Increased risk of severe infections in cancer patients treated with vascular endothelial growth factor receptor tyrosine kinase inhibitors: a meta-analysis

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Background: Vascular endothelial growth factor receptor tyrosine kinase inhibitors (VEGFR-TKIs) have been widely used in a variety of solid malignancies. Concerns have arisen regarding the risk of severe infections (≥grade 3) with use of these drugs, but the contribution of VEGFR-TKIs to infections is still unknown.

Methods: The databases of PubMed and abstracts presented at oncology conferences’ proceedings were searched for relevant studies from January 2000 to December 2014. Summary incidences, Peto odds ratio (Peto OR), and 95% confidence intervals (CIs) were calculated by using either random-effects or fixed-effects models according to the heterogeneity of included studies.

Results: A total of 16,488 patients from 27 randomized controlled trials were included. The risk of developing severe (Peto OR 1.69, 95% CI: 1.45–1.96, \(P = 0.001\)) and fatal infections (Peto OR 1.78, 95% CI: 1.13–2.81, \(P = 0.013\)) was significantly increased in patients treated with VEGFR-TKIs when compared to controls. Exploratory subgroup analysis showed no effect of tumor types, phase of trials, or agent used on the Peto OR of severe infections. When stratified according to specific infectious events, the risks of high-grade febrile neutropenia, pneumonia, fever, and sepsis were increased compared with controls, with Peto ORs of 1.57 (95% CI: 1.30–1.88, \(P = 0.001\)), 1.79 (95% CI: 1.29–2.49, \(P = 0.001\)), 5.35 (95% CI: 1.47–19.51, \(P = 0.011\)), and 3.68 (95% CI: 1.51–8.99, \(P = 0.004\)), respectively. Additionally, VEGFR-TKIs significantly increased the risk of fatal sepsis (OR 3.66, 95% CI: 1.47–9.13, \(P = 0.005\)) but not fatal pneumonia (OR 1.34, 95% CI: 0.80–2.25, \(P = 0.26\)).

Conclusion: The use of VEGFR-TKIs significantly increases the risk of developing severe and fatal infectious events in cancer patients. A close monitoring for any signs of infections is recommended for patients treated with VEGFR-TKIs.

Keywords: VEGFR-TKIs, infections, cancer, randomized controlled trials, meta-analysis

Introduction

Tumor angiogenesis is a complex process that is crucial for tumor growth, invasion, and metastasis. During the past decades, many new agents targeting vascular endothelial growth factor (VEGF) and its receptors (VEGFRs) have proven to be a successful strategy in patients with cancer. Until now, the US Food and Drug Administration has approved a number of VEGFR tyrosine kinase inhibitors (TKIs) in multiple indications: sunitinib, sorafenib, pazopanib, and axitinib have been approved for patients with metastatic renal cell carcinoma (RCC). Moreover, sunitinib has been approved for pancreatic neuroendocrine tumors and refractory gastrointestinal stromal tumors.
(GISTs), and sorafenib has been approved for advanced hepatocellular carcinoma (HCC) and radioiodine-refractory differentiated thyroid carcinoma. Additionally, vandetanib has been approved for symptomatic or progressive medullary thyroid cancer, and regorafenib has been approved for refractory advanced colorectal cancer and GISTs.

However, the toxicity profiles of VEGFR-TKIs are unique compared with the adverse effects typically associated with traditional cytotoxic anticancer therapies. They include mucocutaneous adverse events, liver dysfunction, gastrointestinal perforation, and cardiovascular toxicities. Additionally, severe infections (≥grade 3) associated with VEGFR-TKIs have been reported in randomized controlled trials (RCTs). However, the incidence has varied substantially among clinical trials, and there has been no systematic attempt to synthesize the data in order to define the overall incidence and risk of infections associated with these drugs. Therefore, we conducted a systematic review and meta-analysis of RCTs to determine the overall risk of developing severe infection in cancer patients treated with these drugs.

Methods

Data sources

Studies were identified by searching the Cochrane Central Register of Controlled Trials (CENTRAL), PubMed (up to December 2014), Web of Science, and EMBASE, including abstracts from the leading conference proceedings. The search was limited to prospective RCTs published in English. Keywords were sorafenib, nexavar, BAY43-9006, sunitinib, sutent, SU11248, pazopanib, votrient, GW786034, vandetanib, caprelsa, ZD6474, axitinib, AG-013736, cediranib, AZD2171, tivozanib, regorafenib, BAY 73-4506, caboza
tinib, brivanib, ramucirumab, IMC-1121B, nintedanib, BIBF 1120, motesanib, randomized controlled trials, and cancer. The search strategy also used text terms such as angiogenesis inhibitors and vascular endothelial growth factor receptor-tyrosine kinase inhibitors to identify relevant information (Supplementary material). When more than one publication or presentation was identified from the same clinical trial, the most recent report with complete information about infectious events was included for analysis. The quality of reports of clinical trials was assessed and calculated using the five-item Jadad scale including randomization, double-blinding, and withdrawals.

Study selection

The purpose of this study was to determine whether VEGFR-TKIs contribute to the development of severe and fatal infectious events in patients with cancer. Therefore, we only selected those randomized clinical trials that directly compared patients with cancer treated with and without VEGFR-TKIs for analysis. Clinical trials that met the following criteria were included: (1) prospective randomized controlled Phase II and III trials in cancer patients, (2) randomized assignment of patients to VEGFR-TKIs treatment or control in addition to current chemotherapy and/or biological agent, and (3) events or event rate and sample size available for high-grade (grade 3–4) and fatal (grade 5) infectious events. Phase I trials were excluded because of interstudy variability in drug dosing as well as the small number of patients in these trials. Study selection was conducted according to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses statement.

Data extraction and clinical endpoints

Two investigators independently performed data extraction. Agreement between the two data extractors was assessed with the Kappa statistic test. The following information was recorded for each study: first author’s name, year of publication, trial phase, number of patients enrolled, treatment arms, number of patients in treatment and controlled groups, underlying malignancy, median age, median progression-free survival and overall survival, adverse outcomes of interest (infectious events), and name and dosage of VEGFR-TKIs. The following adverse outcomes were considered as infectious events and were included in the analyses: infections (not specified), febrile neutropenia, sepsis, septic shock, fever, bacterial peritonitis, and pneumonia. Adverse events of severe infections (≥grade 3), as assessed and recorded according to the National Cancer Institute’s common terminology criteria (version 2 or 3; http://ctep.cancer.gov), were extracted for analysis, which has been widely used in cancer clinical trials.

Statistical analysis

The principal summary measures were incidence, Peto odds ratio (Peto OR), and corresponding 95% confidence intervals (CIs). For the calculation of incidence, the number of patients experiencing infections and total number of patients treated with VEGFR-TKIs were extracted from the safety profiles of all selected clinical trials; the proportion of patients with infections and 95% CIs were derived for each study.

We also calculated the Peto ORs and 95% CIs of infections in patients assigned to VEGFR-TKIs vs control treatment. For one study that reported zero events in the treatment or control arm, we applied the classic half-integer correction.
to calculate the relative risk (RR) and variance. We also conducted the following prespecified subgroup analyses to find the potential risk factor of infections: tumor types, VEGFR-TKIs, and phase of trials. For each meta-analysis, the Cochran Q statistic and F score were first calculated to determine heterogeneity among the proportions of the included trials. For $P < 0.10$ values of the Cochran Q statistic, the assumption of homogeneity was deemed invalid, and a random-effects model was reported. Otherwise, results from the fixed-effects model were reported. Additionally, we also calculated the number needed to harm from the absolute difference of the pooled estimates between the two groups. Finally, potential publication biases were evaluated with funnel plots for severe infections, which assessed the relative symmetry of individual study estimates around the overall estimate, followed by Begg’s and Egger’s tests. A two-tailed $P$-value of $< 0.05$ without adjustment for multiplicity was considered statistically significant. The leave-one-out procedure was also performed for primary endpoint analysis. A two-tailed $P$-value of $< 0.05$ was considered statistically significant. The results of the meta-analysis were reported as classic forest plots. All statistical analyses were performed by using Version 2 of the Comprehensive MetaAnalysis program (Biostat, Englewood, NJ, USA) and Open Meta-Analyst software version 4.16.12 (Tufts University).

**Trial sequential analysis**

Trial sequential analyses (TSAs) were performed post hoc to assess the risk of random errors and false-positive results, and to help clarify the need for additional trials. RR was used as effect estimate in a DerSimonian and Laird random-effects model. Zero-event trials were handled by adding 0.5 events to the two arms. We used two-sided tests, type I error set at 5% and power set at 80%. In the TSA based on all trials, the boundaries were calculated with a relative risk reduction set at an arbitrary level of 60% and with model variance-based heterogeneity correction. The incidence of severe infections in the control group was set at 3.3% for cancer patients. TSA was performed in TSA V.0.9 B (http://www.ctu.dk/tsa/).

**Results**

**Search results**

Our search yielded 982 clinical studies relevant to VEGFR-TKIs (sunitinib, sorafenib, pazopanib, vandetanib, axitinib, cediranib, tivozanib, regorafenib, cabozantinib, nintedanib, brivanib, ramucirumab, and motesanib). After excluding review articles, Phase I studies, single-arm Phase II studies, case reports, meta-analyses, observation studies, duplicated RCTs, commentaries, letters, and RCTs without adequate infections data (Figure 1), we selected 27 RCTs, including 24 Phase III and three Phase II trials, for the purpose of analysis (Table 1). A total of 16,488 patients from 27 clinical trials were included for analysis. The characteristics of patients and studies are listed in Table 2. Underlying malignancies included nonsmall-cell lung cancer (NSCLC) (ten trials), colorectal cancer (three trials), thyroid cancer (three trials), HCC (two trials), advanced breast cancer (one trial), and urothelial cancer.
Studies, 35 0.001 52/1,732, 84 48 1/381 2/319, 2 9 7 362/4,891 3.57 (0.78–16.33) 211/2,061 2 5 4 2.11 (1.48–3.00) 0.006 174 1.25 (0.92–1.70) 1.99 (1.19–3.33) 129/2,069 336 1.99 (0.57–7.02) 14/653 0.44 15/424 1.71 (1.47–1.99) 0.001 22/680 0.28 23/1,435 1.62 (1.32–2.00) <0.001 25

Vandetanib
Sorafenib
Sunitinib
Cediranib
Regorafenib
Others

274/7,767 1.69 (1.45–1.96) <0.001 53 NA

Abbreviations: VEGFR-TKIs, vascular endothelial growth factor receptor tyrosine kinase inhibitors; RR, relative risk; CI, confidence interval; NSCLC, non-small-cell lung cancer; CRC, colorectal cancer; HCC, hepatocellular carcinoma; NA, not available.

**Table 1** Relative risk of severe infectious events according to tumor types, VEGFR-TKIs, and phases of trials

<table>
<thead>
<tr>
<th>Groups</th>
<th>Studies, n</th>
<th>Severe infectious events, n/total, n</th>
<th>VEGFR-TKIs</th>
<th>Control</th>
<th>RR (95% CI)</th>
<th>P-value</th>
<th>Numbers needed to harm</th>
<th>P-value for group difference</th>
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<td>NSCLC</td>
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<td>362/4,891</td>
<td>210/4,597</td>
<td>1.65</td>
<td>(1.39–1.96)</td>
<td>&lt;0.001</td>
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<td>43/1,389</td>
<td>19/995</td>
<td>1.99</td>
<td>(1.19–3.33)</td>
<td>0.009</td>
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<td>Thyroid cancer</td>
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<td>6/510</td>
<td>1/381</td>
<td>3.57</td>
<td>(0.78–16.33)</td>
<td>0.10</td>
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<td>HCC</td>
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<td>2/293</td>
<td>4/302</td>
<td>0.52</td>
<td>(0.10–2.65)</td>
<td>0.44</td>
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<td>Others</td>
<td>9</td>
<td>71/1,854</td>
<td>40/1,492</td>
<td>1.73</td>
<td>(1.17–2.56)</td>
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<td></td>
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<tr>
<td>Vandetanib</td>
<td>7</td>
<td>111/2,387</td>
<td>69/1,936</td>
<td>1.25</td>
<td>(0.92–1.70)</td>
<td>0.16</td>
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<td>43/1,497</td>
<td>2.11</td>
<td>(1.48–3.00)</td>
<td>&lt;0.001</td>
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<td>23/1,435</td>
<td>2.18</td>
<td>(1.35–3.53)</td>
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<td>Cediranib</td>
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<td>14/653</td>
<td>8/511</td>
<td>1.56</td>
<td>(0.66–3.65)</td>
<td>0.31</td>
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<td>9/637</td>
<td>2/319</td>
<td>1.99</td>
<td>(0.57–7.02)</td>
<td>0.28</td>
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<td>Others</td>
<td>4</td>
<td>211/2,061</td>
<td>129/2,069</td>
<td>1.62</td>
<td>(1.32–2.00)</td>
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<td>Phases of trials</td>
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<td>Phase II</td>
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<td>22/680</td>
<td>15/424</td>
<td>1.21</td>
<td>(0.60–2.44)</td>
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<td>Phase III</td>
<td>23</td>
<td>462/8,257</td>
<td>259/7,343</td>
<td>1.71</td>
<td>(1.47–1.99)</td>
<td>&lt;0.001</td>
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<tr>
<td>Overall</td>
<td>27</td>
<td>484/8,937</td>
<td>274/7,767</td>
<td>1.69</td>
<td>(1.45–1.96)</td>
<td>&lt;0.001</td>
<td>53</td>
<td>NA</td>
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(0 trial),36 pancreatic cancer (one trial),57 gastric cancer (one trial),4,5 melanoma (one trial),39 RCC (one trial),60 acute myeloid leukemia (one trial),61 castration-resistant prostate cancer (one trial),62 and GIST (one trial).13 When examining by agent, sorafenib was investigated in seven trials (2,964 patients), vandetanib in seven trials (4,223 patients), sunitinib in five trials (3,167 patients), cediranib in two trials (1,164 patients), regorafenib in two trials (956 patients), motesanib in one trial (1,072 patients), ramucirumab in one trial (1,253 patients), nintedanib in one trial (655 patients), and brivanib in one trial (502 patients). The Cohen–Kappa statistic for agreement between the two reviewers was 0.866 (95% CI: 0.80–0.93).

**Trial quality**

Randomized treatment allocation sequences were generated in all trials. Twenty-one trials were placebo-controlled and double-blinded. Follow-up time was generally adequate for each trial and included a period of approximately 2–4 weeks after end of therapy on trial. All the trials were of moderately high to high quality (Jadad score 3–5).

**RR of severe and fatal infections**

Severe infections occurred in 484 out of 8,937 (3.8%) patients receiving VEGFR-TKIs. In the non-TKI group, severe infections occurred in 274 out of 7,767 (3.0%) patients. Subjects in the VEGFR-TKI group were at significantly higher risk of severe infections than subjects in the non-TKI group (OR 1.69, 95% CI: 1.45–1.96, P<0.001; Figure 2). There was no evidence of heterogeneity (Q=31.61, P=0.206, F=17.76%). Fatal infections occurred in 52 out of 4,923 (1.0%) patients receiving VEGFR-TKIs. In the non-TKI group, fatal infections occurred in 26 out of 4,111 (0.6%) patients. There was significant difference in risk of fatal infections between subjects in the VEGFR-TKI group and those in the non-TKI group (OR 1.78, 95% CI: 1.13–2.81, P=0.013; Figure 3), and there was no evidence of significant heterogeneity (Q=15.0, P=0.31, F=13.4%). We also did sensitivity analysis to examine the stability and reliability of pooled severe ORs by sequential omission of individual studies. The results indicated that the significance estimate of pooled severe RRs was not significantly influenced by omitting any single study (Figure 4).

**Subgroup analysis of RR of severe infections**

To determine whether the observed increase in ORs of developing severe infections was the result of confounding bias, we performed subgroup analyses according to the underlying malignancy, VEGFR-TKIs, and phase of trials. When stratified by tumor types, a significantly increased risk of severe infections was observed in colorectal cancer (OR 1.69, 95% CI: 1.45–1.96, P<0.001; Figure 2). However, no significant differences in ORs of severe infections were found...
<table>
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<th>Authors</th>
<th>Histology</th>
<th>Phase</th>
<th>Patients enrolled</th>
<th>Treatment arm</th>
<th>Median age (years)</th>
<th>Median PFS (months)</th>
<th>Median OS (months)</th>
<th>Number for analysis</th>
<th>Severe infections</th>
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<td>3.2</td>
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**Abbreviations:** PFS, progression-free survival; OS, overall survival; NSCLC, non-small-cell lung cancer; qd, once daily; po, per oral; Doc, docetaxel; ABC, advanced breast cancer; bid, twice daily; PTX, paclitaxel; CBP, carboplatin; HCC, hepatocellular carcinoma; NR, not reported; q3w, every 3 weeks; CRC, colorectal cancer; FOLFIRI, folinic acid (leucovorin), fluorouracil, and irinotecan; GIST, gastrointestinal stromal tumor; AML, acute myeloid leukemia; TACE, transcatheter arterial chemoembolization; RCC, renal cell carcinoma; CRPC, castration-resistant prostate cancer.
among these tumor types (P=0.85). Clinicians should be cautious when interpreting these results due to the limited RCTs of HCC and colorectal cancer included for the OR calculation.

The risk of severe infections might be related to different VEGFR-TKIs. Our results demonstrated that the use of sorafenib (OR 2.11, 95% CI: 1.48–3.00, P=0.001) and sunitinib (OR 2.18, 95% CI: 1.35–3.53, P=0.001) significantly increased the risk of severe infections, while a nonsignificantly increased risk of severe infections was observed in vandetanib, cediranib, and regorafenib (Table 1). Again, no significant differences in ORs of severe infections were found among these drugs (P=0.48). Then, we also carried out a subgroup risk analysis stratified according to phase of trials (Phase II vs Phase III). Patients from Phase III trials had an OR of 1.71 (95% CI: 1.47–1.99, P<0.001), while patients from Phase II studies had an OR of 1.21 (95% CI: 0.60–2.44, P=0.60; Table 1).

TSA assessed the effect of VEGFR-TKIs on severe infections in cancer patients and showed that the required information size was 3,259 which was less than that in our study (n=16,488), and the cumulative Z-curve crossed the trial sequential monitoring boundary for harm, indicating that further studies are unlikely to change the current conclusion (Figure 5).

![Figure 2](image1.png)

**Abbreviations:** VEGFR-TKIs, vascular endothelial growth factor receptor tyrosine kinase inhibitors; CI, confidence interval.

![Figure 3](image2.png)

**Abbreviations:** VEGFR-TKIs, vascular endothelial growth factor receptor tyrosine kinase inhibitors; CI, confidence interval.
Figure 4 Meta-analysis of severe infections associated with VEGFR-TKIs vs control: “leave-one-out” sensitivity analysis.
Abbreviations: VEGFR-TKIs, vascular endothelial growth factor receptor tyrosine kinase inhibitors; CI, confidence interval.

Figure 5 Trial sequential analysis of 27 trials with lower risk of bias reporting severe infections.
Abbreviation: VEGFR-TKIs, vascular endothelial growth factor receptor tyrosine kinase inhibitors.
Infectious events with VEGFR-TKIs

Risk of severe and fatal infections by specific types

Individual specified and nonspecified causes of severe and fatal infections are listed in Table 3. Of those severe infections that were specified, the most common events for severe infections were febrile neutropenia (61.6%). We then calculated the risk of severe infections stratifying trials according to specific type of severe infections. Our results showed that the use of VEGFR-TKIs significantly increased the risk of severe febrile neutropenia (OR 1.57, 95% CI: 1.30–1.88, \( P = 0.001 \)), pneumonia (OR 1.79, 95% CI: 1.29–2.49, \( P = 0.001 \)), fever (OR 5.35, 95% CI: 1.47–19.51, \( P = 0.011 \)), and sepsis (OR 3.68, 95% CI: 1.51–8.99, \( P = 0.004 \)). We also calculated the risk of fatal infections stratifying trials according to specific types of infections; the use of VEGFR-TKIs significantly increased the risk of fatal sepsis (OR 3.66, 95% CI: 1.47–9.13, \( P = 0.005 \)) but not fatal pneumonia (OR 1.34, 95% CI: 0.80–2.25, \( P = 0.26 \); Table 1).

Publication bias

A funnel plot and both Begg’s and Egger’s tests were performed to assess the publication bias of the selected studies. The shapes of the funnel plots showed no evidence of obvious asymmetry (\( P = 0.61 \) for OR of severe infections; Figure 6). The results from Egger’s test were not significant (\( P = 0.58 \)).

Discussion

During the past decade, identification of the importance of VEGF signal pathway in tumor growth, invasion, and metastasis has led to the development of VEGFR-targeted treatments, which has significantly changed the prognosis of several solid tumors including RCC, HCC, colorectal cancer, and thyroid cancer. Although VEGFR-TKIs are generally well tolerated, infection is an emerging complication with the use of these drugs. Infections can delay treatment or reduce patient compliance with VEGFR-TKI therapies, and the management of infections also increases the cost of cancer treatment. Thus, it is particularly important for all health care practitioners and patients to understand and recognize the risk of infection associated with VEGFR-TKI therapies.

To the best of our knowledge, this is the first and largest meta-analysis evaluating the risk of infections associated with VEGFR-TKIs. In this comprehensive analysis of 16,488 patients, 27 randomized Phase II and III trials using VEGFR-TKIs (sunitinib, sorafenib, pazopanib, axitinib, vandetanib, cediranib, ramucirumab, regorafenib, nintedanib, and motesanib) were included. We did not include Phase I trials in our meta-analysis, since these studies are not randomized and include a wide range of different dosages of drugs. We observed a significant 1.69-fold increase in the risk of high grades of infections with VEGFR-TKIs compared to controls not receiving VEGFR-TKIs. Sensitivity analysis demonstrated that the significance estimate of pooled severe ORs was not significantly influenced by omitting any single study. We also investigated the outcome...
of VEGFR-TKIs-associated severe infections; our results showed that the use of VEGFR-TKIs significantly increased the risk of fatal infections when compared to non-VEGFR-TKI regimens ($P=0.013$). However, given that the absolute risk of fatal infections is low, the use of VEGFR-TKIs should be considered in the context of overall survival benefits. As VEGFR-TKIs are increasingly used in cancer patients, it is particularly important for clinicians to understand and recognize the risk of severe and fatal infection.

Given an increased OR of treatment-related infections, it is clear that proper monitoring, immediate intervention, and effective management are crucial to achieve the maximal therapeutic benefit of VEGFR-TKIs. However, there are no specific guidelines for the treatment of VEGFR-TKIs-induced infections because there is a lack of controlled studies addressing the subject. Based on our findings, the following approaches may be considered to reduce the association of VEGFR-TKIs with risk of infections. Before the initiation of VEGFR-TKIs, clinicians must fully treat patients with any active infection and must monitor patients during the course of VEGFR-TKIs treatment. Clinicians should be cautious when adding VEGFR-TKIs, especially sorafenib and sunitinib, to the first-line or second-line therapies for the treatment of NSCLC and colorectal cancers.

Despite the size of this meta-analysis, our study has some limitations. First, this was a trial-level meta-analysis, and confounding variables at the patient level, such as comorbidities, younger age, prior cardiac history, and previous chemotherapeutic exposure, could not be incorporated into the analysis. However, all of the included trials exhibited moderate- or good-quality Jadad scores, and a report suggests that trial-level and patient-level meta-analyses yield similar results. Second, although most of these trials carried out the randomization process adequately, the infectious events are retrospectively collected. Therefore, these data should be interpreted cautiously because the extracted data used for this analysis could not be considered randomized, which somehow compromised the evidence level. Finally, our literature search is limited to articles published in English creating some selection bias. However, our research detects no publication bias using Begg and Egger tests for ORs of severe and fatal infections.

**Conclusion**

The use of small-molecule VEGFR-TKIs is associated with an increase in the risk of developing severe and fatal infectious events in cancer patients. Close monitoring for any sign of infections is recommended, especially in NSCLC and colorectal cancer patients. Nevertheless, for the average patient, these approved drugs do improve clinical outcomes in their respective indications, and the benefits generally outweigh the risks.

**Authors’ contributions**

QM and D-SZ designed research. L-YG, YYR, and L-LZ conducted research. TG, D-SZ, and YYR analyzed data. QM wrote the draft. All authors read, reviewed, and approved the final manuscript. D-SZ had primary responsibility for final content. All authors contributed toward data analysis, drafting and revising the paper and agree to be accountable for all aspects of the work.

**Disclosure**

All authors declare that they have no potential conflicts of interests.

**References**


Supplementary materials
Search strategy for meta-analysis of association between use of vascular endothelial growth factor receptor tyrosine kinase inhibitors (VEGFR-TKIs) and the risk of severe infections in cancer patients

PubMed
1. VEGFR-TKIs [ALL]
2. “VEGFR-tyrosine kinase inhibitors” [ALL]
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ALL = all fields, MH = MeSH Terms, NM = substance name

Limitation: humans

Date of Search: December 2014 (1974–December 2014)

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35. [embase]/lim OR [embase classic]/lim
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/de = Mapped terms, /lim = Limitation
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CENTRAL database (The Cochrane Library)
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