Red cell alloimmunization may induce severe hemolytic side effects. Identification of risk-modifying conditions will help tailor preventative strategies. This study aims to quantify the associations of hematologic malignancies and solid cancers with red cell alloimmunization in patients receiving red cell transfusions. We performed a nested multicenter case-control study in a source population of 24,063 patients receiving their first and subsequent red cell transfusions during an 8-year follow-up period. Cases (n=505), defined as patients developing a first transfusion-induced red cell alloantibody, were each compared with 2 non-alloimmunized controls (n=1010) who received a similar number of red cell units. Using multivariate logistic regression analyses, we evaluated the association of various malignancies and treatment regimens with alloimmunization during a delineated 5-week risk period. The incidence of alloimmunization among patients with acute (myeloid or lymphoid) leukemia and mature (B- or T-cell) lymphoma was significantly reduced compared to patients without these malignancies: adjusted relative risks (RR) with 95% confidence interval (CI) 0.36 (range 0.19-0.68) and 0.30 (range 0.12-0.81). Associations were primarily explained by immunosuppressive treatments [RR for (any type of) chemotherapy combined with immunotherapy 0.27 (95%CI: 0.09-0.83)]. Alloimmunization risks were similarly diminished in allogeneic or autologous stem cell transplanted patients (RR 0.34, 95%CI: 0.16-0.74), at least during the six months post transplant. Alloimmunization risks of patients with other hematologic diseases or solid cancers, and their associated treatment regimens were similar to risks in the general transfused population. Our findings suggest that, in contrast to malignancies in general, hemato-oncological patients treated with dose-intensive regimens have strongly diminished risk of red cell alloimmunization.
Introduction

Transfusion of red cells exposes recipients to non-self antigens and, consequently, may induce alloantibody formation. Although prior alloimmunization requires the exclusive administration of donor blood that is negative for the cognate antigen, accidental re-exposure may induce severe hemolytic transfusion reactions. Prevention of alloimmunization and its consequences is promoted by transfusion of ABO/RhD compatible units to all red cell recipients. In addition, matching beyond those antigens is recommended for certain patients considered to be at high risk of alloimmunization due to repeated exposure, since the number of transfusions is strongly associated with the likelihood of alloimmunization. As such, in several high-income countries, patients with hemoglobinopathies and with myelodysplastic syndrome (MDS), who often face regular transfusions over long periods of time, receive red cell units matched for the most immunogenic and clinically relevant antigens C, c, E, e, and K. The ability of the recipient’s immune system to evoke a humoral alloimmune response upon red cell alloantigen exposure is likely modulated by his or her clinical condition. In this regard, while oncological patients were suggested to have a similar alloimmunization risk to the general transfused population, some studies reported high incidences of alloimmunization among MDS patients. Importantly, apart from the study by Sanz et al., these reports did not take into account the cumulative red cell exposure, which in the oncological patient population is often considerable and a main determinant of alloimmunization. Therefore, the possible influence of disease-specific features remains to be clarified. In addition, cancer types differ from one another in their intrinsic immunobiological characteristics as well as in the immunosuppressive nature of their treatments. Therefore, alloimmunization rates observed in a heterogeneous oncological patient population cannot be extrapolated to specific diseases.

Here we report the results of a nested case-control study quantifying the associations of various hematologic malignancies and solid cancers with the risk of red cell alloimmunization in a cohort of red cell transfusion recipients.

Methods

Study design and setting

We performed a nested case-control study within a mainly Caucasian source population of patients receiving their first and subsequent red cell transfusion between 2005 and 2013 at one of six Dutch hospitals. All six hospitals treat patients diagnosed with oncological pathologies; treatment includes standard remission-induction chemotherapy for acute leukemia patients. Allogeneic hematopoietic stem cell transplantation (HSCT) is performed at four of these centers.

Details of the source population, including eligibility criteria, study period per hospital, and the methods adopted have been published previously (see the Online Supplementary Appendix for details).

Briefly, cases were all patients who developed a first transfusion-induced alloantibody against C, c, E, K, C, Fy, Fy, Jk, Jk, Lu, Lu, M, N, S, or s. For all cases, we assumed the last antigen mismatched transfusion preceding the first positive screen (the ‘Nth’ transfusion) to have been likely to elicit alloimmunization and defined this as the implicated transfusion. If, due to incomplete donor typing, this last mismatched transfusion could not be identified, the last non-tested unit preceding the first positive screen was considered as the implicated transfusion. For each case, we then randomly sampled 2 non-alloimmunized controls on the pre-condition that these patients received at least N or more transfusions at the same hospital, hereby following an ‘incidence-density sampling strategy’. After marking the Nth transfusion in the 2 matched controls, we subsequently constructed a so-called ‘alloimmunization risk period’ in both the case and the 2 controls, which stretches from 30 days before to seven days after this Nth (implicated) transfusion (Figure 1). Next, hospital electronic laboratory information systems and patient medical charts were consulted to record the presence of various clinical conditions during this period.

The study protocol was approved by the Ethical Review Board in Leiden and by the board of each participating center.

Malignancies and their treatments

We used internationally approved response criteria to define the remission state of various hematologic malignancies. Malignancies in complete remission during the alloimmunization risk period were considered as absent. The presence of minimal residual disease was not taken into account. All medication under subcategory L01 in the World Health Organization’s Anatomical Therapeutic Chemical (ATC) classification index was defined as chemotherapy, with the exception of agents in the pharmacologically subgroup L01XC, as these involve monoclonal antibodies. Within subgroups L01XC and L04AA, we defined rituximab, alemtuzumab, and rabbit- or horse-derived anti-thymocyte globulin (ATG) as anti-lymphocyte immunotherapy.

Statistical analysis

Multiple imputation was used to account for missing data. Potential confounders were identified on the basis of their association with the assessed determinant among the source population (i.e. the non-alloimmunized controls).

Using multivariate logistic regression analyses conditioning on the matched variables and on the identified potential confounders, we evaluated the associations of various hematologic malignancies and solid cancers, treatment modalities, and degree of leukopenia with the development of red cell alloimmunization. As we used an incidence-density sampling procedure to select controls, all odds ratios are presented as relative risks (RRs).

Further details on the statistical analytical methods adopted are provided in the Online Supplementary Appendix.

Results

Among 54,347 newly-transfused patients, 24,063 met all study criteria. The majority of excluded patients were ineligible due to the absence of an antibody screen following a single transfusion episode (n=25,057).

First-formed red cell alloantibodies were identified in 505 patients (2.1%) (Online Supplementary Table S1). Thirty-seven of those patients (7.3%), including 21 of 32 (65.6%) who formed anti-Lu, only received units for which testing of the cognate antigen had not been performed; we assumed the last non-tested unit preceding the first positive screen to have elicited alloimmunization. General and clinical characteristics of the 505 alloimmunized patients and their 1010 matched control subjects are presented in Online Supplementary Tables S1 and S2.
Malignancies present during the alloimmunization risk period

A total of 606 patients (40.0%) had at least one type of malignancy (270 had a hematologic malignancy and 338 a solid tumor; 2 patients presented with both types of malignancies). Online Supplementary Table S3 presents types and subtypes of malignancies.

The presence of a malignancy could not be confirmed for 12 patients: 4 patients with a clinical condition suspected for a malignancy that was not further evaluated, 4 patients with a suspected malignancy in whom a malignancy was later confirmed, and 4 patients receiving treatment for a solid tumor for whom the remission status at the time of the risk period was unclear. These 12 patients were not included in the corresponding analyses.

Online Supplementary Tables S4 and S5 show identified confounders for each type of malignancy. Control patients with acute leukemia and lymphoma, as compared to control patients without these diseases, were younger and had less comorbidity (including renal insufficiency and presence of other malignancies). They more frequently received chemotherapy and immunosuppressant medication and more frequently had decreased leukocyte counts.

Maximum frequency of missing data per identified confounder was 2.7% (Online Supplementary Table S6).

The association between types of malignancies and red cell alloimmunization

Table 2 presents the number of cases and controls according to various types of malignancies. Acute leukemia was present in 14 cases (2.8%) compared to 74 (7.3%) controls. There was a reduced incidence of red cell alloimmunization in patients with acute (myeloid or lymphoblastic) leukemia and in patients with mature (B- or T-cell) lymphoma [adjusted RR 0.36 (95%CI: 0.19-0.68) and 0.30 (95%CI: 0.12-0.81), respectively]. Conversely, patients with chronic lymphocytic leukemia (CLL) showed a modest, albeit statistically non-significant, increased risk [adjusted RR 1.20 (95%CI: 0.36-3.93)]. No association between the other types of malignancies and red cell alloimmunization was observed, including MDS and solid malignancies. Similarly, subtypes of solid tumors were not associated to red cell alloimmunization, although some RRs presented with wide 95% CIs (Online Supplementary Table S7). As extensive matching recommendations have only been introduced in the Netherlands since 2011,1 only one of 64 patients (1.6%) with MDS received CcEe- and K-matched units.

Effects were similar in all six hospitals (data not shown).

The association between treatment modalities and red cell alloimmunization

A total of 290 patients received chemo- and/or (anti-)lymphocyte immunotherapy during the implicated risk period. Use of any type of chemotherapy without immunotherapy was not associated with red cell alloimmunization. However, when regimens included lymphocyte-targeted monoclonal antibodies the adjusted RR was 0.27 (95%CI: 0.09-0.83) (Table 3). Twenty-five of the 49 patients (51%) treated with monoclonal antibodies received ATG (with or without alemtuzumab) for in vivo depletion of T cells in the context of an allogeneic HSCT (n=21), aplastic anemia (n=3), or combined pancreas-kidney organ transplant (n=1).

Table 1. Patients’ characteristics during the alloimmunization risk period.

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Cases (N=505)</th>
<th>Controls (N=1010)</th>
<th>RR (CI)*</th>
<th>Adjusted RR (CI)†</th>
<th>Excluded from analysis</th>
</tr>
</thead>
<tbody>
<tr>
<td>Men</td>
<td>237 (46.9)</td>
<td>568 (56.2)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age in years (median, IQR)</td>
<td>67.0 (55.9-75.9)</td>
<td>65.3 (51.6-75.1)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cumulative number of red cell units received (median, IQR)</td>
<td>4 (2-8)</td>
<td>4 (2-8)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Over lifetime*</td>
<td>4 (2-8)</td>
<td>4 (2-8)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>During risk period</td>
<td>3 (2-6)</td>
<td>4 (2-8)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Days transfused during risk period (median, IQR)</td>
<td>1 (1-3)</td>
<td>2 (1-3)</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Values are number (N) (%), unless otherwise stated. IQR: interquartile range. *Up until the first positive screen for cases and up until the last available (negative) screen for controls.

Table 2. Association between various malignancies and red cell alloimmunization.

<table>
<thead>
<tr>
<th>Cases (N=505)</th>
<th>Controls (N=1010)</th>
<th>RR (CI)*</th>
<th>Adjusted RR (CI)†</th>
<th>Excluded from analysis</th>
</tr>
</thead>
<tbody>
<tr>
<td>Hematologic malignancies</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Acute leukemia</td>
<td>14 (2.8)</td>
<td>74 (7.2)</td>
<td>0.31 (0.17-0.58)</td>
<td>0.36 (0.19-0.68)</td>
</tr>
<tr>
<td>Myeloid</td>
<td>14 (2.8)</td>
<td>62 (6.1)</td>
<td>0.38 (0.20-0.71)</td>
<td>0.41 (0.22-0.79)</td>
</tr>
<tr>
<td>Lymphoblastic</td>
<td>0 (0)</td>
<td>12 (1.2)</td>
<td>0.00 (NC)</td>
<td>0.00 (NC)</td>
</tr>
<tr>
<td>Myelodysplastic syndrome</td>
<td>18 (3.6)</td>
<td>46 (4.6)</td>
<td>0.76 (0.43-1.36)</td>
<td>0.75 (0.41-1.36)</td>
</tr>
<tr>
<td>Multiple myeloma</td>
<td>10 (2.0)</td>
<td>26 (2.6)</td>
<td>0.77 (0.36-1.62)</td>
<td>0.79 (0.36-1.71)</td>
</tr>
<tr>
<td>Myeloproliferative neoplasm</td>
<td>9 (1.8)</td>
<td>29 (2.9)</td>
<td>0.62 (0.29-1.33)</td>
<td>0.64 (0.29-1.41)</td>
</tr>
<tr>
<td>Chronic lymphocytic leukemia</td>
<td>5 (1.0)</td>
<td>7 (0.7)</td>
<td>1.45 (0.45-4.67)</td>
<td>1.29 (0.36-3.93)</td>
</tr>
<tr>
<td>Lymphoma*</td>
<td>5 (1.0)</td>
<td>35 (3.5)</td>
<td>0.27 (0.10-0.69)</td>
<td>0.30 (0.12-0.81)</td>
</tr>
<tr>
<td>(Mature) B-cell lymphoma</td>
<td>4 (0.8)</td>
<td>28 (2.8)</td>
<td>0.27 (0.09-0.77)</td>
<td>0.30 (0.10-0.89)</td>
</tr>
<tr>
<td>T-cell lymphoma</td>
<td>1 (0.2)</td>
<td>6 (0.6)</td>
<td>0.33 (0.04-2.75)</td>
<td>0.37 (0.04-3.15)</td>
</tr>
<tr>
<td>Non-hematologic malignancies</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Carcinoma</td>
<td>112 (22.3)</td>
<td>183 (18.2)</td>
<td>1.30 (0.99-1.70)</td>
<td>1.01 (0.75-1.37)</td>
</tr>
<tr>
<td>Other</td>
<td>12 (2.4)</td>
<td>31 (3.1)</td>
<td>0.77 (0.39-1.53)</td>
<td>0.83 (0.41-1.68)</td>
</tr>
</tbody>
</table>

Values are expressed as number (N) (%). *Adjusted for the matched variables: number of transfused red cell units and hospital. Additionally adjusted for other potential confounders (for details, see Online Supplementary Table S5). Acute lymphoblastic leukemia and acute lymphoblastic lymphoma. Six patients were diagnosed with a myelodysplastic syndrome in combination with another hematological disorder including polycythemia vera, essential thrombocytosis, primary myelofibrosis, juvenile and chronic myelomonocytic leukemia. *One patient was diagnosed with an undifferentiated mature lymphoma. RR: relative risk; NC: not computable.
Patients receiving chemotherapeutic agents for acute leukemia or lymphoma during the implicated risk period had substantially reduced alloimmunization incidences [RR 0.29 (95%CI: 0.14-0.60) and 0.08 (95%CI: 0.01-0.57), respectively]. This reduction in risk did not seem to be further influenced to any great extent by the time interval between the initial diagnosis and the period of risk (data not shown). In contrast, non-treated patients with these disorders demonstrated risks comparable to the remainder of the patient population (Table 4). Sixty-two of the 74 treated patients (84%) with acute leukemia received induction therapy during the alloimmunization risk period. Analogous to acute leukemia and mature lymphoma, the 22 patients who received treatment for MDS (including 13 patients receiving induction therapy and 7 receiving hypomethylating agents), demonstrated a trend towards reduced alloimmunization incidences [RR 0.31 (95%CI: 0.09-1.06)] (Table 4). Chemotherapy did not have any impact on risks in patients with other types of hematologic malignancies or carcinoma (Table 4 and Online Supplementary Table S8).

A total of 54 patients received radiotherapy (of any dose and frequency), including 10 patients who received total body irradiation in the setting of an allogeneic HSCT. Radiotherapy was not associated with red cell alloimmunization (Table 3).

Respectively 51, 13, and 10 patients underwent an allogeneic HSCT, an autologous HSCT, or both before or during the risk period. In 51 patients, a reduced-intensity allogeneic HSCT conditioning regimen was followed (including 8 patients who received a double cord transplant), while 10 patients received a myeloablative conditioning regimen. Alloimmunization incidences were substantially decreased in these allogeneic or autologous stem cell transplant recipients [RR 0.34, (95%CI: 0.16-0.74)], at least during the first six months after transplant (Table 3). There was no difference in alloimmunization risk between recipients of an autologous or allogeneic HSCT (data not shown).

Finally, the degree of leukopenia was strongly associated with diminished red cell alloimmunization (Table 5). Here, patients with leukocyte counts of less than 1.0x10^9/L demonstrated an adjusted RR of 0.38 (95%CI: 0.20-0.55). Similar results were obtained when we restricted these analyses to leukocyte counts determined within the week following the implicated transfusion (Table 5). The degree of leukopenia was associated with the type of malignancy and whether or not the patient received chemotherapy. In this regard, minimum leukocyte counts of less than 1.0x10^9/L were observed respectively in 66.2%, 75.9%, and 13.8% of patients with acute leukemia, lymphoma, and carcinoma receiving chemotherapy during the risk period (P<0.0001 for carcinoma vs. acute leukemia and for carcinoma vs. lymphoma).

Discussion

In this nested case-control study, we evaluated whether patients diagnosed with hematologic malignancies and solid cancers differed from the general transfused patient population with regards to the risk of forming red cell alloantibodies. Patients treated for acute leukemia (of either myeloid or lymphoblastic origin) and patients with mature (B- or T-cell) lymphomas demonstrated a 3-fold decrease in the incidence of clinically relevant alloantibodies against red cell alloantigens. In contrast, the alloimmunization incidence among patients treated for other hematologic malignancies or solid tumors was similar to those among the non-malignant patient population.

Although earlier reports only observed similar or even increased red cell alloimmunization frequencies in the oncological patient population,9-11 these prevalence-based studies did not adjust for the substantial number of transfusions these patients usually receive. However, it is well known that the cumulative transfusion dose is an important determinant of alloimmunization.5 Consequently, the

[Figure 1. Illustration of the alloimmunization risk period. For each case, the last antigen mismatched transfusion preceding the first serological detection of an alloantibody was defined as the ‘implicated (Nth) transfusion’ since this transfusion most likely triggered alloimmunization. Alloimmunizations within seven days of the first antigen mismatched transfusion were not taken into consideration as these most likely represented boosting rather than primary alloimmunizations. An alloimmunization risk period was then constructed starting 30 days before and finishing seven days after the defined implicated transfusion. Subsequently, for each case, 2 controls who received at least the same number of red cell units were randomly selected and a similar alloimmunization risk period was constructed around the Nth transfusion. In this example, as the fourth red cell unit most likely elicited red cell alloimmunization, the alloimmunization risk period in both the case and control was constructed around the fourth transfusion. Figure adapted from: Evers et al.15]
observed positive associations might have been due to the quite intensive red cell transfusion support that is generally needed in the treatment of certain malignancies rather than to disease-specific characteristics. So far, no studies have compared specific oncological diseases for alloimmunization risks.

Our findings suggest that especially the dose-intensive immunosuppressive therapy influences alloimmunization. This seems biologically plausible. Several classical cytotoxic agents frequently used in the treatment of acute leukemia and lymphoma, including cyclophosphamide, purine nucleoside analogs, and anthracyclines, are known to induce prolonged (mainly naïve) CD4+ T-cell and B-cell depletion.25-28 Moreover, chemotherapeutic regimens often include corticosteroids, a class of immunosuppressants which we earlier reported to protect against red cell alloimmunization.1 Therefore, we cannot exclude the possibility that part of the observed effects could be directly related to the diseases themselves, i.e. induction of an immunosuppressive but tumor tolerant state via host immune evasion mechanisms of malignant cells.47,48

Furthermore, as patients received a wide range of different chemotherapeutic regimens at varying times before the alloimmunization risk period, it is not possible to come to any firm conclusions as to whether or to what extent patients in complete remission of their treated malignancy should be considered to be significantly immunosuppressed. As such, our RRs might underestimate true effects and our results do not preclude the possibility that these patients have a diminished red cell alloimmunization risk.

In contrast to some other studies,2,13 our incidence-based analysis did not demonstrate an enhanced alloimmunization susceptibility with a diagnosis of MDS. However, and similar to intensively treated patients with acute leukemia and mature lymphoma, patients who received treatment for their MDS tended to show incidence of reduced alloimmunization. Consequently, the decision to transfuse extended donor-matched products to this patient population should not be based on the MDS diagnosis itself, but on other factors associated to an increased alloimmune response, e.g. a high transfusion burden.

Finally, the alloimmunization RR in patients with chronic lymphocytic leukemia (CLL) independent of their treatment seemed to be increased compared to lymphoma patients, although we acknowledge that the number of CLL patients in the current study is not sufficient to con-
firm such a hypothesis. However, CLL is characterized by profound immune disturbances including non-clonal formation of IgG auto-antibodies directed against blood cell antigens. Observations seem to suggest that the disease disturbs normal regulatory potential. Seemingly in contrast with these findings, antimicrobial vaccination responses are often compromised in CLL patients.

Some final comments regarding our methods are appropriate. First, the use of an incidence-density sampling strategy guaranteed that controls were exposed to at least the same number of red cell units as their matched cases. Given this adjustment for cumulative number of red cell exposures, our RRs reflect relative risks independent of exposures. Our alloimmunization risk period was defined specifically to provide a comprehensive study of the influential effect of conditions present around the time of red cell exposure. As the immunosuppressive effects of various treatment regimens are slow to wear off, we preferred to use a relatively long period of risk to precede the implicated transfusion.

Second, our strategies do not fully guarantee the exclusion of all boosting events. Actual ‘lag periods’, i.e. the time needed before antibody levels become detectable after primary antigen encounter, are currently unknown and may even differ according to the antigen used. For our chosen lag period of seven days, we cannot, therefore, fully exclude the possibility that our study included patients whose antibody titers became undetectable over time and who quickly demonstrated recall responses upon re-exposure to the alloantigen. However, we had thought that a substantial amount of boosting reactions as primary alloimmunization events would have biased our RRs towards the null-effect. However, a sensitivity analysis, in which we excluded the 53 patients in whom alloantibodies were discovered during the second week after their first antigen-incompatible transfusion, showed no change in RRs (data not shown). We are confident, therefore, that any possible bias deriving from our choice of lag period is small.

Third, we observed no associations with red cell alloimmunization other than the above mentioned hematologic malignancies and specific types of solid malignancies.

### Table 4. Chemotherapy and red cell alloimmunization risks.

<table>
<thead>
<tr>
<th>Type of malignancy</th>
<th>Chemotherapy</th>
<th>Cases (N=505)</th>
<th>Controls (N=1010)</th>
<th>RR (CI)*</th>
<th>Adjusted RR (CI)†</th>
</tr>
</thead>
<tbody>
<tr>
<td>Acute leukemia</td>
<td>-</td>
<td>489</td>
<td>931</td>
<td>ref</td>
<td>ref</td>
</tr>
<tr>
<td></td>
<td>+</td>
<td>4</td>
<td>10</td>
<td>0.77 (0.22-2.66)</td>
<td>0.88 (0.25-3.09)</td>
</tr>
<tr>
<td></td>
<td>++</td>
<td>10</td>
<td>64</td>
<td>0.25 (0.12-0.51)</td>
<td>0.29 (0.14-0.60)</td>
</tr>
<tr>
<td>Myelodysplastic syndrome</td>
<td>-</td>
<td>484</td>
<td>959</td>
<td>ref</td>
<td>ref</td>
</tr>
<tr>
<td></td>
<td>+</td>
<td>15</td>
<td>28</td>
<td>1.06 (0.54-2.07)</td>
<td>1.04 (0.52-2.06)</td>
</tr>
<tr>
<td></td>
<td>++</td>
<td>3</td>
<td>18</td>
<td>0.32 (0.09-1.12)</td>
<td>0.31 (0.08-1.06)</td>
</tr>
<tr>
<td>Lymphoma</td>
<td>-</td>
<td>498</td>
<td>969</td>
<td>ref</td>
<td>ref</td>
</tr>
<tr>
<td></td>
<td>+</td>
<td>4</td>
<td>7</td>
<td>1.08 (0.31-3.76)</td>
<td>1.26 (0.35-4.51)</td>
</tr>
<tr>
<td></td>
<td>++</td>
<td>1</td>
<td>28</td>
<td>0.07 (0.01-0.49)</td>
<td>0.08 (0.01-0.57)</td>
</tr>
<tr>
<td>Carcinoma</td>
<td>-</td>
<td>390</td>
<td>821</td>
<td>ref</td>
<td>ref</td>
</tr>
<tr>
<td></td>
<td>+</td>
<td>85</td>
<td>141</td>
<td>1.28 (0.95-1.73)</td>
<td>0.99 (0.71-1.38)</td>
</tr>
<tr>
<td></td>
<td>++</td>
<td>26</td>
<td>39</td>
<td>1.40 (0.84-2.35)</td>
<td>1.14 (0.67-1.94)</td>
</tr>
</tbody>
</table>

* + = present; - = absent. Only numbers of patients for whom the presence or absence of a given malignancy and the use of chemotherapy during the alloimmunization risk period could be determined are presented. * Adjusted for the matched variables: number of transfused red cell units and hospital. † Additionally adjusted for other potential confounders (for details, see Online Supplementary Table S5). N: number; RR: relative risk; CI: confidence interval.

### Table 5. Leukopenia and red cell alloimmunization risks.

<table>
<thead>
<tr>
<th>Minimum leukocyte counts (x10^9/L) during:</th>
<th>Cases (N=505)</th>
<th>Controls (N=1010)</th>
<th>RR (CI)*</th>
<th>Adjusted RR (CI)†</th>
</tr>
</thead>
<tbody>
<tr>
<td>Alloimmunization risk period‡</td>
<td>307</td>
<td>524</td>
<td>ref</td>
<td>ref</td>
</tr>
<tr>
<td>2-&lt;4</td>
<td>61</td>
<td>128</td>
<td>0.82 (0.58-1.15)</td>
<td>0.87 (0.61-1.24)</td>
</tr>
<tr>
<td>1-&lt;2</td>
<td>14</td>
<td>43</td>
<td>0.52 (0.27-0.99)</td>
<td>0.59 (0.31-1.13)</td>
</tr>
<tr>
<td>0.5-1</td>
<td>26</td>
<td>142</td>
<td>0.27 (0.17-0.44)</td>
<td>0.33 (0.20-0.55)</td>
</tr>
<tr>
<td>≤1 week following implicated transfusion</td>
<td>273</td>
<td>485</td>
<td>ref</td>
<td>ref</td>
</tr>
<tr>
<td>2-&lt;4</td>
<td>44</td>
<td>107</td>
<td>0.72 (0.47-1.10)</td>
<td>0.80 (0.52-1.23)</td>
</tr>
<tr>
<td>1-&lt;2</td>
<td>15</td>
<td>41</td>
<td>0.60 (0.30-1.23)</td>
<td>0.75 (0.36-1.58)</td>
</tr>
<tr>
<td>0.5-1</td>
<td>19</td>
<td>119</td>
<td>0.24 (0.13-0.44)</td>
<td>0.34 (0.17-0.66)</td>
</tr>
</tbody>
</table>

Minimum leukocyte counts as measured during the alloimmunization risk period and as measured during the week following the implicated transfusion. Values are expressed as number (N) (%). Cumulative numbers of presented cases and controls do not necessarily equal the total number of cases and controls, as patients with leukocytosis are not presented. * Adjusted for the matched variables: number of transfused red cell units and hospital. † Additionally adjusted for other potential confounders (for details, see Online Supplementary Table S5). P<0.02 for trend analysis. RR: relative risk; CI: confidence interval.
although the low numbers of some of these subgroups and the consequent wide CIs per RR prevent any firm conclusions to be made. A much larger study or a meta-analysis of similar studies is needed to assess whether these malignancies are indeed not associated to red cell alloimmunization. Also, due to the fact that remission evaluations available during the alloimmunization risk period were not always complete, we were unable to assess whether the disease stage itself is associated to cell alloimmunization. Finally, since patients treated with chemotherapy received a wide range of chemotherapeutic agents and combinations, as well as varying dose intensities, we were not able to quantify risks according to each single agent.

In conclusion, risks associated with red cell alloimmunization are significantly reduced in patients treated for acute leukemia and mature lymphomas, as well as in recipients of an autologous or allogeneic HSCT. These diminished immune responses most likely reflect the intensity of treatment-associated immunosuppression. In contrast, alloimmunization risks in patients with other hematologic diseases and in patients with solid cancers are similar to those in the general, non-oncological transfused patient population. These findings clearly indicate that, in addition to cumulative red cell exposure, disease-specific conditions should be taken into account when considering the risk of red cell alloimmunization in order to select those patients who would most benefit from extended matched red cell transfusions.

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References
1. TRIP Hemovigilance report 2014. Available at: https://www.tripnet.nl/pages/nl/docu-
mants/TRIP2014Hemovigilantie.pdf.
menten/en/prod-en-dienst/287294/blood-
transfusion-guideline.pdf.
4. Handbook of Transfusion Medicine, United Kingdom Blood Services, 5th edition, 2013. Available at: http://www.transfusionguide-
lines.org.uk/transfusion-handbook.
5. Evers D, Middelburg RA, de Haas M, et al. Red-blood-cell alloimmunisation in relation to antigens’ exposure and their immuno-
6. Bauer MF, Wiersum-Osselton J, Schipperus M, Vandenhoute JC, Briet E. Clinical pre-
dictions of alloimmunisation after red blood cell transfusion. Transfusion. 2007;
7. Fasano RM, Booth GS, Miles M, et al. Red blood cell alloimmunization is influenced by recipient inflammatory state at time of trans-
19. Chenos BD, Greenberg PL, Bennett JM, et al. Clinical application and proposal for modifi-
25. Ferrano C, Guemeneur L, Frigent AF, Taverner C, Revillard JP, Bonnefoi-Berard N. Anthracyclines trigger apoptosis of both GO-
G1 and cycling peripheral blood lympho-
cytes and induce massive deletion of mature T and B cells. Cancer Res. 2000; 60(7):1901-
1907.
26. Gaftter-Guili A, Pollack A. Bendamustine associated immune suppression and infec-
tions during therapy of hematological malign-
ancies. Leuk Lymphoma. 2016; 57(5):512-
519.8.
29. Remmerberger M, Sundberg B. Rabbit-
immunoglobulin G levels in patients receiv-
ing thymoglobulin ... as part of con-
30. Mohity M. Mechanisms of action of antithy-
31. Zand MS, Vo T, Huggins J, et al. Polyclonal rabbit antithymocyte globulin triggers B-cell and plasma cell apoptosis by multiple path-
ways. Transplantation. 2005; 79(11):1507-
1515.
32. Bedognetti D, Zoppioli G, Massucco C, et al. Impaired response to influenza vaccine associ-
ciated with persistent memory B cell deple-
47. Meirow Y, Kanterman J, Baniyash M. Paving the Road to Tumor Development and Spreading: Myeloid-Derived Suppressor Cells are Ruling the Fate. Front Immunol. 2015;6:523.