# Association between acute graft versus host disease and lung injury after allogeneic haematopoietic stem cell transplantation

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**Objective:** To investigate the characteristics of chest high-resolution computed tomography (HRCT) and pathogenesis of acute graft versus host disease (acute GVHD)-induced lung injury after allogeneic haematopoietic stem cell transplantation (allo-HSCT).

**Methods:** A study of 47 patients with acute GVHD of grades II–IV describes the clinical manifestations and characteristics of chest HRCT of acute GVHD-induced lung injury. Detection of serum interferon  $\gamma$  (IFN $\gamma$ ) and tumour necrosis factor  $\alpha$  (TNF $\alpha$ ) were performed before the treatment for acute GVHD. Transbronchial biopsy was performed in four patients whose chest HRCT did not recover completely after treatment for acute GVHD. Pulmonary function was measured in patients who survived more than 6 months in every 3 months.

Results: Chest HRCT scans were performed in 47 cases and 20 cases showed abnormal in which 17 cases were suspected of acute GVHD-induced lung injury. In 17 patients with acute GVHDinduced lung injury, HRCT revealed diffused interstitial infiltrate in five cases, diffused interstitial and alveolar infiltrate in seven cases, diffused interstitial and segmental lobar alveolar infiltrate in five cases accompanied by bilateral pleural effusion and hydropericardium in nine patients. There was no statistical significance between the levels of serum IFN $\gamma$  and TNF $\alpha$  in cases with and without lung injury, but the levels of serum IFN<sub> $\gamma$ </sub> and TNF $\alpha$  in patients were significantly higher than the healthy group (IFN<sub>2</sub>: p=0.000, TNF $\alpha$ : p=0.000). The histopathology of the lung tissue was characterized by disorganization, epithelial cell damage, interstitial fibroplasias and interstitial T lymphocyte or macrophage infiltrate. Forty-seven cases all attained the treatment for acute GVHD, and the total effective rate and the rate of completely remission (CR) were 74.47 and 55.32%, respectively. The total effective rate and the rate of CR in the treatment for acute GVHDinduced lung injury were 94.12 and 58.82%, respectively. The effective rate of treatment for acute GVHD-induced lung injury positively correlated with that for acute GVHD (r=0.771, p=0.001). Three cases in nine cases with lung injury and three cases in 15 cases without lung injury who survived more than 6 months developed late-onset non-infectious lung injury. Eleven patients of 24 patients who survived more than 6 months had abnormal pulmonary function, including seven patients in nine patients with acute GVHD-induced lung injury and four patients in 15

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patients without acute GVHD-induced lung injury. There was no difference in the incidence of late-onset non-infectious lung injury, but significance in the incidence of abnormal pulmonary function between cases with and without lung injury (p=0.033, cross-tabs).

**Conclusions:** These results suggested that the lung might be one of the target organs of acute GVHD and participation of T lymphocyte, macrophage and cytokines such as IFN $\gamma$  and TNF $\alpha$  might play a role in the pathogenesis of acute GVHD-induced lung injury. Acute GVHD-induced lung injury may progress to late-onset non-infectious lung injury.

Keywords: allogeneic haematopoietic stem cell transplantation, acute graft versus host disease, lung injury

#### Introduction

Allogeneic haematopoietic stem cell transplantation (allo-HSCT) has been used with increasing frequency for the malignant or non-malignant haematological diseases. Acute graft versus host disease (acute GVHD) and lung injury remain common complications and are significant causes of mortality after allo-HSCT. Acute GVHD usually involves the liver, skin and gut. In recent years, other organs such as thymus and lung have also been reported to be involved by acute GVHD.<sup>1-3</sup> In 1978, Beschorner et al.<sup>4</sup> reported a group of patients who developed lymphocytic bronchitis associated with acute GVHD after allo-HSCT. Although many studies reported that early lung injury was related to acute GVHD, whether the lung is one of the target organs of acute GVHD remains controversial.<sup>1,2</sup> The pathogeneses of acute GVHDinduced lung injury is unknown. The mechanisms of T-cell axis and gut-liver-lung inflammatory cytokine axis were proposed. Some studies proposed that cytokines such as interferon  $\gamma$  (IFN $\gamma$ ) and tumour necrosis factor  $\alpha$  (TNF $\alpha$ ) might play a critical role in the process of acute GVHD-induced lung injury.<sup>3,5,6</sup> Clinical reports of imaging characteristics of lung at the onset of acute GVHDinduced lung injury are rare.

In this report, the characteristics of chest highresolution computed tomography (HRCT) and pathogenesis of acute GVHD-induced lung injury were investigated in 47 patients with acute GVHD of grades II–IV after allo-HSCT. Chest HRCT of the cases suspected of acute GVHD-induced lung injury was characterized by diffused interstitial, alveolar and segmental lobar infiltrates. The histopathology of lung tissue showed T lymphocyte or macrophage infiltrates. Increased levels of serum IFN $\gamma$  and TNF $\alpha$  were found in patients with acute GVHD. The association between acute GVHD and lung injury, and the association between acute GVHD-induced lung injury and lateonset non-infectious lung injury, are discussed.

# Patients and methods

### Patients

Forty-seven patients with a median age of 32 years (range: 14–49 years) who were diagnosed of acute GVHD (grades II–IV) after allo-HSCT from February 2003 to June 2007 were enrolled in this study. The clinical characteristics of the cases are shown in Table 1.

#### Conditioning regimens

Twenty-four patients received total-body irradiation (TBI) and cyclophosphamide conditioning regimens (TBI + CY) and 19 patients were treated with busulfan, cyclophosphamide and cytarabine conditioning regimens (modified BUCY). Four patients received increased intensity conditioning regimens (fludarabine: 50 mg once daily i.v. on -10 to -6 day; cytarabine 1.0 to 2.0 g once daily i.v. on -10 to -6 day; TBI 4.5–5.0 GY/day, on -5, -4 day; cyclophosphamide 50 mg/kg once daily i.v. on -3 to -2 day).<sup>7</sup>

#### Diagnosis criteria of acute GVHD-induced lung injury

Diagnosis criteria of acute GVHD-induced lung injury are as follows: (1) patients have manifestations of acute GVHD involving at least one organ, e.g. skin, liver and gut;<sup>8</sup> (2) patients have abnormal chest HRCT; (3) lung injury induced by infection and heart diseases can be excluded; (4) chest HRCT scans are improved after treatment for acute GVHD.

#### Treatment and therapeutic effect

Methylprednisolone at a dose of 2 mg/kg/day was used for the treatment of acute GVHD. Antihuman thymocyte globulin (ATG) or ATG combined with CD25 monoclonal antibody and other immunosuppressants were used for steroid-resistant acute GVHD cases. Acute GVHD-induced lung injury was treated according to acute GVHD protocols. Therapeutic effect was stratified into complete remission (CR), partial response (PR) and non-remission (NR) (CR: chest HRCT scans recover completely; PR: chest HRCT improved after treatment; NR:

Case	Age (years)	Gender	Diagnosis	Donor	Stem cell source	HLA-matching (serology)	HLA-matching (genotype)	Conditioning regimens	Acute GVHD	Chronic GVHD	GVHD prophylaxis	Lung injury before HSCT	CMV	Survival time
-	34	Male	CML-CP -NB	Unrelated	BMT	Matched	1 genetic locus mismatched	Modified BUCY	≡	I	MTX + CsA + ATG	I	+	86 d
N	47	Female	MDS	Related	PBSCT	2 antigen mismatchad	1	Modified BUCY	=	I	MTX + CsA	I	I	>3 years
Ю	28	Female	AML-CR	Unrelated	BMT	Matched	Matched	Modified BUCY	=	I	MTX + CsA	I	+	82 days
4	48	Male	CML-BP	Related	PBSCT	Matched	I	Over intensity	=	Intensive	H MTX + CsA	I	+	273 days
Q	29	Male	CML-CP	Related	PBSCT	1 antigen mismatchad	I	Improved BUCY	=	Limited	MTX + CsA	I	+	149 days
9	22	Male	ALL-CR	Related	PBSCT	1 antigen mismatched	I	Modified BUCY	≡	Limited	MTX + CsA + ATG	Ι	I	170 days
7	33	Female	AML-CR	Related	PBSCT	Matched	I	TBI + CY	=	Limited	MTX + CsA	I	+	916 days
00	29	Female	AML-CR	Related	PBSCT	Matched	I	Modified BUCY	=	Intensive		I	+	811 days
ი	49	Male	AML-CR	Related	PBSCT	Matched	I	TBI + CY	=	I	MTX + CSA + ATG	I	+	167 days
10	41	Male	AML-CR	Unrelated	PBSCT	Matched	2 genetic locus mismatched	TBI + CY	=	Intensive	MTX+CsA + ATG	I	+	762 days
11	36	Male	AML-NR	Unrelated	BMT	Matched	2 genetic locus mismatched	Modified BUCY	≡	Limited	MTX + CsA + ATG + MMF	I	+	58 days
10	50	Eamalo			PRCCT	Matchad	I	Modified BLICV	=	Limitod		I	I	9/10/10/10
10	45	Male	CML-BP	Unrelated	PBSCT	Matched	1 genetic locus	TBI + CY	= =	Limited	MTX+CSA MTX+CSA	Ι	+	613 days
7	00			Dototo Doto	RMT	a antiae	mismatched -		=	1	HAIGHMW		-	aver Og
<u>†</u>	0	Maid		וופומופת		mismatched		5 +	=		+ ATG + MMF	-	F	oo days
15	39	male	CML-CP -NR	Unrelated	PBSCT	Matched	matched	Modified BUCY	≥	I	MTX + CsA + ATG + MMF	I	+	74 days
16	21	Female	ALL-CR	Related	PBSCT	matched	I	TBI + CY	=	Limited	T IVIIVI MTX + CsA	Ι	Ι	>3 years
17	34	Female	AML-CR	unrelated	BMT	Matched	2 genetic locus mismatched	Over intensity	=	Intensive	MTX + CsA + ATG	I	Ι	855 days
18	32	Female	AUL-NR	Related	PBSCT	2 antigen mismatched			2	I	CsA + ATG	Pulmonary tuberculosis	I	32 days
19	18	Female	AML-CR	Unrelated	BMT	Matched	1 genetic locus mismatched	TBI + CY	=	I	MTX + CsA + ATG	Ē	+	86 days
20	14	Male	ALL-NR	Unrelated	PBSCT	Matched	3 genetic locus	TBI + CY	$\geq$	Limited	T MINI MTX + CsA + ATG	I	+	65 days
21	17	Male	ALL-CR	Unrelated	BMT	Matched	1 genetic locus mismatched	TBI + CY	$\geq$	Limited	MTX + CSA + ATG	Ι	+	177 days
22	36	Male	ALL-CR	Related	PBSCT	Matched	1	TBI + CY	=	Intensive	MTX + CsA	I	+	3 years

Table 1 The clinical characteristics of the 47 cases

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22       Female AML-CR       Unrelated PBSCT       Matched       1 genetic locus       Modified BUCY       I. Limited       MTX + CsA + MX + CsA + MX + CsA         39       Female AML-NR       Related PBSCT       Matched       TBI + CY       IV       -       MTX + CsA + MX + CsA + MX + CsA + MX + M	Case (yea	(years)	Gender	Diagnosis	Donor	source	HLA-matching (serology)	(genotype)	Conditioning regimens	GVHD	GVHD	GVHD prophylaxis	Derore HSCT	CMV	Survival time
39         Famale         Rearied         RBSCT         Matched         -         TBI + CY         IV         -         MTX + CSA         ATG           14         Fernale ALL-RR         Related         PBSCT         Matched         -         TBI + CY         II         -         MTX + CSA         + ATG           15         Fernale ALL-RR         Related         PBSCT         Matched         -         MTX + CSA         + ATG           22         Male         AUL-CR         Related         PBSCT         Matched         -         TBI + CY         II         -         MTX + CSA         + ATG           38         Male         AUL-CR         Related         PBSCT         Matched         1         genetic locus         Modified BUCY         II         Immession         MTX + CSA         + ATG           38         Male         AUL-CR         Related         PBSCT         Matched         1         genetic locus         Modified BUCY         II         Immession         MTX + CSA         + ATG           38         Male         AUL-CR         Related         PBSCT         Matched         1         genetic locus         Modified BUCY         II         Immession         MTX + CSA         ATG <td></td> <td>22</td> <td>Female</td> <td>AML-CR</td> <td>Unrelated</td> <td></td> <td>Matched</td> <td>1 genetic locus mismatched</td> <td>Modified BUCY</td> <td>=</td> <td>Limited</td> <td>+ CsA +</td> <td></td> <td>+</td> <td>105 days</td>		22	Female	AML-CR	Unrelated		Matched	1 genetic locus mismatched	Modified BUCY	=	Limited	+ CsA +		+	105 days
14         Female ALL-NF         Related PBSCT         Matched PBSCT		39	Female	AML-NR	Related	PBSCT	Matched		TBI + CY	$\geq$	I	+	I	I	76 davs
19         Male Fenale         AML-CR MIC         Unrelated BSCT         Matched Matched         matched Fenale         matched MIX         Cost intensity (MIX         1         -         MIX         Cash ATG MIX         Cash ATG MIX </td <td></td> <td>14</td> <td>Female</td> <td>ALL-NR</td> <td>Related</td> <td>PBSCT</td> <td>Matched</td> <td>I</td> <td>TBI + CY</td> <td>≡</td> <td>Intensive</td> <td>+</td> <td>I</td> <td>+</td> <td>202 davs</td>		14	Female	ALL-NR	Related	PBSCT	Matched	I	TBI + CY	≡	Intensive	+	I	+	202 davs
45       Female       CML-CP       Related       PBSCT       Matched        Modified BUCY       II        MTX + CSA + ATG         22       Male       AUL-CP       Unrelated       BMT       Matched       1        MTX + CSA + ATG         38       Male       AUL-CP       Unrelated       BMT       Matched       1       TBI + CY       III       Initiated       MTX + CSA + ATG         38       Male       AUL-CR       Related       PBSCT       1 antigen        MTX + CSA + ATG         31       Hate       AUL-CR       Related       PBSCT       1 antigen        MTX + CSA + ATG         32       Male       AUL-CR       Related       PBSCT       1 antigen        MTX + CSA + ATG         32       Male       AUL-CR       Related       PBSCT       1 antigen        MTX + CSA + ATG         32       Male       CML-CP-PR       Unrelated       PBSCT       matched       1 genetic locus       Modified BUCY       I        MTX + CSA + ATG         32       Male       CML-CP-PR       Related       PBSCT       matched       0 over modified       I        MTX + CSA + ATG		19	Male	AML-CR	Unrelated	BMT	Matched	matched	Over intensity	=		+ CsA +	۱ گ	- 1	117 davs
22       Male       CML-CP       Unrelated       BMT       Matched       1 genetic locus       Modified BUCY       II       Imited       MTX + CsA + ATG         38       Male       AUL-CR       Related       PBSCT       Matched       TBI + CY       II       Imited       MTX + CsA + ATG         14       Fernale       ANL-CR       Related       PBSCT       Matched       TBI + CY       II       Limited       MTX + CsA + ATG         32       Male       AUL-CR       Related       PBSCT       Zamigen       -       MC4ified BUCY       II       Limited       MTX + CsA + ATG         32       Male       CML-CP-PR       Unrelated       PBSCT       Matched       -       Modified BUCY       II       Intensive       MTX + CsA + ATG         32       Male       CML-CP-PR       Unrelated       PBSCT       Matched       -       Modified BUCY       II       Intensive       MTX + CsA + ATG         32       Male       ALL-NR       Unrelated       PBSCT       Matched       -       MC4 field BUCY       II       Intensive       MTX + CsA + ATG         33       fermale       ALL-NR       Unrelated       PBSCT       Matched       -       Over modified BUCY		45	Female	CML-CP -NB	Related	PBSCT	Matched	I	Modified BUCY	≡	I	+ CsA +	- 5	I	3 years
38         Male         AUL-CR         Related         PBSCT         Immatched         TBI + CY         II         Limited         MTX + CsA         ATG           14         Male         AML-CR         Related         PBSCT         1 antigen         -         Modified BUCY         II         Immersive         MTX + CsA         + ATG           32         Male         CML-CP-PR         Unrelated         PBSCT         2 antigen         -         Modified BUCY         II         -         MTX + CsA         + ATG           28         Male         CML-CP-PR         Unrelated         PBSCT         mismatched         -         Modified BUCY         II         -         MTX + CsA         + ATG           28         Male         CML-CP-PR         Unrelated         PBSCT         mismatched         -         MOdified BUCY         II         -         MTX + CsA         + ATG           20         Male         ALL-NR         Unrelated         PBSCT         Matched         -         MOdified BUCY         II         Intensive         MTX + CsA         + ATG           21         Male         ALL-NR         Unrelated         PBSCT         Matched         -         MOdified BUCY         II         - <td></td> <td>22</td> <td>Male</td> <td>CML-CP</td> <td>Unrelated</td> <td>BMT</td> <td>Matched</td> <td>1 genetic locus</td> <td>Modified BUCY</td> <td>=</td> <td>limited</td> <td>+ CsA +</td> <td></td> <td>+</td> <td>938 days</td>		22	Male	CML-CP	Unrelated	BMT	Matched	1 genetic locus	Modified BUCY	=	limited	+ CsA +		+	938 days
14         Male         AML-CR         Related         PBSCT         1 antigen		38	Male	AUL-CR	Related	PBSCT	Matched		TBI + CY	≡	Limited	+	Ι	I	255 days
11       Female       ALL-CR       Related       PBSCT       2 minutached mismatched       -       TB + CY       I       -       MTX + CsA + ATG         32       Male       CML-CP-PR       Unrelated       BMT       Matched       1       emotified BUCY       I       intensive       MTX + CsA + ATG         28       Male       CML-CP-PR       Unrelated       PBSCT       matched       -       MOdified BUCY       I       intensive       MTX + CsA + ATG         20       Male       CML-CP-PR       Unrelated       PBSCT       matched       -       Modified BUCY       I       intensive       MTX + CsA       ATG         20       Male       ALL-NR       Related       PBSCT       matched       Over modified BUCY       I       intensive       MTX + CsA       ATG         22       female       CML-CP       Related       PBSCT       matched       Over modified BUCY       I       intensive       MTX + CsA       ATG         23       Male       CML-CR       Intensive       MTX + CsA       I       -       MTX + CsA       ATG         34       Male       CML-CR       Intensive       MTX + CsA       I       -       MTX + CsA       ATG		14	Male	AML-CR	Related	PBSCT	1 antigen	I	Modified BUCY	≡	Intensive	+ CsA +		+	906 days
32       Mate       CML-CP-PR       Unrelated       BMT       Matched       1 genetic locus       Modified BUCY       In Intensive       MTX + CsA + ATG         28       Male       CML-CP-PR       Unrelated       PBSCT       mismatched       Modified BUCY       In       Intensive       MTX + CsA + ATG         20       Male       ALL-NR       Unrelated       PBSCT       matched       Modified BUCY       In       Intensive       MTX + CsA + ATG         20       Male       ALL-NR       Unrelated       PBSCT       Matched       Over modified       WTX + CsA + ATG         21       Female       CML-CP       Related       PBSCT       Matched       3genetic locus       TBI + CY       IV       Intensive       MTX + CsA + ATG         22       female       CML-CP       Related       PBSCT       Matched       3genetic locus       TBI + CY       IV       Intensive       MTX + CsA + ATG         23       Male       AML-NR       related       PBSCT       Matched       1genetic locus       TBI + CY       II       Intensive       MTX + CsA + ATG         36       Male       AML-NR       related       PBSCT       Matched       1genetic locus       TBI + CY       II       III <td></td> <td>41</td> <td>Female</td> <td>ALL-CR</td> <td>Related</td> <td>PBSCT</td> <td>2 antigen</td> <td>I</td> <td>TBI + CY</td> <td>=</td> <td>I</td> <td>+ CsA +</td> <td></td> <td>I</td> <td>59 days</td>		41	Female	ALL-CR	Related	PBSCT	2 antigen	I	TBI + CY	=	I	+ CsA +		I	59 days
28       Male       CML-CP       related       PBSCT       manatched       Modified BUCY       Intensive       MTX + CSA         20       Male       ALL-NR       Unrelated       PBSCT       Matched       -       Modified BUCY       IV       -       MTX + CSA         20       Male       ALL-NR       Unrelated       PBSCT       Matched       -       Modified BUCY       IV       -       MTX + CSA         22       female       CML-CP-PR       Unrelated       PBSCT       1 antiched       Over modified BUCY       IV       -       MTX + CSA       +       +       MTX + CSA         17       Female       CML-CP       Related       PBSCT       1 antiched       Over modified BUCY       II       -       MTX + CSA       +       +       MTX<+		32	Male	CML-CP-PR		BMT	Matched	1 genetic locus	Modified BUCY	=	Intensive	+		I	855 days
28       Male       CML-CP       related       PBSCT       matched       -       MTX + CsA         20       Male       AML-NR       Related       PBSCT       Matched       -       MTX + CsA         20       Male       AML-NR       Related       PBSCT       Matched       -       MTX + CsA         20       Male       AML-NR       Unrelated       PBSCT       Matched       -       MCTX + CsA         22       female       CML-CP-PR       Unrelated       PBSCT       Matched       -       MCTX + CsA       +         17       Female       CML-CP-PR       Unrelated       PBSCT       1 antigen       -       MCTX + CsA       +         16       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA       +         36       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA       +         36       Male       AML-PR       Natched       1 genetic locus       TBI + CY       II       -       MTX + CsA       +         37       Male       AML-								mismatched				+ MMF			
40       Male       AML-NR       Related       PBSC1       Matched       - MIX + CsA         20       Male       ALL-NR       Unrelated       PBSC1       Matched       - Modified       V       - MIX + CsA         22       female       CML-CP-PR       Unrelated       PBSC1       Matched       - Modified       V       - MIX + CsA         22       female       CML-CP-R       Unrelated       PBSC1       1 antigen       - Modified       - MIX + CsA         17       Female       CML-CP-R       Related       PBSC1       1 antigen       - MIX + CsA       + MMF         16       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       - MIX + CsA         36       Male       AML-NR       related       PBSC1       Matched       - Mix + CSA       + MMF         37       Male       AML-NR       related       PBSC1       Matched       - TBI + CY       II       Limited       MIX + CSA         33       Male       AML-PR       Matched       - TBI + CY       III       Limited       MIX + CSA         33       Male       AML-PR       Matched       - TBI + CY       III		28	Male	CML-CP	related	PBSCT	matched	I	Modified BUCY	=	Intensive	+	I	+	854 days
20       Male       ALL-NR       Unrelated       PBSCT       Matched       matched       Cover modified       IV       Limited       MTX + CsA + MMF         17       Female       CML-CP-PR       Unrelated       PBSCT       Matched       3 genetic locus       TBI + CY       IV       Limited       MTX + CsA + MMF         17       Female       CML-CP-PR       Unrelated       PBSCT       1 antigen       -       MTX + CsA + MMF         16       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + MMF         36       Male       AML-NR       related       PBSCT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + MMF         32       Male       AML-NR       related       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA + MMF         32       Male       AML-PR       Intensive       MTX + CsA       -       MTX + CsA       +         33       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       III       Limited       MTX + CsA       +         33		40	Male	AML-NR	Related	PBSCT	Matched		Modified BUCY	2		+ CsA	I	+	166 days
22       female       CML-CP-PR       Unrelated       PBSCT       Matched       3 genetic locus       TBI + CY       IV       Intensive       MTX + CsA + + MMF         17       Female       CML-CP-PR       Unrelated       PBSCT       1 antigen       -       Modified BUCY       II       -       + MMF         16       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + + MMF         36       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + + MMF         36       Male       AML-PR       Unrelated       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA + + MMF         37       Male       AML-PR       Unrelated       PBSCT       Matched       -       TBI + CY       III       -       MTX + CsA + MMF         32       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       III       Intensive       MTX + CsA + MMF         33       Male       MDS       Related       PBSCT       Matched       -       TBI + CY		20	Male	ALL-NR		PBSCI	Matched	matched	Over modified	2	Limited	+ CsA +	۱ ت	+	122 days
17       Female       CML-CP       Related       PBSCT       1 antigen       -       Modified BUCY       II       -       HMMF         16       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + + MMF         36       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + + MMF         36       Male       AML-NR       related       PBSCT       Matched       1 genetic locus       TBI + CY       II       -       MTX + CsA + + MMF         32       Male       AML-PR       Related       PBSCT       Matched       1       -       TBI + CY       III       -       MTX + CsA + + MMF         32       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       III       -       MTX + CsA + MMF         33       Male       MDS       Related       PBSCT       Matched       -       TBI + CY       III       Intensive       MTX + CsA         42       male       CLL-NR       Related       PBSCT       Matched       -       TBI + CY       III		22	female	CML-CP-PR		PBSCT	Matched	3 genetic locus	TBI + CY	$\geq$	Intensive	CsA +	- 5	I	682 days
16       Male       ALL-CR       Unrelated       BMT       mismatched       + MMF       + MMF         36       Male       ALL-CR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       I       Limited       MTX + CsA + MMF         36       Male       AML-NR       related       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA + MMF         32       Male       AML-PR       Unrelated       BSCT       Matched       -       TBI + CY       II       -       MTX + CsA + MMF         32       Male       AUL-PR       Related       PSCT       Matched       -       TBI + CY       II       Intensive       MTX + CsA + MMF         33       Male       AUL-PR       Related       PSCT       Matched       -       TBI + CY       II       Intensive       MTX + CsA + MIX + CsA + MIX + CsA         34       Male       CUL-NR       Related       PSCT       Matched       -       TBI + CY       II       Intensive       MTX + CsA + MIX		17	Female	CML-CP	Related	PBSCT	1 antigen	mismatched -	Modified BUCY	≡	Ι	+	- D	+	672 days
36       Male       AML-NR       related       PBSCT       Matched       + MMF         24       Female       AML-PR       Unrelated       BNT       Matched       -       TBI + CY       III       -       MTX+CsA         32       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       III       -       MTX+CsA         32       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       III       Limited       MTX + CsA         32       Male       MDS       Related       PBSCT       Matched       -       TBI + CY       II       Intensive       MTX + CsA         42       male       CLL-NR       Related       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA         43       Male       CML-BP       Unrelated       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA         43       Male       CML-CP-NR       Related       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA         43       Male       CML-CP-NR       Related       PBSCT		16	Male	ALL-CR	Unrelated	BMT	mismatched Matched	1 genetic locus	+	=	Limited	+	ا ا	+	652 days
24       Female       AML-PR       Unrelated       BMT       Matched       1 genetic locus       TBI + CY       III       Limited       MTX + CsA       +         32       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       III       Limited       MTX + CsA       +         32       Male       AUL-PR       Related       PBSCT       Matched       -       TBI + CY       II       Intensive       MTX + CsA         34       Male       MDS       Related       PBSCT       Matched       -       TBI + CY       II       Limited       MTX + CsA         42       male       CLL-NR       Related       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA         43       Male       CML-BP       Unrelated       PBSCT       Matched       -       TBI + CY       II       -       MTX + CsA         18       Male       CML-CP-NR       Related       BMT       2 antigen       -       MTX + CSA       +       MMF         18       Male       CML-CP-NR       Related       BMT       2 antigen       -       MTX + CSA       +       MMF         18		36	Male	AML-NR	related	PBSCT	Matched	mismatched -	+	≡	I	+ MMF MTX+CsA	I	I	46 davs
<ul> <li>32 Male AUL-PR Related PBSCT Matched</li> <li>33 Male MDS Related PBSCT Matched</li> <li>34 Male MDS Related PBSCT Matched</li> <li>35 Male CLL-NR Related PBSCT Matched</li> <li>36 male CLL-NR Related PBSCT Matched</li> <li>37 mismatched</li> <li>38 Male CML-BP Unrelated PBSCT Matched</li> <li>39 mismatched</li> <li>30 mismatched</li> <li>31 mismatched</li> <li>31 mismatched</li> <li>32 mismatched</li> <li>33 mismatched</li> <li>34 mismatched</li> <li>35 mismatched</li> <li>36 mismatched</li> <li>37 mismatched</li> <li>38 matched</li> <li>39 mismatched</li> <li>30 modified BUCY II</li> <li>31 mismatched</li> <li>31 mismatched</li> <li>32 mismatched</li> <li>33 mismatched</li> <li>34 modified BUCY II</li> <li>34 mMF</li> <li>34 mMF</li> <li>34 mMF</li> </ul>		24	Female	AML-PR	Unrelated	BMT	Matched	1 genetic locus	+	: ≡	Limited	+	آ آ	+	245 davs
<ul> <li>Male AUL-FR Related PBSCT Matched - TBI + CY III Intensive MIX + CSA</li> <li>Male MDS Related PBSCT Matched - TBI + CY II Limited MTX + CSA</li> <li>42 male CLL-NR Related PBSCT Matched - TBI + CY III - MTX + CSA</li> <li>43 Male CML-BP Unrelated PBSCT Matched 2 genetic locus TBI + CY III - MTX + CSA + mismatched</li> <li>18 Male CML-CP-NR Related BMT 2 antigen - Modified BUCY II - MTX + CSA + MMF</li> </ul>		0						mismatched		Ξ					
<ul> <li>Male MUS Helated PBSCT Matched - TBL + CY II LIMIted MIX + CSA</li> <li>male CLL-NR Related PBSCT Matched - TBL + CY III - MIX + CSA + 43 Male CML-BP Unrelated PBSCT Matched 2 genetic locus TBL + CY II Intensive MTX + CSA + mismatched</li> <li>Male CML-CP-NR Related BMT 2 antigen - Modified BUCY II - MTX + CSA + + MMF</li> </ul>		N N	Male	AUL-PH	Kelated		Matched	I	+ •	= =	Intensive	+ •	I	•	5/2 days
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18 Male CML-CP-NR Related BMT 2 antigen – Modified BUCY II – MTX + CsA + mismatched + MMF								mismatched		= :			2	F	0,0 days
		18	Male	CML-CP-NH		BMI	2 antigen	I	Modified BUCY	=	I	+	- 5	+	500 days
15 Male CML-BP-NR Unrelated PBSCT Matched - TBI + CY IV 0 MTX + + MMF	46	15	Male	CML-BP-NR	Unrelated	PBSCT	Matched	I	TBI + CY	≥	0	CsA +	- 51	+	28 days
47 36 Male AML-CR Related PBSCT Matched – Modified BUCY II –		36	Male	AML-CR	Related	PBSCT	Matched	I	Modified BUCY	=	Ι	-	I	I	428 days

chest HRCT aggravates or has no improvement after treatment).

#### Chest HRCT

Chest HRCT scans were performed in 45 patients within 3 days after the onset of symptoms of acute GVHD. Another two patients had chest HRCT scans after the body temperature was over  $38.5^{\circ}$ C for over 12 h and the occurrence of dyspnea, respectively.

#### Histopathology and immunohistochemistry

Transbronchial biopsy was performed in four patients whose chest HRCT did not recover completely after treatment for acute GVHD. Samples of lung tissue were obtained using fibreoptic bronchoscopy and immersed in 10% neutral formalin. Formalin-preserved specimens were then embedded in paraffin, cut into  $1-2 \mu m$  thick sections, and stained with haematoxylin and eosin for histopathology examination. For immunohistochemistry examination, paraffin-embedded sections were dewaxed, rehydrated and incubated with 0.5% hydrogen peroxide in methanol to quench endogenous tissue peroxidase. Sections were incubated with pepsin for 45 min for antigen retrieval. After blocking nonspecific sites with 1% BSA in PBS, sections were treated with primary anti-CD3 and anti-CD68 (DAKO) and with appropriate horseradish peroxidase-conjugated secondary antibodies for 90 and 45 min, respectively.

#### Pulmonary function

Pulmonary function was measured three times monthly in 24 patients who survived more than 6 months. A decrease of less than 80% of the total lung capacity was defined as a new restrictive physiological defect. A new obstructive defect was defined by a 15% or more reduction in forced expiratory volume in one-second with a concomitant decrease in the forced expiratory volume in one-second to forced vital capacity ratio. A certified technician performed all pulmonary function tests.

#### The levels of serum IFN $\gamma$ and TNF $\alpha$

The levels of serum IFN $\gamma$  and TNF $\alpha$  were measured with enzyme linked immunosorbent assay (ELISA) in 47 patients before the treatment for acute GVHD. Venous peripheral blood from each patient was drawn into tubes containing EDTA and into anticoagulant-free tubes. After centrifugation, serum samples were stored at  $-20^{\circ}$ C until required. ELISA was performed according to the manufacturer's instruction (IFN $\gamma$  and TNF $\alpha$  ELISA kit, Jingmei Corporation, China). The healthy group consisted of 10 healthy adults (five males and five females) who were enrolled in the study.

#### Prophylaxis of infection

All patients were treated with compound sulfamethoxazole and norfloxacin by oral administration. Ganciclovir was used as the fundamental prophylaxis for cytomegalovirus (CMV) infection. Antifungal agents were used 5 days before allo-HSCT. Patients who did not have invasive fungal infection (IFI) history took fluconazole at a dose of 0.3 g once daily until peripheral white blood cells count was over  $2.0 \times 10^{9}$ /l. Patients who had IFI history used itraconazole at a dose of 0.4 g once daily or voriconazole at a dose of 0.4 g once daily or AmBisome at a dose of 2 mg/kg once daily by intravenous administration. Oral application of itraconazole and voriconazole were started after peripheral white blood cells count was over  $2.0 \times$ 10<sup>9</sup>/1 and discontinued for 90 days, except one case with Mucor infection receiving AmBisome by intravenous administration intermittently.

#### Results

#### Onset time and characteristics of acute GVHD

In 47 patients, 25 had acute GVHD of grade II, 13 had acute GVHD of grade III and nine had acute GVHD of grade IV. The onset time of acute GVHD was from +17 to +122 days with the median day of +31 days after allo-HSCT. Thirty-two patients had skin, liver and gut involved, 10 patients had two organs involved and five patients had only skin or liver involved. At the onset of acute GVHD, 42 patients had fever, 28 patients had dry cough and 11 patients presented with shortness of breath. Two patients had respiratory symptoms and abnormal chest HRCT without skin, liver and gut involved early at the onset of acute GVHD-induced lung injury. The first patient had fever and her HRCT showed diffuse interstitial and alveolar infiltrate accompanied by bilateral pleural effusion. Antiinfection treatment was discontinued and methylprednisolone was added when her clinical condition and chest HRCT worsened after anti-infection for 1 week. Respiratory symptoms were improved and the temperature returned to normal by the third day after methylprednisolone was added. The repeat chest HRCT showed pleural effusion and hydropericardium disappeared with lung diffuse infiltrate improved obviously after anti-GVHD for 10 days (Fig. 1A-F). Rechecked chest HRCT showed thickening and deranged interstitial markings 95 days after the onset (Fig. 1G). The second had only

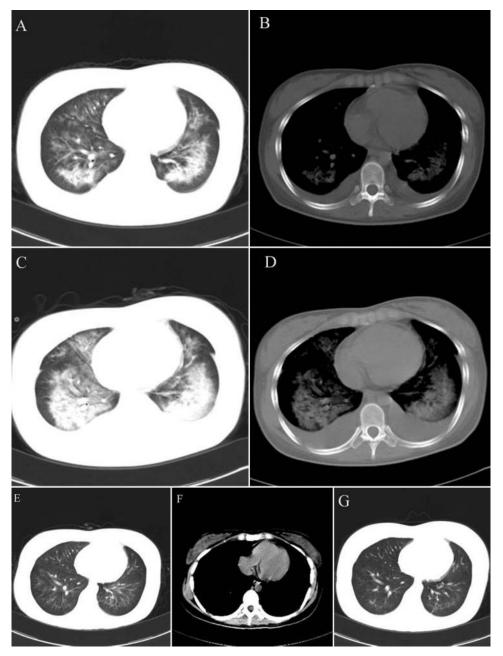


Figure 1 This patient had only fever, respiratory symptoms and abnormal chest HRCT without skin, liver and gut involvement on +89 days post-transplantation. Chest HRCT showed diffused interstitial and alveolar infiltrate (A) and bilateral pleural effusion accompanied by hydropericardium (B) at the onset. Chest HRCT showed diffused interstitial and alveolar infiltrate worsened (C) and worsened bilateral pleural effusion (D) after anti-infection for 1 week. Chest HRCT showed that intraparenchymatous infiltrate was obviously improved (E), hydropericardium and pleural effusion disappeared (F) after anti-GVHD for 10 days. Chest HRCT showed thickening interstitial markings and deranged vascular shadow after anti-GVHD for 95 days (G)

respiratory symptoms without fever when chest HRCT showed diffuse interstitial accompanied by bilateral pleural effusion and hydropericardium. Thoracentesis was performed and pathogen culture of hydrothorax was negative. Her cough and shortness of breath resolved and rechecked chest HRCT showed that interstitial infiltrate and hydropericardium were obviously improved and pleural effusion disappeared after anti-GVHD for 11 days (Fig. 2A–D). Chest HRCT recovered completely 88 days after the onset (Fig. 2 E). These two patients presented with manifestations of acute GVHD (grade II) on skin and liver 9 and 11 days after the occurrence of respiratory symptoms, respectively, and evolved to limited chronic GVHD of skin and liver, respectively.

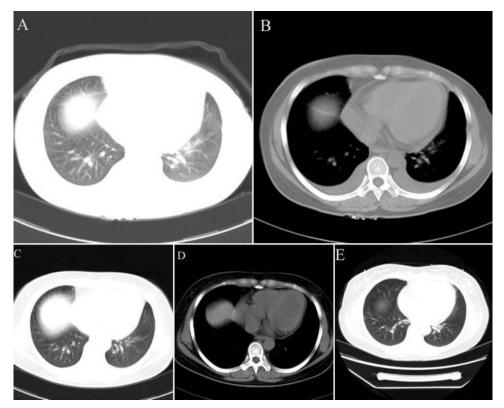


Figure 2 This patient had only respiratory symptoms and abnormal chest HRCT without skin, liver and gut involvement on +122 day post-transplantation. Chest HRCT showed diffused interstitial infiltrate (A), bilateral pleural effusion and hydropericardium (B) at the onset. Chest HRCT showed that interstitial infiltrate was obviously improved (C), pleural effusion disappeared and hydropericardium obviously improved (D) after treatment for GVHD for 11 days. Chest HRCT returned to normal after anti-GVHD for 88 days (E)

#### Characteristics of chest HRCT

Chest HRCT scans were performed in 47 patients and 20 patients had imaging abnormalities. Seventeen cases were suspected of acute GVHD-induced lung injury and three cases were diagnosed with IFI relapse, invasive *Aspergillus* pneumonia and bacterial pneumonia, respectively. In 17 cases suspected of acute GVHD-induced lung injury, HRCT revealed diffused interstitial infiltrate in five cases, diffused interstitial and alveolar infiltrate in seven cases, and diffused interstitial and segmental lobar infiltrate in five cases

accompanied by bilateral pleural effusion and hydropericardium in nine patients (Fig. 3). HRCT showed that 14 cases had the whole lung involved and three cases had the median and inferior lobe involved.

#### Levels of serum IFN $\gamma$ and TNF $\alpha$ and analysis of hydrothorax

The levels of serum IFN $\gamma$  were 6.901  $\pm$  1.751 ng/ml in the cases with lung injury, 6.280  $\pm$  1.150 ng/ml in the cases without lung injury and 3.343  $\pm$  0.737 ng/ ml in the healthy group. The levels of serum TNF $\alpha$ were 399.514  $\pm$  101.598 pg/ml in the cases with lung

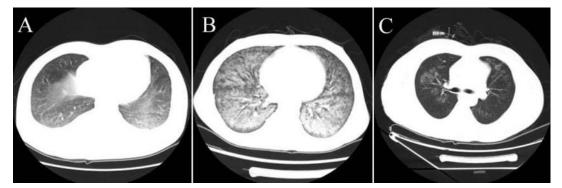


Figure 3 (A) Diffused interstitial infiltrate; (B) diffused interstitial and alveolar infiltrate; (C) diffused interstitial and segmental lobar infiltrate

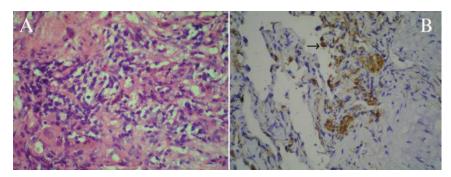


Figure 4 (A) H&E-stained histological section of the transbronchial lung biopsy. There is lymphocytic infiltrate which consists mostly of T lymphocytes (HE, ×200). (B) Immunohistochemistry. There is macrophage infiltrates (CD68+, ×200)

injury,  $427.871 \pm 83.287$  pg/ml in the cases without lung injury and  $79.298 \pm 14.555$  pg/ml in the healthy group. No statistical significance was shown in the levels of serum IFN $\gamma$  and TNF $\alpha$  between cases with and without lung injury (IFN $\gamma$ : p=0.202, TNF $\alpha$ : p=0.306), but the levels of serum IFNy and TNF $\alpha$  in patients were higher than in the healthy group (IFN<sub>γ</sub>: p=0.000, TNF $\alpha$ : p=0.000). Four of nine patients with bilateral pleural effusion had thoracentesis. Examination of hydrothorax showed that white blood cell count 1200  $\pm$  517·204/µl (800–2200/µl) and the quantity of protein  $38.00 \pm 13.06$  g/l (25.00-62.00 g/l) increased. Bacterium or fungus cultures of hydrothorax were negative.

#### Therapeutic effect

Forty-seven cases all attained the treatment for acute GVHD and 26 of them received CR: nine were PR and 12 were NR. The total effective rate was 74.47% and the CR rate was 55.32% in the treatment for acute GVHD. The total effective rates of treatment for acute GVHD were 88.24 and 66.67% in patients with (CR: 10 cases, PR: five cases and NR: two cases) and without lung injury (CR: 16 cases, PR: four cases and NR: 10 cases), respectively. There was no statistical significance in the effective rate of treatment for acute GVHD between patients with and without lung injury (p=0.200). Seventeen patients suspected of acute GVHD-induced lung injury rechecked chest HRCT during 7-14 days after anti-GVHD. Rechecked HRCT scans showed that 10 patients received CR: six patients were PR and one patient aggravated. The total effective rate of treatment for acute GVHD-induced lung injury was 94.12% and the CR rate was 58.82%. Comparison between the effective rates of treatment for acute GVHD and acute GVHD-induced lung injury showed no statistical significance (p=0.169). The effective rate of treatment for acute GVHD-induced

lung injury positively correlated with that for acute GVHD (r=0.771, p=0.001).

#### Histopathology and immunohistochemistry

Transbronchial biopsy was performed in four patients whose HRCT showed PR during 26–52 days after the onset of acute GVHD. The histopathology of the lung tissue was characteristic by disorganization, epithelial cell damage, interstitial fibroplasias and cells infiltrate. Immunohistochemistry showed that three cases had predominately T lymphocyte infiltrate and one case had macrophage infiltrate (Fig, 4).

#### Pulmonary function

Seven patients had abnormal pulmonary function in nine patients with acute GVHD-induced lung injury who survived more than 6 months, including six patients with obstructive defect and one patient with restrictive defect. Four patients had abnormal pulmonary function in 15 patients without acute GVHD-induced lung injury who survived more than 6 months, including two patients with obstructive defect, one patient with restrictive defect and one patient with both. There was significance in the incidence of abnormal pulmonary function between patients with or without acute GVHD-induced lung injury (p=0.033, cross-tabs).

#### The relationship between acute GVHD-induced lung injury and late-onset non-infectious lung injury

Three patients developed late-onset non-infectious lung injury in nine patients with acute GVHD-induced lung injury and three had late-onset non-infectious lung injury in 15 patients without acute GVHD-induced lung injury who survived more than 6 months, according to diagnostic criteria of clinical manifestation, imaging characteristics and pulmonary function in references of late-onset non-infectious lung injury.<sup>9,10</sup> The incidence of late non-infectious lung injury between these two groups had no significant difference (p=0.635).

#### Survival status

Until now, 18 patients still survived and disease free survival rate for 3 years was 38·30%. Six patients died of acute GVHD, three secondary infections and one late-onset non-infectious lung injury, in 17 patients with acute GVHD-induced lung injury. Ten cases died of acute GVHD, five secondary infection, one lateonset non-infectious lung injury, one renal inadequacy, one leukaemia relapse and one lymphoproliferative diseases in 30 patients without lung injury. Disease free survival rate for 3 years was  $41\cdot18 \pm 11\cdot94$  and  $26\cdot48 \pm 12\cdot36\%$  in cases with and without lung injury, respectively. There was no statistical significance of disease free survival rate between patients with and without lung injury (p=0.707).

#### Discussion

Lung injury is a major complication of allo-HSCT that occurs in 25-50% of recipients and can account for approximately 50% of transplant-related mortality.<sup>2,3</sup> It is a major prognostic factor following allo-HSCT. Lung injury after allo-HSCT can be divided into infectious and non-infectious lung injury according to the pathogeneses and can be either early lung injury which occurs within 100 days or late-onset lung injury that occurs over 100 days after allo-HSCT. Historically, about half of all lung injury after allo-HSCT has been secondary to infection, but the judicious use of broad-spectrum antimicrobial prophylaxis in recent years has altered the balance of lung injury from infectious to non-infectious causes. A large body of literature demonstrated that late-onset non-infectious lung injury after allo-HSCT was immunologically mediated disease and shared similar pathogenesis with chronic GVHD.<sup>1-3,11</sup> On the issue about the relationship of acute GVHD and early lung injury, in 1978, Beschorner et al. reported a group of patients who developed lymphocytic bronchitis associated with acute GVHD after allo-HSCT.<sup>4</sup> Many studies suggested that the lung was the target organ of acute GVHD and some early lung injury was related to acute GVHD confirmed by animal experiment and clinical cases, but whether the lung is one of the target organs of acute GVHD remains controversial.<sup>1,2,11</sup> Some reports suspected the association between early lung injury and acute GVHD based on the absence of specific epithelial apoptosis in the lung tissue which is considered as histological feature for acute GVHD that existed in the skin, liver and gut, and there is no consistency of the occurrence of early lung injury and acute GVHD. However, the thymus is a known target of GVHD and displays extensive cytolytic damage in

the course of GVHD, but epithelial cell apoptosis is not a prominent histological feature.<sup>12</sup> Recently, Cooke et al. reported that lung damage was also detectable in mice after T lymphocyte depleted allo-HSCT when signs of clinical and histological acute GVHD were absent.<sup>2</sup> Bolanos-Meade et al. and Knox et al. reported patients who presented with pneumonia as the early manifestation of acute GVHD without skin, liver and gut involvement after allogeneic peripheral haematopoietic stem cell transplantation and liver transplantation, respectively.<sup>1,13</sup> In this report, 20 of 47 patients with grades II-IV acute GVHD after allo-HSCT had abnormal chest HRCT. Three cases were diagnosed of infectious lung injury and 17 cases were suspected of acute GVHD-induced lung injury. In total, 94.12% of patients' lung injury was improved in17 cases suspected of acute GVHDinduced lung injury after anti-GVHD treatment. The efficacy of treatment for acute GVHD-induced lung injury paralleled with that for GVHD. Additionally, two of these 17 patients had no any stigmata of acute GVHD in skin, liver and gut early at the onset of acute GVHD-induced lung injury. Clinical condition and chest HRCT improved after anti-GVHD treatment. These two patients presented with manifestations of acute GVHD (grade II) on skin and liver 9 and 11 days after the onset of lung injury, respectively, and evolved to limited chronic GVHD. These confirmed that lung injury correlated to acute GVHD and lung injury might be the initial manifestation of acute GVHD.

There are few reports about chest imaging characteristics of acute GVHD-induced lung injury. Early idiopathic pneumonia syndrome after allo-HSCT, which is reported to be related to acute GVHD, has multilobar infiltrate as one of the chest imaging diagnosis criteria.<sup>14–17</sup> Recently, Bolanos-Meade et al. reported a case of acute GVHD-induced lung injury after allo-HSCT. Chest imaging characteristics of the reported case included diffuse centrilobular nodules, lung parenchymal and interstitial infiltrate.<sup>1</sup> In this study, 17 cases suspected of acute GVHD-induced lung injury had 14 cases with the whole lung involved and three cases with the median and inferior lobe involved. HRCT characteristics including diffused interstitial and alveolar infiltrate were similar to the previous reports. Furthermore, nine cases suspected of acute GVHD-induced lung injury in this study had bilateral pleural effusion and hydropericardium.

The pathogenesis of acute GVHD-induced lung injury is not clear. Yanik and Cooke<sup>3</sup> proposed the mechanisms of T-cell axis and gut–liver–lung inflammatory cytokine axis. The lymphocyte activation pathway depends on interactions between donor T cells and host antigen-presenting cells. Donorderived T cells are activated by host antigen presenting cells and secrete inflammatory cytokines like IFN $\gamma$ , IL-2 and IL-1 which subsequently recruit to the lung by specific chemokine receptor, and finally contribute to the early pro-inflammatory events associated with lung injury. Donor macrophages, primed by cytokines produced by gut-liver-lung inflammatory cytokine axis, are recruited to the lung where they are triggered by lipopolysaccharide to secrete inflammatory cytokines like  $TNF\alpha$ , resulting in enhanced chemokine expression, the recruitment of neutrophils to the lung and increased tissue damage.<sup>2,3,5,6,14</sup> In our four patients, immunohistochemistry of lung tissue showed T lymphocyte infiltrate predominately in three cases and macrophage infiltrate in one case. It suggested that T lymphocyte and macrophage might play a role in acute GVHD-induced lung injury. In this study without non-GVHD transplant controls, the difference of the levels of serum IFN $\gamma$  and TNF $\alpha$  in patients with and without lung injury showed no statistical significance, but the levels of serum IFN $\gamma$ and TNF $\alpha$  in patients were significantly higher than that in healthy group. The results suggested that IFN $\gamma$  and TNF $\alpha$  might be involved in lung injury. The comparison of levels of serum IFN $\gamma$  and TNF $\alpha$ in GVHD and non-GVHD patients might further demonstrate the participation of IFN $\gamma$  and TNF $\alpha$  in acute GVHD-induced lung injury.

Reports about the association between early noninfectious lung injury and late-onset non-infectious lung injury after allo-HSCT are rare. It was reported that the incidence of late-onset non-infectious lung injury was higher in patients with acute GVHD than in those without acute GVHD, which suggested that acute GVHD might be related to late-onset non-infectious lung injury.<sup>10,17–19</sup> Twenty-four of 47 patients with grades II-IV of acute GVHD who survived more than 6 months were followed up. Three cases each developed late-onset non-infectious lung injury in cases with and without lung injury. Although the difference of the incidence of late non-infectious lung injury was not significant, there was significance in the incidence of abnormal pulmonary function between cases with and without lung injury. These suggested that early acute GVHD-induced lung injury could evolve to late-onset non-infectious lung injury.

These results suggested that lung might be one of the target organs of acute GVHD and participation of T lymphocyte, macrophage and cytokines like IFN $\gamma$  and TNF $\alpha$  might play a role in the pathogenesis of acute GVHD-induced lung injury. GVHD-induced early lung injury might progress to late-onset non-infectious lung injury.

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