

2098: CMV INFECTION FOLLOWING HEMATOPOIETIC CELL TRANSPLANTATION WITH PTCY: OUTCOMES IN HAPLOIDENTICAL VERSUS MATCHED OR MISMATCHED DONOR ALLOGRAFTS

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INTRODUCTION

- Post-transplant cyclophosphamide (PTCy) is increasingly being used for prevention of graft-versus-host disease (GVHD) in matched unrelated donor (MUD), mismatched unrelated donor (MMURD), and haploidentical donor hematopoietic cell transplantation (HCT).
- PTCy has been associated with increased risk of viral infections including respiratory viruses, herpesviruses, and cytomegalovirus (CMV) infection after HCT.
- It remains unclear if the increased risk of viral infections is independent of donor allograft selection.
- This study evaluated the incidence of clinically significant CMV infection (CS-CMVi) following PTCy in patients who received MUD/MMURD allografts compared to haploidentical donor allografts.

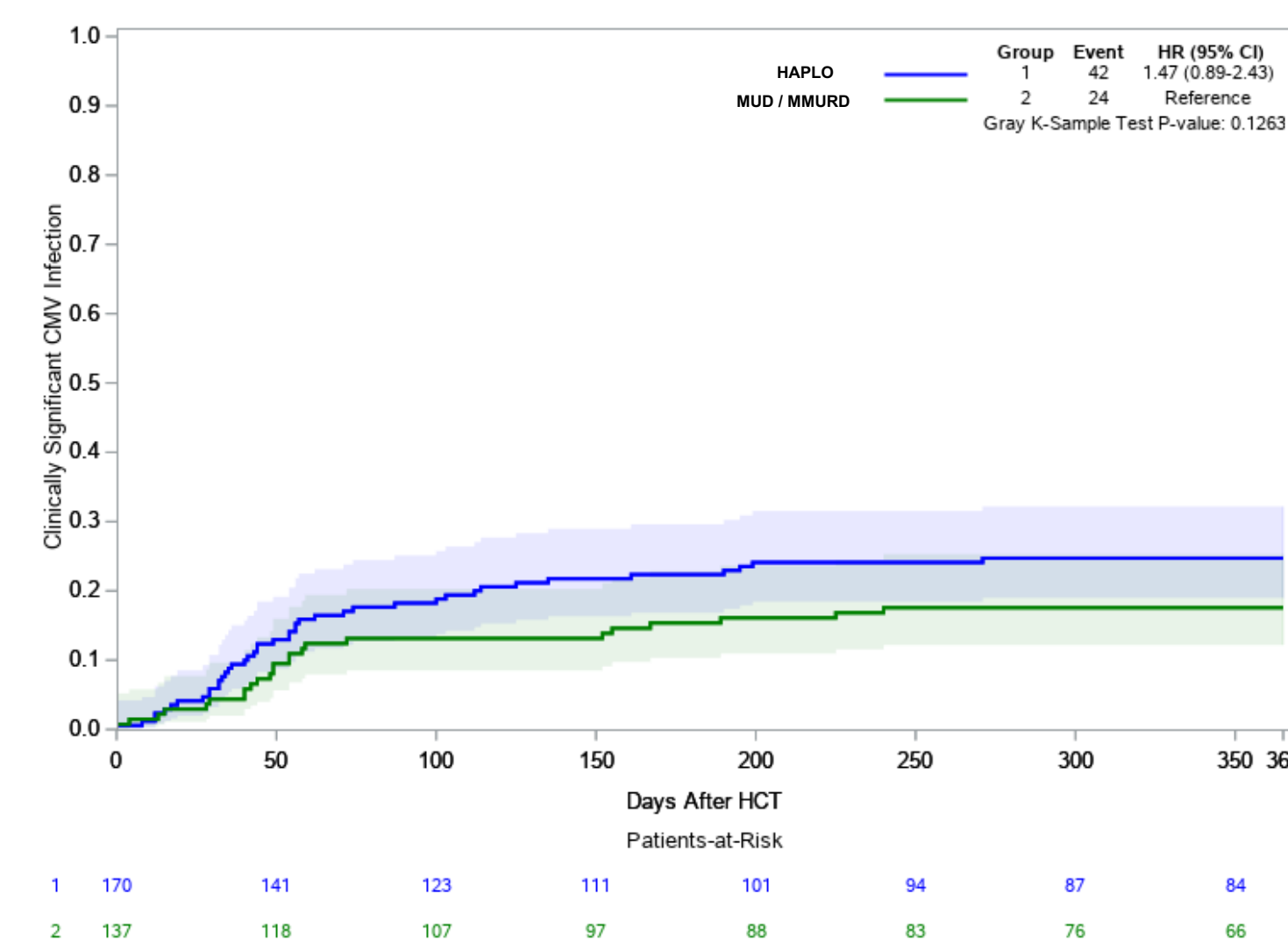
METHODS

- We performed a single-center retrospective study of adults undergoing HCT with PTCy between January 1, 2015 and July 1, 2021. Per institutional protocol, patients who were CMV seropositive (R+) received letermovir prophylaxis.
- CS-CMVi, defined as CMV viremia or disease requiring initiation of CMV therapy, was evaluated in two groups: haploidentical (n=170) and MUD/MMURD (n=137) allograft HCT recipients.
- CMV disease was defined according to criteria published by the Disease Definitions Working Group of the Cytomegalovirus Drug Development Forum.
- We assessed the incidence of CMV viremia, CS-CMVi, and CMV disease, as well as incidence of letermovir breakthrough infections and late CMV events after cessation of prophylaxis for one year following HCT.
- One-year Cumulative Incidence Functions were calculated based on time to CS-CMVi using dates of infection-free death, disease relapse, and repeat HCT as competing risks.

RESULTS

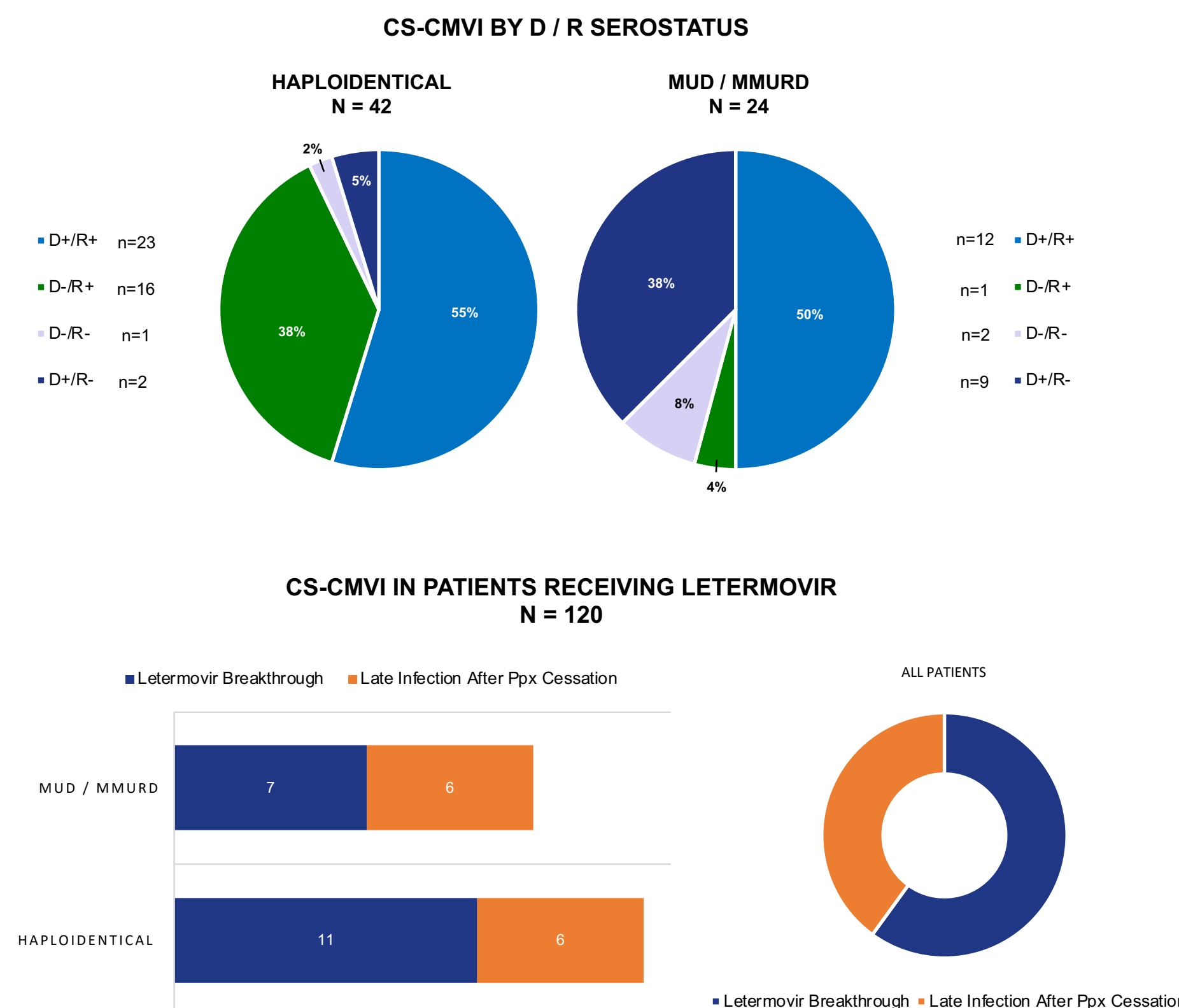
| Baseline Characteristics | Haploidentical HCT n=170 | MUD / MMURD HCT n=137 |
|---------------------------------------|-----------------------------|--------------------------|
| Demographics | | |
| Age, median (range) | 59 (19 – 77) | 58 (21 – 78) |
| Male sex, no. (%) | 103 (61) | 74 (54) |
| Underlying Disease | | |
| AML, no. (%) | 69 (41) | 51 (37) |
| MDS/MPN, no. (%) | 39 (23) | 35 (26) |
| ALL, no. (%) | 19 (11) | 24 (18) |
| NHL, no. (%) | 19 (11) | 10 (7) |
| HL, no. (%) | 4 (2) | 6 (4) |
| Other, no (%) | 20 (12) | 11 (8) |
| Transplant Characteristics | | |
| Prior HCT | | |
| Allogeneic | 19 (11) | 19 (14) |
| Autologous | 9 (5) | 4 (3) |
| Graft | | |
| Peripheral blood | 99 (58) | 111 (81) |
| Bone Marrow | 71 (42) | 26 (19) |
| Conditioning | | |
| Reduced Intensity | 127 (75) | 97 (71) |
| Myeloablative | 43 (25) | 40 (29) |
| HLA Match | | |
| Fully Matched (10 of 10) | - | 62 (45) |
| Mismatched (7, 8, or 9 of 10) | - | 75 (55) |
| CMV Serostatus | | |
| CMV D positive / R positive | 59 (35) | 47 (34) |
| CMV D positive / R negative | 14 (8) | 12 (9) |
| CMV D negative / R positive* | 40 (24) | 7 (5) |
| CMV D negative / R negative | 55 (32) | 71 (52) |
| Letermovir Primary Prophylaxis | | |
| CMV D positive / R positive | 66 (39) | 54 (39) |
| CMV D positive / R negative | 37 (22) | 47 (34) |
| CMV D negative / R positive | 26 (15) | 7 (5) |
| Other** | 3 (2) | 0 (0) |
| Transplant Outcomes | | |
| CRS Diagnosis | 102 (60) | 29 (21) |
| Acute GVHD | 85 (50) | 58 (42) |
| Grade 3 – 4 | 6 (4) | 11 (8) |
| Tx ≥ 1 mg/kg Corticosteroids | 48 (28) | 29 (21) |
| Death | 50 (29) | 36 (26) |
| Days to Death, median (range) | 223 (9 – 1795) | 152 (10 – 1026) |

*Two patients were donor seronegative with indeterminate recipient serostatus
**Two CMV D negative / R negative patients and one CMV D positive / R negative patients received letermovir prophylaxis



| CMV Outcomes | Haploidentical HCT n=170 | MUD / MMURD HCT n=137 | P-value |
|--------------------------------------|-----------------------------|--------------------------|---------|
| CMV Events | | | |
| CMV disease | 1 (0.6) | 1 (0.7) | - |
| CMV Viremia | 68 (40) | 50 (36) | 0.557 |
| Time to CMV Viremia, median (range) | 29 (2 – 337) | 29 (0 – 252) | - |
| CS-CMVi | 42 (25) | 24 (18) | 0.162 |
| Time to CS-CMVi, median (range) | 44 (0 – 315) | 49 (0 – 240) | - |
| CS-CMVi Risk group | | | |
| CS-CMVi in seropositive recipients | 39 (23) | 13 (9) | 0.002 |
| CS-CMVi in seronegative recipients | 3 (2) | 11 (8) | 0.012 |
| CMV Donor Status | | | |
| CS-CMVi with seropositive donor | 25 (15) | 21 (15) | 0.874 |
| CS-CMVi Infection Type | | | |
| Letermovir Breakthrough Infection | 11 (6) | 7 (5) | 0.808 |
| Infection After Letermovir Cessation | 6 (4) | 6 (4) | 0.772 |
| Not Receiving Letermovir | 25 (15)* | 11 (8) | 0.077 |

*One patient with CS-CMVi was on the Letermovir versus Placebo study



DISCUSSION

- Despite the use of letermovir prophylaxis, high-risk (CMV R+) haploidentical recipients had a higher incidence of CS-CMVi than MUD / MMURD recipients (23% versus 9%; p=0.002), suggesting that haploidentical HCT may have increased risk for viral infections independent of PTCy, possibly related to impaired antigen presentation in the setting of HLA mismatch.
- For low-risk (CMV R-) recipients, there was a notably increased incidence of CS-CMVi in the MUD/MMURD cohort (8% versus 2%; p=0.012), that was driven primarily by D+/R- recipients. This raises the question of whether letermovir may benefit (CMV R-) patients when the donor is seropositive.
- Out of 120 patients who received letermovir, 25% developed CS-CMVi, with 60% of those cases due to breakthrough infection and 40% following cessation of letermovir. Further data is needed on the impact of delayed CMV-specific immunity related to letermovir use.

KEY REFERENCES

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