

S. S. Agarwala<sup>1</sup>, S. O'Day<sup>2</sup>, Y. Zakharia<sup>3</sup>, B. Voorhies<sup>4</sup>, M. Milhem<sup>3</sup>; <sup>1</sup>Medical Oncology, St. Luke's Hospital & Health Network, Bethlehem, PA, United States of America, <sup>2</sup>Medical Oncology, John Wayne Cancer Institute, Santa Monica, CA, United States of America, <sup>3</sup>Medical Oncology, University of Iowa Hospitals and Clinics, Iowa City, IA, United States of America, <sup>4</sup>Huntsman Cancer Institute, University of Utah, Utah, UT, United States of America

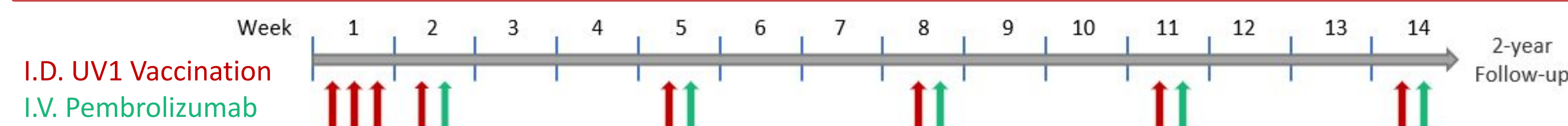
**BACKGROUND:**

**UV1** (developed by Ultimovacs ASA) consists of three peptides (15,15 and 30 amino acids) representing fragments of the human reverse transcriptase subunit of telomerase (hTERT). Telomerase activation is the major mechanism implicated in human cell immortalization and cancer cell pathogenesis [1]. Telomerase is expressed in all cancer cells at every stage of tumor evolution, from the cancer stem cell to circulating tumor cells. A CD4+ Th1 response against telomerase has recently been implicated as a positive prognostic factor in cancer [2]. Thus, telomerase represents a unique cancer antigen as a basis for immunotherapy [3]. UV1 contains both CD4 and CD8 epitopes and has been shown to be immunogenic in 78% (40/52) of HLA unselected patients across three completed phase I studies. The vaccine mainly induces Th1 reactivity (i.e. secretion of IFN-γ, TNFα, and IL-2), and an immune response against the UV1 peptides is associated with epitope spreading within hTERT and prolonged survival [4].

**Study rationale:** Efficacy of pembrolizumab depends on the presence of spontaneously induced T cell responses against relevant tumor antigens. Patients who lack or have few T cells in their tumor are less likely to obtain durable benefit from pembrolizumab alone. UV1 has the potential to increase the efficacy of pembrolizumab by amplifying the pool of hTERT-specific, tumor-reactive CD4 T cells and by increasing the breadth and diversity of the tumor-reactive T cell response (epitope spread). Reciprocally, the efficacy of UV1 vaccination may be enhanced since pembrolizumab augments the effector activity of UV1 induced T cells that are otherwise restricted by intrinsic immune regulatory and tumor induced suppressor mechanisms. Thus, the addition of UV1 to pembrolizumab has the potential to produce synergistic immunological activity which may transfer into increased clinical benefit compared to pembrolizumab monotherapy.

**TRIAL DESIGN:**

This is an ongoing Phase I, open-label, multicenter study investigating UV1 vaccination in combination with pembrolizumab in 30 patients with untreated unresectable or metastatic melanoma. Patients receive 8 UV1 vaccinations (300 μg) with GM-CSF as adjuvant. 20 patients will receive 37.5 μg while 10 patients will receive 75 μg GM-CSF. Five doses of pembrolizumab are given during the UV1 treatment. Pembrolizumab continues per label after completed UV1 treatment.



**KEY ENTRY CRITERIA:**

**Key Inclusion Criteria**

- Unresectable histological confirmed malignant melanoma (Stage IIIB-C, IV)
- Measurable and evaluable disease per RECIST v.1.1
- Previous untreated and eligible for pembrolizumab (prior BRAF and MEK inhibitors permitted)
- ECOG 0-1

**Key Exclusion Criteria**

- Uveal or ocular melanoma
- Prior therapy with anti-CTLA4, anti-PD-L1/L2 or oncolytic virus
- Active brain metastases (MRI)

**OBJECTIVES:**

**Primary**

- Safety and tolerability of the combination of UV1 and pembrolizumab

**Secondary**

- Immunological response against the vaccine
- Progression Free Survival (PFS) (RECIST1.1 and iRECIST)
- Overall Survival (OS)
- Duration of pembrolizumab treatment
- Time to subsequent anticancer therapies

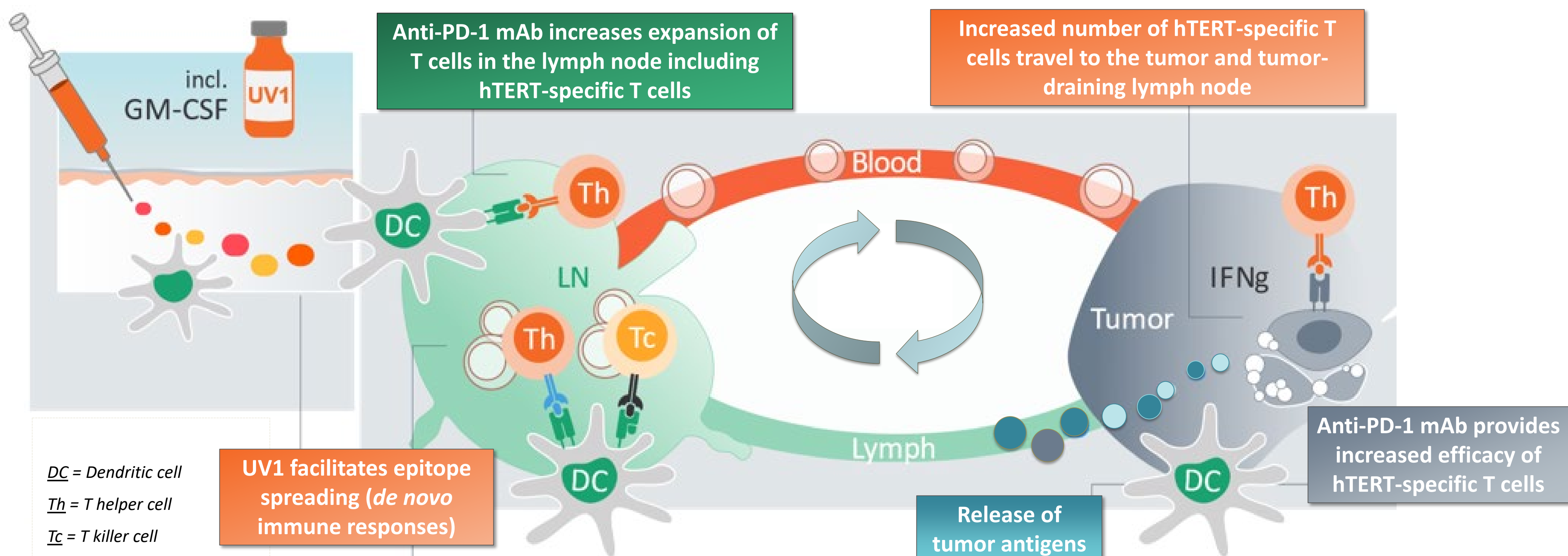
**SUMMARY:**

- The study is open and actively accruing at sites in the US
- Accrual is expected to be completed early Q12020
- For additional information please contact [wenche.rasch@ultimovacs.com](mailto:wenche.rasch@ultimovacs.com)

**REFERENCES:**

- 1.Hanahan, D. and R.A. Weinberg, Hallmarks of cancer: the next generation. Cell, 2011. 144(5): p. 646-74.
- 2.Laheurte, C., et al., Distinct prognostic value of circulating anti-telomerase CD4(+) Th1 immunity and exhausted PD-1(+)/TIM-3(+) T cells in lung cancer. Br J Cancer, 2019. 121(5): p. 405-416.
- 3.Zanetti, M., A second chance for telomerase reverse transcriptase in anticancer immunotherapy. Nat Rev Clin Oncol, 2017. 14(2): p. 115-128.
- 4.Inderberg-Suso, E.M., et al., Widespread CD4+ T-cell reactivity to novel hTERT epitopes following vaccination of cancer patients with a single hTERT peptide GV1001. Oncoimmunology, 2012. 1(5): p. 670-686.

**PROPOSED MECHANISM OF SYNERGY BETWEEN IMMUNE ACTIVATION AND CHECKPOINT INHIBITION**



DC = Dendritic cell  
 Th = T helper cell  
 Tc = T killer cell  
 IFNγ = Interferon gamma  
 LN = Lymph node

**UV1 facilitates epitope spreading (de novo immune responses)**

**Relevance of the hTERT-specific CD4 cells**

- hTERT is present throughout the tumorigenesis and the tumor cannot escape from dependence on telomerase.
- Due to the temporo-spatial presence of the target, the CD4 T cells induced by UV1 can provide a pro-immunological environment in the tumor and tumor-draining lymph node, continuously and over time, regardless of the genetic makeup of the individual tumor cell.
- The vaccine mediated CD4-immune response optimize for priming of de novo immune responses (epitope spreading) and tumor cell killing in an otherwise highly dynamic tumor.

**TRANSLATIONAL RESEARCH OBJECTIVES:**

- Correlation between UV1 specific immune responses in vitro (proliferation/ELISPOT) and in vivo (DTH\*)
- Correlation between exploratory biomarkers and clinical response
- Whole-exome sequencing for TMB estimation and neoantigen prediction
- Evolution of the TCR repertoire in response to treatment by TCR sequencing of PBMCs
- Correlation of mutations seen on liquid and solid tissue biopsies\*\*
- Correlation of fecal microbiome and response to treatment

\*DTH = Delayed type hypersensitivity \*\* Cell-free DNA in blood will be sequenced and analyzed for tumor-specific mutations