.64	0.47	5/11	3/11
Monocytes	=	16.49	15.09	0.58	4/11	3/11
Refined Subsets	Direction	Pre	D29	P value	# Increasing	# Decreasing
	of Change	(median)	(median)		>25%	>25%
CD8 ki67	1	0.28	0.76	0.003	9/11	0/11
CD8 EM ki67	\uparrow	0.15	0.50	0.003	9/11	0/11
NK ki67	1	0.46	0.81	0.02	8/11	1/11
Int Monocyte	1	0.48	0.84	0.01	9/11	0/11

Median frequency of PBMC subsets before and after 1 month of treatment with SOC (A) or SOC + IO (B). Subsets with a potentially biologically relevant change (bold type) were defined as those with p < 0.05, most patients having a > 25% change, and difference in medians > 0.05% of PBMCs.

*Although p<0.05, most patients did not change by > 25%.

Abbreviations: Int, intermediate; IO, immuno-oncology; SOC, standard of care.

Supplemental Table 3. Effect of treatment on serum cytokines/soluble factors and complete blood count parameters

Serum Factor	Direction of Change	Pre (median)	D29 (median)	P value	# Increasing >25%	# Decreasing >25%
sCD27	=	80.00	93.00	0.23	1/7	0/7
sCD40L	\checkmark	18.00	10.00	0.03	0/7	6/7
Ratio sCD27:sCD40L	1	5.000	8.000	0.03	6/7	0/7
sPD-L1	=	78.80	79.00	0.94	0/7	0/7
IL-8	=	26.30	20.90	0.69	3/7	3/7
TGFβ	\checkmark	27779	17646	0.02	0/7	6/7
IFNg	=	0.000	0.000	1.00	0/7	1/7
CBC Measure	Direction of Change	Pre (median)	D29 (median)	P value	# Increasing >25%	# Decreasing >25%
ALC	=	1.440	1.170	0.03*	0/7	2/7
NLC	\checkmark	4.850	2.130	0.02	0/7	7/7
NLR	\checkmark	3.292	1.821	0.02	0/7	6/7
Platelet	\checkmark	327.0	185.0	0.02	0/7	7/7

A. SOC

B. SOC + IO

Serum Factor	Direction of Change	Pre (median)	D29 (median)	P value	# Increasing >25%	# Decreasing >25%
sCD27	=	104.5	129.0	0.25	4/10	1/10
sCD40L	\checkmark	23.50	16.50	0.02	0/10	6/10
Ratio sCD27:sCD40L	1	5.000	8.000	0.03	7/10	0/10
IL-8	=	29.65	18.10	0.32	3/10	5/10
TGFβ	\checkmark	32893	21308	0.002	0/10	6/10
IFNg	=	0.000	0.000	0.50	0/10	2/10
CBC Measure	Direction of Change	Pre (median)	D29 (median)	P value	# Increasing >25%	# Decreasing >25%
ALC	=	1.535	1.635	0.81	1/8	2/8
NLC	=	4.960	2.745	0.06	1/8	5/8
NLR	=	2.746	2.015	0.46	2/8	4/8
Platelet	=	323.0	223.5	0.004*	0/10	4/10

Median frequency of measure before and after 1 month of treatment with SOC (A) or SOC + IO (B). Factors with a potentially biologically relevant change (bold type) were defined as those with p < 0.05, and most patients having a > 25% change.

*Although p<0.05, most patients did not change by > 25%. sPD-L1, TGF- β , IFN- γ , and IL-8 were measured in pg/mL, sCD27 in U/mL, and sCD40L in ng/mL.

Abbreviations: IO, immuno-oncology; SOC, standard of care; SOC + IO, AdCEA vaccine plus avelumab.

Supplemental Table 4. Frequency of patients with anti-Ad5 neutralizing antibodies pre- and post-SOC + IO treatment

Anti-Ad-5 Neutralizing Antibody Titer	Pre	~D29	~D56	~D111	~D167
Positive at <u>></u> 1:100	0/10	4/9	6/10	5/6	5/5
serum dilution	(0%)	(44%)	(60%)	(83%)	(100%)
Positive at <u>></u> 1:500	0/10	4/9	4/10	2/6	1/5
serum dilution	(0%)	(44%)	(40%)	(33%)	(20%)
Positive at <u>></u> 1:1,000 serum dilution	0/10	2/9	2/10	1/6	1/5
	(0%)	(22%)	(20%)	(17%)	(20%)

Abbreviations: IO, immuno-oncology; SOC, standard of care; SOC + IO, AdCEA vaccine plus avelumab.