

Version 1.5

**Study #:** 20-614

**Study ID:** Generation and characterization of T cells expressing PACT391C

**Responsible Scientist:** Andrew Conroy

**Date:** 30th November, 2020

Study Objective:

The goal of this study is to generate and characterize T cells expressing TCRs pulled down from patient PACT391C by the imPACT technique of the study of the

tetramer group. The edited T cells were assessed for TCR expression and IFNy secretion in the comPACT coating assay.

Study Sheet Initiated by: Oghene Efagene / 30th November, 2020

Report generated by: Pankaj Tiwari / 2nd July, 2021

Initial Study reviewed by: Study Report Approval / 14th March, 2021Final Study reviewed by: Study Report Approval / 9th August, 2021

Day 0: Activation
Day 3:Nucleofection

Day 4-14: Cytokine Treatment and Expansion

**Readouts:** Gene editing Dextramer binding

IFNγ secretion in the comPACT coating assay

#### 1.0 MATERIALS AND METHODS

#### **Reagent IDs:**

Reagent ID	Reagent	Description	Vendor/Study/ Experiment	Container	
PACT497	Cells	CD4/CD8 cells from Prodigy	EXP20002315	N/A	
p001-020	HR DNA	pUCu-Kan TRAC(1k)_P2A.Neo12TRBC2opt.f-P2A.TRA(Va)opt	Nature Technology	ep15236	
rnp001-063	RNPα	TRAC-1.Aldevron.sNLS-SpCas9-sNLS.Research Grade	20-614 /EXP20000119	ep15411	
rnp002-060	RNPβ	TRBC-2.Aldevron.sNLS-SpCas9-sNLS.Research Grade	20-614 /EXP20000119	ep15412	

#### **Study Groups:**

On day 0, cells were activated with TransACT in the presence of IL-7 and IL-15 (12.5 ng/mL each). On Day 2, cells were transfected.

Grp	n	Cells	Cuvette/Condition	<b>Transfection Conditions</b>	Expansion
1	1	7000000.00	Lonza X-Unit 100 µL	Neo12 + RNPα + RNPβ	24-well G-rex
2	1	7000000.00	Lonza X-Unit 100 μL	p959-PACT391C_TCR792 + RNPα + RNPβ	24-well G-rex
3	1	7000000.00	Lonza X-Unit 100 µL	p960-PACT391C_TCR793A01 + RNPα + RNPβ	24-well G-rex
4	1	7000000.00	Lonza X-Unit 100 μL	p961-PACT391C_TCR793A02 + RNPα + RNPβ	24-well G-rex

### **Experiment List:**

Date	Experiment	Description	Initials
30th November, 2020	20002249	[PD-010] Receive isolated T cells and T cell activation	OE
3rd December, 2020	20002250	[PD-010] Nucleofection	OE
7th December, 2020	20002293	PACT391C: neoTCR T cell comPACT Dextramer screening	TM
7th December, 2020	20002294	PACT391C: comPACT coating assay for cytokine secretion	TM

### **Key Identifiers:**

Clinical Study ID	Clinical Patient ID	Patient Date of Birth	PACT ID		
PACT-0101	0401	20th March, 1974	PACT391C		

#### **Notes:**

This test conclusion.

1



Version 1.5

#### 2.0 RESULTS:

2.1 Dextramer Screening:

Table 1: Summary of the dextramer/2a data for all the neoTCRs tested

Plasmid-PACT-TCR Name	Matched comPACT ID tested	neoE Sequence	Gene	% Gene Edited (2a+) CD8 T cells	% Gene Edited (2a+) CD 4 T cells	% cognate comPACT Dex+2a+CD8 T cells	% cognate comPACT Dex+2a+CD4 T cells	Pass CD8 binding	Pass CD4 binding	% mis match comPACT Dex + 2a + CD 8 T cells	% mis match comPACT Dex + 2a + CD 4 T cells
N eo1 2	PRO004	YLYHRVDVI	USP7	45.1	34.4	44.8	33.6	Yes	Yes	0.7	0.0
p959-PACT391C_TCR792	comPACT28036	SNNPLFQYTY	RNF145	37.2	34.2	35.9	20.3	Yes	Yes	0.8	0.0
p960-PACT391C_TCR793A01	comPACT27877	IQIETVLPAE	NUP210	14.8	11.9	0.7	0.1	No	No	1.0	0.0
p961-PACT391C_TCR793A02	comPACT27877	IQIETVLPAE	N UP 2 1 0	13.8	9.7	0.7	0.0	No	No	0.8	0.1

#### **Results Summary:**

 $3\ \text{of}\ 3\ \text{PACT}\,391\text{C}\ \text{neoT}\,\text{CRs}\ \text{received}$  were tested in the functional assays.

- 1 of the 3 PACT 391C TCRs tested in this study exhibited dextramer binding in the CD4 T cell population (CD8-independent binding): TCR792.
- 2 of the 3 neoTCRs was negative for specific binding to CD8 and CD4 T cells: TCR793A01,TCR793A02

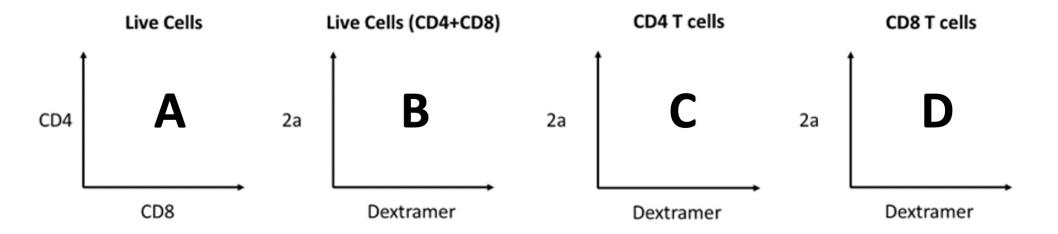
**Definition of CD8-independent TCR:** % Dextramer binding to CD4 T cells has to be at least 50% of Dextramer staining observed on CD8 T cells stained under same conditions for the TCR to pass this criterium.



Version 1.5

2.2 Dextramer Screening Raw data:

#### Schema of the flow plots for Figures 1 to 4:



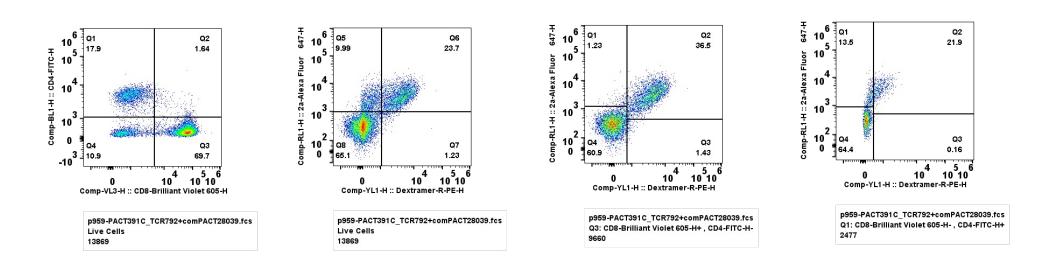
For the above gating schema, the "Live Cells" gate (plots A and B) are derived from parental gates (not shown) "Intact T cells  $\rightarrow$  doublet discrimination  $\rightarrow$  live cells".

- Plot A illustrates the CD4 and CD8 composition of the live cell population.
- Plot B illustrates the dextramer and 2a peptide binding of the live cell population (combined CD4 and CD8 T cells).
- Plot C illustrates the dextramer and 2a peptide binding in the CD4 T cell population
- Plot D illustrates the dextramer and 2a peptide binding in the CD8 T cell population.

Version 1.5

### p959-PACT391C\_TCR792

### - Cognate comPACT: comPACT28039 (RNF145: SNNPLFQYTYL - HLA-A29:02) 11 mer



#### - Mismatch comPACT: comPACT6050 (ARHGAP44: TTDNMMLEFY- HLA-A29:02- HLA Allele Control)

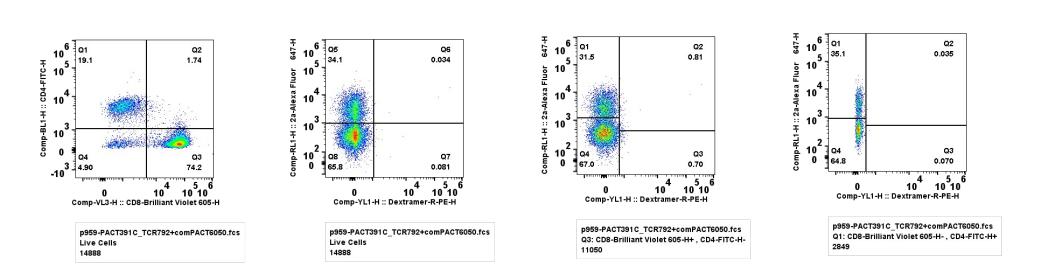
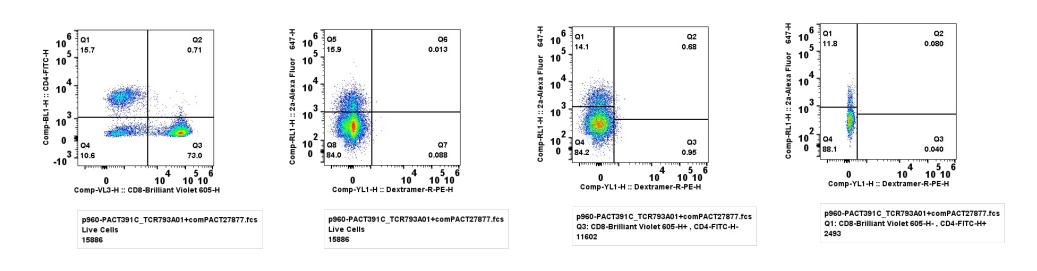


Figure 1: Flow plots for TCR792.

Version 1.5

#### p960-PACT391C TCR793A01

#### - Cognate comPACT: comPACT27877 (NUP210: IQIETVLPAE - HLA-B15:01) 10 mer



#### - Mismatch comPACT: comPACT1130 (ELF4: KQIFRTSEM- HLA-B15:01- HLA Allele Control)

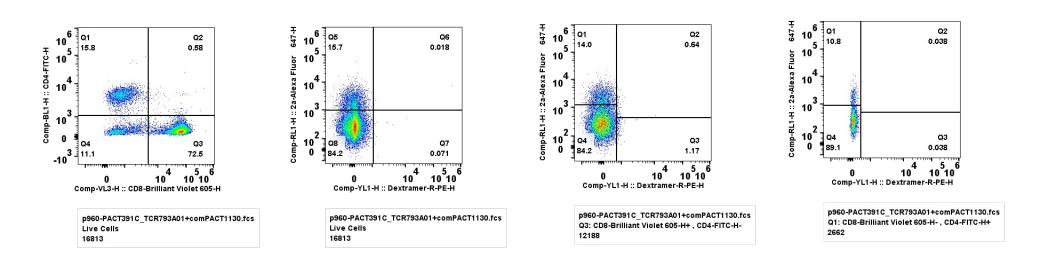
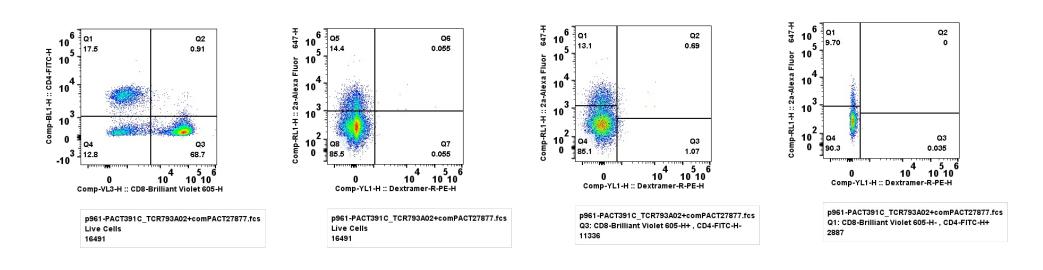


Figure 2: Flow plots for TCR793A01.

Version 1.5

### p961-PACT391C\_TCR793A02

#### - Cognate comPACT: comPACT27877 (NUP210: IQIETVLPAE - HLA-B15:01) 10 mer



#### - Mismatch comPACT: comPACT1130 (ELF4: KQIFRTSEM- HLA-B15:01- HLA Allele Control)

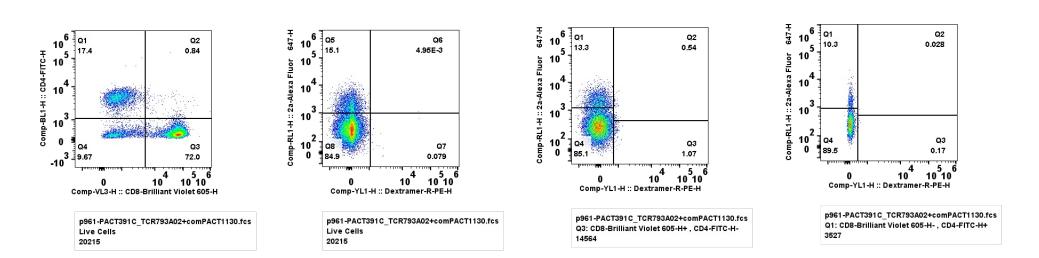


Figure 3: Flow plots for TCR793A02.



Version 1.5

### Neo12 Control TCR

#### - Cognate comPACT: PRO004 (USP7: YLYHRVDVI-HLA-A02:01) 9mer

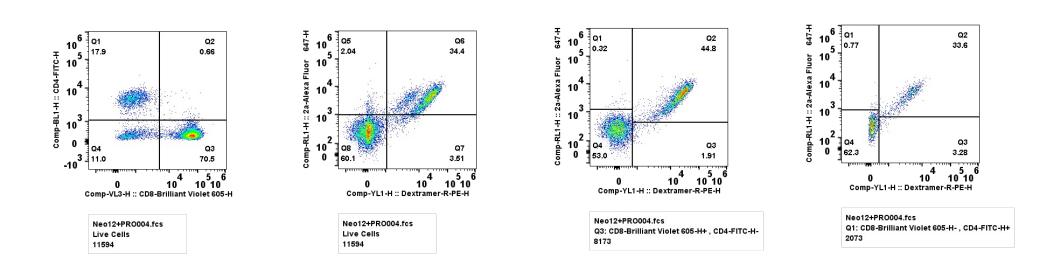


Figure 4: Flow plots for neo12 control TCR.

2.3 Results of functional characterization IFN $\gamma$  secretion

Table 2: Summary of the IFNy secretion

					IFNg(pg/10^3 edited cells)												
comPACT ID	comPACT Name	neoE Sequence	Gene	TCR ID	1000	333	111	37.0	12.3	4.12	1.37	0.46	0.15	0.05	0	EC50 ng/ML	Mismatch pass(Y/N)
PRO004		YLYHRVDVI	USP7	Neo12	25.9	24.7	25.4	25.3	24.6	21.5	15.4	5.1	0.6	0.0	0.0	1.1	Yes
comPACT28036	PACT391C_T_PP001370_O49_CP_HLA-A29:02	SNNPLFQYTY	RNF145	p959-PACT391C_TCR792	23.9	24.2	21.6	15.2	4.4	0.0	0.0	0.0	0.0	0.0	0.0	28.5	Yes
comPACT27877	PACT391C_T_PP001370_O150_CP_HLA-B15:01	IQIETVLPAE	NUP210	p960-PACT391C_TCR793A01	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	ND	Yes
comPACT27877	PACT391C_T_PP001370_O150_CP_HLA-B15:01	IQIETVLPAE	NUP210	p961-PACT391C_TCR793A02	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	ND	Yes

 $(EC_{50} \text{ with cognate comPACT and whether IFN} \gamma \text{secretion was observed in presence of mismatched comPACT}) \text{ for all the neoTCR candidates tested.}$  ND = Not determined

Version 1.5

### Functional characterization (IFNysecretion) raw data:

Figures 5 to 7 illustrate the potency data graphs illustrating the amount of IFN $\gamma$  secreted by neoTCR expressing T cells stimulated with increasing concentration of the cognate comPACT or a single concentration of negative control mismatch comPACT.

IFNy secretion upon stimulation with comPACT27877 (NUP210: IQIETVLPAE - HLA-B15:01)

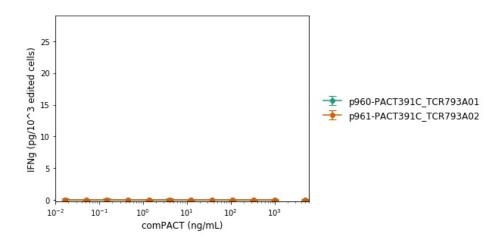


Figure 5: Potency data for TCR793A01, TCR793A02.

IFNy secretion upon stimulation with comPACT28036 (RNF145: SNNPLFQYTY - HLA-A29:02)

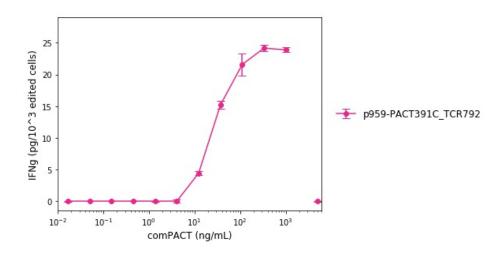


Figure 6: Potency data for Missing Data.

Version 1.5

#### Positive control

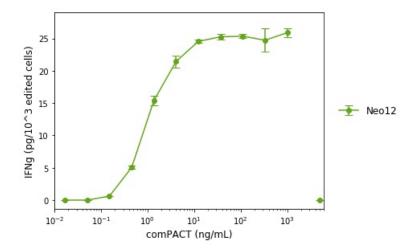


Figure 7: Potency data for the neoTCR positive control neo12.

#### **3.0 CONCLUSIONS**

Table 3: Summary of the dextramer screening data and the IFNy secretion potency assay.

Plasmid-PACTTCR Name	Matched comPACT ID tested	ne o E Se que nce	Gene	% Gene Edited (2a+) CD8 T cells	% Gene Edited (2a+) CD4 T cells	% cognate comPACT Dex+2a+CD8 T cells	% cognate comPACT Dex+2a+CD4 T cells	Pass CD8 binding	Pass CD4 binding	EC50 ng/ML	Pass mismatch
N eo1 2	PRO004	YLYHRVDVI	USP7	45.1	34.4	44.8	33.6	Yes	Yes	1.1	Yes
p959-PACT391C_TCR792	comPACT28036	SNNPLFQYTY	RNF145	37.2	34.2	35.9	20.3	Yes	Yes	28.5	Yes
p960-PACT391C_TCR793A01	comPACT27877	IQIETVLPAE	NUP210	14.8	11.9	0.7	0.1	No	No	ND	Yes
n961-PACT391C TCR793A02	comPACT27877	IOIETVI.PAE	NUP210	13.8	9.7	0.7	0.0	No	No	ND	Yes

#### The summary of the functional screening results for PACT-0101 0401 (PACT391C) are as follows:

3 out of the 3 PACT 391C neoTCRs were screened in both the dextramer binding and potency assays. Among the 3 neoTCRs screened:

- 1/3 passed all functional criteria for selection as TCR: CD8 independent dextramer binding and IFN $\gamma$  secretion.
- 2/3 neoTCRs did not meet criteria for either cognate dextramer binding or IFN $\gamma$  secretion.

The neo12 control neoTCR performed as expected in both the dextramer binding and IFN  $\!\gamma$  secretion assays.

### **Supporting Documents:**

• 2541372\_163633b3-89a9-4994-8b68-21f9b3cfcaf8.pdf