

Phase I trials of personalized cancer vaccine TG4050 in surgically treated high-risk head and neck squamous cell carcinoma (HNSCC) and relapsing ovarian cancer (OvC) patients

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BACKGROUND

Cancer mutations acquired during carcinogenesis are not subject to central tolerance and thus are immunologically attractive targets for vaccination.. The use of mutationderived neoepitopes to design cancer vaccines is an unprecedented opportunity to target efficient antigens, but the stochastic occurrence of mutations implies that a tailored vaccine needs to be designed and manufactured for each patient. We used next generation sequencing, machine learning based genomic data analysis and advances in genome editing to systematically identify private neoepitopes in high-risk HPV negative HNSCC and relapsing OvC patients.

METHODS

Mutations were identified by DNA and RNA sequencing and putative epitopes were selected for vaccine design based on their likelihood to elicit a class I or II response using data on HLA binding, allelic frequency, prediction of processing and expression at RNA level. The vaccine was a recombinant virus of the modified Vaccinia Ankara (MVA) strain encoding for up to 30 sequences of 29 amino acids, centered on the predicted epitopes and flanked with the normal germline sequence in case of missense mutations, and of the mutated protein sequence in case of frameshift mutations. The viral vaccine was amplified under GMP conditions and administered to patients after completion of SOC treatment given with curative intent in two phase I trials. Monotherapy with the vaccine was started for OvC patients at relapse defined as elevation of CA-125 or onset of suspicious radiological findings and for locally advanced HNSCC patients following upfront surgery and adjuvant therapy. Head and Neck patients were randomized to receive the vaccine immediately after first line treatment in monotherapy (Arm A) or at relapse (Arm B) in conjunction with standard of care. In both studies, the vaccine was administered weekly for 6 weeks and a booster dose every three weeks over a year.

TRANSLATIONAL STUDY PLAN Plasma, Serum and T-cell response ex vivo ELISPOT

DESIGN OF PATIENT SPECIFIC VACCINE

HLA Typing

Variant Calling

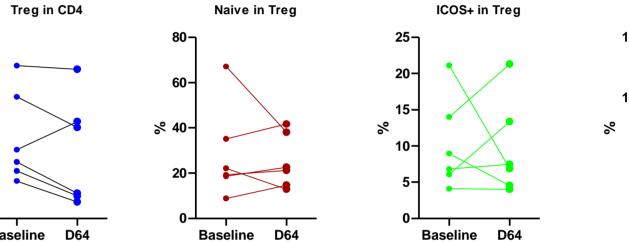
Sequencing data

RNA -

CLINICAL FOLLOW-UP

PHENOTYPING OF CIRCULATING CELLS PBMC were analysed by flow cytometry to quantify relevant immune cell subpopulations, in particular monocytes, DC, NK cells, subcells of CD8, CD4, Treg and expression of immune checkpoints ICOS and PD1. They were no striking changes in the absolute counts of cell populations; but an

cells was observed for CD4 and CD8 T cells in all patients between baseline and D64. Additionally, CD16⁻ subset in CD56^{dim} NK cells increased in all patients over the treatment period.



STUDY POPULATION

Ovarian cancer patients

Key Inclusion Criteria

- Stage IIIC or stage IVA (FIGO staging) high grade serous ovarian, fallopian or primary peritoneal carcinoma
- Complete response maintained at least 6 months after debulking surgery and first-line chemotherapy
- Asymptomatic relapse (elevated CA-125 or radiological findings)
- ECOG Performance status 0 or 1

Key Exclusion Criteria

- Prior exposure to cancer immunotherapy including anti-cancer vaccines, any antibody targeting T cell co-regulatory proteins such as anti-PD L1, anti-PD 1, or anti-CTLA-4 antibodies
- Chronic treatment with systemic corticosteroids

HNSCC patients

Key Inclusion Criteria

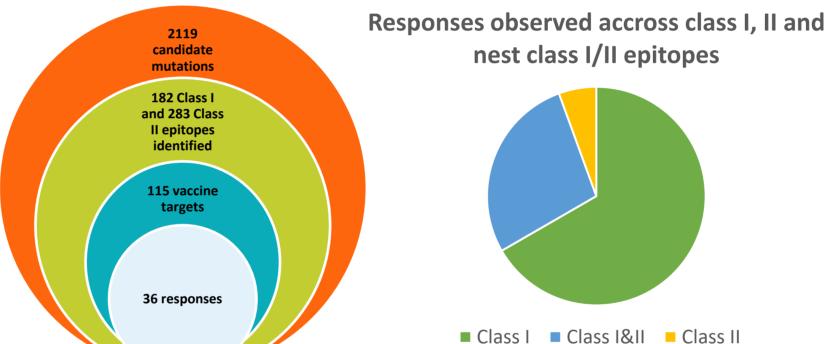
- Newly diagnosed stage III or IVA squamous-cell carcinoma of the oral cavity, oropharynx, hypopharynx or larynx eligible for gross total resection and adjuvant therapy
- Complete response 3 months after completion of adjuvant therapy
- ECOG Performance status 0 or 1

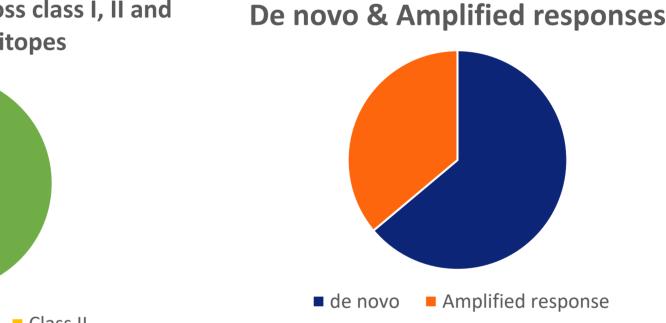
Key Exclusion Criteria

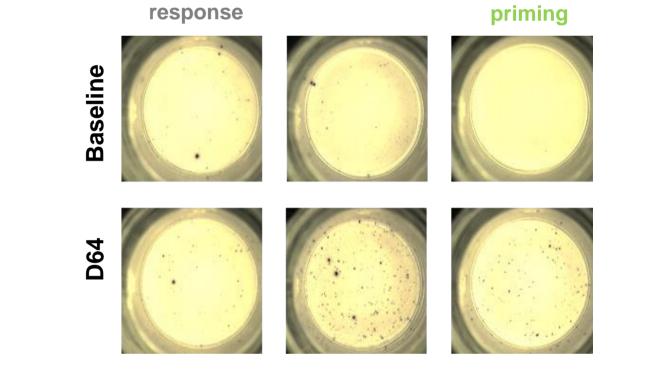
- HPV-positive oropharynx primaries, carcinoma of the nasopharynx, squamous cell-carcinoma of unknown primary, squamous cell carcinoma that originates from the skin and salivary gland or paranasal sinus, nonsquamous histologies
- Prior exposure to cancer immunotherapy including anti-cancer vaccines, any antibody targeting T cell co-regulatory proteins such as anti-PD L1, anti-PD 1, or anti-CTLA-4 antibodies
- Chronic treatment with systemic corticosteroids

ADAPTIVE T-CELL RESPONSES BY EX VIVO ELISPOTS IFNV

increase in the percentage of effector cells and a corresponding reduction in naïve T

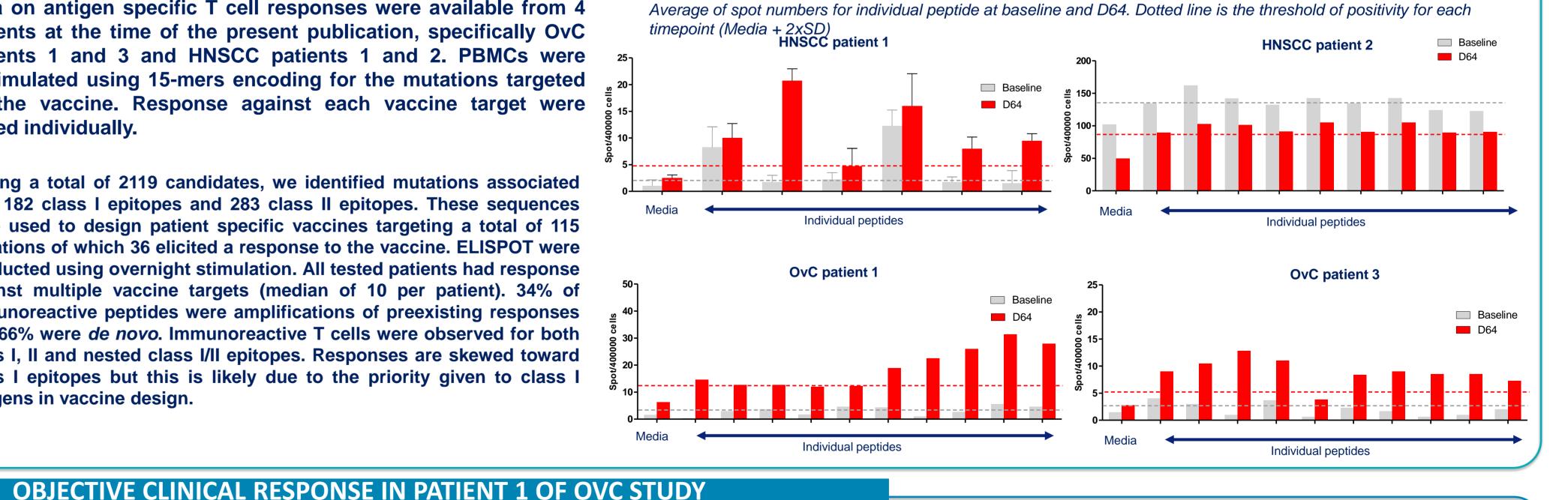






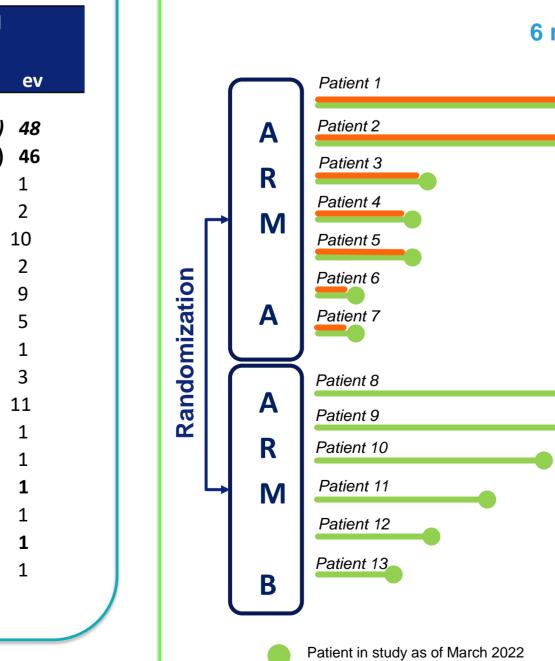
Data on antigen specific T cell responses were available from 4 patients at the time of the present publication, specifically OvC patients 1 and 3 and HNSCC patients 1 and 2. PBMCs were restimulated using 15-mers encoding for the mutations targeted by the vaccine. Response against each vaccine target were tested individually.

Among a total of 2119 candidates, we identified mutations associated with 182 class I epitopes and 283 class II epitopes. These sequences were used to design patient specific vaccines targeting a total of 115 mutations of which 36 elicited a response to the vaccine. ELISPOT were conducted using overnight stimulation. All tested patients had response against multiple vaccine targets (median of 10 per patient). 34% of immunoreactive peptides were amplifications of preexisting responses and 66% were de novo. Immunoreactive T cells were observed for both class I, II and nested class I/II epitopes. Responses are skewed toward class I epitopes but this is likely due to the priority given to class I antigens in vaccine design.





	Ovarian cancer (N=4)		HNSCC (N=6)		Overall	
					(N=10)	
	N (%)	ev	N (%)	ev	N (%)	ev
Patient with at least one AE related to TG4050	4 (100.0%)	29	6 (100.0%)	19	10 (100.0%)	48
GENERAL DISORDERS AND ADMINISTRATION SITE CONDITIONS	4 (100.0%)	28	6 (100.0%)	18	10 (100.0%)	46
Chills	1 (25.0%)	1	0 (0.0%)	0	1 (10.0%)	1
Fatigue	1 (25.0%)	1	1 (16.7%)	1	2 (20.0%)	2
Injection Site Erythema	3 (75.0%)	8	1 (16.7%)	2	4 (40.0%)	10
Injection Site Induration	0 (0.0%)	0	1 (16.7%)	2	1 (10.0%)	2
Injection Site Inflammation	0 (0.0%)	0	4 (66.7%)	9	4 (40.0%)	9
Injection Site Mass	1 (25.0%)	5	0 (0.0%)	0	1 (10.0%)	5
Injection Site Nodule	1 (25.0%)	1	0 (0.0%)	0	1 (10.0%)	1
Injection Site Oedema	0 (0.0%)	0	3 (50.0%)	3	3 (30.0%)	3
Injection Site Pain	2 (50.0%)	10	1 (16.7%)	1	3 (30.0%)	11
Injection Site Pruritus	1 (25.0%)	1	0 (0.0%)	0	1 (10.0%)	1
Injection Site Swelling	1 (25.0%)	1	0 (0.0%)	0	1 (10.0%)	1
NVESTIGATIONS	0 (0.0%)	0	1 (16.7%)	1	1 (10.0%)	1
Blood Alkaline Phosphatase Increased	0 (0.0%)	0	1 (16.7%)	1	1 (10.0%)	1
MUSCULOSKELETAL AND CONNECTIVE TISSUE DISORDERS	1 (25.0%)	1	0 (0.0%)	0	1 (10.0%)	1
Musculoskeletal Chest Pain	1 (25.0%)	1	0 (0.0%)	0	1 (10.0%)	1

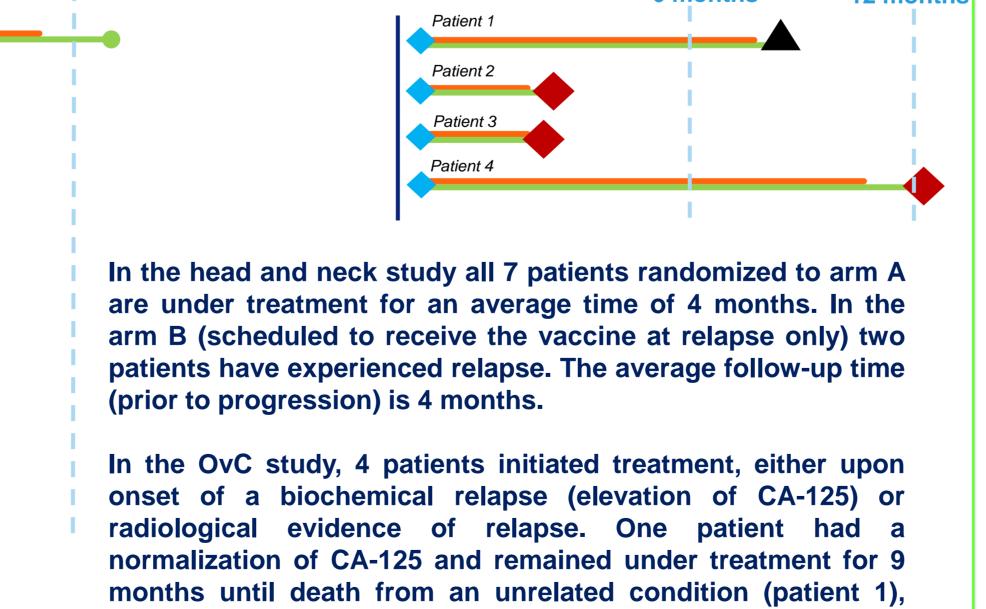


Relapse/Progressive disease

Death from non-cancer cause

Asymptomatic relapse

HNSCC patients



Ovarian cancer patients

DATA PROCESSIN

ARTIFICIAL

INTELLIGENCE

Selection of 30 neoantigens

RNA Expression

GENE EDITING

MANUFACTURING

Vaccine

generation and

Vaccine

NEC'S NEOANTIGEN PREDICTION

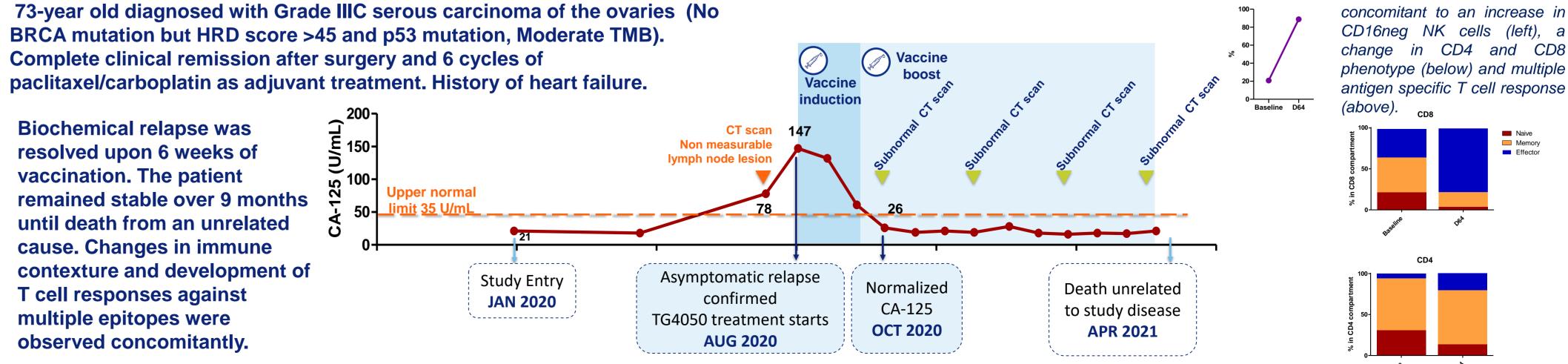
Epitope Ranking

NEC

Overall Ranking

relational learning

paclitaxel/carboplatin as adjuvant treatment. History of heart failure. **Biochemical relapse was** resolved upon 6 weeks of vaccination. The patient remained stable over 9 months until death from an unrelated cause. Changes in immune contexture and development of T cell responses against multiple epitopes were observed concomitantly.



ACKNOWLEDGEMENTS

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CONCLUSIONS

The viral vectored personalized vaccine approach was feasible: relevant targets could be identified in all patients, and time of manufacturing and drug release was compatible with the clinical course of treatment. Administration of the vaccine was safe and induced tumor specific T cell response against multiple targets. Early signs of clinical activity are encouraging with changes of tumor markers observed in a treated patient under vaccine monotherapy.

another patient was treated upon onset of radiological

evidence of relapse and was stable for 11,4 months (patient 4).

Patients 2 and 3 were treated upon onset of abnormal

radiological findings and were progressors at D43.