

HHS Public Access

Author manuscript Med (N Y). Author manuscript; available in PMC 2021 May 24.

Published in final edited form as:

Med (N Y). 2021 April 9; 2(4): 423-434. doi:10.1016/j.medj.2021.02.002.

Increased Incidence of Venous Thromboembolism with Cancer Immunotherapy

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Abstract

Background: Cancer immunotherapy is associated with several immune-related adverse events, but the relationship between immunotherapy and venous thromboembolism has not been thoroughly studied.

Methods: We conducted a retrospective cohort study of 1,686 patients who received immunotherapy for a variety of malignancies to determine the incidence of venous thromboembolism and the impact of venous thromboembolism on survival. To examine the potential role of inflammation in venous thromboembolism, we also profiled immune cells and plasma cytokines in blood samples obtained prior to initiation of immunotherapy in a sub-cohort

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AUTHORSHIP

Author Contributions

JR, AK, LG and BT performed chart reviews on study patients. JR and SS wrote the manuscript. KRM and AAK edited and revised the manuscript and contributed to experimental design. SS, CMDM, PR, PGP and SK performed the correlative studies. BH, WW, SS, TJA and NS performed the statistical analysis. PF referred patients to the study. JB reviewed and edited the manuscript.

Inclusion and Diversity Statement:

We worked to ensure gender balance in the recruitment of human subjects. One or more of the authors of this paper self-identifies as an underrepresented ethnic minority in science. The author list of this paper includes contributors from the location where the research was conducted who participated in the data collection, design, analysis, and/or interpretation of the work.

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of patients treated on clinical trials who subsequently did (N = 15), or did not (N = 10) develop venous thromboembolism.

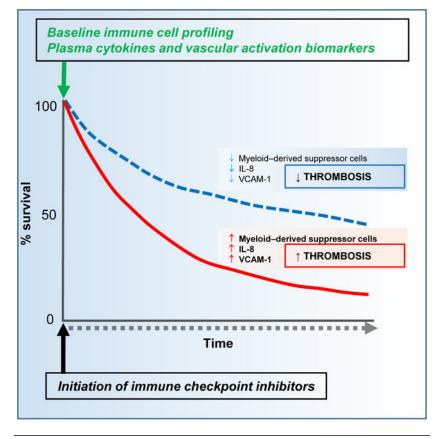
Findings: Venous thromboembolism occurred while on immunotherapy in 404/1686 patients (24%) and was associated with decreased overall survival [HR=1.22 (95% CI 1.06–1.41), p<0.008]. Patients that developed venous thromboembolism had significantly higher pretreatment levels of myeloid-derived suppressor cells (5.382 ± 0.873 vs. 3.341 ± 0.3402 , mean \pm SEM; p=0.0045), interleukin 8 (221.2 ± 37.53 vs. 111.6 ± 25.36 , mean \pm SEM; p=0.016), and soluble vascular cell adhesion protein 1 (1210 ± 120.6 vs. 895.5 ± 53.34 , mean \pm SEM; p=0.0385).

Conclusions: These findings demonstrate that venous thromboembolism is an underappreciated and important immune-related adverse event associated with cancer immunotherapy, and may implicate an interleukin 8 and myeloid-derived suppressor cell-driven pathway in pathogenesis.

eTOC blurb

Venous thrombosis is an underappreciated adverse event associated with cancer immunotherapy. Roopkumar et al describe the incidence of venous thrombosis in cancer patients receiving immunotherapy and suggest that elevated levels of myeloid-derived suppressor cells, interleukin 8 and soluble vascular cell adhesion molecule 1 before immunotherapy correlates with development of venous thrombosis.

Graphical Abstract



Introduction

Cancer immunotherapy disables pathways that regulate the immune response, offering a novel approach to the treatment of malignancy ^{1, 2}. The immune response is normally regulated by binding of CD80/CD86 and programmed death-ligand 1 (PD-L1) to their cognate receptors, cytotoxic T-lymphocyte-associated-4 (CTLA-4) and programmed cell death protein 1 (PD-1) expressed on the surface of cytotoxic T cells ^{3, 4}. Immune checkpoint inhibitors (ICIs) block these ligand-receptor interactions, releasing an immune response toward the tumor². There are currently six ICIs approved for treatment of advanced malignancies: ipilimumab, which targets CTLA-4, nivolumab and pembrolizumab, directed against PD-1, and atezolizumab, avelumab and durvalumab, which bind PD-L1. While these agents have demonstrated substantial efficacy in some cases, ICIs are associated with a variety of immune-related adverse effects (irAEs) that differ from those associated with traditional cytotoxic chemotherapeutic agents ^{5, 6}. These irAEs may involve the skin, gastrointestinal tract, lungs, endocrine, musculoskeletal, renal, nervous, hematologic, cardiovascular and ocular systems. Hematologic irAEs include neutropenia ⁷, immune thrombocytopenia⁸, acquired hemophilia⁹, autoimmune hemolytic anemia^{10, 11} and aplastic anemia ¹². While the mechanisms underlying these toxicities are not well understood, evidence implicates a role for cellular immune responses, inflammatory cytokines and complement-mediated inflammation ^{5, 13, 14}. Interestingly, some studies suggest that occurrence of irAEs may be associated with improved cancer prognosis ^{15–17}.

Thromboembolism, including venous and arterial thromboembolism (VTE and ATE, respectively) is a leading cause of death in cancer patients ¹⁸. Patients with cancer that develop a VTE such as deep-vein thrombosis (DVT), pulmonary embolism (PE) or visceral vein thrombosis (VVT) may experience decreased survival ^{19, 20}. The incidence of VTE is increased in patients with certain types of malignancies ²¹, as well as in those treated with specific cytotoxic chemotherapies and targeted agents ^{22–25}. However, despite the expanding use of ICIs, there is little information available concerning their association with thrombosis ^{26, 27}. Moreover, whether the occurrence of VTE in patients receiving ICIs is associated with longer or shorter survival is unknown. We hypothesized that the pro-inflammatory effects of immunotherapy might initiate a thromboinflammatory response that augments the already-elevated risk of VTE in malignancy ^{22, 28}, and therefore conducted a retrospective cohort study to determine the incidence of VTE in cancer patients receiving immunotherapy. We also assessed the association of VTE with overall survival, and to obtain insight into potential mechanisms, probed for biomarkers predictive of VTE.

Results

Patient characteristics

The study population comprised 1,686 patients with a mean age of 64.5 years (range, 18–93 years; Table 1). Most patients were male (n = 1014, 60.1%), white (n = 1464, 86.8%), non-Hispanic (n = 1610, 96.4%), and had metastatic disease (n = 1523, 90.3%). The most common types of primary cancers were lung (49.6%) and melanoma (13.2%) (Table 1 and Table S1). Nivolumab was the most commonly used immunotherapy, followed by pembrolizumab, atezolizumab, ipilimumab, avelumab and durvalumab. The most common

combination regimen was nivolumab and ipilimumab. The median patient follow-up was 438 days from the initiation of immunotherapy (range: 7–1,971 days). Data concerning concurrent chemotherapy was not collected, but it is unlikely that many patients were receiving chemotherapy given the types of malignancies included in this cohort and the fact that chemotherapy is not often used for these types of tumors per current guidelines. We attempted to define a concurrent control group with similar cancers not receiving immunotherapy, however this was not possible given that immunotherapy represents the current standard of care.

VTE events

VTE occurred in 404 of 1,686 patients (24%). Of patients with VTE, 42.6% developed DVT (n=172), 33.2% PE (n=134), 15.3% both DVT and PE (n=63) and 5.4% VVT (n=24) (Table 2). However, the incidence of concurrent PE and DVT may have been higher, since patients with clinically-diagnosed PE did not routinely undergo lower extremity ultrasound. There were no significant differences observed between the rates of VTE in patients treated with different immunotherapies (Table 1). Likewise, the incidence of VTE in patients receiving single versus combination immunotherapy did not differ.

The six-month and one-year cumulative VTE incidence rates for all immunotherapies were 7.13% and 10.86%, respectively (time to event analysis is shown in Table S2). We did not discriminate between symptomatic versus asymptomatic events, since many patients with incidental and apparently asymptomatic thrombi might have experienced symptoms that would be apparent only upon careful questioning, but cannot be captured in a retrospective study.

Clinical predictors of VTE

In univariate analysis, age and the presence of advanced tumor stage and/or metastatic disease was significantly associated with VTE (p < 0.05 for each), while gender, race, ethnicity, number of immunotherapy treatments and type of immunotherapy were not. In multivariate analysis, only younger age at diagnosis [odds ratio (OR) per one year increase in age = 0.99 (95% CI, 0.980–0.998), p=0.015] and metastatic disease [OR 1.71 (95% CI 1.12–2.61), p=0.013] were significantly associated with VTE. Given the high incidence of irAEs and the retrospective nature of our study, we were unable to assess the relationship between thrombosis and other irAEs.

Relationship between VTE and overall survival

Of the 1,686 study patients, the median overall survival was 416 days (95% CI, 380–471 days), and 906 patients died by the end of the first year of follow-up. The median survival in patients with VTE was significantly decreased: 365 days (95% CI 288, 438 days) compared to 453 days (95% CI 398, 529 days) for patients without VTE (p= 0.002) by log rank test (Figure 1). Decreased survival in patients with VTE was also observed when the two most common tumors in the study, lung cancer and melanoma, were examined individually (Figure S1).

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In a multivariate Cox proportional hazard analysis for overall survival, occurrence of VTE remained significantly associated with decreased survival [HR = 1.22 (95% CI 1.06–1.41), p < 0.008] (Table 3). Other factors significantly associated with decreased survival included stage IV or metastatic disease [HR = 6.16 (95% CI 1.54–24.71) p < 0.001], older age at diagnosis [HR = 1.01 (95% CI 1.00–1.01) p < 0.006] and single agent immunotherapy [HR = 1.37 (95% CI 1.01–1.85), p = 0.04].

Biomarkers Predictive of VTE

Given the association of inflammation with ICI-associated irAEs as well as thrombotic disease, we hypothesized that thrombosis in immunotherapy-treated patients may have an inflammatory etiology. Therefore, we analyzed the phenotype of circulating peripheral blood mononuclear cells (PBMC) in blood samples obtained prospectively from a cohort of 25 patients (15 of whom subsequently developed VTE and 10 who did not, matched for age, sex, site and stage of cancer) at the time of entry into immunotherapy clinical trials. Of these individuals, 7 had bladder carcinoma, 6 had renal cell carcinoma, 8 had melanoma, and 4 had lung cancer (Table 4). In terms of PBMC, the only phenotypic difference observed between these cell populations was a significant elevation in numbers of total MDSCs (% frequency of live PBMC) in patients who subsequently developed VTE compared to those who did not (mean \pm SEM: 5.382 \pm 0.873 vs. 3.341 \pm 0.3402, p = 0.0045) (Figure 2A). There were also higher levels of the polymorphonuclear (mean \pm SEM: 2.720 \pm 0.6119 vs. 1.641 ± 0.3520 , p = 0.6), monocytic (mean \pm SEM: 1.363 ± 0.3561 vs. 0.8114 ± 0.3184 , p = 0.15) and early (mean \pm SEM: 1.163 \pm 0.1624 vs 0.8406 \pm 0.1647, p = 0.28) MDSC subtypes in patients who developed VTE (Figure 2B–D), though these differences were not statistically significant.

We also assessed levels of 51 plasma cytokines and chemokines associated with inflammation and/or vascular damage in the same pretreatment samples (Table S3). Significantly elevated levels of interleukin 8 (IL-8) (p = 0.016), and soluble vascular cell adhesion molecule 1 (sVCAM-1) (p = 0.038) were present in patients who developed VTE (Figure 2 E, F). Other biomarkers including granulocyte-macrophage colony-stimulating factor (GM-CSF) and interleukin-1 receptor antagonist (Figure 2G, H) (Table S3) were also elevated in VTE patients. When we examined correlations between levels of different inflammatory cytokines using correlation matrices, a large cluster of correlated cytokines that included IL-8, TNF α , IL-6, IL-1 β and several others was detected in the group of patients that developed VTE, but not in those who did not (Figure 3). This cytokine cluster also correlated positively with PMN-MDSCs and negatively with CD4+ and CD8+ T cells, although given the small sample size across several tumor types these correlations were not statistically significant.

We also used non-metric multi-dimensional analysis was used to visualize the distributions of the 55 biomarker assays (plasma analytes and MDSC populations) across the three sample populations. Using the PERMANOVA approach, significant differences in the multivariate distributions across the three populations were found (p=0.001, R2=0.32) (Figure S2).

Discussion

The association of VTE with standard cancer therapies is well-established ^{21, 24, 29, 30}. However, in this retrospective cohort study, we report that VTE also developed in nearly one-fourth of all cancer patients treated with immunotherapy. These VTE were clinically important, particularly since nearly one-half were due to pulmonary emboli. Moreover, VTE was associated with decreased overall survival. Hence, although immune checkpoint inhibitors have emerged as powerful agents in the treatment of cancer ^{2, 31–33}, VTE should be added to the expanding list of immune-related adverse events associated with these therapies ^{5, 13}.

These findings are novel since there is little information in the literature concerning the association of cancer immunotherapy with VTE. Kunimasa et al first described a single case of VTE in a patient treated with pembrolizumab ²⁷. Ando expanded on this report, describing VTE and arterial thromboembolism in 8.2% of a retrospective cohort of 122 patients treated with pembrolizumab or nivolumab ²⁶. Gutierrez-Sainz et al described VTE in 16 of a cohort of 229 patients with primarily lung cancers and melanoma over a median follow-up period of 9.8 months ³⁴, and Moik et al. described a cumulative incidence of VTE and ATE in 12.9 and 1.8% of a cohort of 662 patients with a median follow up of 8.5 months ³⁵. In our substantially-larger analysis, we have observed a similar incidence of VTE when comparable follow-up periods are examined, although our studies demonstrate that the risk of VTE extends beyond the initial 6–12 months of therapy.

Our study is the first to define potential biomarkers of thrombotic risk in ICI-treated patients. These immune biomarkers differ from those that have been identified in thrombosis associated with standard cancer therapies, such as tissue factor-bearing microparticles ^{36, 37}, D-dimer and P-selectin ^{38–40}; suggesting a novel mechanism of thrombosis in ICI-treated patients. However, it remains possible that some of the same biomarkers identified as risk factors for cancer thrombosis in patients treated with standard cancer therapies may also contribute to thrombosis in immunotherapy-treated patients, a hypothesis that we did not address in this study. Likewise, the biomarkers and mechanisms identified here have yet to be thoroughly tested in patients treated with standard chemotherapy approaches.

The six-month and one-year cumulative incidence of VTE in our population is similar to that observed in cancer patients at high risk for VTE, such as those with pancreas cancer ^{41, 42}. This is remarkable given the fact that the primary sites of cancer in most patients in this study are associated with a low or intermediate VTE risk ²², suggesting that the high incidence of VTE was indeed immunotherapy-related.

The biomarkers observed in this study are potentially informative concerning the pathogenesis of VTE in immunotherapy-treated patients. MDSCs are associated with tumor progression, metastasis, and reduced survival ^{43–46} and are key targets of immunotherapy ^{47, 48}, but have not been linked to cancer thrombosis. However, an association of MDSCs with activation of coagulant pathways is suggested by the observation that these cells may impair anti-tumor immunity through modulation of procoagulant activity ⁴⁹. Moreover,

MDSC exposed to metastatic tumors in vitro secrete proteins that promote platelet aggregation 50 .

Another intriguing mechanism that our studies suggest may contribute to immunotherapyassociated thrombosis involves the IL-8/CXCR1/2 axis, an important inflammatory pathway. IL-8 released by tumor and other cells in the tumor microenvironment is strongly linked to accumulation of MDSCs within tumors ⁵¹, and elevated plasma levels of IL-8 have been associated with resistance to atezolizumab ⁵² as well as nivolumab and ipilimumab ⁵³ in large retrospective analyses. Moreover, IL-8, through activation of CXCR1/2, induces release of neutrophil extracellular traps (NETs) by tumor-infiltrating MDSCs, and these NETs may shield tumor cells from the cytotoxic effects of CD8+ T cells and NK lymphocytes ^{52, 53}. However, NETs also play a critical role in thromboinflammatory disorders through activation of the complement, contact activation ^{54, 55}, and other systems, and have been implicated in the prothrombotic mechanisms underlying disorders such as heparin-induced thrombocytopenia ⁵⁶, antiphospholipid syndrome ⁵⁷, and SARS-COV2 infection ⁵⁸, among others. We hypothesize that like these non-malignant disorders, immunotherapy-induced thrombosis is largely an immune-mediated thromboinflammatory disorder ²⁸, and that NETs may play a central role.

In addition to NETs, our data suggests that other inflammatory pathways may contribute to immunotherapy-associated thrombosis; indeed, a correlation matrix (Figure 4) taken together with the NMDA analysis (Figure 5) suggests that in immunotherapy patients who develop thrombosis, elevated levels of IL-8 are associated with elevations of a number of other inflammatory cytokines and identify a patient subgroup with increased thrombotic risk. These cytokines may act directly on endothelial and other cells to induce proinflammatory and procoagulant responses ^{59, 60}. Indeed the elevated levels of sVCAM-1 (Figure 2 and Table S3) observed in immunotherapy-treated patients who develop thrombosis support the argument that systemic endothelial activation may be present in these individuals ^{61–63}. Moreover, IL-6, IL-8 and IL-1 β increase the viscoelasticity of blood, and the combination of these three cytokines induce platelet activation and spreading ⁶⁴.

These results have significant implications for patient care, since recent studies demonstrate that VTE in cancer patients can be prevented by prophylactic anticoagulation ^{65, 66}. Confirmation of our findings might suggest that thromboprophylaxis should be considered for patients receiving immunotherapy. However, the safety of prophylactic anticoagulation in immunotherapy-treated patients must be established before such treatment can be routinely recommended.

Limitations of study

Our study has several limitations, including the fact that our data is derived from a singlehealth system-derived retrospective cohort. The retrospective nature of the study precluded us from thoroughly assessing other comorbidities that may have contributed to thrombotic risk in individual patients. However, the large sample size and detailed patient follow-up enhance the fidelity of our observations. We also did not include arterial events in our analysis, though this may have led us to underestimate the overall thrombotic rate. Due to the evolution of cancer therapy, we were not able to identify a contemporary control cohort

treated with non-immunotherapy approaches. However, studies performed prior to the immunotherapy era suggest that the odds ratio for thrombosis in patients with melanoma was 0.78 (95% CI 0.29–2.09), for GU cancer 2.72 (1.80–4.12) and for lung cancer 3.87 (2.73–5.49) ⁶⁷; though it is difficult to translate these odds ratios into incidence rates, these historical results suggest that the thrombosis incidence in our immunotherapy patients is substantially increased compared to patients with similar cancers not treated with immunotherapy. Finally, the predictive biomarkers reported here must be considered exploratory given the relatively small sample size, may be potentially prone to artifacts related to false discovery rates with multiple testing, and require confirmation in a prospective study.

STAR ★ METHODS

RESOURCE AVAILABILITY

Materials Availability—The study did not generate any new materials or reagents

Data and Code Availability—Data involved in this research is available from the corresponding authors upon request. The request should be in the form of a proposal with aims, a statistical approach and other information that will provide assurance of data confidentiality. Patient data will be shared after review and approval of the request by the Cleveland Clinic Institutional Review Board. No data that includes patient names, birthdates, or other potentially identifying information can be shared.

EXPERIMENTAL MODEL AND SUBJECT DETAILS

Study Design and participants—We performed a retrospective single-institution cohort study of cancer patients treated with immunotherapy at the Taussig Cancer Institute, Cleveland Clinic, between July, 2015 and December, 2017. Inclusion criteria included 1) a biopsy proven diagnosis of malignancy, and 2) active treatment with an FDA-approved immunotherapy agent. Patients were identified through a pharmacy database and clinical data were collected directly from the electronic medical record by manual review and stored in a secure web-based server. We included all adult patients who received immunotherapy for their cancer diagnosis and were followed at Cleveland Clinic.

We identified and confirmed all VTE events (including DVT, PE and VVT) through physician notes in the electronic medical record, and confirmed that all VTE events were documented by imaging studies. Since this was a retrospective study in which all included patients received their care through the Cleveland Clinic health system, no patients were lost to follow-up. The study protocol was approved by the Cleveland Clinic Institutional Review Board.

METHOD DETAILS

Biomarker Analysis—For biomarker analysis, we identified a sub-cohort of patients that had been treated on clinical trials of cancer immunotherapies for whom pretreatment analysis of peripheral blood leukocyte populations, including myeloid-derived suppressor cells (MDSC), had been performed (n=25). Pre-treatment plasma from these patients was

also available for cytokine analysis. Of these individuals, 7 had bladder carcinoma, 6 had renal cell carcinoma, 8 had melanoma, and 4 had lung cancer. Fifteen of these patients developed VTE during administration of immunotherapy (cases) and ten did not. The third quartile cutoff for time on immunotherapy was 26.75 months for the no VTE group and 10 months for patients with VTE (p = 0.945). Nine healthy individuals without cancer were included as controls.

Phenotyping of MDSCs was performed on fresh peripheral blood samples immediately after phlebotomy. Briefly, peripheral blood mononuclear cells (PBMC) were purified from whole blood using Ficoll density centrifugation (10 minutes, 1200 x g), then stained using a panel of nine fluorochrome-conjugated monoclonal antibodies that included anti-CD14-APC H7, anti-CD15-PE CF594, anti-CD33-PE, and anti-HLA-DR-APC (Supplementary Table 4). Stained cells were analyzed using an LSR Fortessa flow cytometer (BD Biosciences) with compensation for spectral overlap and auto-fluorescence determined using single-stained and unstained PBMC. The compensation matrix was calculated using FACSDiva software (BD Biosciences), with manual fine-tuning to align the median fluorescence intensities of negative and positive populations. Analysis of cell populations was performed using FlowJo v.10.4.1 (FlowJo LLC, Ashland, OR), with PBMC gating strategies as previously described ⁶⁸. Total MDSC were defined as CD33+HLADR–. MDSC subsets were defined as: monocyte (M-MDSC) = CD14+CD15–, polymorphonuclear (PMN-MDSC) = CD14–CD15+, and early-stage (e-MDSC) = CD14–CD15–⁴⁴.

Analysis of soluble plasma biomarkers was performed using clinical-grade electrochemiluminescence assays (V-PLEX Human Biomarkers). Briefly, pre-coated electrochemiluminescence assay plates were incubated with a blocking solution. Plates were then washed, and incubated with diluted plasma samples and/or calibrators. Bound analytes were detected using sulfo-TAG detection antibodies and plates were analyzed on a SECTOR S 120 plate reader (MSD)⁶⁹. Data was analyzed using MSD Discovery Workbench Version 4.0.

Quantification and statistical analysis—Statistical analysis including overall survival (OS) was performed using the Kaplan-Meier method, and its association with VTE following immunotherapy was estimated using Cox proportional hazard regression adjusted for age at diagnosis and presence/absence of metastases. We compared the incidence of VTE following immunotherapy between single and combination therapy cohorts using the Fisher's exact test.

A multivariate logistic regression model was constructed to identify clinical variables associated with VTE. The cumulative incidence rate of VTE was estimated using the method of Fine and Gray ⁷⁰. Survival and logistic regression analyses were performed using SAS Studio version 3.7 (SAS Institute, Cary, NC) and R version 3.6.3 (R Foundation, Vienna, Austria). All tests were two-sided and P-values of 0.05 or less were considered statistically significant.

Correlative analysis was performed using GraphPad Prism. A non-parametric factorial Kruskal-Wallis (KW) rank-sum test was used to categorize statistical differences between

three groups (i.e. VTE, no VTE and control). Pair-wise, two group analysis was performed using the Wilcoxon rank-sum test ⁷¹. Kruskal-Wallis and Wilcoxon rank-sum tests were adjusted by the Benjamini and Hochberg method ⁷². V-PLEX assay data was normalized using the varstab method implemented in R. The Bray-Curtis method was used to calculate dissimilarities between samples ⁷³.

Non-metric multidimensional scaling (NMDS) plots were generated in the ggplot2 package using Bray-Curtis dissimilarity matrices ⁷⁴. PERMANOVA with the Bray-Curtis dissimilarities was used to compare the multivariate assay distributions across the 3 sample groups (Vegan R package). We also generated correlation plots using R version 3.6.3 (https://cran.case.edu/bin/macosx/R-3.6.3.nn.pkg). The R package corrplot was used to generate each correlation matrix. Hierarchical clustering⁷⁵ was determined using hclust (http://search.r-project.org/R/library/stats/html/hclust.html).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgements:

This work was supported by U01 HL143402 to KRM and AAK, and a Cleveland Clinic VeloSano award.

Funding: NIH U01 HL143402

Declaration of Interest Statement:

JR, LG, BT, SS, JR, WW, NS, MDR, PR, PGP, SK and PF have no competing interests. KRM served on a data safety monitoring board on a study sponsored by Halozyme and received fees from Paraxel for this service. AK reports personal fees and non-financial support from Janssen, personal fees and non-financial support from Bayer, personal fees and non-financial support from Sanofi, personal fees from Parexel, personal fees and non-financial support from Sanofi, personal fees from Parexel, personal fees and non-financial support from Sanofi, personal fees from Parexel, personal fees and non-financial support from Sanofi, personal fees from Parexel, personal fees and non-financial support from Seattle Genetics, personal fees from Pharmacyclics, personal fees from Pharmacyte, personal fees and non-financial support from AngioDynamics, personal fees from Leo Pharma, personal fees from TriSalus, personal fees from Medscape, grants from Merck, grants from Array, grants from Bristol Myers Squibb, and grants from Leap, outside the submitted work. BH has research funds from Amgen, is a consultant to SimulStat, and a scientific advisor to Presagia.

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Highlights

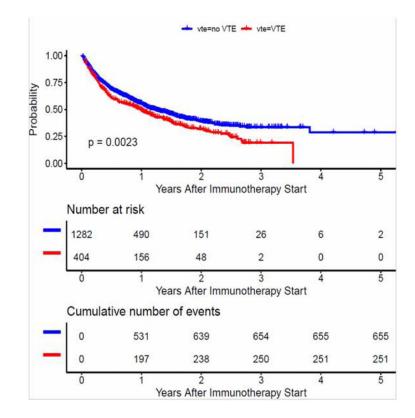
• Cancer immunotherapy blocks physiologic immune regulatory pathways

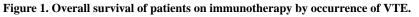
- Immune regulatory blockade is associated with immune-related adverse events
- Venous thrombosis is an underappreciated immune-related adverse event
- Novel biomarkers may predict thrombosis in patients receiving immunotherapy

Context and Significance

Immunotherapy is a promising new treatment approach for cancer. Researchers from the Cleveland

Clinic demonstrate that venous thrombosis (the formation of blood clots in veins) occurs in up to 25% of patients receiving this treatment, and thus must be considered to be a common immune-related adverse event in immunotherapy-treated cancer patients. The authors also identified biomarkers that may recognize patients receiving immunotherapy who are at highest risk of thrombosis. These findings suggest that venous thrombosis is an underappreciated toxicity of cancer immunotherapy, and may guide future studies that focus on the validation of the biomarkers demonstrated here, as well as the role of prophylactic anticoagulation to prevent thrombosis in high-risk cancer patients considered for immunotherapy.





Kaplan-Meier curve for landmark analysis (overall survival) for patients on immunotherapy by occurrence of venous thromboembolism demonstrating reduced survival in those with VTE.

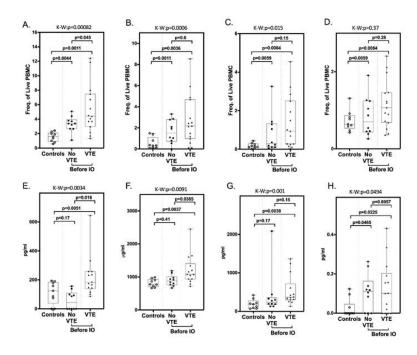


Figure 2. Elevation of total MDSC and soluble biomarkers correlate with VTE after immunotherapy.

MDSC within peripheral blood mononuclear cell fractions from healthy, age-matched donors (controls), and cancer patients before initiation of immunotherapy were quantified by flow cytometry analysis. (A) Total MDSC (CD33+/HLADR–), (B) polymorphonuclear MDSC (CD14–/CD15+), (C) monocytic MDSC (CD14+CD15–), (D) early stage MDSC (CD14–CD15–). Plasma concentrations of immune/inflammatory biomarkers from healthy, age-matched donors "controls" and cancer patients before initiation of immunotherapy. (E) IL-8 (pg/ml), (F) sVCAM-1 (µg/ml), (G) IL-1ra (pg/ml), (H) GM-CSF (pg/ml). Box-and whisker plots show median and 25th and 75th percentiles, minimum and maximum. Kruskal-Wallis (K-W) rank-sum test was used to categorize statistical differences seen between the three groups (VTE, no VTE, controls) with results shown above each graph. Pair-wise two-group analysis was performed using the Wilcoxon rank-sum test. *p* values are indicated. These studies demonstrate a significantly increased rate of thrombosis in patients with elevated levels of total MDSC, IL-8, sVCAM-1, and GM-CSF compared to normal controls and patients with no VTE.

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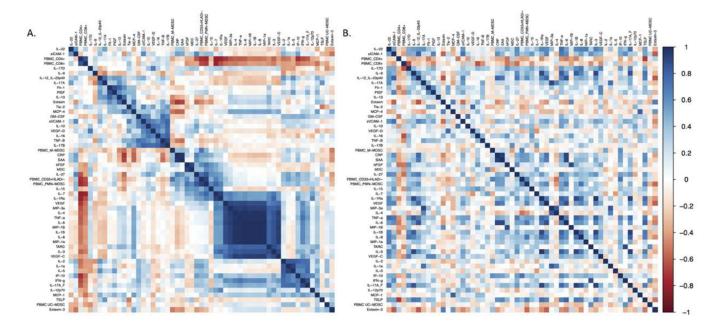


Figure 3: Correlation matrix of MDSCs and soluble plasma analytes associated with VTE in immunotherapy-treated patients with (A) and without (B) VTE. Pairwise Pearson's correlations with hierarchical clustering among the absolute plasma analytes and MDSC populations (n=55 parameters in total). Key to the right of the

analytes and MDSC populations (n=55 parameters in total). Key to the right of the correlation matrix represents the correlation coefficient [ρ = –1 negative correlation (red) to ρ =1 positive correlation (blue)]. The prominent blue square in the right lower quadrant of the matrix in the VTE patients (A) indicates that elevated levels of IL-8 correlate with elevated levels of several other inflammatory cytokines including IL-1 β , IL-6 and TNF α . This clustering is not observed in patients without VTE.

Table 1.

Patient characteristics by VTE status

Characteristic	VTE (N = 404)	No VTE (N = 1282)	Total (N =1686)	P value
Age				0.02*
Median	63.5	65	64.5	
Range	23 - 90	18 - 93	18 – 93	
Sex - no. (%)				0.054
Male	226 (55.9)	788 (61.5)	1014 (60.1)	
Female	178 (44.1)	494 (38.5)	672 (39.9)	
Race – no. (%)				0.1
Asian	3 (0.7)	5 (0.4)	8 (0.5)	
African American	46 (11.4)	108 (8.4)	154 (9.1)	
Caucasian	341 (84.5)	1123 (87.6)	1464 (86.9)	
Multiracial / Other	5 (1.2)	21 (1.7)	26 (1.5)	
Unknown	9 (2.2)	25 (1.9)	34 (2.0)	
Stage – no. (%)				0.03
Stage I - III	28 (6.9)	135 (10.5)	163 (9.6)	
Stage IV	376 (93.1)	1147 (89.5)	1523 (90.4)	
Number of IO drug – no. (%)				>0.99
>1	8 (24.2)	25 (75.8)	33 (2.0)	
1	396 (24.0)	1257 (76.0)	1653 (98.0)	
Type of Immunotherapy – no. (%)				0.58
Ipilimumab	8 (2.0)	40 (3.1)	48 (2.8)	
Nivolumab	214 (53.0)	655 (51.1)	869 (51.5)	
Pembrolizumab	106 (26.2)	350 (27.3)	456 (27.1)	
Atezolizumab	40 (9.9)	135 (10.6)	175 (10.4)	
Durvalumab	5 (1.2)	14 (1.1)	19 (1.1)	
Avelumab	8 (2.0)	12 (0.9)	20 (1.2)	
Nivolumab + ipilimumab	23 (5.7)	76 (5.9)	99 (5.9)	
Current status – no. (%)				
Alive	153 (37.9)	627 (48.9)	780 (43.36)	
Dead	251 (62.1)	655 (51.1)	906 (53.74)	

* p-value by Wilcoxon rank sum test

Table 2.

Types of venous thromboembolic events

Site(s) of VTE	N (%)
Deep-vein thrombosis	172 (42.6)
Pulmonary embolism	137 (33.2)
Visceral vein thrombosis	24 (5.9)
Deep vein thrombosis with pulmonary embolism	
Deep vein thrombosis with visceral vein thrombosis	
Pulmonary embolism with visceral vein thrombosis	
Deep vein thrombosis with pulmonary embolism and visceral vein thrombosis	

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Table 3.

Multivariate Cox model analysis for overall survival

Clinical characteristic	Comparison	Hazard Ratio	95% LCL	95% UCL	P value
VTE	VTE vs. No VTE	1.22	1.06	1.41	0.008
Stage	Stage II vs. I	4.89	1.06	22.65	0.01
	Stage III vs. I	1.58	0.37	6.72	0.49
	Stage IV vs. I	6.16	1.54	24.71	< 0.0001
Use of combination immunotherapy	No vs. Yes	1.37	1.01	1.85	0.04
Age at diagnosis	1 year increase	1.01	1.00	1.01	0.006

Table 4:

Characteristics of patients in the subcohort used to define VTE biomarkers

Characteristic	VTE (N = 15)	No VTE (N = 10)	Total (N =25)
Age			
Median	69	63	66
Range	42-84	51-85	42-85
Sex – no. (%)			
Male	10	7	17
Female	5	3	8
Type of Immunotherapy – no. (%)			
Nivolumab	3 (20) PE	0(0)	
Pembrolizumab	6 (40) 2 DVT, 4 PE	2 (20)	7
Atezolizumab	2 (13.3) DVT	2 (20)	5
Durvalumab	1 (6.6) DVT	5(50)	6
Nivolumab + ipilimumab	3 (20) DVT	1 (10)	4
Current status – no. (%)			
Alive	3	9	12
Dead	12	1	13

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Key Resources Table

Resource	Source	Identifier	
Antibodies			
anti-CD14-APC H7 (MΦP9)	BD Biosciences (Franklin Lakes, NJ)	Catalog #560180	
anti-CD15-PEC F594 (W6D3)	BD Biosciences (Franklin Lakes, NJ)	Catalog #562372	
anti-HLA-DR-APC (G46-6)	BD Biosciences (Franklin Lakes, NJ)	Catalog #559866	
anti CD33-PE (WM53)	BD Biosciences (Franklin Lakes, NJ)	Catalog #555450	
anti-CD4-PE (RPA-T4)	BD Biosciences (Franklin Lakes, NJ)	Catalog #555347	
anti-CD8-APC (RPA-T8)	BD Biosciences (Franklin Lakes, NJ)	Catalog #555369	
anti-PD1-FITC (MIH4)	eBioscience (San Diego, CA)	Catalog #11-9969-42	
anti-PDL1-PerCPeF710 (MIH1)	eBioscience (San Diego, CA)	Catalog #46-5983-42	
anti-human VISTA/B7-H5/PD-1H AF 488	R&D Systems (Minneapolis, MN)	Catalog #FAB71261RG-100UG	
Assay kits			
Human biomarker 46-PLEX assay (V-PLEX)	MesoScale Discovery	K15088D	
V-PLEX vascular injury panel 2 (V-PLEX)	MesoScale Discovery	K15198D	
Software			
FACS-Diva	Becton-Dickinson	https://www.bdbiosciences.com/en-us/instruments/ research-instruments/research-software/flow- cytometry-acquisition/facsdiva-software	
Flow-Jo	Becton Dickinson	https://www.flowjo.com/	
R 3.6.3 (including corrplot and hclust))	R Foundation for statistical computing	https://www.r-project.org/	
Graph Pad Prism	GraphPad.com	https://www.graphpad.com/scientific-software/ prism/	