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| Reference | Study design | Follow -up, (mo) | N | Study objective | Age range, (years) | MRD detection method | Time of MRD assessment | ISS | Statistics | Regimen | Mainte nance | Depth of response |
|-----------------------|--|-------------------------|----|--|--------------------------|----------------------------|--|---|--|--|-----------------|---|
| Rawstron 2002 | Prospective | up to 39 | 45 | Whether MFC results (levels of malignant vs normal plasma cells) predict outcomes after HDT and SCT | 41-65 | MFC | 3 months after Tx; 3–6 month intervals thereafter | NS | Univariate (log-rank test) and multivariate (Cox- regression) analysis | C-VAMP followed by MEL + HD ASCT | None | Following induction: 22% CR, 78% PR Following HD: 73% CR: 42% MRD ⁺ |
| San Miguel 2002 | Prospective, randomized, multicenter PETHEMA trial | 65 (PFS), 53 (OS) | 87 | Determine whether changes in the plasma cell compartment (MRD using MFC) could predict disease outcome | 31-70 | PCR | 3 months after ASCT, 1 month after 12 cycles chemotherapy | 58–61% stage II 39–42% stage III | Mann- Whitney U, Wilcoxon for between- group differences. Kaplan-Meier for survival curves | VBMCP/VBAD followed by ASCT or 8 cycles chemotherapy | IFNα + DEX | MRD ⁻ : 36% of ASCT patients vs 15% CT patients (p=0.04) |
| Ferrero 2014 | GIMEMA trial | 93 | 39 | Impact of MRD kinetics on survival when using VTD consolidation | 42–69 | PCR | After 2 VTD courses, end of treatment, every 6 months until relapse | NS | Univariate Cox proportional hazards model | VTD, MEL, ASCT | None | Full MRD: 18% Major MRD: 67% |
| Bakkus 2004 | NS | NS | 67 | Whether post- SCT tumor load predicts duration of response | 30–65 | PCR | 3–6 months post-HDT | 12% stage IIA, 47% stage IIIA, 8% stage IIIB | Log-rank test | VAD, MEL± TBI with single or tandem autologous PBSCT | None | 28% CR |
| Dal Bo 2013 | Prospective | 18 | 44 | Whether presence of MRD 3 months post- | 52.2– 64 | MFC | 3 months | NS | Log-rank test | MEL, ASCT | None | 32.6% CR, 40% MRD⁻ |

| | | | | SCT predicts relapse or death | | | | | | | | |
|------------------|--|----|--------------------------------------|--|-------|-----|---|---|---|---|----------------------------|--|
| Paiva 2011 | Prospective | 32 | 102 | Prognostic value of MFC vs IF vs SFLC | 65–84 | MFC | After 6 cycles of induction | 29% stage I, 38% stage II, 33% stage III | Two-sided log-rank test | VMP vs VTP for 6 cycles | VT vs VP for max 3 y | CR 43%, MRD 30% |
| Paiva 2008 | Prospective | 57 | 295 | Prognostic value of post-SCT MFC remission | 29–70 | MFC | 100 days | 39% stage I, 41% stage II, 20% stage III | Log-rank test | VBMCP/VBAD, MEL, ASCT | None | 50% CR, 42% MRD⁻ |
| Korthals 2012 | NS | 61 | 53 | Whether pre- and post-SCT MRD status predicts EFS/OS | 31–75 | PCR | 3–6 months after SCT | 11% stage I+II, 89% stage III | Kaplan-Meier plots and the log-rank test | Idarubicin/dex amethasone induction, MEL, ASCT | IFN or THAL | 25% nCR, 21% MRD⁻ |
| Korthals 2013 | Retrospective | 45 | 42 | Whether MRD status in PB predicts remission status | 31–66 | PCR | 3 mo | 12% Stage I+II, 88% Stage III | Kaplan-Meier plots and the log rank test. | Idarubicin/dex amethasone induction, MEL, ASCT | IFN or THAL | 28% CR |
| Swedin 1998 | NS | 29 | 36 | Utility and clinical value of ASO-PCR to evaluate MRD | 31–60 | PCR | 3 + 6 months after ASCT, 6 months thereafter | NS | Log rank test | VAD, MEL, ASCT | IFN | 50% CR |
| Rawstron 2013 | Prospective | 71 | 397 (INT) and 245 (nINT) | Prognostic value of MRD, measured using MFC, on outcomes | NS | MFC | 100 days after ASCT (intensive pathway only) | NS | Fisher's exact test | CTD vs CVAD (INT) or CTDa vs MP (nINT), MEL and ASCT | THAL vs no THAL | MRD ⁻ 62% (INT) and 15% (nINT) |
| Roussel 2014 | Prospective, multicenter, single-arm, open-label, phase II study | 39 | 31 | Response with RVD induction/ consolidation | 33–65 | MFC | Baseline, post- induction/pre- ASCT, post- ASCT, post- consolidation, end of | 48% stage I, 36% stage II, 16% stage III | Kaplan-Meier | RVD, MEL, ASCT | LEN for 1 year | 58% CR, 68% MDR⁻ |

| | | | | | | | treatment | | | | | |
|---------------------|--|------|----|---|---------------------|-----|--|--|---|---|--------------|---|
| Fukomoto 2016 | Retrospective | 40.9 | 78 | Impact of immunophenoty pic CR (MFC) on survival outcomes | 44–87 | MFC | Bone marrow samples taken at presentation, and at VGPR/CR | 53% stage III | Univariate analysis and multivariate analysis using a Cox proportional hazards model | 87% IMiD- based regimens and 94% BORT- based therapies | BORT+ DEX | 44% iCR |
| Sarasquet e 2005 | Prospective (GEMM2000) | NS | 32 | Compare ASO real-time qPCR vs MFC for MRD monitoring | 59 ± (SD) 9.7 | PCR | 3 months after transplant | NS | Mann- Whitney U and Kruskal– Wallis tests | VBCMP/VBAD, MEL, ASCT | None | 58% IF⁻ CR |
| Ludwig 2015 | Randomized, open-label, multicenter, phase II | 33 | 98 | Response rates after VTD vs VTDC induction | 33–68 | MFC | 40–269 days after SCT | 18–24% stage I, 45–47% stage II, 31–35% stage III | NS | VTD vs VTDC, single or double ASCT | None | MRD ⁻ : 53% (VTD) 33% (VTDC) |

Abbreviations: ASCT = autologous stem cell transplantation; BORT = bortezomib; C-VAMP = cyclophosphamide, vincristine, adriamycin plus methylprednisolone; CR = complete response; CT = chemotherapy; CTD = cyclophosphamide-thalidomide-dexamethasone; CTDa= attenuated CTD; CVAD = cyclophosphamide-vincristine-doxorubicin-dexamethasone; DEX = dexamethasone; HD = high-dose; IF = immunofixation; IFNα = interferon alfa; INT = intensive pathway; MEL = melphalan; MFC = multiparameter flow cytometry; mo = months; MP = melphalan-prednisolone; MRD = minimal residual disease; nINT = non-intensive pathway; NS = not specified; PFS = progression-free survival; OS = overall survival; PBSCT = peripheral blood stem cells transplant; PR = partial response; (q)ASO-PCR = (quantitative) allele-specific oligonucleotide polymerase chain reaction; SCT = stem cell transplantation; sFLC = serum free light chain; TBI = total body irradiation; THAL = thalidomide; Tx = transplantation; VBAD = vincristine-bis-chloroethylnitrosourea-doxorubicin-dexamethasone; VBMCP = vincristine-bis-chloroethylnitrosourea-melphalan-prednisolone; VP = bortezomib-prednisolone; VT = bortezomib-thalidomide; VTD = bortezomib-thalidomide-dexamethasone; VTP = bortezomib-thalidomide; VTP = bortezomib-thalidomide-dexamethasone; VTP = bortezomib-thalidomide-dexamethasone; VTP = bortezomib-thalidomide

Supplementary Figure. Overall survival in patients achieving CR according to cytogenetic risk category (FISH) and MRD status. CR, complete response; FISH, fluorescent in situ hybridization; MRD, minimal residual disease; OS, overall survival.

