

Supplementary Methods

Definition

Acute graft-versus-host disease (GVHD) was diagnosed and graded clinically according to the Glucksberg's criteria [1], and chronic GVHD according to the National Institutes of Health Consensus criteria [2]. Relapse was defined by any evidence of the disease after the hematopoietic cell transplantation (HCT), and non-relapse mortality (NRM) was defined as death from any cause except for relapse. The HCT comorbidity index (HCT-CI) was applied according to a previous study by Sorror *et al.* [3] The disease risk index (DRI) consisted of the disease and stage risk, each of which was derived from a diagnosis with the cytogenetics data and remission status at the time of the HCT, respectively, and has been shown to successfully risk stratify heterogeneous allogeneic transplant recipients [4]. For the

current study, the DRI was collapsed into a 2-group system of low/intermediate and high/very high risk, as proposed by the original study [4].

Statistical endpoints included GVHD-free, relapse-free survival (GRFS), disease-free survival (DFS), and overall survival (OS). GRFS is a composite endpoint encompassing ongoing morbidity from GVHD in addition to relapse and death [5, 6]. GRFS events were defined as grade III-IV acute GVHD, chronic GVHD requiring systemic immunosuppressive treatment, relapse, or death. DFS was calculated as the time from the HCT to relapse or death, and OS was defined as the time from the HCT to death.

Statistical methods

Patient and treatment characteristics were compared with Chi-square test for categorical variables and t-test for continuous variables. Survival curves were plotted using the

Supplementary Table 1. Patients and treatment characteristics.

| Variables | Overall (N=130) | By ATG doses | | P ^{a)} |
|---|--------------------|-------------------|--------------------|-----------------|
| | | 6 mg/kg (N=43) | 9 mg/kg (N=87) | |
| Patient age, median (range) | 52 (16–68) | 55 (18–67) | 52 (16–68) | 0.219 |
| Patient sex, N (%) | | | | 0.264 |
| Male | 68 (52.3%) | 19 (44.2%) | 49 (56.3%) | |
| Female | 62 (47.7%) | 24 (55.8%) | 38 (43.7%) | |
| HCT-CI total scores, median (range) | 0 (0–5) | 0 (0–4) | 0 (0–5) | 0.411 |
| Disease, N (%) | | | | 0.073 |
| AML | 78 (60.0%) | 28 (65.1%) | 50 (57.5%) | |
| MDS | 19 (14.6%) | 2 (4.7%) | 17 (19.5%) | |
| ALL | 33 (25.4%) | 13 (30.2%) | 20 (23.0%) | |
| Disease risk index | | | | 1.000 |
| Low/intermediate | 99 (76.2%) | 33 (76.7%) | 66 (75.9%) | |
| High/very high | 31 (23.8%) | 10 (23.3%) | 21 (24.1%) | |
| Time from diagnosis to HCT in mo, median (range) | 5.92 (2.07–197.43) | 5.46 (2.07–31.50) | 6.07 (2.57–197.43) | 0.450 |
| Female donor-to-male patient, N (%) | 21 (16.2%) | 5 (11.6%) | 16 (18.4%) | 0.464 |
| Donor type | | | | 0.008 |
| HID | 55 (42.3%) | 25 (58.1%) | 30 (34.5%) | |
| PUD | 31 (23.8%) | 4 (9.3%) | 27 (31.0%) | |
| MUD | 44 (33.8%) | 14 (32.6%) | 30 (34.5%) | |
| ABO incompatibility | | | | 0.592 |
| Matched | 48 (36.9%) | 16 (37.2%) | 32 (36.8%) | |
| Major mismatch | 36 (27.7%) | 9 (20.9%) | 27 (31.0%) | |
| Minor mismatch | 27 (20.8%) | 10 (23.3%) | 17 (19.5%) | |
| Bidirectional | 18 (13.8%) | 8 (18.6%) | 10 (11.5%) | |
| Infused CD34 ⁺ cells in ×10 ⁶ /kg, median (range) | 5.14 (0.97–12.68) | 5.59 (1.80–12.68) | 4.97 (0.97–12.09) | 0.093 |
| CNI use | | | | < 0.001 |
| Cyclosporin A | 68 (52.3%) | 12 (27.9%) | 56 (64.4%) | |
| Tacrolimus | 52 (40.0%) | 30 (69.8%) | 22 (25.3%) | |
| Switch | 10 (7.7%) | 1 (2.3%) | 9 (10.3%) | |
| Methotrexate use | | | | 0.005 |
| Yes | 79 (60.8%) | 34 (79.1%) | 45 (51.7%) | |
| No | 51 (39.2%) | 9 (20.9%) | 42 (48.3%) | |

^{a)}P-value by χ^2 test for categorical variables and t-test for continuous variables.

Abbreviations: ALL, acute lymphoblastic leukemia; AML, acute myeloid leukemia; ATG, antithymocyte globulin; CNI, calcineurin inhibitor; HCT, hematopoietic cell transplantation; HCT-CI, hematopoietic cell transplantation comorbidity index; HID, haploidentical familial donors; MDS, myelodysplastic syndrome; MUD, matched unrelated donors; PUD, partially-matched unrelated donors.