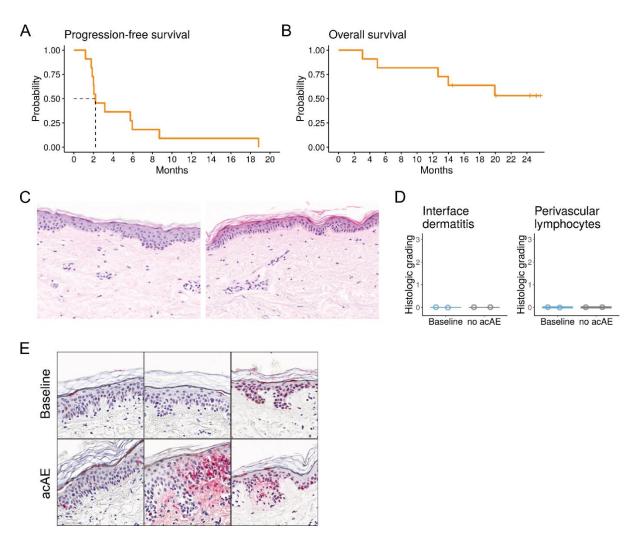
Supplementary Figure 1



A) Kaplan-Meier curve of progression-free survival (PFS) and **B)** overall survival of patients with uveal melanoma treated with tebentafusp (n=11). Tick marks indicate censoring. **C)** Representative histology images of on-treatment skin biopsies from patients without acAE. **D)** Histologic grading of interface dermatitis and perivascular lymphocytes in baseline and non-acAE on-treatment skin samples (2 patients, paired). **E)** Representative TUNEL (Terminal deoxynucleotidyltransferase mediated dUTP Nick End Labeling) staining of baseline and acAE skin samples.

Α В Vascular EC Fibroblasts T cells Macrophage/DC Pericytes Keratinocytes Melanocytes Smooth muscle cells Lymphatic EC Glial cells Glial cells Percent Expressed Lymphatic EC . 0 • 25 • 50 • 75 Smooth muscle cells UMAP_2 Melanocytes Keratinocytes Average Expression

2.5
2.0
1.5
1.0
0.5
0.0 Pericytes Macrophage/DC T cells Fibroblasts -10 Vascular EC -UMAP_1

15

16

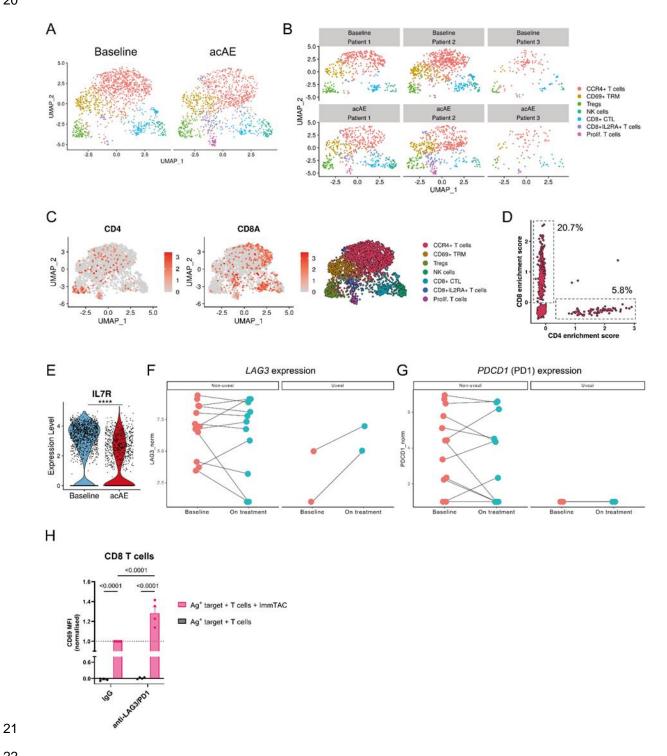
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- **A)** UMAP representation of all cell types in integrated baseline and acAE skin samples analyzed by single cell RNA-sequencing (3 patients, paired). **B)** Dotplot of lineage marker expression.
- 18 EC: Endothelial cells



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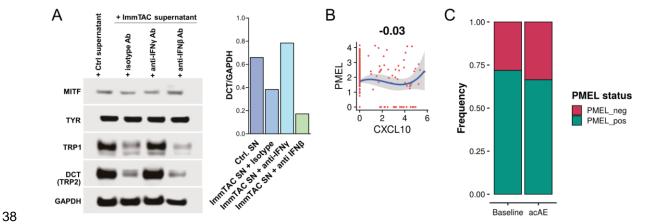
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A) UMAP of T and NK cell subclusterss split by timepoint. B) UMAP split by timepoint and patient. C) UMAP feature plot showing CD4 and CD8A expression. D) Feature scatter plot showing the percentage of CD4 and CD8A/CD8B expressing cells in the CCR4+ T cell-subcluster. E) Violin plot of IL7R

expression in T and NK cells on scRNA-seq. **F)** Log2-normalized expression of *LAG3* and **G)** *PDCD1* (PD1) in baseline and on treatment tumor biopsies from patients with uveal (n=2) or non-uveal (n=11) melanoma treated with tebentafusp, derived from a published RNA dataset (*Middleton, M.R., et al. (2020). Tebentafusp, A TCR/Anti-CD3 Bispecific Fusion Protein Targeting gp100, Potently Activated Antitumor Immune Responses in Patients with Metastatic Melanoma. Clin. Cancer Res. 26, 5869–5878.). H) CD8 T cell activation as measured by CD69 when co-cultured with antigen expressing target cells with or without ImmTAC and anti-LAG3/anti-PD1 antibodies or IgG isotype control.*

Ag: Antigen

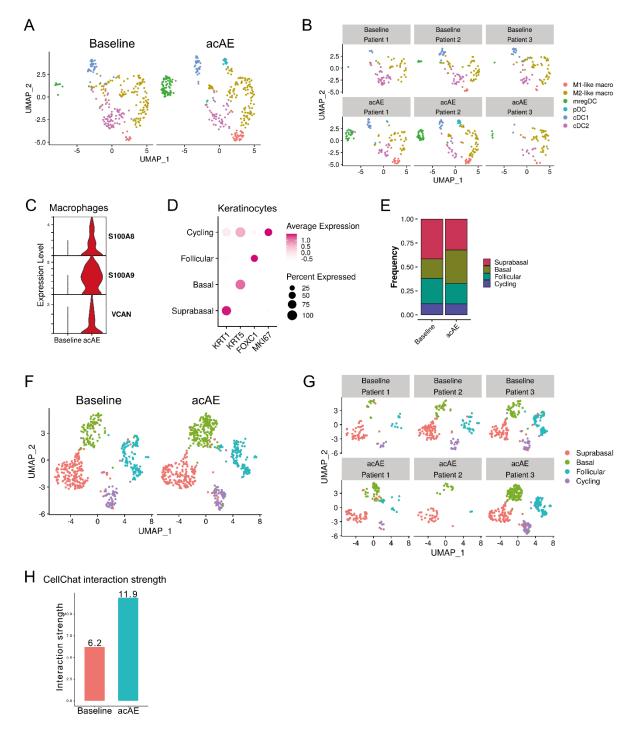


- -

A) WB of the melanocytic proteins MITF (Melanocyte Inducing Transcription Factor), TYR (Tyrosinase), TRP1 (Tyrosinase Related Protein 1), and DCT (Dopachrome Tautomerase) derived from melanocytes treated with supernatant derived from gp100-ImmTAC co-culture experiments with the addition of an isotype control, anti-IFN-γ, or anti-IFN-β antibody. The normalized DCT values are shown as barplots. **B)** Correlation of *PMEL* and *CXCL10* expression levels in melanocytes on scRNA-seq. Pearson correlation is indicated. **C)** Frequency of *PMEL*-positive melanocytes on scRNA-seq.

Ctrl: Control, Ab: Antibody, ImmTAC: Immune Mobilizing Monoclonal T cell Receptor Against Cancer, GAPDH: Glyceraldehyde-3-Phosphate Dehydrogenase, SN: Supernatant, acAE: acute cutaneous adverse event





A) UMAP of myeloid cell subclusters split by timepoint and **B)** split by timepoint and patient. **C)** Violin plot of *S100A8*, *S100A9* and *VCAN* expression in macrophages on scRNA-seq. **D)** Marker gene dotplot and **E)** cell type composition barplot of keratinocyte subclusters on scRNA-seq. **F)** UMAP of keratinocyte subclusters split by timepoint and **G)** split by timepoint and patient. **H)** Cell-cell communication interaction strength as inferred from CellChat analysis.

- 58 acAE: acute cutaneous adverse event, cDC1: classical dendritic cell type 1,cDC2: classical dendritic
- 59 cell type 2, mregDC: mature DC enriched in immunoregulatory molecules, pDC: plasmacytoid DC

Supplementary Table 1: Skin-related information

| Patient | Sex | Age | Cutaneous adverse events | Follow-up skin biopsy | Biopsy from vitiligo-like pigmentation disorder | Assays |
|---------|-----|-----|---|--------------------------|--|--------------------------------|
| 1 | m | 58 | Diffuse erythema, pruritus, vitiligo-like pigmentation disorder | Day 15 | Month 7 (tebentafusp treatment ongoing) | scRNA-seq mIHC histology |
| 2 | m | 74 | Diffuse erythema, pruritus, vitiligo-like pigmentation disorder | Day 8 | Month 12 (tebentafusp treatment ongoing) | scRNA-seq mIHC histology |
| 3 | f | 62 | Diffuse erythema, macular exanthema, facial edema, pruritus, vitiligo-like pigmentation disorder | Day 2 | | scRNA-seq mIHC histology |
| 4 | m | 57 | Maculopapular exanthema, partly diffuse erythema on face and neck, vitiligo-like pigmentation disorder | Day 2 | Month 8 (tebentafusp treatment ongoing) | mIHC histology |
| 5 | m | 66 | Diffuse erythema, pruritus, vitiligo-like pigmentation disorder | Day 15 | Month 9 (tebentafusp treatment ongoing) | mIHC histology |
| 6 | m | 56 | Diffuse erythema, vitiligo-like pigmentation disorder | Day 8 | Month 7 (tebentafusp treatment ongoing) | mIHC histology |
| 7 | f | 47 | Diffuse erythema, single fluid-filled vesicle | Day 17 | | mIHC histology |
| 8 | f | 67 | Diffuse erythema, pruritus | Day 1 | | mIHC histology |
| 9 | m | 66 | Diffuse erythema, hand and facial edema, vitiligo- like pigmentation disorder | Day 8 | | mIHC |
| 10 | m | 67 | None | Day 14 | | histology |
| 11 | m | 62 | None | Day 8 | | histology |

m: male, f: female, scRNA-seq: single cell RNA-sequencing, mIHC: multiplex immunohistochemistry

Supplementary Table 2: Tumor-related information

| Patient # | Sex | Age | Uveal melanoma disease stage | Site of metastases | Previous treatments | Cytokine- release syndrome (CRS) on tebentafusp | Treatment of CRS |
|--------------|-----|-----|---------------------------------------|--|--|---|--|
| 1 | m | 58 | Stage IV | Lymph nodes, liver, bone, retroperitoneal space, pleura | Dacarbazine, Ipilimumab/Nivolumab, Temozolamid, Denosumab | CRS grade 2 | fluids, paracetamol, metamizole, morphine |
| 2 | m | 74 | Stage IV | Liver | lpilimumab/Nivolumab | CRS grade 2 | fluids, paracetamol, metamizole |
| 3 | f | 62 | Stage IV | Liver, lung | Pembrolizumab, Temozolamid, Ipilimumab/Nivolumab | CRS grade 2 | fluids, paracetamol, metamizole |
| 4 | m | 57 | Stage IV | Liver | none | CRS grade 2 | fluids, paracetamol |
| 5 | m | 66 | Stage IV | Liver | Ipilimumab/Nivolumab | CRS grade 1 | paracetamol, metamizole |
| 6 | m | 56 | Stage IV | Liver, lymph nodes | none | no CRS | none |
| 7 | f | 47 | Stage IV | Liver, bone | none | CRS grade 2 | fluids, ibuprofen |
| 8 | f | 67 | Stage IV | Liver | none | no CRS | none |
| 9 | m | 66 | Stage IV | Liver | none | no CRS | none |
| 10 | m | 67 | Stage IV | Liver, brain, lymph node, pleura, pericardium, lung, muscle, thyroid, pancreas, colon, abdominal soft tissue, kidney, bone | Ipilimumab/Nivolumab, Temozolamid, Mekinist, Denosumab | CRS grade 1 | paracetamol, metamizole |
| 11 | m | 62 | Stage IV | Liver, lung, lymph node, subcutaneous tissue, muscle | Ipilimumab/Nivolumab, Temozolamid, Pembrolizumab, Lenvatinib, Carboplatin | CRS grade 1 | paracetamol, domperidone |

CRS: Cytokine-release syndrome (according to CTCAE v5)

Supplementary Table 3: Vitiligo-like pigmentation disorder

| Patient | Fitzpatrick Phototype | Distribution pattern of VLPD | Signs of activity (trichrome, confetti) | Diagnosis of VLPD | Biopsy timepoint | Biopsy site | VLPD dynamic at time of biopsy |
|---------|--------------------------|---|--|----------------------|---------------------|----------------|---|
| 1 | IV | Generalized, symmetrical, mottled with confluent patches. Generalized leukotrichia. | confetti and trichrome | Month 3 | Month 7 | Upper arm | Expanding |
| 2 | III | Generalized, symmetrical, mottled. Generalized leukotrichia. | confetti | Month 9 | Month 12 | Lower arm | Stable |
| 4 | II | NA | confetti | Month 6 | Month 8 | Lower arm | Stable |
| 5 | III | Leukotrichia of eye brows and lashes. | confetti | Month 7 | Month 9 | Face | Expanding |
| 6 | II | Generalized, symmetrical, mottled with confluent patches. Generalized leukotrichia. | confetti | Month 5 | Month 7 | Face | Expanding |

VLPD: vitiligo-like pigmentation disorder