

## Infections in lymphoma patients treated with bispecific antibodies: A systematic review and meta-analysis.

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### Abstract:

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45

## 46 **Key Points**

47 A high proportion of fatal and severe infections post bispecific antibody treatment were viral  
48 in etiology, highlighting the importance of prophylaxis, vaccination and investigation for viral  
49 reactivation.

50 Improved elucidation of risk factors for infections in bispecific-treated patients will help define  
51 periods of infection risk and guide prophylactic approaches.

## 52 **ABSTRACT**

53 Bispecific antibodies (BsAb) have rapidly entered the treatment paradigms of both indolent  
54 and aggressive lymphomas. Infection is a commonly reported toxicity of BsAb therapy,  
55 however characterization of infection patients among patients treated with BsAbs is poorly  
56 understood. A systematic review and meta-analysis of published studies of CD20-directed  
57 BsAb-treated lymphoma patients was performed. Twenty-seven studies, including 2100  
58 patients, were included. Median follow-up was 12 months; 17% of patients had received  
59 prior cellular therapies. The pooled prevalence of infections of any-grade was 44%  
60 (95%CI:37-50%), with a prevalence of grade  $\geq 3$  infections of 20% (95CI:15-21%) and 3% of  
61 patients experiencing fatal infections (95%CI:2-5%). Infection rates did not differ between  
62 patients with aggressive or indolent lymphomas, or between BsAb monotherapy and  
63 combination therapy. Viral infections constituted a significant proportion (41%) of fatal  
64 infections. Future studies should closely examine the incidence of, and risk factors for,  
65 severe infections classically associated with T-cell depletion. As BsAbs are progressively  
66 incorporated into lymphoma treatment paradigms, the risk of infection needs to be  
67 comprehensively profiled, monitored and proactively managed with multidisciplinary  
68 approaches.

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77 Bispecific antibodies (BsAb) have demonstrated efficacy in newly diagnosed and  
78 relapsed/refractory (R/R) lymphomas. BsAbs form immune synapses between effector cells  
79 (T-cells) and target cell surface markers; typically, CD19 and CD20 in B-cell lymphoma.  
80 BsAb use in myeloma has raised concerns about increased infection risk;<sup>1</sup> less is known  
81 about infection rates following BsAb-treatment for lymphoma. While some CD20-directed  
82 BsAb are now available for lymphoma as standard-of-care, approvals across jurisdictions  
83 such as the US, Canada and Europe are evolving rapidly. <sup>2</sup> FDA approvals were largely  
84 based on small phase Ib/II studies, in heterogenous populations, with different dosing  
85 schedules, treatment durations and supportive care protocols. Despite regulatory approvals,  
86 there are no guidelines for anti-infective prophylaxis and an incomplete understanding of the  
87 rates, timing and types of infections experienced. Therefore, a systematic review and meta-  
88 analysis were undertaken to better characterize infection risks associated with BsAb therapy.

89 We conducted a PRISMA systematic review and meta-analysis (PROSPERO:  
90 CRD42023433207).<sup>3</sup> The search strategy is detailed in the supplement. Studies reporting  
91 infection outcomes following CD20-directed BsAb therapy for the treatment of B-cell  
92 lymphoma, in adult patients, were included. Study identification, data-extraction and bias-  
93 assessment,<sup>4</sup> were performed independently by two authors (GR/MM).

94 The primary outcome was the proportion of BsAb-treated lymphoma patients treated on  
95 clinical trial who experienced  $\geq 1$  infection of any grade. Secondary outcomes included the  
96 rate of severe (grade  $\geq 3$ ) and fatal (grade 5) infections. Sub-group analyses of severe  
97 infections were performed according to aggressive versus indolent lymphoma (defined by  
98 WHO diagnostic criteria),<sup>5</sup> monotherapy versus combination therapy, newly-diagnosed  
99 versus relapsed/refractory (R/R) disease, and bispecific agent. Observational studies were  
100 described separately.

101 Meta-analysis of proportions estimated the pooled infection incidence. Cochran's Q test  
102 examined heterogeneity. Secondary outcomes and subgroup analyses were performed  
103 using random effects models (Mantel-Haenszel). Institutional Review Board approval was  
104 not sought as this study did not constitute human participant research.

105 Of 1133 studies screened, 27 studies (2228 patients, 58% male) were included (Figure S1).  
106 Twenty-three clinical trials (2100 patients) and four observational studies (128 patients)  
107 reported infection outcomes following receipt of one of four CD20-targeting BsAbs (Table 1).

108 The median cohort age was 65 years (IQR:61.2–67). Patients received a median of 3 prior  
109 therapies (IQR:1–3) and had infrequently undergone autologous SCT (16%) or prior  
110 chimeric-antigen receptor T-cell (CAR-T) therapy (17%). The pooled prevalence of grade  $\geq 3$   
111 cytokine release syndrome (CRS), immune-effector cell associated neurotoxicity (ICANS),  
112 and grade  $\geq 3$  neutropenia was: 3.3% (95%CI:0 – 2%), 1% (95%CI: – 2%), and 22%  
113 (95%:6–27%), respectively. Grade  $\geq 3$  leukopenia was 17% (6 studies, 490 patients).  
114 Treatment-emergent hypogammaglobulinemia was not routinely reported. Median follow-up  
115 was 12 months (IQR:6–15). Additional extracted variables are presented in Table S1.

116 Our primary outcome of any-grade infections occurred in 44% patients treated with a CD20-  
117 BsAb (21 studies, 1961 patients, 95%CI:37-50%,  $I^2=88\%$ ). Twenty percent of patients  
118 experienced a grade  $\geq 3$  infection (19 studies, 1791 patients, 95%CI:15 – 21%). The causes  
119 of severe infections were incompletely reported (12/19 studies, Table S2); four studies  
120 reported the cause of all Grade 3 or 4 infections,<sup>6-9</sup> two studies reported only the proportion  
121 of severe infections attributable to COVID-19,<sup>10,11</sup> and six studies reported the aetiology of at  
122 least one severe infection in addition to COVID-19.<sup>12-16</sup> Aggregated across studies, just 133  
123 of 319 (42%) grade  $\geq 3$  infections had an aetiology reported. Among these 133, the  
124 commonly reported causes of severe infection were COVID-19 (32%), clinically-diagnosed  
125 pneumonia (26%) and sepsis (12%), Severe opportunistic infections were reported  
126 specifically in six studies (included in Table S2), including HSV/VZV reactivations (9  
127 patients), CMV, EBV and severe influenza (2 patients each), toxoplasmosis (1  
128 patient), *Pneumocystis pneumonia* (1 patients), and fungal pneumonia (fungus not specified,  
129 1 patient)

130 Fatal infections occurred in 79/1774 patients (3%, 95%CI:2-5%). The cause of fatal infection  
131 was reported in 67% (54/79, Table 2). Of reported infections, microbiologically-confirmed  
132 fatal infections (53%) were more common than clinically-defined fatal infections (15%). Viral  
133 infections were the most common microbiological cause of fatal infections (41%, 32/79),  
134 largely reflecting COVID-19 mortality (91% of viral infections), followed by fungal infections  
135 (6% of total fatal infections, predominately *Pneumocystis*, and one case of systemic  
136 mycosis) and bacterial infections (5% of total fatal infections). One case of fatal  
137 toxoplasmosis was reported.<sup>16</sup> In the included observational studies, viral infections were  
138 also the most common cause of fatal (73%) and severe infections (100%), which detailed 75  
139 infections (11 fatal) (Table S3).

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141 Several planned sub-analyses were then performed. The rate of all-grade (47 vs. 48%),  
142 grade  $\geq 3$  (20 vs. 21%) and fatal infections (4% vs. 3%) did not differ significantly between  
143 patients with DLBCL versus follicular lymphoma (Table S4).

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145 Additionally, in DLBCL, the rates of all-grade (41 vs. 49%,  $p = 0.17$ ) and severe infections  
146 (19 vs. 20%,  $p=0.91$ ) did not differ between patients with DLBCL who received BsAb for first-  
147 line therapy (4 studies, 164 patients), or for relapsed-refractory disease (8 studies, 715  
148 patients,  $p=0.17$ ). In follicular lymphoma, the pooled rates of all-grade infections (59% vs.  
149 26%) and grade  $\geq 3$  infections (25% vs. 5%,  $p<0.01$ , Figure S1) were significantly higher in  
150 patients receiving BsAb for R/R follicular lymphoma (4 studies, 372 patients) compared to  
151 the single published study examining first-line FL treatment with epcoritimab (39 patients).

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153 When considering monotherapy with BsAb, there were no differences in all-grade (49% vs.  
154 50%) or grade  $\geq 3$  infections (18 vs. 27%) in R/R DLBCL patients receiving a BsAb as  
155 monotherapy compared to combination therapy (FigureS2). The impact of combination  
156 therapy could not be analysed in patients receiving upfront treatment because all BsAb were  
157 administered in combination with chemotherapy.

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159 In agent-specific analyses, in DLBCL patients receiving BsAb as first-line therapy, there were  
160 no significant differences in severe infection rates between BsAb products (Figure S3a). In  
161 R/R DLBCL, the significant differences observed between products was driven by single  
162 studies of mosunetuzumab (9%, 95%CI:4–17%) and odronextamab (37%, 95%CI:29–45%,  
163 Figure S3b), In follicular lymphoma, rates of grade  $\geq 3$  infection did not differ significantly  
164 between patients treated with mosunetuzumab (2 studies, 133 patients, 25%, 95%CI:7-42%)  
165 and epcoritimab (1 study, 111 patients, 13%, 95%CI:7-20%,  $p>0.05$ ).

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167 Our systematic review of infections in lymphoma patients receiving bispecific antibodies  
168 reveals a notable rate of all-grade (44%) and grade  $\geq 3$  (20%) infections, with highly variable  
169 reporting of infection type. The etiologies of severe (grade  $\geq 3$ ) and fatal infections were  
170 underreported. Yet, infection remains a prominent cause of treatment interruption, non-  
171 disease-related treatment discontinuation, and non-relapse mortality.<sup>7,17</sup> Preventative  
172 strategies including antimicrobials, infection screening and vaccination require a detailed  
173 understanding of the types of infections they aim to prevent; likewise for diagnostic  
174 investigation. Infection reporting is crucial to the safe implementation of these approaches.

175 A significant proportion of serious and fatal infections were viral, in contrast to other R/R  
176 lymphoma treatments for R/R lymphoma like CAR T-cell therapy, where fatal bacterial  
177 infections predominate.<sup>18</sup> Fatal viral infections were frequently due to COVID-19, which  
178 highlights the importance of understanding both antibody and T-cell specific responses to  
179 COVID-19 vaccine in the context of B-cell depletion, as well as optimal vaccine timing in this

180 cohort.<sup>19</sup> Additionally, reporting of centre-specific prophylaxis regimens could help elucidate  
181 whether the severe viral reactivations observed (e.g. CMV, VZV and HSV) – less commonly  
182 seen in lymphoma patients - would benefit from more intensive preventative approaches. For  
183 clinicians, our results highlight the importance of investigating reactivated or disseminated  
184 viral infection in the context of clinically compatible syndromes in a patient who has or is  
185 receiving BsAb.

186 Analysis of infection rates by advanced disease, aggressive disease, and combination  
187 therapy, did not identify any specific BsAb cohorts at higher infection risk. These results  
188 contrast some observational studies highlighting advanced disease and extensive pre-  
189 treatment as risk factors for infection in lymphoma patients, which reflect the cohorts of  
190 patients who are typically treated on clinical trials.<sup>20</sup> Based on currently available information,  
191 some risk factors for infection cannot be separated by subgroup meta-analysis, which is a  
192 limitation of our study. For example, first-line BsAb-regimens were administered with CHOP  
193 chemotherapy, likely confounding the effect of disease stage and combination therapy on  
194 infection outcomes. Agent-specific effects were likely also confounded with follow-up  
195 duration; odronextamab demonstrated the highest any-grade infection rate in the context of  
196 the longest median follow-up.<sup>21,22</sup> Comprehensive registries and accelerated public access to  
197 individual patient data will help reduce inter-study heterogeneity, and rapidly identify cohorts  
198 at higher infection risk.<sup>23,24</sup>

199 Additionally, evaluation of host- and treatment-related risk factors for opportunistic infections,  
200 may be critical to understanding infection in BsAb-treated patients, given the small but  
201 notable occurrence these infections in a haematological population where these infections  
202 are relatively uncommon. The duration of neutropenia, lymphopenia and  
203 hypogammaglobulinemia were infrequently reported; insufficient data precluded regression  
204 analysis. Cumulative steroid exposure was also underreported, differed notably between  
205 treatment regimens and correlated with BsAb-therapy duration. Recent studies suggesting  
206 that continuous T-cell reduction with BsAbs may induce functional T-cell exhaustion may  
207 provide a further mechanistic explanation for the occurrence of opportunistic infections in  
208 BsAb-treated patients.<sup>25</sup> Future studies should report more detail around the timing of  
209 infections to help compare the relative contribution of early CRS (and its treatment), steroids  
210 and drug ramp-up with the effects of long-term exposure to BsAb. Prospectively collected  
211 minimum dataset of validated risks for opportunistic infection, such as depth of cytopenia,  
212 steroid burden, and infection prophylaxis may help better define these periods of infection  
213 risk.<sup>23</sup> It also provides impetus for consideration and investigation of time-limited and/or  
214 response-adapted BsAb therapy, especially in curative contexts such as DLBCL.<sup>14</sup>

215 The analysis presented is limited by incomplete reporting of infections across different  
216 grades and incomplete reporting of infection etiologies. Similarly, information specific to  
217 COVID-19 risk, such as vaccination status, predominant viral strain, and timing of COVID-19  
218 infection was not available to provide further comment on the high proportion of COVID-19  
219 deaths observed. Significant heterogeneity across product, lymphoma sub-type and  
220 combination therapy was observed, and while addressed by planned sub-group analyses,  
221 resulted in small groups available for pooled analysis.

222 As BsAbs are increasingly integrated into a broader range of treatment paradigms for  
223 lymphoma, the risk of infection needs to be fully characterized, monitored and managed. The  
224 fatal and severe viral and fungal infections in this cohort contrast to the higher rates of  
225 bacterial infections following other anti-lymphoma therapies and highlights the potential for  
226 rapid induction of B- and T-cell dysfunction as a CD20-BsAb class-effect. Comprehensive  
227 registries and enhanced reporting as part of clinical trials are required to design and  
228 implement careful strategies to minimize morbidity and mortality associated with increased  
229 utilization of BsAb.

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Table 1 – Summary of included clinical trials by malignant target and bispecific product.

Malignant Target	Bispecific antibody	Number of Trials	Lymphoma subtype (No. of trials)	Number of patients	All Grade Infection (% , 95%CI)	Median length of follow-up (IQR)
CD20	Epcoritamab	7	Aggressive (5), Indolent (1), B-cell NHL NOS (1)	470	39 (29 - 47)	11.4 (6.1-17.1)
	Glofitamab	7	Aggressive (6), B-cell NHL NOS (1)	618	42 (30 - 53)	10.6 (6 - 15)
	Mosunetuzumab	6	Aggressive (3), Indolent (2), B-cell NHL NOS (1)	599	43 (47 - 50)	12.5 (8 - 28.5)
	Odrone tamab	3	Aggressive (1), Indolent (1), B-cell NHL NOS (1)	414	59 (48 - 69)	21 (N/A)

Table 2 – Aetiology of Fatal Infections

	N, %
<b>Fatal Infections</b>	79
<b>Microbiologically Confirmed</b>	<b>42 (53% of fatal infections)</b>
<i>Viral</i>	32 (41% of fatal infections)
SARS-COV-2	> 29*
Epstein-Barr Virus	1
CMV	1**
PML	1
<i>Bacterial</i>	4 (5% of fatal infections)
Gram negative bacteraemia	4
<i>Fungal</i>	5 (6% of fatal infections)
Candidemia	1
<i>Pneumocystis jirovecii</i> pneumonia	3
Systemic mycoses	1
<i>Protozoan</i>	
Toxoplasmosis	1
<b>Clinically Diagnosed</b>	<b>12 (15% of fatal infections)</b>
<i>Sepsis</i>	4
<i>Pneumonia</i>	8
<b>Aetiology not reported</b>	<b>25 (32% of total infections)</b>

\* Some studies reported > the reported number of SARS-COV-2 infections

\*\* Following first-line therapy