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ORIGINAL ARTICLE

Clinical features and outcome of 2009-influenza A (H1N1) after allogeneic hematopoietic SCT

B Mohty^{1,5}, Y Thomas^{2,3}, M Vukicevic^{1,5}, M Nagy^{1,5}, E Levrat^{1,5}, M Bernimoulin^{1,5}, L Kaiser^{2,3,4}, E Roosnek^{1,5}, J Passweg^{1,5} and Y Chalandon^{1,5}

¹Division of Hematology, University Hospital, Geneva, Switzerland; ²Laboratory of Virology, University Hospital, Geneva, Switzerland; ³Swiss National Centre for Influenza, University Hospital, Geneva, Switzerland; ⁴Division of Infectious Diseases, University Hospital, Geneva, Switzerland and ⁵Division of Hematology, Geneva University Hospital Blood and Marrow Transplant Program, Geneva, Switzerland

The impact of the 2009 H1N1-Influenza A (H1N1) pandemic in allogeneic hematopoietic SCT recipients (allo-HSCT) is not yet well defined. Between May 2009 and May 2010, all allo-HSCTs who presented with respiratory symptoms were screened for the presence of the H1N1 virus. Oseltamivir resistance was assessed and chart reviews were performed for all cases. In all, 51 of 248 (20%) allo-HSCT recipients followed at our outpatient clinic were screened. We identified 10 patients with H1N1 infection. Close contact with children was the most commonly suspected mode of transmission. Upper and lower respiratory tract infections were present in eight and five patients, respectively. Lymphopenia (<1 G/L) was the most frequent biological abnormality. High immunosuppression was responsible for severe infection requiring mechanical ventilation associated with prolonged viral shedding in three patients who had significant comorbidities and GvHD. Two of them developed an oseltamivirresistant strain and both patients died subsequently despite intensive therapy, resulting in a case fatality rate of 20%. In conclusion, although most allo-HSCTs had mild symptoms from H1N1 infection, severe immunosuppression and emergence of oseltamivir resistance were likely responsible for a substantial morbidity, further supporting the need for vaccination and monitoring of close contacts, especially children.

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Introduction

Respiratory virus infections (RVIs) are common after allogeneic hematopoietic SCT (allo-HSCT),¹⁻³ and influenza may account for up to 30% of all RVIs.^{4.5} Influenza pneumonia-related mortality can reach 25%, particularly in those patients with chronic GvHD.^{1.2} Recently, a new type of influenza A (H1N1) virus, characterized by antigenically distant surface Ags compared with human viruses,⁶ has been involved in a worldwide pandemic outbreak that began in Mexico in March 2009.⁷ On 11 June 2009, WHO (World Health Organization) declared it as the first pandemic of the twenty-first century and >17700 deaths were reported as of March 2010.⁸

Although several large studies have characterized 2009-H1N1 as a novel infection in the general population,^{9–11} the true extent of 2009-H1N1 infection is not yet well defined in allo-HSCT recipients. To better define H1N1 infection in this highly immunocompromised patient population, we reviewed our recent experience at the Geneva University Hospital. This observational study describes the incidence, clinical features and outcome of H1N1 infection among allo-HSCT recipients seen in our department during the pandemic.

Patients and methods

Study population

The Hematology Division at the Geneva University Hospital (Geneva, Switzerland) is a referral center for adults receiving allo-HSCT (population 1.8 million) and patients are closely followed up for many years post transplant. Between May 2009 and May 2010, 248 adult allo-HSCT recipients were followed up. Overall, 19 (8%) patients were within 6 months post transplant and 33 (13%) within 1 year. A total of 46 had an active GvHD (acute grade ≥ 2 or chronic extensive). According to predefined institutional guidelines, all allo-HSCT recipients who presented with respiratory symptoms (such as sore throat, cough, rhinorrhea, nasal congestion or dyspnea) with or without fever were investigated for the presence of

Correspondence: Dr B Mohty, Service d'Hématologie, Hopital Universitaire de Genève, Rue Gabrielle-Perret-Gentil 4, 1211 Geneva-14, Switzerland.

E-mail: bilal.mohty@hcuge.ch

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H1N1 and other community-acquired respiratory viruses (such as adenovirus, metapneumovirus, seasonal influenza A and B, parainfluenza, picornavirus, respiratory syncytial virus). All specimens for microbiological confirmation were taken from combined nasopharyngeal and throat swabs at the time of initial screening. In addition, bronchoalveolar lavages were performed in patients requiring mechanical ventilation. Serial specimens were collected at the discretion of the treating physician. All patients gave their informed consent and the study was approved by the Institutional Review Board.

Virological methods

All analyses were performed according to standardized protocols¹² running in our virology laboratory, which is a WHO referral center for influenza. In brief, H1N1 virus was detected in samples using a real-time reverse transcription-PCR assay in accordance with the protocol from the US Centres for Disease Control and Prevention.¹³ Whenever possible, H1N1 virus isolates were analyzed to determine the presence of the H275Y NA (neuraminidase) mutation using a nucleic acid sequencing assay.

Data collection and analysis

Chart review was performed for all patients who received a PCR-documented diagnosis of H1N1 infection during the study period. Transplant-related characteristics, immunosuppressive medication regimen, H1N1 infection-related features, copathogens and outcomes were assessed. Death due to H1N1 infection was defined as patients dying of respiratory failure after H1N1 infection or its complications. Respiratory tract infections were classified according to Ljungman et al.²: upper respiratory tract infection (URTI) was defined as detection of H1N1 from upper respiratory secretions together with symptoms from the upper respiratory tract. Lower respiratory tract infection (LRTI) was defined as hypoxia, pulmonary infiltrates or new abnormal chest auscultation findings, together with identification of the H1N1 virus in bronchoalveolar lavage or upper respiratory secretions. Hospital-acquired infection was defined as symptom onset >7 days after admission. The immunodeficiency status was graded as published elsewhere,4 either as severe (SID) or MID (moderate). Lymphopenia was defined by lymphocyte counts < 1 G/L. Data were analyzed with descriptive statistics and proportions were compared with a Fisher's exact two-tailed test using SPSS Statistics 13.0 software (SPSS, Chicago, IL, USA).

Results

Baseline characteristics

During the study period, 51 (20%) of 248 allo-HSCT recipients followed up at our outpatient clinic were screened. We identified 10 patients with H1N1 infection between 18 August 2009 and 29 December 2009, giving a frequency of 20% (10/51) (95% confidence interval, 9–30%) based on the number of patients screened or 4% (10/248) (95% confidence interval, 2–7%) based on the total number of patients followed up. Close contact with

children ≤ 12 years or adolescents in the previous 7 days was the most frequently suspected mode of transmission (50% of cases), whereas one case was considered nosocomial (UPN 10).

The median age of H1N1-infected patients was 50.5 (range, 23–62) years, and the median time since transplant was 15 (range, 6-109) months. Eight donors were HLAidentical siblings and all patients, but one (UPN 9), received PBSCs. Half of the patients received a so-called reduced-intensity conditioning regimen. At the time of infection, all patients were in CR for their underlying disease. Six patients had received a partial T cell-depleted transplant and four an unmanipulated graft. Most of the patients were heavily pre-treated before allo-HSCT, and one patient (UPN 4) was on lenalidomide as posttransplant maintenance therapy for multiple myeloma. Four patients had a Karnofsky score $\leq 80\%$ and presented various comorbid conditions. With respect to their immunodeficiency status, five patients were classified as having SID and five as MID. Five patients had GvHD (one acute grade 4 and four chronic extensive) and were receiving immunosuppressive drugs and corticosteroids at a mean dose of 46 mg prednisone equivalent daily (range, 30-80). Immunological reconstitution was incomplete in most of the patients as assessed by mean CD3. CD4 and CD8 lymphocyte count measured within few weeks before diagnosis: 0.7 G/L (range, 0.037–1.4), 0.26 G/L (0.011–0.74) and 0.43 G/L (0.02-1.07), respectively (UPN 9 had large granular lymphoproliferation (LGL) and was not included). All patients, except UPN 4, had been immunized against seasonal influenza and three patients had been vaccinated against H1N1 within a median of 19 days (range, 10-24) before symptom onset (UPN 2 and 3 were vaccinated 1 and 7 days, respectively, after the onset of symptoms) (Table 1).

Clinical features

All patients presented with fever and cough. Eight patients presented an URTI and five an LRTI (three patients had concomitant URTI and LRTI). The median duration of symptoms before virological diagnosis was 2.5 (range, 1-15) days. Commonly reported clinical manifestations were runny nose (n=8, 80%), myalgia (n=6, 60%) and dyspnea (n=6, 60%). Less common signs included sore throat (40%), gastrointestinal symptoms (20%), rarely sweating and fatigue. Abnormal physical examination findings were wheezes or rales on pulmonary auscultation (UPN 4, 5, 7, 8 and 10) and pharyngeal erythema (40% of patients). Chest X-rays were performed in seven patients. Alveolar or interstitial infiltrates were detected in two patients at presentation. Complete blood counts and serum chemistries were drawn in all patients at diagnosis, and revealed normal neutrophil counts in all cases and lymphopenia (<1/L) in six patients. Associated respiratory pathogens were present at diagnosis or developed during the disease course in six patients as shown in Table 1.

Serial virological testing was performed in three patients who required mechanical ventilation and in one patient who had persistent respiratory signs for 23 days, despite having received oseltamivir treatment (UPN 4, 5, 6 and 7).

Table 1	Baseline	e and clinical	features of H1N	[1-infected]	patients at prese	entation								
Patients	Sex/age malignancy	Donor type Reg	Karnofsky score/chronic conditions	GvHD IS/Cs	Baseline CD3/CD4/CD8 (μL)	Sampling site/H1N1 immunization- to-symptoms onset interval (D)	Time since transplant (M)/symptom duration before diagnosis (D)	Rhinorrhea/sore throat/dyspnea/ myalgia	URTI LRTI	Associated respiratory copathogens	LOS (D)/ mechanical ventilation duration (D)	Oseltamivir/ zanamivir duration (D)	H275Y mutation/ viral shedding (D)	Outcome
UPN 1	F/23	Sib	100/None	None	1423/310/1076	-/SdN	8/4	$\mathbf{Y}/\mathbf{N}/\mathbf{N}$	\mathbf{Y}/\mathbf{N}	None	5/-	5/-	NA	Full
UPN 2	M/26	MUD	100/None	None	399/164/246	NPS/+1	6/1	Y/N/N/Y	\mathbf{Y}/\mathbf{N}	RSV		5/-	N/NA	Kecovery Full
UPN 3	ALL F/32	MAC/T-dep Sib	100/None	None	1039/744/264	NPS/+7	109/2	Y/N/Y	\mathbf{Y}/\mathbf{N}	Adenovirus None		5/-	N/NA	Recovery Full
UPN 4	CML M/49	MAC/T-dep MUD	70 /None	Extensive	37/11/22	NPS + BAL/-	12/4	$\mathbf{Y}/\mathbf{N}/\mathbf{Y}$	Y/Y	Klebsiella	50/43	15/15	$\mathbf{Y}/12$	Recovery Expired D + 54
UPN 5	MM F/62	MAC/UM Sib	80/Diabetes	Y/30 Extensive	718/352/381	$NPS/-24^{a}$	8/15	$N/\lambda/N/N$	N/N	Uxytoca None	20/6	13/8	N/19	Full
9 NAN	AML M/57	RIC/T-dep Sib DIC/IIM	100/Esophageal	Y/80 None	NA	NPS/-19	86/3	${\bf X}/{\bf X}/{\bf X}$	\mathbf{Y}/\mathbf{N}	Picornavirus		-/9	N/23	Recovery Full Decovery
UPN 7	M/40 AML	Sib MAC/T-dep	70/Diabetes, Renal,	Extensive Y/30	861/121/691	NPS + BAL/-	52/1	$\mathbf{Y}/\mathbf{N}/\mathbf{Y}/\mathbf{N}$	\mathbf{Y}/\mathbf{Y}	CMV EBV	00/06	20/20	$\mathbf{Y}/21$	Expired D + 94
UPN 8	M/56 ▲I T _ Dh: →	Sib	pulmonary 80/pulmonary	Extensive	1005/376/646	$\rm NPS/-10^a$	22/4	$N/\lambda/\lambda/\lambda$	Y/Y	None		5/-	N/NA	Full
6 NAU	M/52	Sib MACT don	100/LGL,	None	6630/580/5967	NPS/-	18/2	N/A/A/A	\mathbf{Y}/\mathbf{N}	RSV		-/L	NA	Full Boomorery
UPN 10	CMML M/53 AML	MAC/1-dep Sib RIC/T-dep	spienectomized 70/Diabetes, renal, pulmonary	Acute Y/60	141/16/115	NPS/-	8/2	X/N/N/N	\mathbf{Y}/\mathbf{N}	Picornavirus	15/-	15/15	NA	Full Recovery
Abbrevi lymphor conditio RSV = r "UPN 5	ations: ALL na; IS/C = in ning; MM = 1 espiratory sy and 8 had 2	Phi + = ALL munosuppress multiple myeloi ncytial virus; S vaccine doses (Ph positive; BAJ sive drug/corticos ma; MUD = matc ib = identical sibl (only time since fi	L = bronchc iteroid (pred ched unrelat ling; T-dep = îrst dose is	balveolar lavage; Inisone equivalen ed donor; N = no = T-cell depletion specified).	CMML = chroi it mg/day); LOS i; NA = not avai i; URTI = upper	uic myelo-monoo S = length of sta, lable; NHL = no respiratory trac	ytic leukemia; C ;; LRTI =lower n-Hodgkin's lymp t infection; UM =	ond Re respiratc homa; h	g = conditioni ory tract infec NPS = nasophi pulated graft;	ng regimen; tion; M = má aryngeal swal Y = Yes.	D = days; F ale; M = mon b; RIC = redt	= female; ths; MAC iced-intens	HL = Hodgkin's = myeloablative ity conditioning;

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The median period of symptomatic viral shedding in those four patients was 22 (range, 19–81) days. Sequencing of the NA gene was performed on seven patients (not possible on the three remaining isolates mainly because of low viral loads). An oseltamivir-resistant strain appeared 7 and 6 days after initiation of oseltamivir in two patients (UPN 4 and 7), respectively, who later expired despite mechanical ventilation and i.v. zanamivir (Table 1).

Treatment and outcome

All patients were treated with oseltamivir at a dose of 75 mg two times a day. Treatment was started within a median of 3 (range, 0–8) days from symptom onset for a median duration of 6.5 (range, 5–20) days. Eight patients (80%) received broad spectrum antibiotics in addition to antiviral treatment.

Five patients of whom four (UPN 4, 5, 7 and 10) had significant comorbidities and active GvHD were hospitalized for a median of 20 (range, 5–90) days. Of the latter, three patients (UPN 4, 5 and 7) required mechanical ventilation for a median time of 43 (range, 6–70) days. Zanamivir was administered i.v. for these four patients (UPN 4, 5, 7 and 10) in addition to oseltamivir for a median duration of 15 (range, 8–20) days. Patients UPN 4 and 7 died of respiratory failure within 54 and 94 days of symptom onset, respectively, resulting in a case fatality rate of 20%. All patients tolerated oseltamivir and zanamivir, and none discontinued treatment because of adverse effects (Table 1).

Discussion

Several studies have described H1N1 infection in allo-HSCT (Table 2),^{14–19} but its clinical spectrum is still being defined. Here, we report the impact of the H1N1 epidemic in a population of allo-HSCT recipients who were followed up in the long term in a clinic using standardized guidelines.²⁰

H1N1 incidence peaked in November 2009, similar to the general Swiss population.²¹ Overall, 4% of our total patient population had a proven H1N1 infection. When considering only those screened for a respiratory illness, $\sim 20\%$ were infected. Redelman-Sidi *et al.*¹⁷ found that 22% of their screened patients were positive for H1N1 during the New York city outbreak. In fact, the denominator of H1N1-infected patients is difficult to establish.^{22,23} Whereas a frequency of 20% is most likely too high, 4% is certainly an underestimate if patients with mild illness may not have sought medical care and therefore may not have been diagnosed.

Nosocomial infection rates may range from 11 to 19% in allo-HSCTs.^{15,18} Close contact with children was the most common presumed mode of transmission in our series, which is not surprising as this has been previously reported as the most significant risk factor of developing RVIs.¹ Indeed, approximately one child in every three was infected with H1N1 in England,²⁴ whereas in the United States, the rate of secondary outbreaks in households was 13% with children at increased risk for infection by a factor of 4.²⁵

Similarly, Kumar *et al.*²⁶ reported that pediatric solid organ transplant recipients were substantially more likely to have fever, rhinorrhoea, sore throat and headache at presentation than were adult patients. Notably in this study, 31% of patients had ill household contacts.²⁶ Taken together, these observations should alert physicians involved in the care of HSCT patients about the high risk of viral transmission through close contact with children, underlining the importance of education and vaccination of patients and their family households. Close contact with symptomatic children should be avoided whenever possible during periods of RVI epidemics.

Fever and cough were the most common symptoms observed. H1N1-related LRTI developed in 50%, of patients, which is in the range of 31–68% of almost all other series.^{14–16,18,19,26,27} The 30% mechanical ventilation and 20% case fatality rates observed in our study correspond to those found in other small series,^{16,18,19,27–29} but are higher than the frequencies reported in larger series (4.0–13.5% and 3.0–7.3%, respectively).^{14,22,26} Although this may reflect a reporting bias if patients with more severe illness were more likely to come to medical attention, we believe that allo-HSCT recipients are at higher risk for severe complications, notably in case of development of H1N1 oseltamivir-resistant strains^{15,19} or concomitant respiratory pathogens.^{16,19,26,27,30–32}

The average duration of H1N1 viral shedding in the general population was ~6 days.^{9–11} Redelman-Sidi *et al.*¹⁷ failed to demonstrate a significant prolonged viral shedding in HSCT patients. However, we along with others^{18,19,30} have found evidence of prolonged viral shedding in severely immunosuppressed allo-HSCTs and in patients with oseltamivir resistance. Although it remains uncertain whether these are representative of the entire population of allo-HSCTs, protracted infections are expected to occur in this patient population.

Half of our patients were classified as SID. Interestingly, LRTI, prolonged viral shedding, hospitalization, longer antiviral treatment, ventilator requirement and H1N1 resistance were documented only in SID patients. Lymphopenia, which has been previously reported as a risk factor for LRTI, both for community-acquired RVI1-4,33 and for H1N1 influenza9,14-18,26 was observed in six patients at the time of diagnosis. Chronic GvHD and its corollary of long-term immunosuppression has also been associated with the risk of developing RVI and LRTI.¹ Recently, Taplitz *et al.*¹⁸ showed that the use of >20 mg of prednisone equivalent daily was significantly associated with the development of LRTI and with 30-day mortality. In our study, 11% (5/46) of patients with active GvHD contracted an H1N1 infection as compared with 2.5% (5/202) of patients without GvHD (P < 0.03). Furthermore, 5 of 33 (15%) patients within the first year post transplant had an H1N1 infection, whereas only 5 of 215 (2.3%) patients after the first year were infected (P < 0.005). Taken together, these observations suggest that highly immunocompromised allo-HSCT recipients are at risk for severe complications from influenza and deserve close monitoring.

There have been conflicting data as to the efficacy of influenza immunization.^{34–36} In our study, three patients had H1N1 infection despite having been vaccinated. In

Table 2 E	3aseline an	nd clinical feature	s of H1N1	l-infected	patients in tl	ne literature								
Author country	No. of patients (% of allo)	Donor type/ Cond Reg	Median time since transplant (range)	Median age (y) (range)	GvHD (%) IS or Cs	Vaccine	Rhinorrhea/ Sore throat/ dyspnea/ myalgia	Fever/ cough/GI symptoms/ CNS symptoms	URTI LRTI (%)	Antiviral treatment	Hospital/ mechanical ventilation	H275Y mutation/ PVS	Outcome/ mortality rate (%)	Comments
This study	10	2 MUD, 8 Sib/5	15 M	50.5	5 (50%) 5	H1N1: 5	8/4/6/6	10/10/2/0	8	O: 10	s, c	ς, τ	Two deaths	
Taplitz	(100%) 27	MAC, 5 KIC 12 MUD,	(0-109) 379 D	(23-02) 46 i	17 (63%) 12	Seasonal: 9 H1N1: 3	12/10/16/14	25/26/	(%0c) c 13	Z: 4 O: 25	s NA	ۍ د 1	(20%) Nine deaths	19% Nosocomial
et al., ¹⁸ USA	$(80\%)^{a}$	10 Sib/NA	(5–2895)	(20-67)		Seasonal: NA		7/2	14 (52%)	Z: 5, P: 2	٢	NA	(33%)	
Redelman-Sidi et al., ¹⁷ USA	21 (57%)	NA/19 MAC, 2 RIC	44 M (5 M to 15 Y)	36 (5–72)	4 (19%) NA	NA	NA	NA	9° 5 (40%)	0: 19 Z: 1	× 0	00	Full recovery	None nosocomial and none <5 M
Tramontana <i>et a</i> l., ¹⁹ Australia	16 (50%) ^d	7 Sib, 1 MUD/NA	3 allo < 100 D 8 auto > 2 V	56.5 (30–72)	5 (31%) 12	H1N1: 0 Seasonal: NA	NA	NA	$ \begin{array}{c} 16 \\ 11 \\ (70\%) \end{array} $	0:15 Z: 2	16 9	. 1	Six deaths (37,5 %)	Three nosocomial and four with copathogens (aspergilus, CMV, Staph, HXV HHV6 FRV)
George <i>et al.</i> ,'' Australia	6 13 (92%)	3 MUD, 8 Sib, 1 Haplo/2 MAC, 10 RIC 1 auto	425 D (2–1541)	53.6 (37–63)	7 (54%) 5	H1N1: NA Seasonal: 9	NA/13/5/NA	13/11/ NA/NA	13 5 (40%)	O: 13 Z: 1	v 4	ΝA	Four deaths (30%)	Five with copathogens and four had H1N1 hefore HSCT
Ditschkowski et al., ²⁸ Germony	10 (NA)	NA NA	10 M	NA	5 (50%) 10	NA	NA	NA	NA	O: 10	с	NA	Two deaths (20%)	
Garland et al. ²⁹ UK	9 (44%)	3 MUD, 1 Sib/1 MAC, 3 RIC, 50000	4 M (8 D to	62 (35–75)	3 NA	H1N1: 1 Seasonal: NA	NA	NA	NA	0: 9 Z: 3	6 K	ΝA	Three deaths (20%)	No copathogen
Patel <i>et al.</i> , ²⁷ USA	5 (80%)	1 MUD, 3 Sib/3 MAC, 1 RIC, 1 anto	(3–19) (3–19)	51 (23–56)	4 (80%) 3	H1N1: 0 Seasonal: NA	NA	NA	5 3 (60%)	O: 5 P: 1	1 1	NA 1	One death (20%)	One with parainfluenza, aspergillosis and rhizopus
Lalayanni <i>et a</i> l., ⁴⁰ Greece	3 (100%)	NA	18 M (15 M to	40 (35–54)	$^{1}_{ m AA}$	H1N1: 1 Seasonal: NA	NA	NA	(00.00) 3 2 (66%)	O: 3 Z: 1	ω α	NA 1	Two deaths (66%)	
CDC, ³⁰ USA	2 (NA)	NA	24 D	10 and 40	NA	NA	1/NA/NA/NA	2/2/ NA/NA	2 NA	O: 2, Z:1 Rib: 1_R:1	- 17	~ ~	Full	One with meumocvetic and stanh
Kharfan-Daba _{, et al} ³¹ 11SA	ja 2 (100%)	2 MUD/	191 D (171–711)	46.5 (41–52)	00	NA	NA/1/1/2	2/2/ 1/NA	- 10	0: 2	- 0 -	² V o	One death	One with Aspergillus
Bastos <i>et al.</i> , ⁴¹ Brazil	(100%)	CB/RIC	3 D	12	10-	NA	1/NA/1/NA	1/1/ NA/NA		O and Z		°A −	Full	No copathogen
Rozovski et al ³² Israel	(100%)	1 Sib/RIC	118 D	62		NA	NA/NA/1/1	1/1/NA/1	- 1	0		NA	Death	Sepsis to Klebsiella meumonia
Frangoul et al. ⁴² USA	(100%)	CB/MAC	Υ	6		NA	NA/NA/1/NA	No/NA/ 1 /NA	00	0	. — —	NA	Death	Infection mimicking gut GVHD
Campbell et al., ⁴³ USA	(0%)	1 Auto	2 D	40	0	NA	NA/NA/No/NA	1/1/ 1/NA	1 1	D, A, Rib, P	1	NA 1	Death	Extrapulmonary H1N1 RNA
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Abbreviations: ARDS = acute respiratory distress syndrome; ATG = anti-thymocyte globulin; Auto = autograft; CB = cord blood; Cond Reg = conditioning regimen; CNS = central nervous system; corticosteroid; LOS = length of stay; LRTI = lower respiratory tract infection; M = months; MAC = myeloablative conditioning; MMF = mycophenolate mofetil; MUD = matched unrelated donor; NA = not Cy = cyclophosphamid; D = days; Fluda = fludarabine; GI = gastrointestinal; Haplo = haplo-identical; HHV 6 = human herpes virus 6; HSCT = hematopoietic SCT; IS/C = immunosuppressive drug/ available; O = oseltamivir; P = peramivir; PVS = prolonged viral shedding; Rib = ribavirine; RIC = reduced-intensity conditioning; R = rimantadine; Ritux = rituximab; Sib = identical sibling; staph = staphy-

lococcus; T-dep = T-cell depletion; URTI = upper respiratory tract infection; Y = year; y = years; Z = zanamivir.

Data are number of patients unless otherwise specified. Only data on HSCT patients (allo and auto) provided in the articles are reported in the table. "Only 17 of 27 patients had a PCR-documented H1N1 infection (probable cases were defined as patients who had a positive test for influenza A in the context of widespread community H1N1 activity, without PCR confirmation).

^bOnly one patient has been tested for H275Y mutation in this study.

^cInformation on URTI or LRTI was lacking for six patients. ^{d6}come patients reported in this study might be common to those reported in the study by George *et al.*¹⁶

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highly immunogenic.³⁷ Ljungman²² failed to demonstrate any protective effect from either the seasonal or the H1N1 vaccines in HSCT patients. In contrast, in the Spanish series,¹⁴ only one of the vaccinated patients against seasonal influenza was diagnosed with pneumonia compared with 41% who were not vaccinated (P = 0.005). Although it is not possible to draw definitive conclusions as to the protective effect of H1N1 vaccine in HSCT patients, preliminary results from a large study (http://Clinical Trials.gov, ID: NCT01022905) conducted in our hospital suggest that severely immunocompromised allo-HSCT patients do not respond to H1N1 vaccination. Therefore, influenza vaccination of family members and close contacts, especially children, is strongly recommended to limit the risk of influenza exposure in HSCT recipients. Additional strategies such as post-exposure oseltamivir prophylaxis for high-risk patients may be warranted.^{19,26,38}

Our study has several limitations. The number of H1N1confirmed cases was low, making the reliable estimation of clinical outcomes, such as mechanical ventilation or mortality rates, difficult. Furthermore, only a limited number of allo-HSCT recipients within 6 months post transplant (19 patients) were included, precluding us to precisely analyze the impact of H1N1 infection in the early post-transplant period.

In conclusion, although most allo-HSCT recipients had mild symptoms from H1N1 infection as does the general population,²³ high immunosuppression and emergence of oseltamivir-resistant strains were responsible for a substantial number of deaths in the allo-HSCT setting. Complications included higher rates of LRTI, prolonged viral shedding and respiratory failure. Our study strongly supports the need for vaccination and monitoring of family households, especially children.³⁹

Conflict of interest

The authors declare no conflict of interest.

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