SUPPLEMENTAL DATA

Supplement to: Tiong IS, et al, Venetoclax induces rapid elimination of *NPM1* mutant measurable residual disease in combination with low intensity chemotherapy in acute myeloid leukaemia.

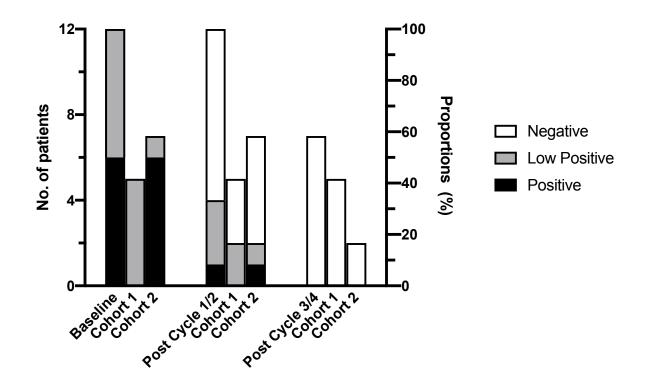
- 1. Supplemental Table SI. Venetoclax regimen, associated toxicities and dose adjustments.
- Supplemental Figure S1. Proportion of NPM1mut MRD response following venetoclax in combination with low-intensity chemotherapy. Five patients (all from cohort 2) did not proceed beyond cycle 2: one patient ongoing post cycle 1 and four patients underwent HSCT (one post cycle 1 and three post cycle 2). Overall molecular response rate was 92% by end of second cycle. Low positive: detectable at >4-log reduction from baseline; positive: detectable at <4-log reduction from baseline.
- 3. **Supplemental Figure S2.** Kaplan-Meier plots of (A) relapse-free and (B) overall survival for patients treated with venetoclax in combination with low-intensity chemotherapy for molecular persistence (cohort 1) and progression (cohort 2).
- 4. **Supplementary Figure S3.** Kaplan Meier plot of relapse-free survival for 6 patients with persistent *NPM1*^{mut}-MRD in the bone marrow post 4 courses of chemotherapy. Data reproduced from Ivey et al, *N Engl J Med* 2016.
- 5. **Supplementary Figure S4.** Kaplan Meier plot of relapse-free survival for 27 patients with molecular relapse who received salvage chemotherapy and subsequent allogeneic stem cell transplantation. Data reproduced from Dillon et al, *Blood* 2020.
- 6. **Case narratives** and serial NPM1mut MRD figures for cases #1 to #12.

Case #	Venetoclax	Concurrent azole	Partner therapy	No. of Cycles	Grade 4 cytopenias	Dose adjustment	Reason for cessation
1	400mg x 28d	No (C5-6)	Aza 75mg/m² x 7d	6	Neut	Aza x5d in cycle 6 Ven 100mg during voriconazole	Invasive fungal infection
	400mg x 28d	No	-	10	-	-	Facial rash
2	100mg x 14d	Yes	LDAC 20mg/m ² x 7d	3+	Both	-	Ongoing
3	100mg x 14d	Yes	LDAC 20mg/m ² x 10d	4	Neut	Ven x 10d from cycle 3	Planned
4	100mg x 14d	Yes	LDAC 20mg/m ² x 10d	4	Both	Ven x 10d from cycle 2	Planned
5	100mg x 14d	Yes	LDAC 20mg/m ² x 10d	10+	-	Ven x 10d + LDAC x 7d from cycle 7; Ven x 7d + LDAC x 5d from cycle 9	Ongoing
6	100mg x 14d	Yes	Aza 75mg/m² x 7d	4+	Both	Aza x 5d from cycle 3 (febrile neutropenia and severe pneumonia requiring admission to intensive care unit)	Ongoing
7	100mg x 28d	Yes	LDAC 20mg/m ² x 10d	2	Both	Ven x 17d in cycle 2	SCT
8	100mg x 14d	Yes	Aza 75mg/m² x 5d DLI	2+	-	-	Ongoing
9	100mg x 14d	Yes	LDAC 20mg/m ² x 10d	2	-	-	SCT
10	100mg x 14d	Yes	LDAC 20mg/m ² x 10d	2	Both	Ven x 28d in cycle 2 (increase)	SCT
11	100mg x 14d	Yes	LDAC 20mg/m ² x 10d	1	Neut	-	SCT
12	600mg x 28d	No	LDAC 20mg/m ² x 10d	7+	-	-	Ongoing

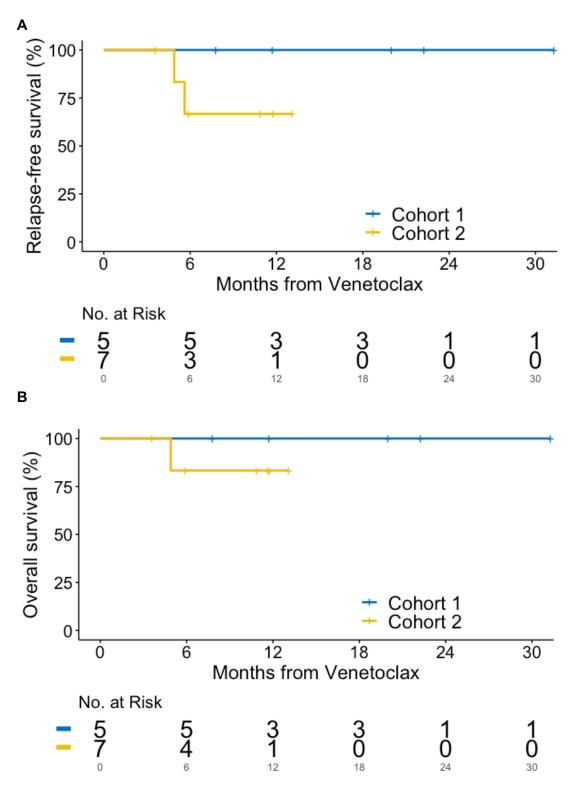
Supplemental Table SI. Venetoclax regimen, associated toxicities and dose adjustments.

Abbreviations: DLI: donor lymphocyte infusion; LDAC: low-dose cytarabine; Neut: neutropenia; SCT: stem cell transplantation.

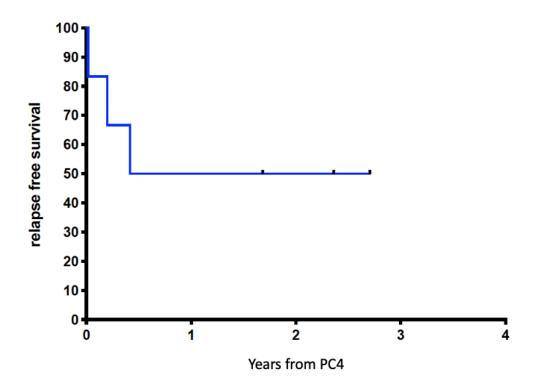
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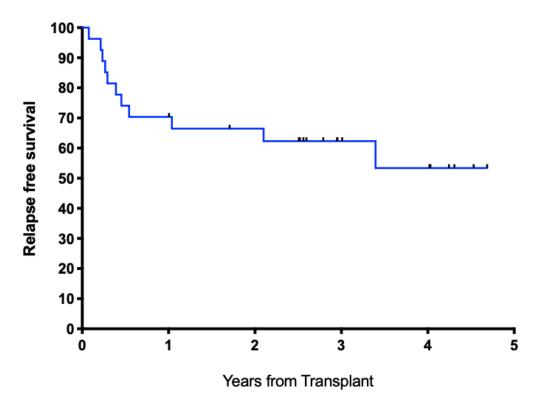
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Supplementary Figure S3. Kaplan Meier plot of relapse-free survival for 6 patients with persistent *NPM1*^{mut}-MRD in the bone marrow post 4 courses of chemotherapy. Data reproduced from Ivey et al, *N Engl J Med* 2016.



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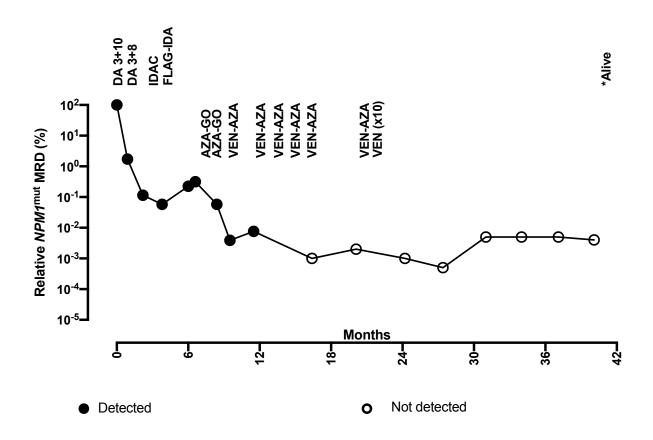
Case Narratives

Case #1

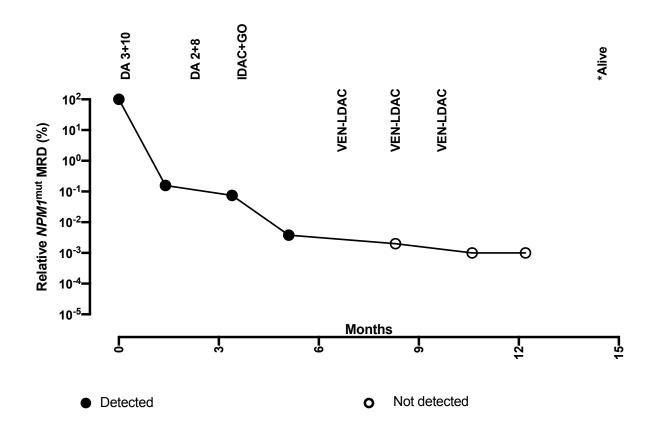
A 71-year-old male was diagnosed with AML with a normal karyotype and co-mutations in *DNMT3A*, *TET2* and *FLT3* (N676K). He was treated with DA3+10 then DA3+8. He achieved CR and a 2-log reduction in *NPM1* mutant transcripts after cycle 1. After cycle 2 there was a further 1 log reduction in *NPM1* mutant transcripts in the bone marrow and these were not detected in the peripheral blood. He received intermediate dose cytarabine ($1.5mg/m^2 \times 6$) for cycle 3 with no further reduction in transcript level and subsequently received FLAG-Ida with a 50% dose reduction for cycle 4 after which there was a 0.6 log increase in expression, confirmed on a second sample consistent with molecular persistence of disease.

He then received two cycles of azacitidine + gemtuzumab ozogamicin resulting in a >2-log reduction in transcript levels which remained positive. This treatment was poorly tolerated and was stopped. He then received venetoclax (400mg daily) + azacitidine (75mg/m² x 7d), achieving molecular complete remission after the second cycle. He received a further five cycles of venetoclax + azacitidine. The 5th cycle was complicated by presumed pulmonary invasive fungal infection requiring admission to hospital. The azacitidine cycle length was reduced to 5 days for the 6th cycle. He then received venetoclax monotherapy for a further 10 cycles following which all therapy was stopped due to the development of a pustular skin rash affecting the face which completely resolved within one month of treatment cessation.

At the time of last follow up 31 months after initiation of venetoclax + azacitidine the patient remains alive and well and in ongoing molecular complete remission.

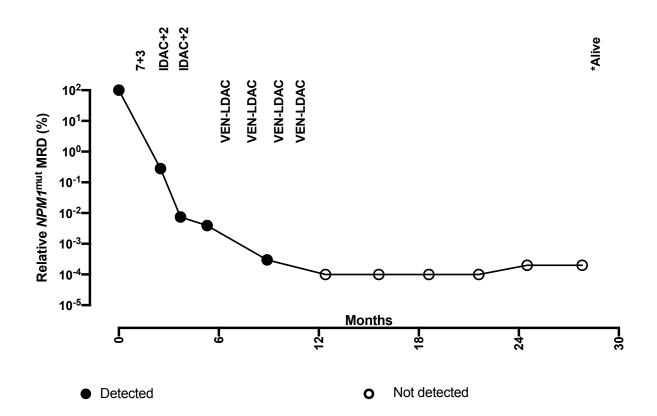


A 79-year-old female was diagnosed with normal karyotype *NPM1* mutated AML. No comutations were detected. She entered the NCRI AML18 clinical trial and received induction therapy with DA3+10 which was extremely poorly tolerated with life threatening bacterial and fungal infections and cardiomyopathy presumed secondary to anthracycline exposure. She achieved CR after cycle 1 and there was a 3-log reduction in *NPM1* mutant transcript levels. She received DA2+8 for the second cycle which was reasonably tolerated but there was no further reduction in transcript levels. Consequently, she received intermediate dose cytarabine (1 g/m² IV once daily x 5 days) with gemtuzumab ozogamicin for cycle 3 which was complicated by marked derangement in liver function tests. *NPM1* mutant transcripts remained detectable at the end of cycle 3 and treatment with venetoclax (100 mg once daily x 14 days with posaconazole) + cytarabine (20 mg/m²/d x 7 days by continuous infusion) was initiated. This was well tolerated without infection or hospitalisation. Molecular complete remission was achieved after the first cycle and at the time of last follow-up she had received two further cycles with a plan for ongoing therapy.

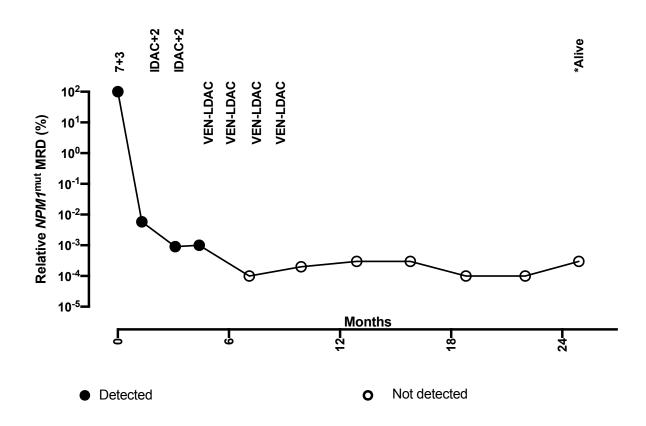


A 67-year-old female was diagnosed with pancytopenia and leucoerythroblastic blood picture. Bone marrow was hypercellular with 9.4% blasts, dysplastic changes with MF-2 reticulin fibrosis, consistent with an MDS/MPN syndrome. Molecular panel demonstrated *NPM1* and *DNMT3A* Met880Val mutations. Patient was treated as AML with 7+3 induction (2.6 log reduction) and 2 cycles of IDAC+2 consolidation (persistent low positive levels at end of therapy). Best response was CR with incomplete platelet recovery (CRp).

She then received 4 cycles of venetoclax (100 mg once daily x 14 days with posaconazole) with low-dose cytarabine, with successful eradication of *NPM1* mutant transcripts. In addition, patient made full hematologic recovery (CR). Treatment course was complicated by oral ulcers but otherwise well tolerated. Patient remained well at 22.2 months post venetoclax initiation.

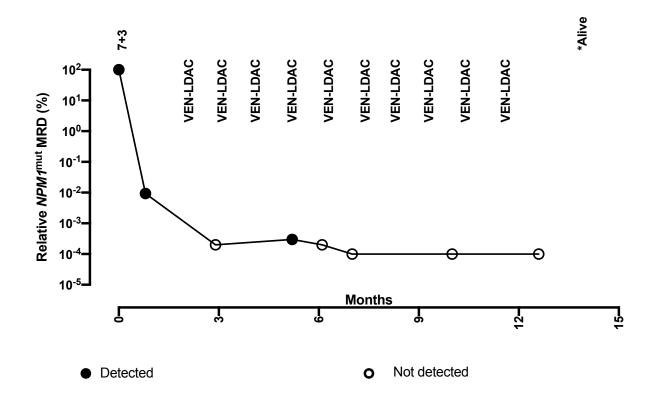


A 61-year old female was diagnosed with normal karyotype AML with mutations in *NPM1* and *IDH2*-R140. This was treated with 7+3 induction with a 4.2-log reduction. However, following further 2 cycles of IDAC+2 consolidation, *NPM1* mutant transcript persisted at low copy numbers. She then received 2 cycles of venetoclax (100 mg once daily x 14 days with posaconazole) with low-dose cytarabine. *NPM1* mutant MRD was eradicated following two cycles of therapy and patient went on to receive two further cycles. She remained well at 20 months post VEN initiation.



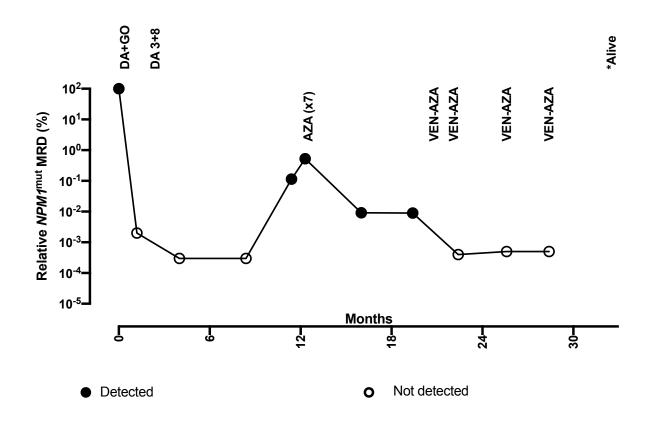
A 62-year-old male presented with WCC > 100×10^{9} /L and diagnosed with AML: trisomy 21, and mutated *NPM1*, *FLT3* Asp839Glu, *DNMT3A* Arg882Cys, *TET2* Arg1354fs and *TET2* Arg1440fs. Induction (7+3 with midostaurin) was complicated by tumour lysis syndrome, oliguric acute kidney injury requiring renal replacement therapy, methicillin-sensitive Staphylococcal aureus sepsis, and intensive care admission. Complete remission was achieved post induction with a 4-log reduction in *NPM1*^{mut} MRD level.

Patient was deemed unfit for further consolidation chemotherapy, but instead was commenced on venetoclax (100 mg once daily x 14 days with posaconazole) with low-dose cytarabine, at 65 days post induction. *NPM1*^{mut} MRD was rendered negative following first cycle of therapy. Post cycle 3 patient was hospitalised for viral upper respiratory tract infection and deranged liver biochemistry. A low level *NPM1*^{mut} MRD was also noted. Posaconazole was ceased from cycle 4-6 and venetoclax increased to 400 mg once daily. Cycle 7 onwards, itraconazole was initiated and venetoclax dose reduced back to 100 mg once daily. Subsequent MRD levels were undetectable and patient remained well with ongoing therapy (cycle 10+).

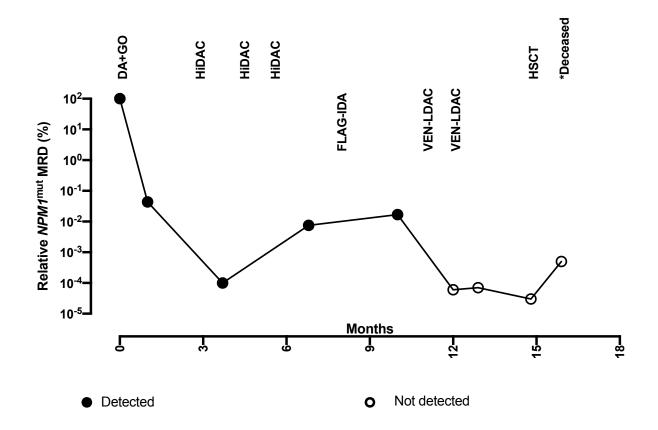


An 80-year-old male was diagnosed with AML with normal karyotype and co-mutations in *IDH1* (R132C) *FLT3* (D835E) and *SRSF2* (P95R). He entered the NCRI AML18 clinical trial and received induction therapy with DA3+10 with two doses of gemtuzumab ozogamicin. He achieved a morphological and molecular complete remission after cycle 1 and received DA3+8 for cycle 2 which was complicated by severe mucositis leading to the decision to withhold further treatment.

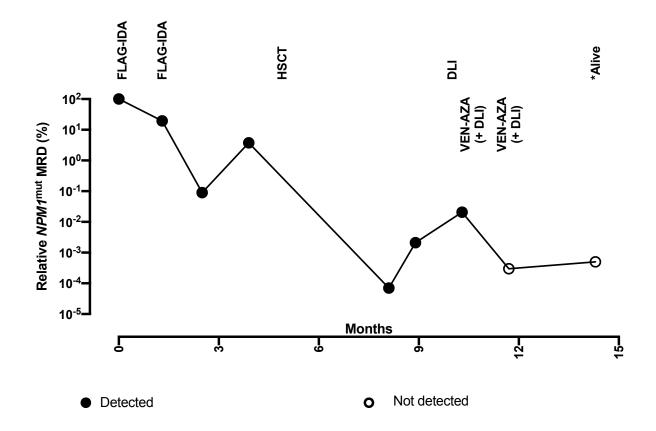
One year after his initial diagnosis, two consecutive bone marrow samples tested positive for *NPM1* mutant transcripts with a rising level and he was diagnosed with molecular relapse. He was treated with azacitidine monotherapy which resulted in an initial reduction in transcript levels which subsequently reached a plateau. He then commenced treatment with venetoclax (100mg od with posaconazole x 14d) + azacitidine (75mg/m² x 7d), achieving a molecular complete remission after the first cycle. The second cycle was complicated by neutropenic sepsis and severe pneumonia requiring intensive care admission for non-invasive ventilation. The length of azacitidine was reduced to 5 days for subsequent cycles and these were well tolerated. At last follow-up 11.8 months after initiation of venetoclax treatment, the patient remains alive and well in molecular complete remission with a plan for ongoing treatment.



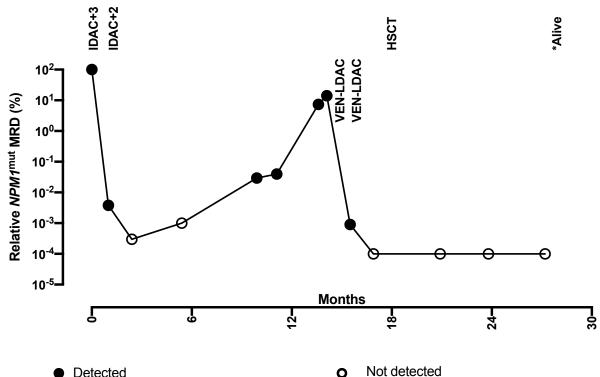
A 40-year-old female was diagnosed with AML with a normal karyotype and mutations in NPM1 and FLT3 (D835Y). She entered the NCRI AML19 clinical trial and received induction therapy with DA3+10 and one dose of gemtuzumab ozogamicin. This was complicated by eosinophilic myocarditis thought to be caused by the gemtuzumab ozogamicin. She achieved CR after the first cycle with a 3.4 log reduction in NPM1 mutant transcript levels and received consolidation with 3 cycles of high dose cytarabine (3 $g/m^2 \times 6$). At the end of treatment, MRD assessment showed a 1.9 log increase in transcript levels consistent with molecular progression. She received FLAG-IDA as salvage therapy; this was poorly tolerated with lifethreatening infection and on regeneration there had been no reduction in transcript levels. She then received venetoclax (100 mg od x 28d + posaconazole) + cytarabine (20 mg/m²/d x 10d) and achieved complete molecular remission, grade 4 neutropenia and thrombocytopenia but no infection. Second cycle of therapy was continued without waiting for neutrophil and platelet recovery to grade <4, with venetoclax ceased at day 18. She went on to receive HSCT and MRD assessment pre-transplant and at D+30 showed ongoing molecular complete remission. Unfortunately, she was readmitted with sepsis after transplant having fully engrafted and subsequently died.



A 28-year-old man presented with relapsed AML 7 years after his original diagnosis, having originally presented in 2012 with AML with fibrosis and an *NPM1* mutation. At relapse, in addition to persistent *NPM1*, he was also found to harbour a *FLT3* TKD (D835) mutation. He received two cycles of FLAG-IDA salvage therapy resulting in CR and a 3-log reduction in transcript levels. Post allogeneic HSCT, the D+100 bone marrow MRD assessment was positive. Despite withdrawal of immunosuppression and donor lymphocyte infusion (DLI; 10^6 cells), *NPM1^{mut}* levels increased in two consecutive samples consistent with molecular progression. Venetoclax (100 mg od x 14d) + azacitidine (75mg/m² for 5 days) was given resulting in molecular complete remission after one cycle of therapy in conjunction with DLI (10^6 cells). Venetoclax and azacitidine was well tolerated without grade 4 cytopenias, infection or hospital admission. Second cycle was delivered with DLI 4.7 x 10^6 cells. The patient remains alive and well with a plan for ongoing therapy.

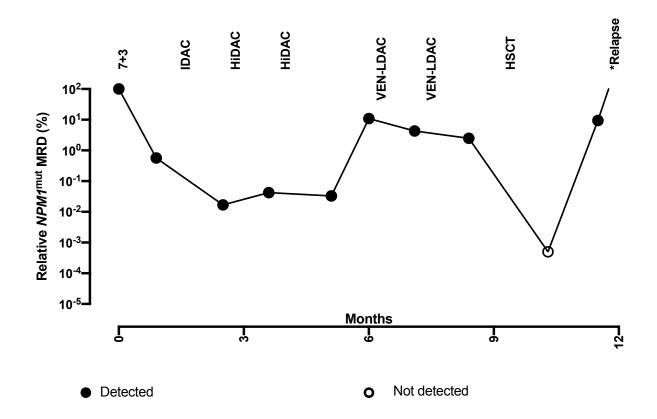


A 52-year-old male was diagnosed with AML with normal karyotype and mutated NPM1 (no other co-mutations). He achieved excellent response following IDAC+3 induction (cytarabine 1.5 g/m² x 8 and idarubicin 12 mg/m² x3) with a 4.4-log reduction in NPM1^{mut} MRD, and molecular complete remission after an IDAC+2 consolidation (cytarabine 1 g/m² x6 and idarubicin 12 mg/m² x2). During surveillance, NPM1^{mut} transcript levels were noted to be rising after 9 months post CR and peaked at 14% (~13 months post CR). Venetoclax (100 mg x 14d with posaconazole) and low-dose cytarabine were initiated with 4.2 log reduction post 1st cycle and molecular complete remission post 2nd cycle. Patient underwent allogeneic HSCT and remained alive at 9.9 months post HSCT.



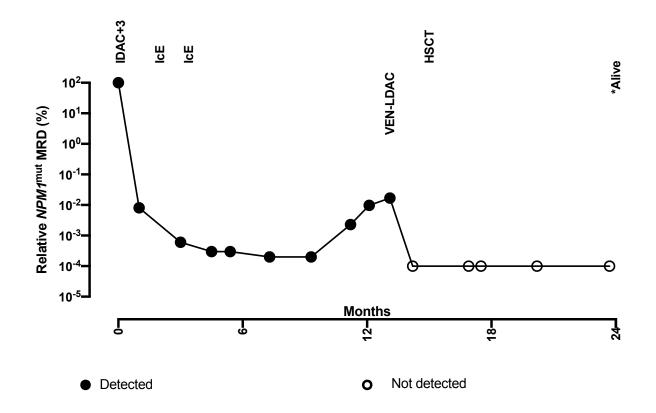
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A 35-year-old female was diagnosed with AML with normal karyotype, mutated *NPM1*, *IDH2* R140 and *DNMT3A* R882. Post induction (7+3) response was suboptimal (2.2 log reduction), followed by three cycles of HiDAC consolidation ($3 \text{ g/m}^2 \times 6$) with persistent low copy numbers at approximately 3.5 log from baseline levels. Two months following completion of chemotherapy, molecular progression was evident. Two cycles of venetoclax (100 mg x 14d during cycle 1 and x 28d during cycle 2) with low-dose cytarabine were attempted with minimal response (0.6 log reduction). Patient went to have a myeloablative allogeneic HSCT but relapsed 78 days post HSCT with a newly acquired *FLT3*-ITD at allelic ratio 0.09. Baseline mutations (*NPM1*, *IDH2* and *DNMT3A*) were all persistent at the time of relapse. She was subsequently treated with gilteritinib.



A 51-year-old male was diagnosed with AML with normal karyotype, *FLT3*-ITD (allelic ratio 0.32) and mutated *NPM1*. He entered ALLG AMLM16 trial and underwent induction with IDAC+3 (cytarabine 1.5 g/m² x8 and idarubicin 12 mg/m² x3) and sorafenib/placebo. Post induction assessment revealed CR with a 4.1-log reduction in *NPM1*^{mut} MRD. This was followed by two cycles of ICE consolidation (idarubicin 9 mg/m² x2, cytarabine 100 mg/m² x5 and etoposide 75 mg/m² x5), with persistent low copy numbers of *NPM1*^{mut} transcripts.

NPM1^{mut} MRD was initially stable during surveillance but molecular progression occurred at six months post completion of chemotherapy. After one cycle of venetoclax (100 mg x 14d with posaconazole) and low-dose cytarabine, *NPM1*^{mut} MRD was rendered undetectable and patient underwent reduced intensity conditioning HSCT. He remained alive and well at 8.9 months post HSCT.



A 69-year-old male was diagnosed with AML with normal karyotype, mutated *NPM1* and *DNMT3A* R882. Post induction (7+3) response was suboptimal (2.8 log reduction), followed by two cycles of IDAC+2 consolidation (cytarabine 1 g/m² x 6 and idarubicin 12 mg/m² x2) with persistent low copy numbers at end of therapy.

NPM1^{mut} transcript levels were initially stable but molecular progression ensued eight months post completion of chemotherapy, followed shortly by early morphological relapse (5% bone marrow blasts) 3.1 months later. Patient entered a clinical trial receiving venetoclax in combination with a MCL1 inhibitor and achieved a 2.1-log MRD reduction but was withdrawn after one cycle due to toxicity. He subsequently received venetoclax (600 mg once daily x28d) and low-dose cytarabine, with a further 2.6-log reduction after one cycle and achieved molecular complete remission after 3 cycles. Patient remained on therapy (7+ cycles) and well at last follow up.

