



2019年终盘点

end of 2019



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1 急性髓系白血病的诊断分层

AML的预后分层

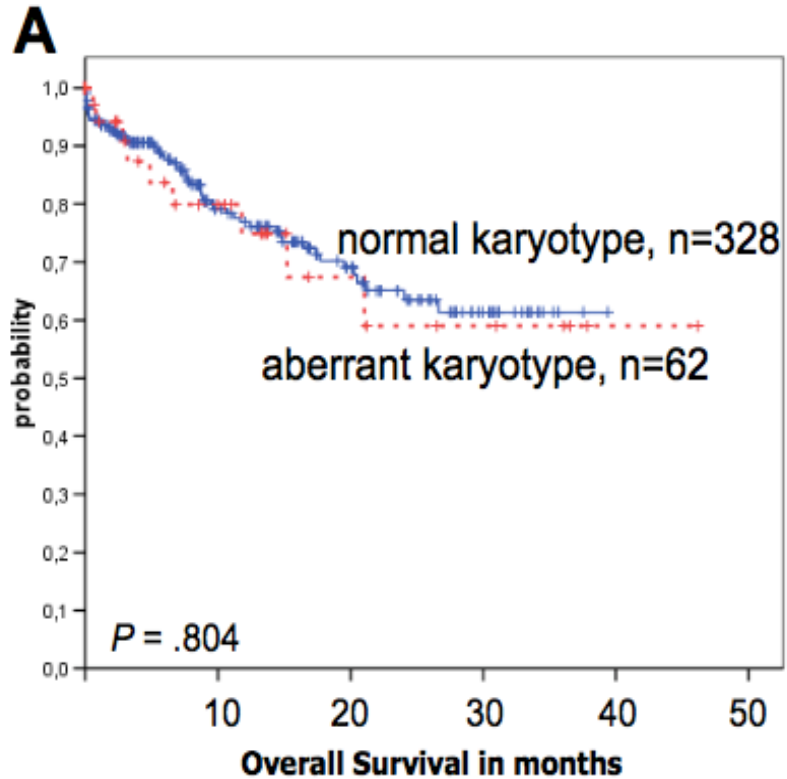
NCCN指南

| 预后分层 | 细胞遗传学 | 分子生物学 |
|------|---|--|
| 预后良好 | Inv(16)(p13q22)或 t(16;16)(p13;q22)或 t(8;21)(q22;q22) | 正常核型： NPM1 突变但不伴有FLT3-ITD 突变，或CEBPA双突变 |
| 预后中等 | 正常核型 +8 t(9;11)(p22;q23) 其他异常 | Inv(16)(p13q22)或 t(16;16)(p13;q22) 伴有c-Kit突变； t(8;21)(q22;q22)伴有c-Kit突变 |
| 预后不良 | 复杂核型(≥3 种) 单体核型 -5, -7, 5q-, 7q- 11q23 染色体易位，除 外t(9;11) inv(3)或t(3;3) t(6;9) t(9;22) | 正常核型： 伴有FLT3-ITD突变 TP53突变 |

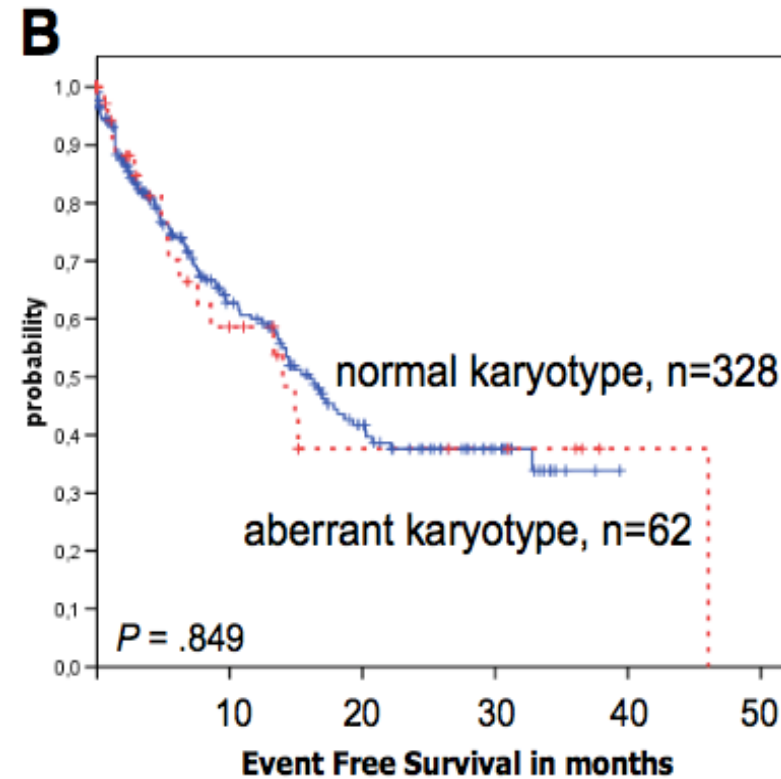
| 预后分层 | 分子遗传学 |
|------|--|
| 预后良好 | t(8;21)(q22;q22.1); RUNX1-RUNX1T1 inv(16)(p13.1q22) 或 t(16;16)(p13.1;q22); CBFB- MYH11 NPM1突变不伴或低水平FLT3-ITD CEBPA双突变 |
| 预后中等 | NPM1 突变伴高水平FLT3-ITD 野生型NPM1 不伴或低水平 FLT3-ITD t(9;11)(p21.3;q23.3); MLLT3-KMT2A 无良好或不良的细胞遗传学异常 |
| 预后不良 | t(6;9)(p23;q34.1); DEK-NUP214 t(v;11q23.3); KMT2A 重排 t(9;22)(q34.1;q11.2); BCR-ABL1 inv(3)(q21.3q26.2) 或 t(3;3)(q21.3;q26.2); GATA2,MECOM(EVI1) -5 或del(5q); -7; -17/abn(17p) 复杂核型或单倍体核型 野生型NPM1 伴高水平FLT3-ITD RUNX1突变 ASXL1突变 TP53突变 |

ELN指南

AML预后分层: NPM1突变与核型



rather than the previous 4-group system (Table 5). A few other changes have been made. Recent studies have shown that in AML with *NPM1* or biallelic *CEBPA* mutations, the presence of coexisting chromosomal abnormalities does not appear to modify the prognostic effect of the mutations^{16,25,115}; prognosis may be more influenced by concurrent gene mutations.³⁷ Accordingly, and as in CBF-AML, the categorization of these cases is now based on the primary leukemia-defining genetic subsets irrespective of the karyotype. The higher relapse rate and poorer OS associated with *FLT3*-ITD largely depends on the ITD



Blood. 2009;114: 3024-3032
Blood. 2017;129(4):424-447

研究背景: 伴 NPM1^{mut} 和 FLT3-ITD^{neg/low} 的AML患者, 是预后较好的分子学异常, 且被认为不管细胞遗传学分型如何。此研究验证合并细胞遗传学异常的 NPM1^{mut} 和 FLT3-ITD^{neg/low} 的AML患者预后情况

研究人群:

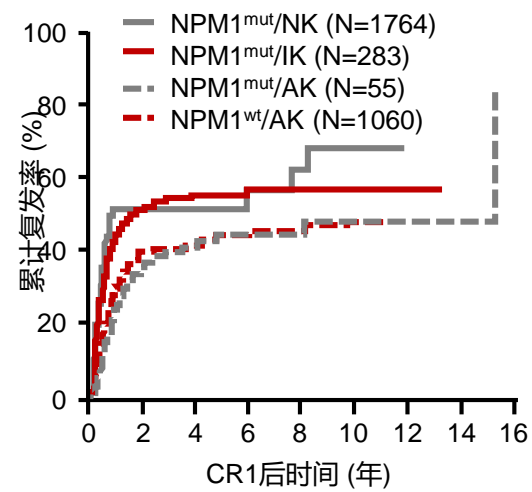
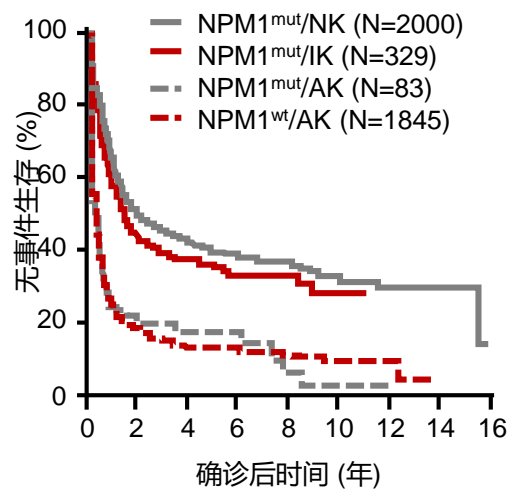
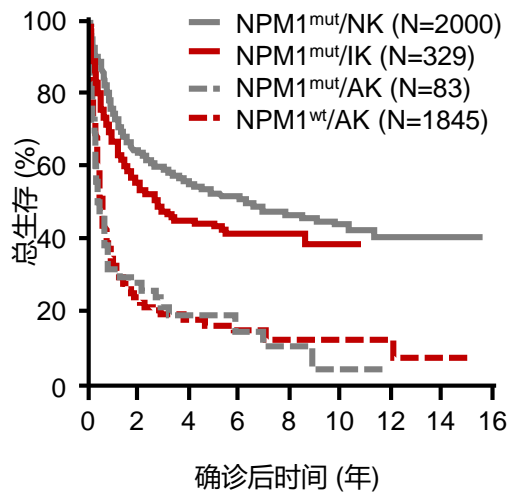
NPM1^{mut}/FLT3-ITD^{neg/low}和染色体核型

9个国际队列

- ≥18岁高强度治疗的AML患者
- 核型根据2017 ELN 进行分类
- **2426例NPM1^{mut}/FLT3-ITD^{neg/low}患者**
 - 2000例正常核型
 - 329例中等异常核型
 - 83例不良核型

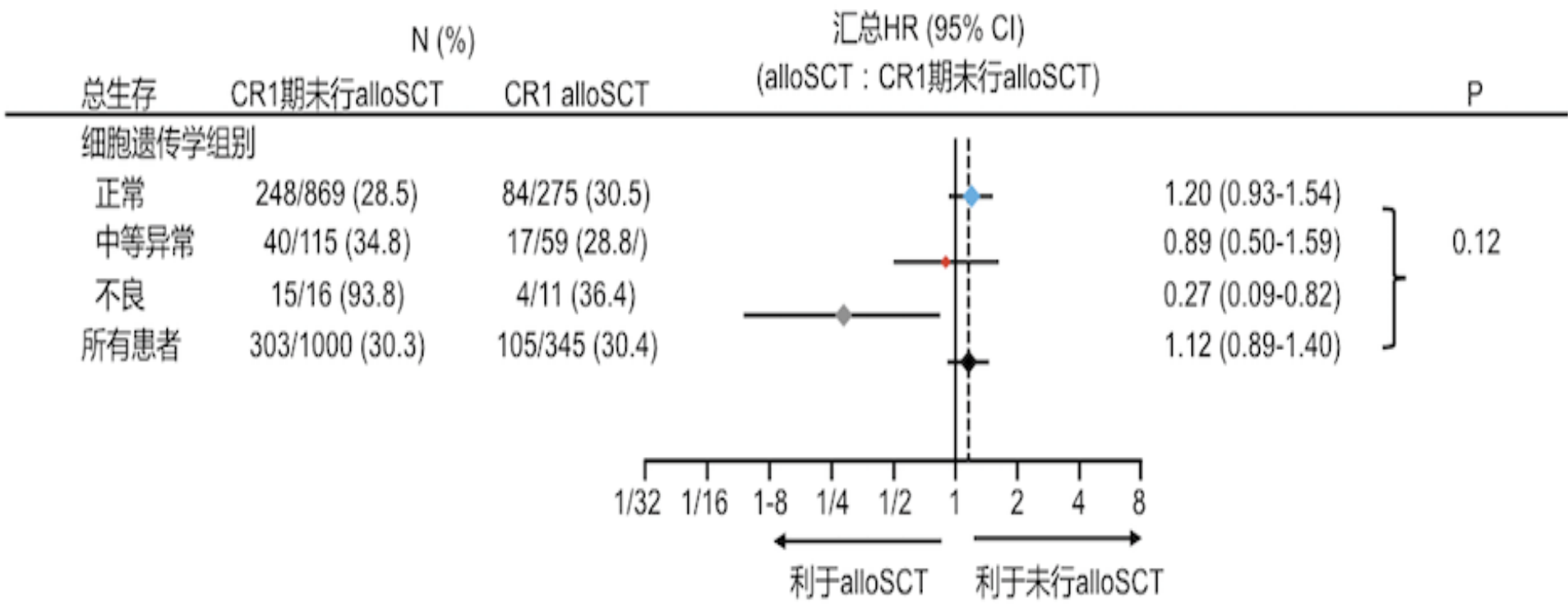
1845例NPM1^{wt}/FLT3-ITD^{neg/low}不良核型患者

研究结果：合并不良细胞遗传学分型的患者CR率更低、OS更差、EFS短，CIR更高



| 多变量 | CR | OS | EFS | CIR |
|----------------|------------------------------|------------------------------|------------------------------|------------------------------|
| 不良核型 vs 正常核型 | 0.29 (0.17-0.48) P<0.0001 | 2.97 (2.29-3.87) P<0.0001 | 2.63 (2.05-3.38) P<0.0001 | 2.52 (1.75-3.62) P<0.0001 |
| 中等异常核型 vs 正常核型 | 0.84 (0.59-1.19) P=0.33 | 1.27 (1.07-1.50) P=0.0060 | 1.21 (1.04-1.41) P=0.014 | 1.18 (0.97-1.44) P=0.10 |

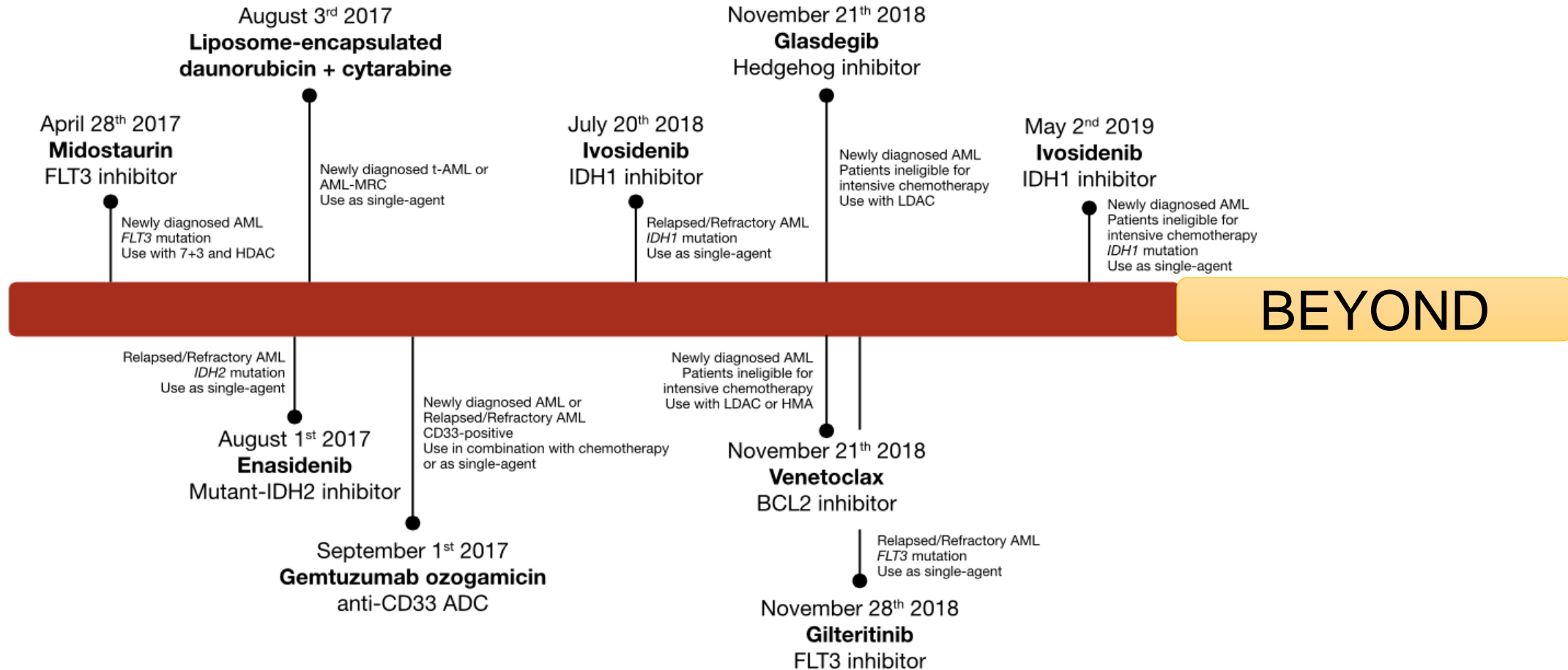
研究结果：细胞遗传学良好或一般患者，CR1后移植与否不影响OS，而细胞遗传学不良患者CR1后行移植明显改善OS



2

急性髓系白血病的诱导、巩固及维持治疗

AML的治疗靶向进展



现有的FLT3抑制剂

| Name | Initial name | Type I or II | Active as monotherapy | Selectivity | Half-life |
|--------------|--------------|--------------|-----------------------|-------------|-----------|
| Midostaurin | PKC412 | I | No | + | 19h |
| Sorafenib | BAY-43-9006 | II | Yes | ++ | 25-48h |
| Quizartinib | AC220 | II | Yes | +++ | ~1.5d |
| Crenolanib | CP-868–596 | I | Yes | ++ | 6-8h |
| Gilteritinib | ASP2215 | I | Yes | ++ | 113h |

FLT3抑制剂相关临床试验

FLT3 Inhibitor Combination Clinical Trials

Newly Diagnosed- Fit

QUIWI
Quizartinib + (7+3)
(FLT3-WT)

AML-18
Multiple
(FLT3-mutated)

QuANTUM First
Quizartinib + (7+3)
(FLT3-ITD)

Quizartinib + CPX-351
(FLT3-ITD/WT/HR MDS)

Quizartinib + CLIA
(FLT3-ITD)

Gilteritinib vs mido + 7+3
(HOVON/SAKK-156)

Gilteritinib vs mido + 7+3
(PrECOG)

Newly Diagnosed- Unfit

Quizartinib + AZA/LoDAC
(FLT3-ITD/WT; MDS, CMML)

LI-1
Multiple
(FLT3-ITD/WT, MDS)

QUIZOM
Quizartinib + Omacetaxine (FLT3-ITD)

Gilteritinib vs Gilteritinib + AZA
vs AZA

MDM2 + Quizartinib
(FLT3-ITD)

LGH447 (Pim kinase inhibitor) + Midostaurin
(FLT3-ITD/WT)

Relapsed/Refractory - Unfit

Quizartinib + Venetoclax
(FLT3-ITD)

Gilteritinib + Venetoclax

Gilteritinib + Atezolizumab

Relapsed/Refractory - Fit

Q-HAM
Quizartinib with HiDAC and
mitoxantrone
(FLT3-ITD)

FLAG – QUIDA
Quizartinib + FLAG-IDA
(FLT3-ITD/WT)




Crenolanib + FLAG-IDA/HAM

Quizartinib + CPX-351
(FLT3-ITD/WT/HR MDS)

Quizartinib + CLIA
(FLT3-ITD/WT)

Pediatric
Quizartinib + Re-induction
(FLT3-ITD)

LEGEND

-  High intensity
-  Low intensity
-  Targeted Therapy

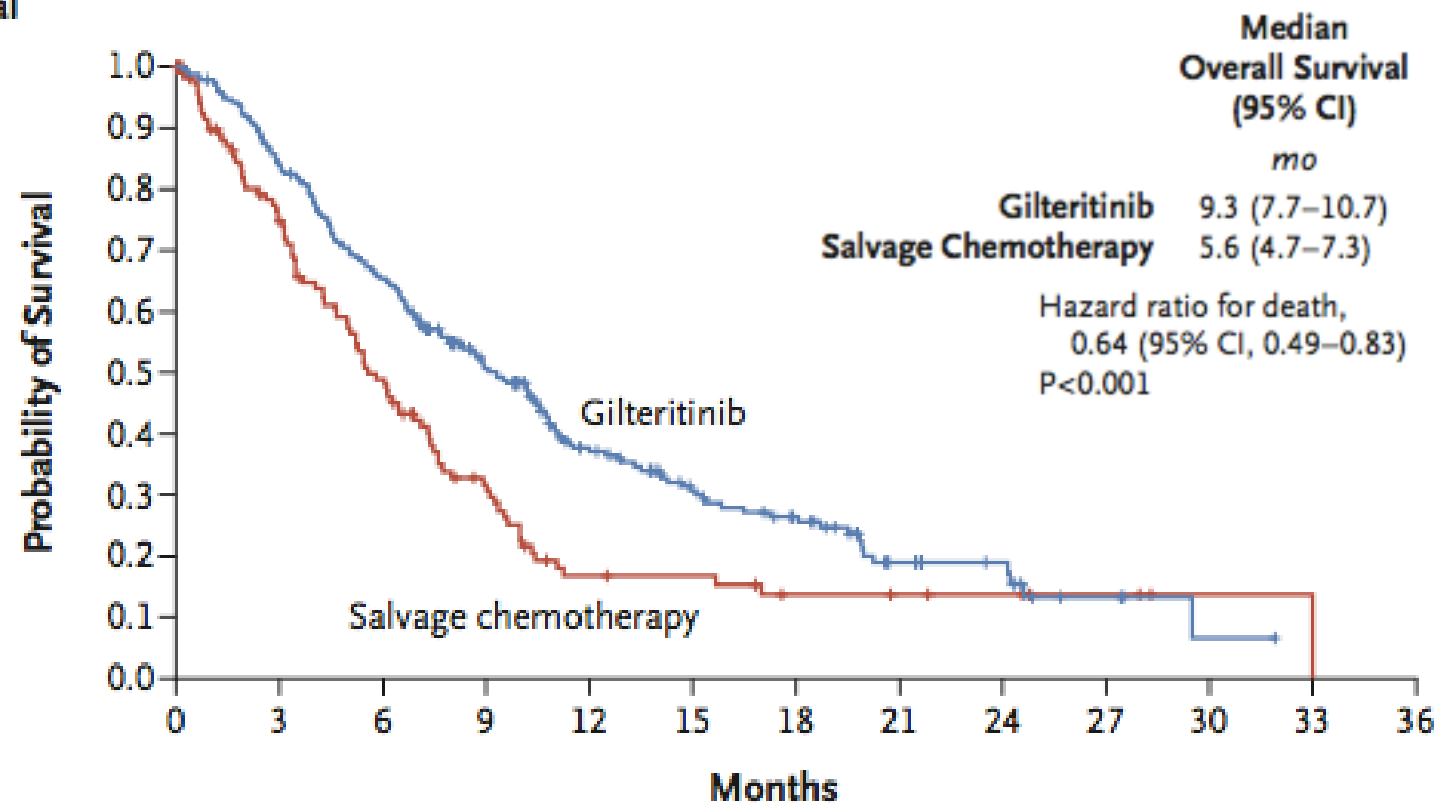
FLT3抑制剂Quizartinib在R/R AML中的疗效

Effect of Co-Mutations and FLT3-ITD Variant Allele Frequency (VAF)
on Response to Quizartinib or Salvage Chemotherapy (SC) in
Relapsed/Refractory (R/R) Acute Myeloid Leukemia (AML)

- Patients with **NPM1wt/DNMT3Amut** had significantly longer median OS with quizartinib vs SC (39.3 vs 19.6 weeks, respectively; HR, 0.239; P = 0.003) while NPM1mut/DNMT3Amut patients had lower and similar median OS between the 2 arms (23.6 vs 23.4 weeks, respectively).
- Quizartinib treatment showed significantly longer median OS vs SC in patients with **high FLT3-ITD** VAF (23.9 vs 17 weeks respectively; HR, 0.689, P = 0.0148), while the median OS in patients with low FLT3-ITD VAF was similar (34.1 vs 26.6 weeks, respectively; HR, 0.857, P = 0.535)

FLT3抑制剂Gilteritinib治疗R/R AML的生存情况

A Overall Survival



No. at Risk

| | | | | | | | | | | | | | |
|----------------------|-----|-----|-----|-----|----|----|----|----|----|---|---|---|---|
| Gilteritinib | 247 | 206 | 157 | 106 | 64 | 44 | 31 | 14 | 11 | 4 | 1 | 0 | 0 |
| Salvage chemotherapy | 124 | 84 | 52 | 29 | 13 | 12 | 8 | 7 | 5 | 3 | 1 | 0 | 0 |

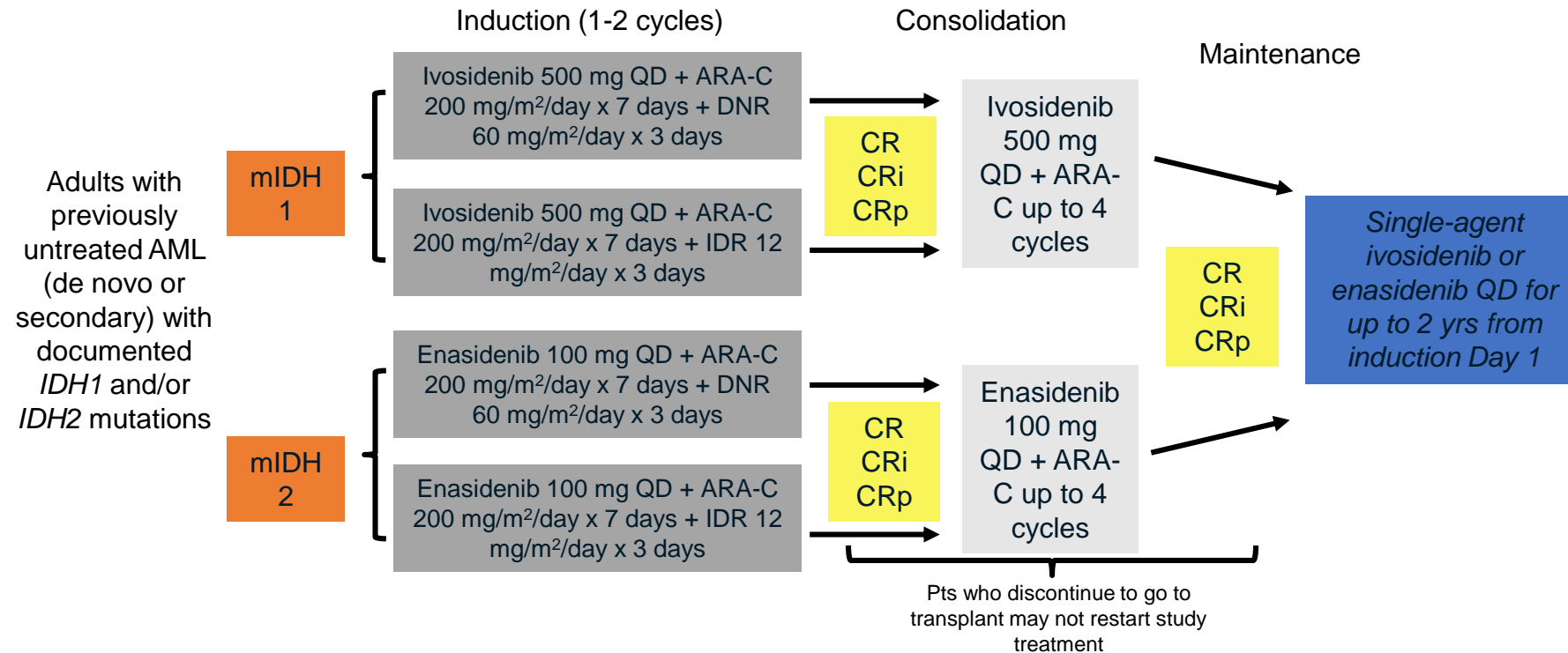
IDH1/2抑制剂在R/R-AML中的疗效

| | Enasidenib | Ivosidenib |
|---------------------------------|------------|---------------------------------|
| CR | 19.3% | 21.6% |
| ORR | 40.3% | 41.6% |
| The median time to a response | 1.9 months | 1.9 months (range, 0.8 to 4.7) |
| The median duration of response | 5.8 months | 6.5 months (95% CI, 4.6 to 9.3) |

N Engl J Med 2018;378:2386-98.
Blood. 2017;130:722-731

IDH1/2抑制剂单药在新诊断AML中的疗效

Open-label, phase I dose-escalation and -expansion trial



Primary objectives: safety and tolerability of ivosidenib and enasidenib

阿扎胞苷联合IVO治疗携带 IDH1 突变的unfit新诊断 AML 患者，安全性良好，ORR达78%

AG221-AML-005 (NCT02677922) 是一项正在进行中的开放标签、多中心 Ib / II 期研究，主要评估 IVO 或 AG221 (enasidenib) 联合 AZA 治疗携带 IDH1 或 IDH2 突变的不适合强化治疗的新诊断AML的安全性和有效性。

此次报告的数据为 IVO + AZA 的数据：中位年龄76岁，继发性 AML (s AML) 患者占 26%。细胞遗传学预后中等和预后不良的比例分别为 65% 和 22%，具体用药为 IVO 500mg qd 口服 + **AZA 75mg/m² 皮下 d1-7, 28 天一个周期**，直至疾病进展。

| IVO+AZA (n=23) | | All-grade AEs regardless of cause in≥30% its,% | | IVO +AZA (n=23) |
|-----------------|---------------|--|--|-----------------|
| CR+CRh,%(n) | 70(16) | 血小板减少 | | 65 |
| ORR,%(n) | 78(18) | 恶心 | | 61 |
| CR,% | 57 | 腹泻 | | 57 |
| CRi/CRp,% | 13 | 贫血 | | 52 |
| MLFS,% | 9 | 便秘 | | 52 |
| 中位至缓解时间, 月 (范围) | 1.8 (0.7-3.8) | 发热性中粒细胞减少 | | 39 |
| 中位至CR时间, 月 (范围) | 3.5 (0.8-6.0) | 发热 | | 39 |
| 中位缓解持续时间, 月 | 未达到 | 呕吐 | | 35 |
| | | 疲乏 | | 35 |
| | | 低钾血症 | | 35 |
| | | 头晕 | | 35 |
| | | 失眠 | | 35 |
| | | 中性粒细胞减少 | | 30 |

IDH2抑制剂+阿扎胞苷在新诊断AML中的疗效

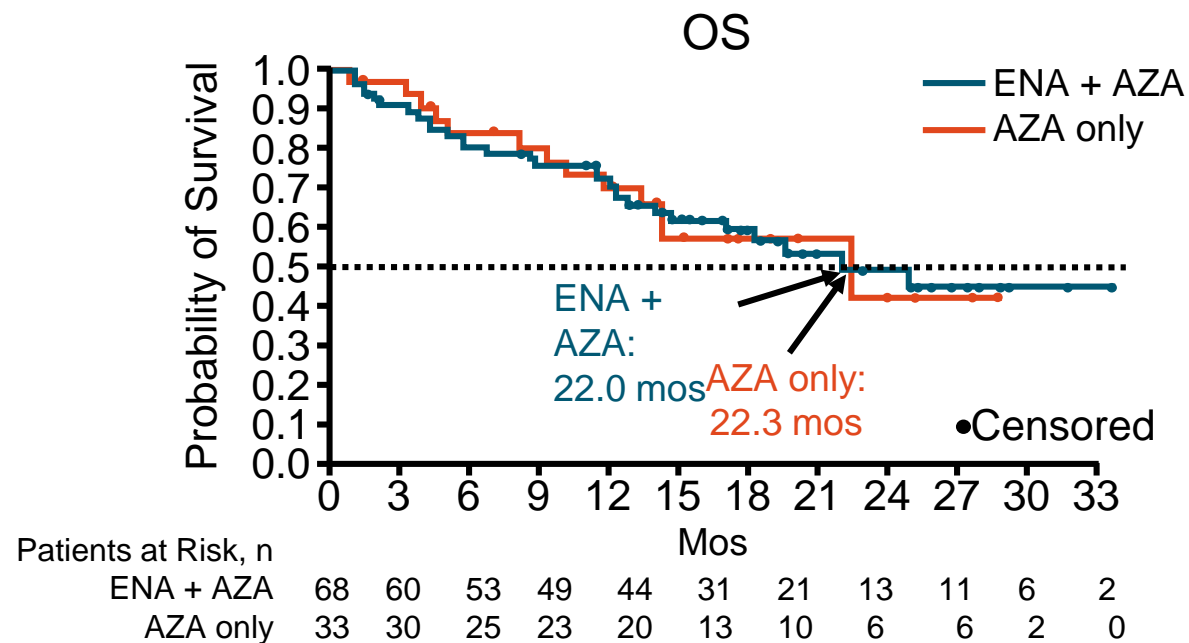
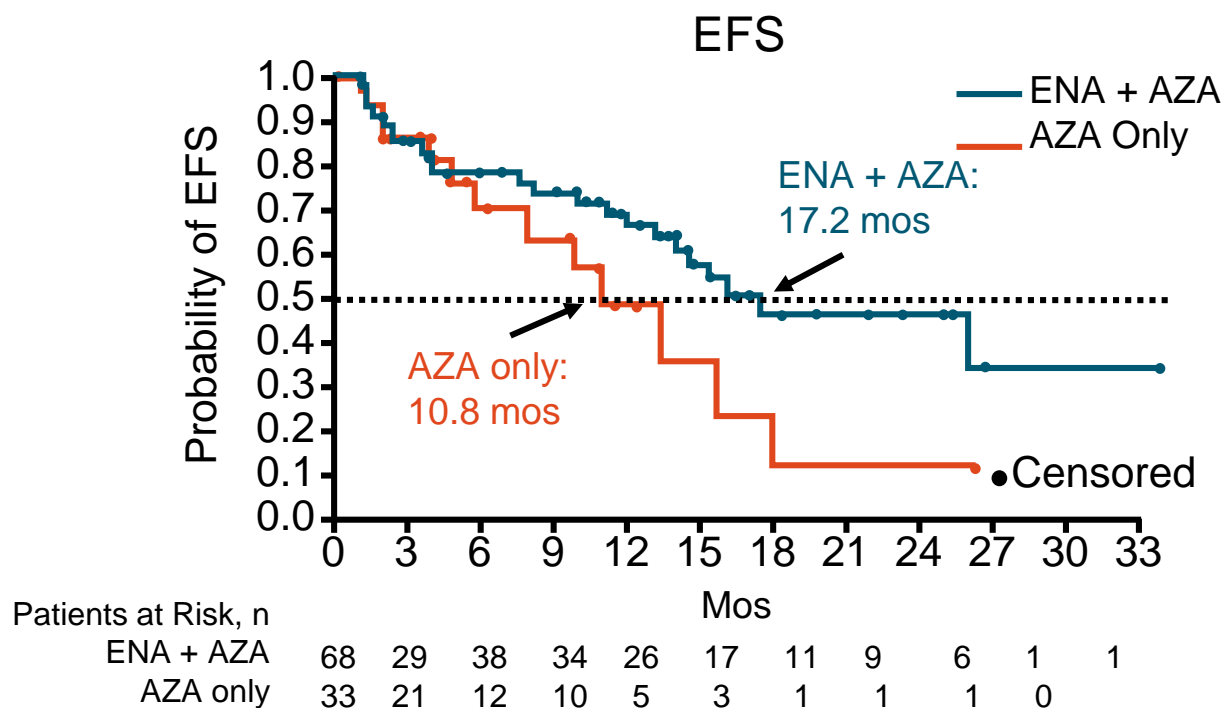
Enasidenib Plus Azacitidine Significantly Improves Complete Remission and Overall Response Compared with Azacitidine Alone in Patients with Newly Diagnosed Acute Myeloid Leukemia (AML) with Isocitrate Dehydrogenase 2 (IDH2) Mutations: Interim Phase II Results from an Ongoing, Randomized Study

ASH2019 Abs643

Table. Clinical efficacy with enasidenib plus azacitidine versus azacitidine monotherapy

| | Enasidenib + Azacitidine (N=68) | Azacitidine Monotherapy (N=33) |
|--|---------------------------------------|--------------------------------------|
| Overall response rate,* n (%) | 46 (68) | 14 (42) |
| [95%CI] | [55, 79] | [26, 61] |
| <i>P</i> value | 0.0155 | |
| Best response, n (%) | | |
| Complete remission (CR) | 34 (50) | 4 (12) |
| [95%CI] | [38, 62] | [3, 28] |
| <i>P</i> value | 0.0002 | |
| CR with incomplete recovery (CRi/CRp) | 6 (9) | 4 (12) |
| Partial remission | 3 (4) | 4 (12) |
| Morphologic leukemia-free state | 3 (4) | 2 (6) |
| Stable disease, [†] n (%) | 15 (22) | 13 (39) |
| Progressive disease, n (%) | 2 (3) | 1 (3) |
| Not evaluable, n (%) | 1 (2) | 0 |
| Missing, n (%) | 4 (6) | 5 (15) |
| Time to first response (months), median (range) | 1.9 (1–9) | 2.0 (1–6) |
| Duration of response (months), median [95%CI] | NR [11, NR] | 10.2 [3, NR] |
| Time to CR (months), median (range) | 5.0 (1–20) | 3.7 (3–4) |
| *Overall response comprises complete remission (CR), CR with incomplete hematologic recovery, CR with incomplete platelet recovery, partial remission, or morphologic leukemia-free state, per IWG 2003 AML response criteria. | | |
| [†] Absence of hematologic response and not meeting criteria for disease progression, sustained for a period of ≥8 weeks. | | |
| <i>P</i> values are from Chi-square test. | | |
| 95%CI, 95% confidence interval; AML, acute myeloid leukemia; CR, complete remission; CRi, CR with incomplete blood count recovery; CRp, CR with incomplete platelet count recovery; IWG, International Working Group; NR, not reached. | | |

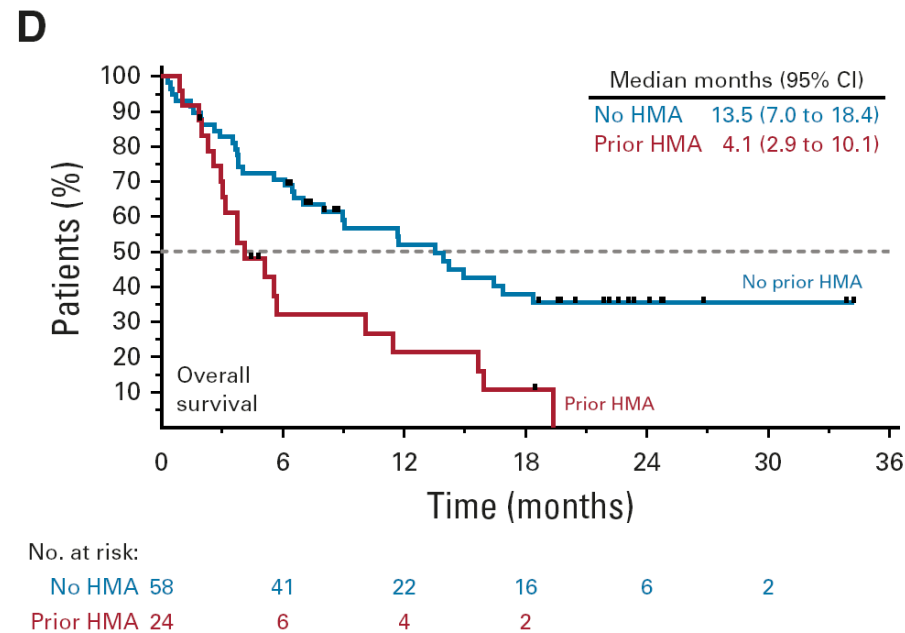
IDH2抑制剂+阿扎胞苷治疗新诊断AML的生存情况



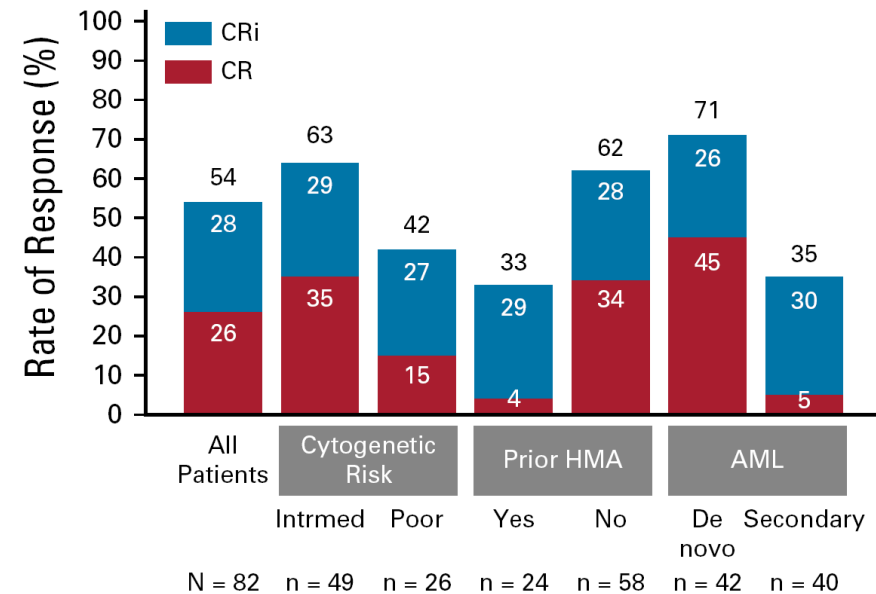
| Endpoint | Enasidenib + Azacitidine (n = 68) | Azacitidine Monotherapy (n = 33) | HR (95% CI) | P Value |
|-----------------|-----------------------------------|----------------------------------|------------------|---------|
| Median OS, mos | 22.0 | 22.3 | 0.99 (0.52-1.87) | .9686 |
| Median EFS, mos | 17.2 | 10.8 | 0.59 (0.30-1.17) | .1278 |

BCL2抑制剂+低甲基化药物在新诊断老年AML中的疗效

Durable Responses



High Response Rate



Venetoclax + LDAC治疗老年新诊断AML

J Clin Oncol. 2019 Mar 20

BCL2抑制剂+ 低甲基化药物在新诊断老年AML中的疗效

Response to venetoclax in combination with LDAC or HMA in untreated patients with AML patients with IDH, FLT3 and other mutations and correlations with BCL2 family expression

| | Responses to Venetoclax | | | |
|--------------------------------|--|-------------------------------|-------------------------------------|----------------------------|
| | Complete Remission/Incomplete Complete Remission Rate, n (%) | Median Overall Survival (mos) | Median Time to First Response (mos) | Duration of Response (mos) |
| Molecular Marker cohort | | | | |
| N = 167 | 109 (65.3%) | 12.5 | 1.2 | 15.0 |
| IDH1/IDH2 | | | | |
| Detected n= 43 | 36 (83.7%) | NR | 1.1 | NR |
| NPM1 | | | | |
| Detected n= 26 | 22 (84.6%) | NR | 1.3 | NR |
| TP53 | | | | |
| Detected n= 37 | 22 (59.5%) | 8.9 | 1.5 | 5.6 |
| FLT3 | | | | |
| Detected n=30 | 16 (53.3%) | 12.4 | 1.8 | 19.9 |

Venetoclax联合IA诱导及大剂量Ara-C巩固在新诊断AML中的I期临床试验

| Pt Age/Gender | Cytogenetics | NGS (VAF%) | Non-Heme Gr 3/4 Adverse events | Outcome (after single induction cycle) |
|---------------|--------------|---|---|--|
| 58F | 9q-, +10 | DNMT3A(40) | DIC after one dose VEN | MRD neg CR |
| 35M | t(1;11), +10 | PTPN11 x 5 (3-9) | Mucositis, elevated ALT | MRD neg CR |
| 62F | Abn 9 | RUNX1 (4) | None | MRD neg CR |
| 54F | 46, XX | DNMT3A (39), NPM1 (31), IDH1 (35), NRAS x 2, (3,20), PTPN11(8) | None | CR |
| 55M | +4, 17q- | DNMT3A (49), NRAS (27), KIT (11), TET2 x 2 (38, 64), U2AF1 (44) | Intracranial hemorrhage/ hemiparesis, B. cereus bacteremia, elevated ALT/AST/bilirubin | CR |
| 73M | 92, XYY | RUNX1 (42), SRSF2 (42) | Typhlitis, heart failure | Early death (day +14, sepsis) |
| 55M | t(X;10), 5q- | CBL (90), ETV6 (43), RUNX1 (49), SETD2 (49) | S. epidermidis and B. cereus bacteremia | MRD neg CR |
| 56F | +8 | PTPN11 (18), IDH1 (4), SRSF2 (13) | Asystolic cardiac arrest, typhlitis, resp failure, acute kidney injury, Klebsiella bacteremia | MRD neg CRi |
| 51F | 7q-, +8 | IDH2 (28), PHF6 (27) | Elevated AST and ALT | MRD pos CR |

- decreasing AraC to 1.5 g/m², and administering VEN for 14 days in ind, and 7 days in cons cycles
- Of 23 R/R pts, **17 pts (74%)** achieved a best response of CR/CRi (12 CR, 5 CRi: 3 CRh + 2 CRp). 12 (52%) pts attained MRD negative status by flow cytometry.
- Of 11 ND pts, **10 pts (91%)** achieved ORR (9 CR, 1 CRh) and all 10 pts became MRD negative by flow cytometry

Venetoclax+去甲基化一线治疗后R/R AML的临床特征

| Treatment Characteristics | All Patients (N = 41) |
|--|--------------------------|
| VEN + HMA regimen, n (%) | |
| ▪ VEN + DEC10 | 18 (44) |
| ▪ VEN + DEC5 | 18 (44) |
| ▪ VEN + AZA | 4 (10) |
| ▪ VEN + DEC10 + FLT3 inhibitor | 1 (2) |
| Response to VEN + HMA, n (%) | |
| ▪ CR | 19 (46) |
| ▪ CRi | 11 (27) |
| ▪ MLFS | 3 (7) |
| ▪ Primary refractory | 8 (20) |
| Median duration of response, mos (range) | 5.3 (0.9-34.1) |
| Median VEN + HMA cycles, n (range) | 4 (1-29) |
| AlloSCT in CR1, n (%) | 4 (10) |

Maiti. ASH 2019. Abstr 738.

Venetoclax+去甲基化一线治疗后R/R AML的治疗仍需进一步探讨

- Outcomes poor for R/R AML after first-line VEN + HMA with median OS of 2.4 mos

| Patient Population | n | Median OS, Mos |
|-----------------------|----|----------------|
| All patients | 41 | 2.4 |
| Subsequent therapy | | |
| ▪ Yes | 24 | 2.9 |
| ▪ No | 17 | 1.3 |
| Disease status | | |
| ▪ Relapse | 33 | 2.3 |
| ▪ Refractory | 8 | 1.7 |
| Initial diagnosis | | |
| ▪ de novo AML | 22 | 2.5 |
| ▪ Secondary AML | 12 | 2.8 |
| ▪ Therapy-related AML | 7 | 1.1 |

Maiti. ASH 2019. Abstr 738.

APR-246+阿扎胞苷在伴TP53突变MDS和AML中的疗效

| Parameter | Overall | MDS | AML | MDS-MPN/CMML |
|------------------------------------|---------|---------|--------|--------------|
| Evaluable patients, n | 45 | 33 | 8 | 4 |
| ORR, n (%) | 39 (87) | 29 (88) | 7 (88) | 3 (75) |
| CR rate, n (%) | 24 (53) | 20 (61) | 4 (50) | 0 (0) |
| Median duration of CR, mos | 7.3 | 7.3 | 7.0 | NE |
| Discontinued for transplant, n (%) | 22 (49) | 17 (52) | 4 (50) | 1 (25) |

- 45/45 (100%) evaluable patients had ≥ 1 mutation in *TP53* DNA-binding domain on sequencing
- Absence of co-mutations at BL predicted higher CR rate (69% vs 25%; $P = .0062$) and ORR (93% vs 69%; $P = .08$)
- $\geq 10\%$ p53 IHC+ BM-MNCs predicted higher CR rate (66% vs 13%; $P = .01$)
- 20/45 (44%) patients obtained NGS negativity at VAF threshold of 5%, with 5 (11%) achieving MRD negativity (0.1% sensitivity) with median MRD VAF at maximum clearance of 0.63%

Blood (2019) 134 (Supplement_1): 676.

CD47单抗Magrolimab (5F9) +阿扎胞苷治疗MDS及AML有效

CD47, a macrophage immune checkpoint and “don’t eat me” signal on cancers

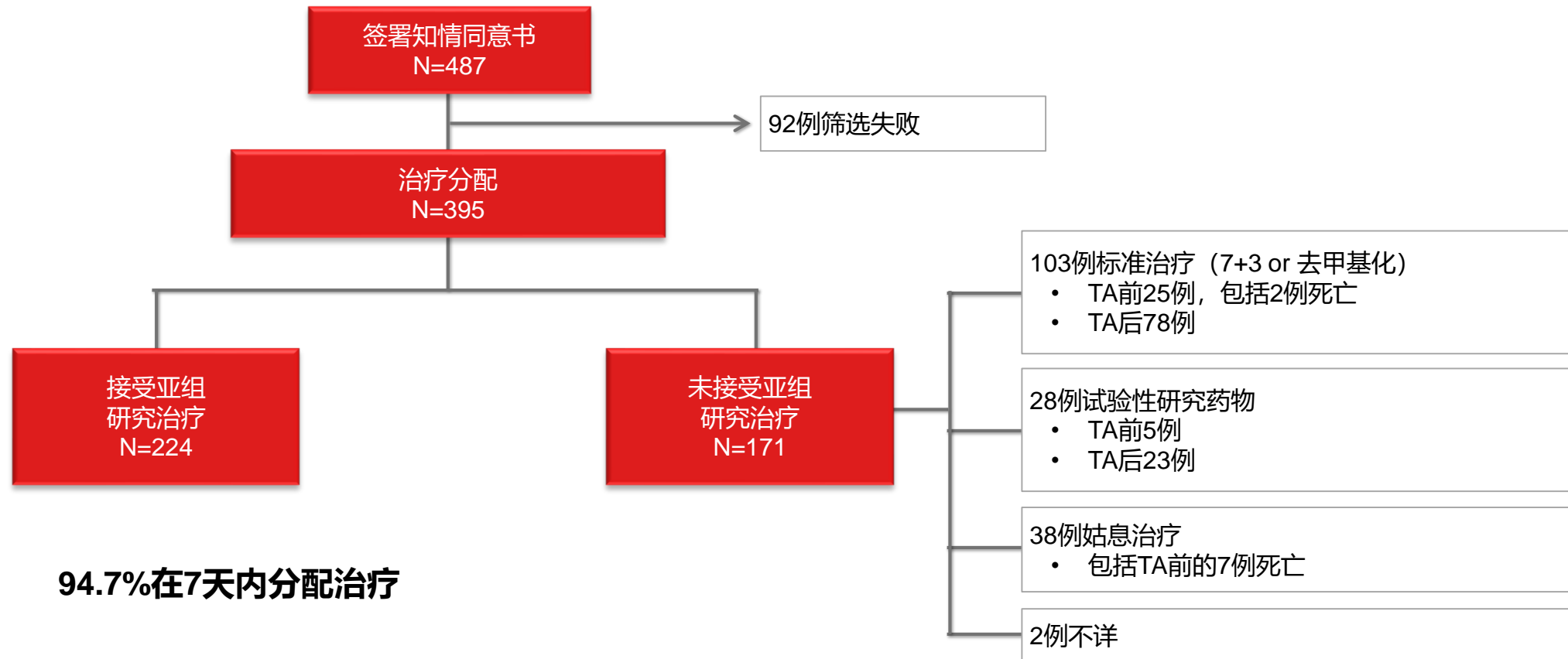
| Magrolimab+AZA (AML: n=16) | |
|----------------------------|----------|
| ORR, n (%) | 11 (69%) |
| CRi/CR, n (%) | 8 (50%) |
| PR, n (%) | 2 (13%) |
| MLFS, n (%) | 1 (6%) |
| SD, n (%) | 5 (31%) |

Magrolimab+AZA is a novel immunotherapy regimen that blocks a key macrophage checkpoint.

ASH Abstract 569

3 老年急性髓系白血病的全程治疗

Beat AML Master Trial



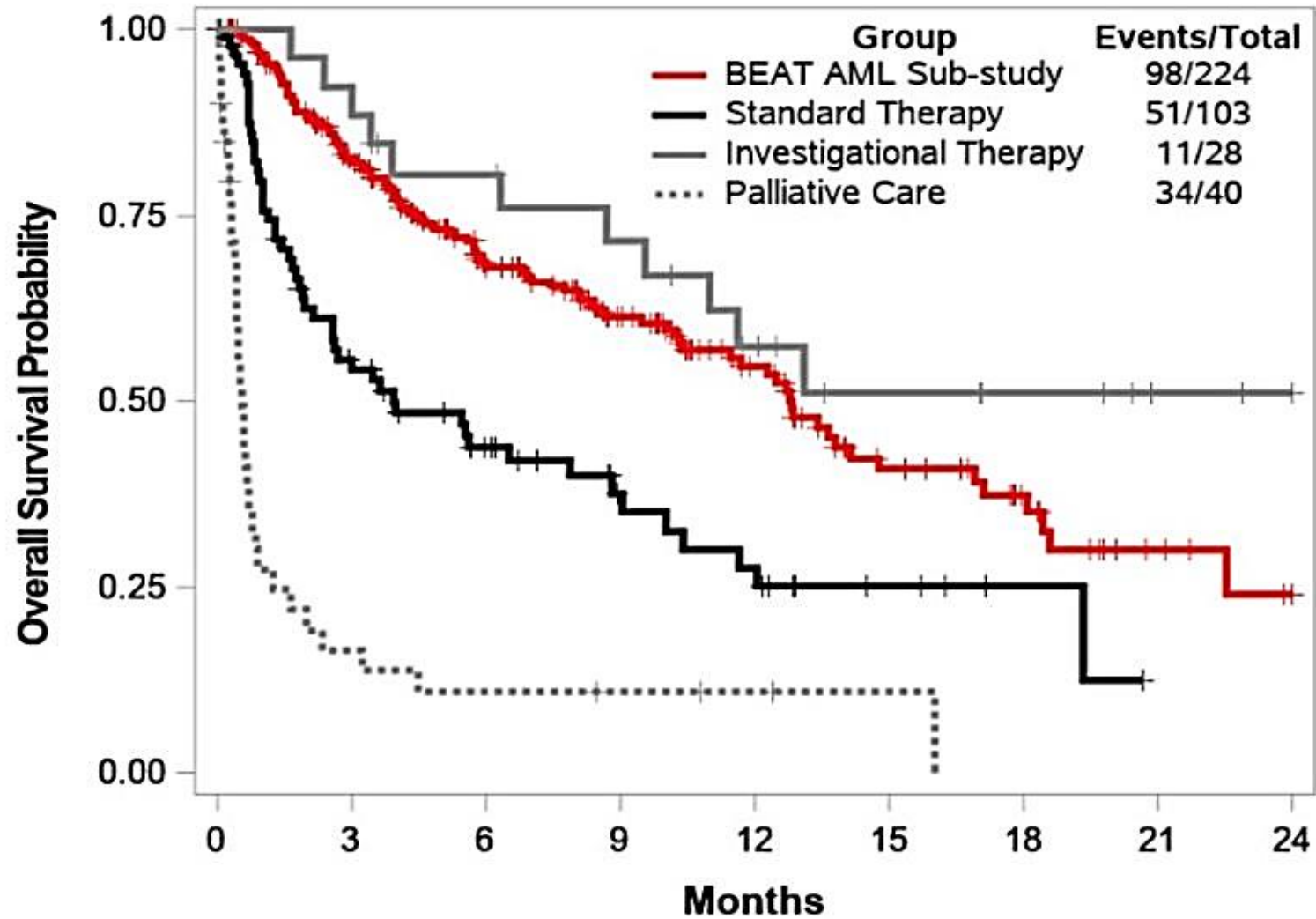
Beat AML Master Trial

| AML Subtype | Drug |
|---|---|
| CBF | Samalizumab (CD200 Ab) + induction |
| NPM1 + FLT3-ITD | Entospletinib (Syk inhibitor) + induction (fit) |
| | Entospletinib (Syk inhibitor) monotherapy (unfit) |
| MLL rearranged | Entospletinib (Syk inhibitor) |
| IDH2 + | Enasidenib |
| IDH1 + | Ivosidenib + Aza |
| TP53+ | Entospletinib (Syk inhibitor) + Decitabine |
| TP53 - Complex Karotype (≥ 3 abn) | Entospletinib (Syk inhibitor) + Decitabine |
| TP53+ | Pevonedistat (Nedd8 inhibitor) + Aza |
| FLT3-ITD+ or FLT3-TKD + | Gilteritinib monotherapy or + Decitabine |
| Tet2/WTI | BI 836858 (CD33 Ab) + Aza |
| Marker Negative | BI 836858 (CD33 Ab) + Aza |

CONCLUSIONS

- Implementation of a rapid treatment assignment umbrella study in elderly AML is feasible with 95% of patients assigned to treatment in ≤ 7 days
- Early death and disease progression prior to treatment assignment is uncommon outside of MLL rearranged AML
- Majority of patients assigned to protocol therapy proceed to trial with increasing frequency as new protocols open
- Promising efficacy observed in several of the treatment arms to date

Beat AML Master Trial中AML的生存情况



HOVON97研究：阿扎胞苷维持治疗可延长老年AML患者DFS

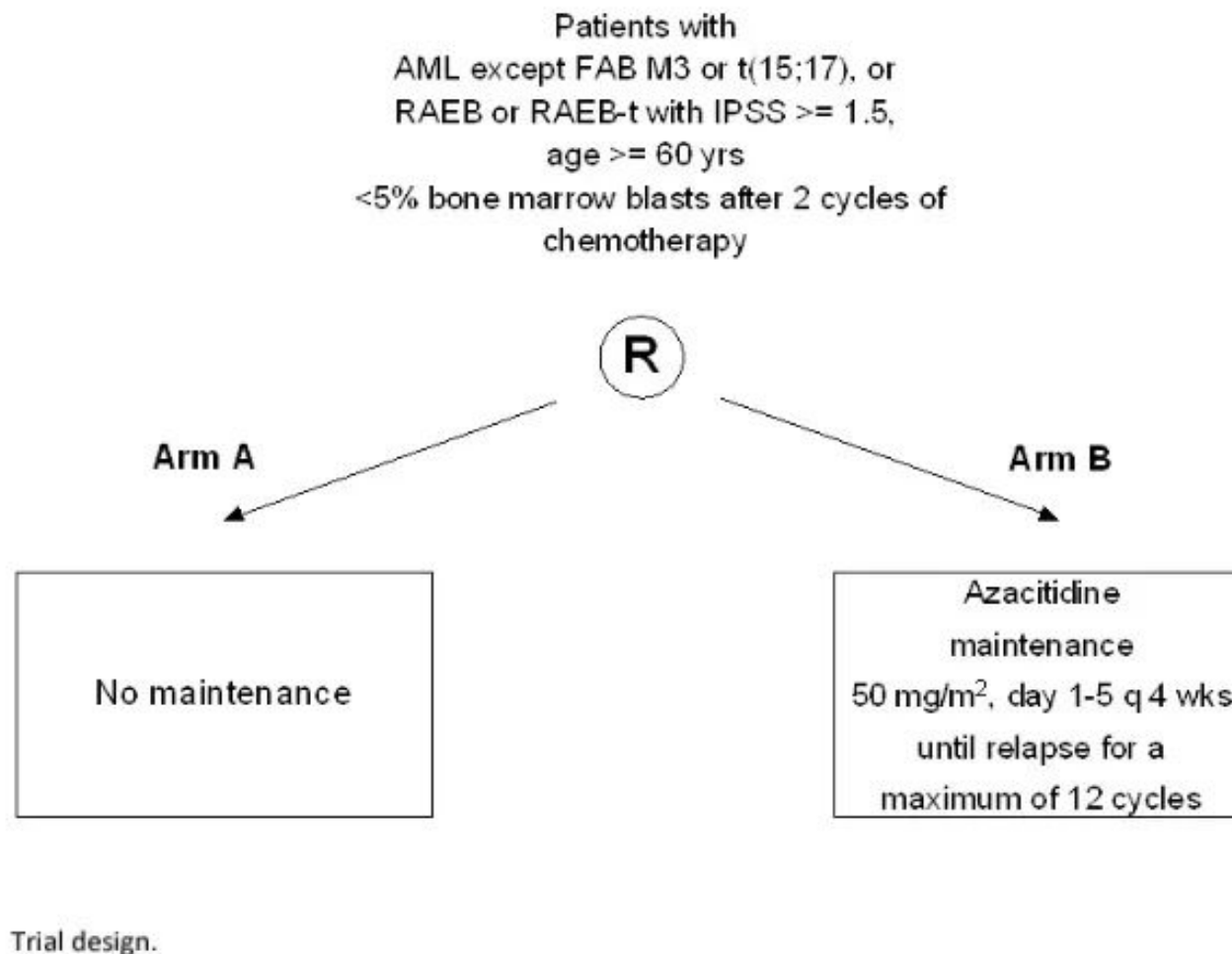
随机性3期临床研究——HOVON-97

N=116，按照1:1分组

A组：观察组

B组：阿扎胞苷组

主要研究终点为DFS，次要研究终点为OS；



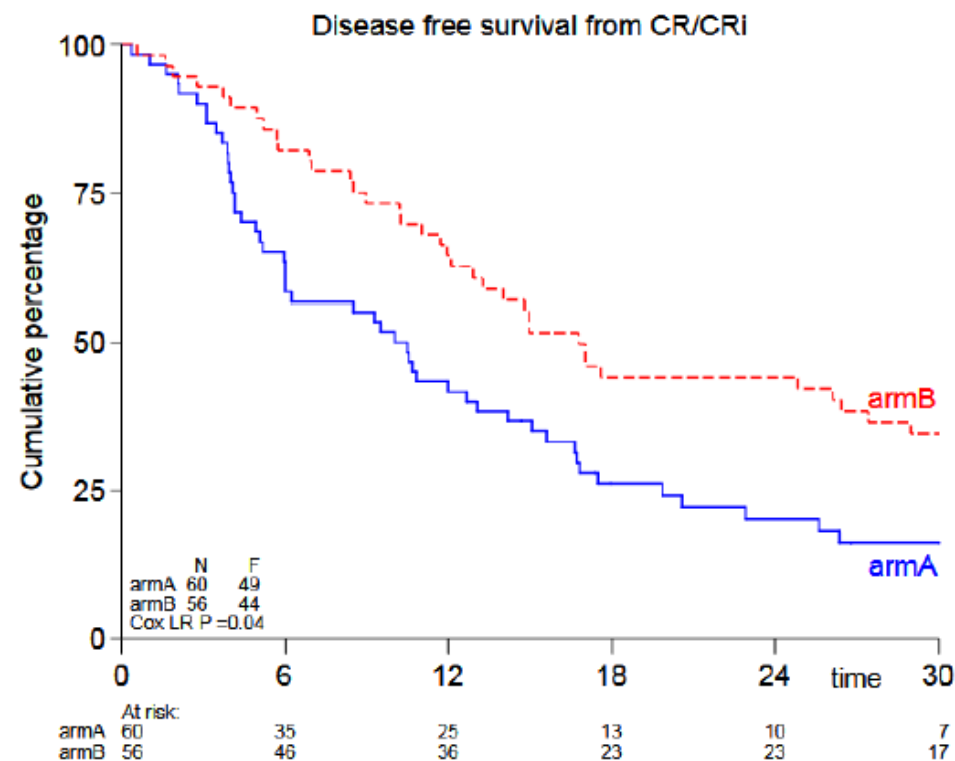
HOVON97研究：阿扎胞苷维持治疗可延长老年AML患者DFS

阿扎胞苷用于这类患者的维持治疗是可行的；

随机化时对诊断时低风险细胞遗传学异常和血小板计数调整后，阿扎胞苷维持治疗组的DFS明显更好

阿扎胞苷组的12个月DFS估计为64%，对照组为42%（ $P = 0.04$ ）；

24个月和36个月DFS估计为44% vs 20%和32% vs 16%。



QUAZAR AML-001研究是口服AZA用于AML维持治疗



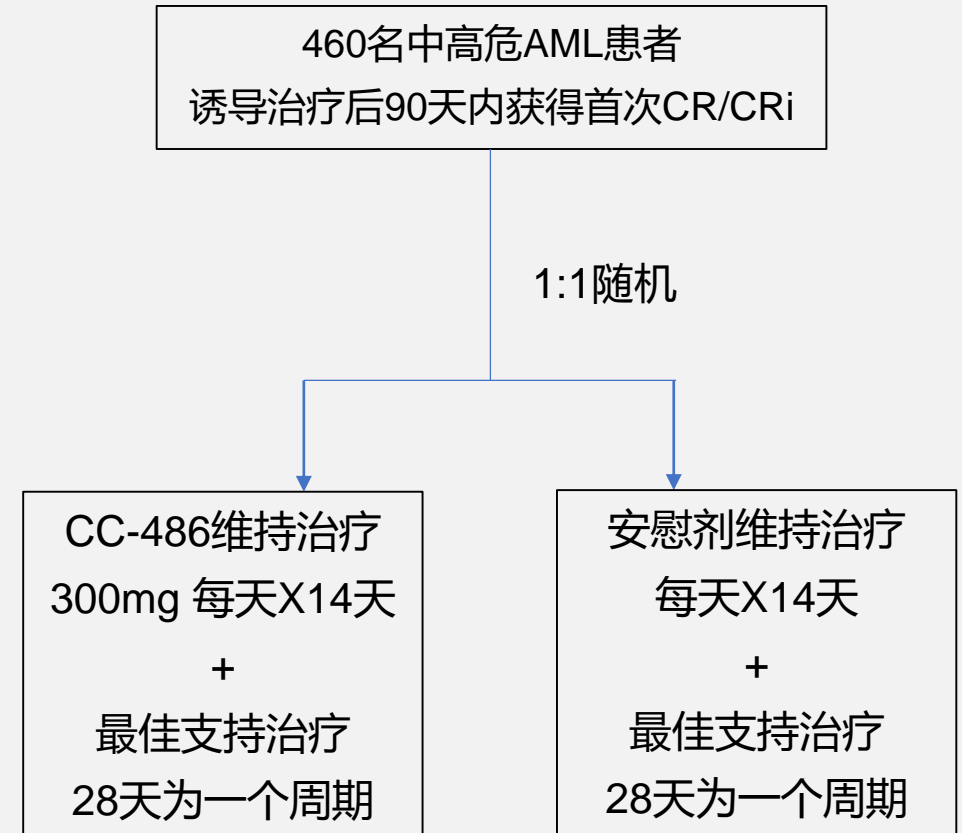
研究设计

1. 含安慰剂对照的III期临床研究，全球13家中心开展。共纳入472名55岁以上中高危AML患者，所有患者均首次通过强烈化疗方案获CR/CRi
2. 患者随机分为CC-486组和安慰剂组，CC-486组接受每周期14天的300mg每天剂量的阿扎胞苷并联合最佳支持治疗，安慰剂组采用同等剂量；每28天为一个周期

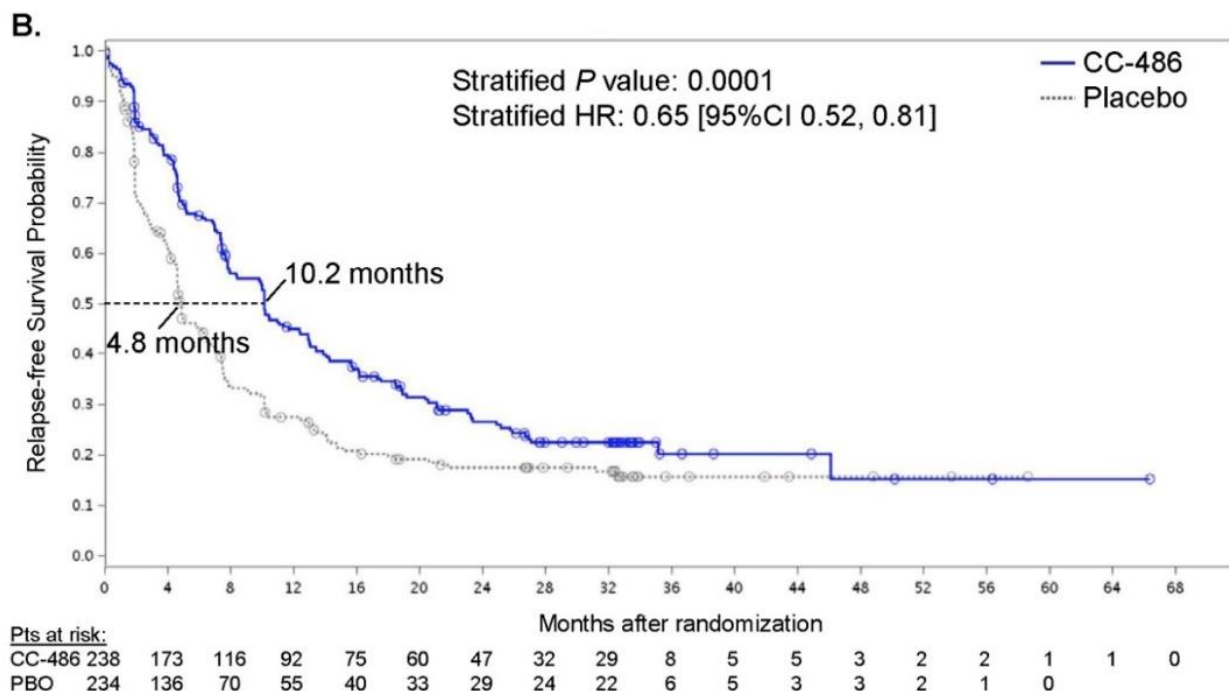


研究目的

OS、RFS、安全性和耐受性、生活质量评分

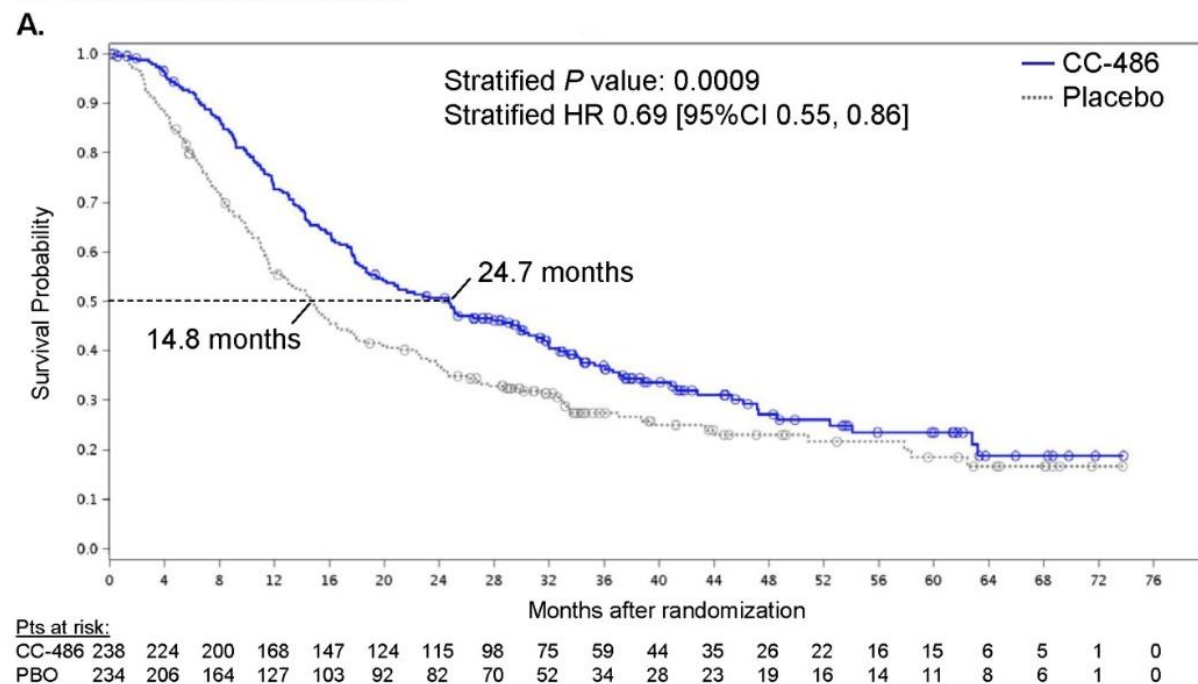


AML的维持治疗：口服阿扎胞苷CC-486用于AML CR1后维持治疗生存获益明显



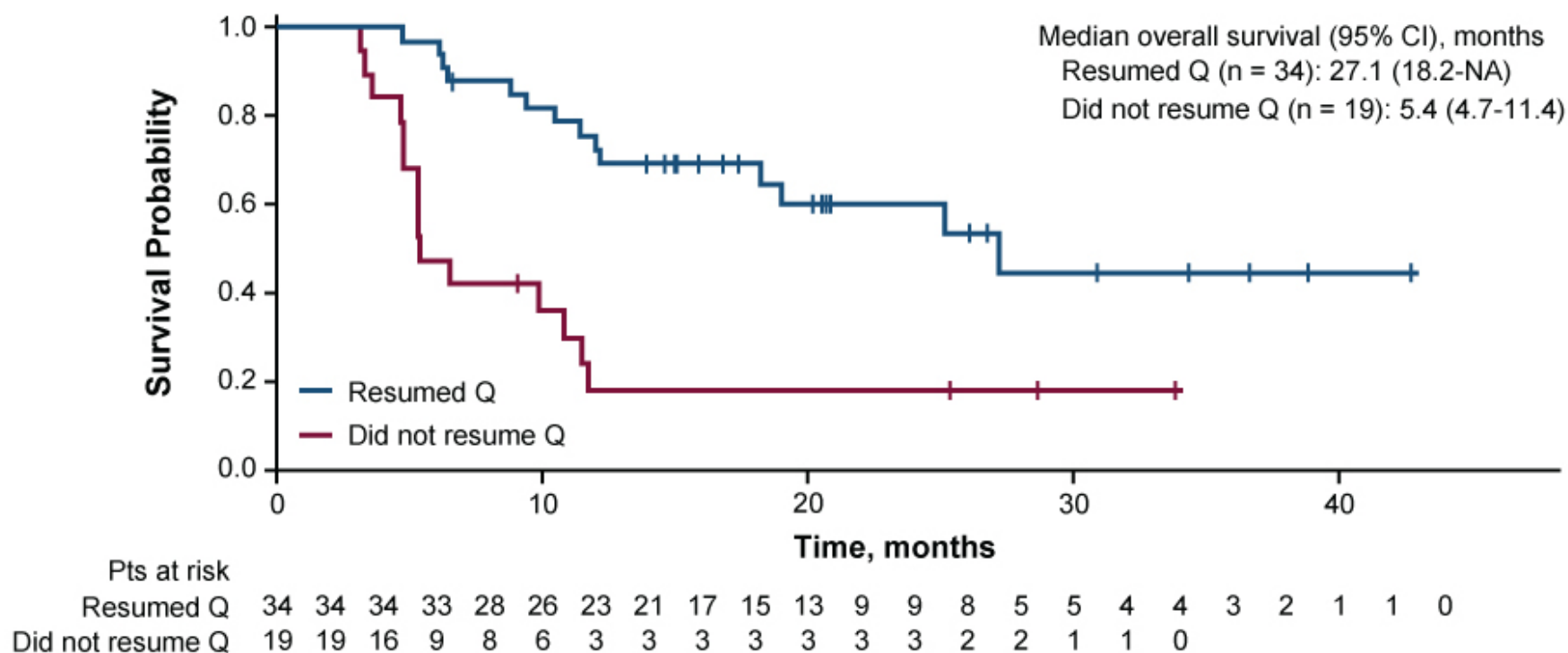
95%CI, 95% confidence interval; HR, hazard ratio; PBO, placebo; Pts, patients

Figure. Kaplan-Meier plots of (A) overall survival and (B) relapse-free survival, from time of randomization



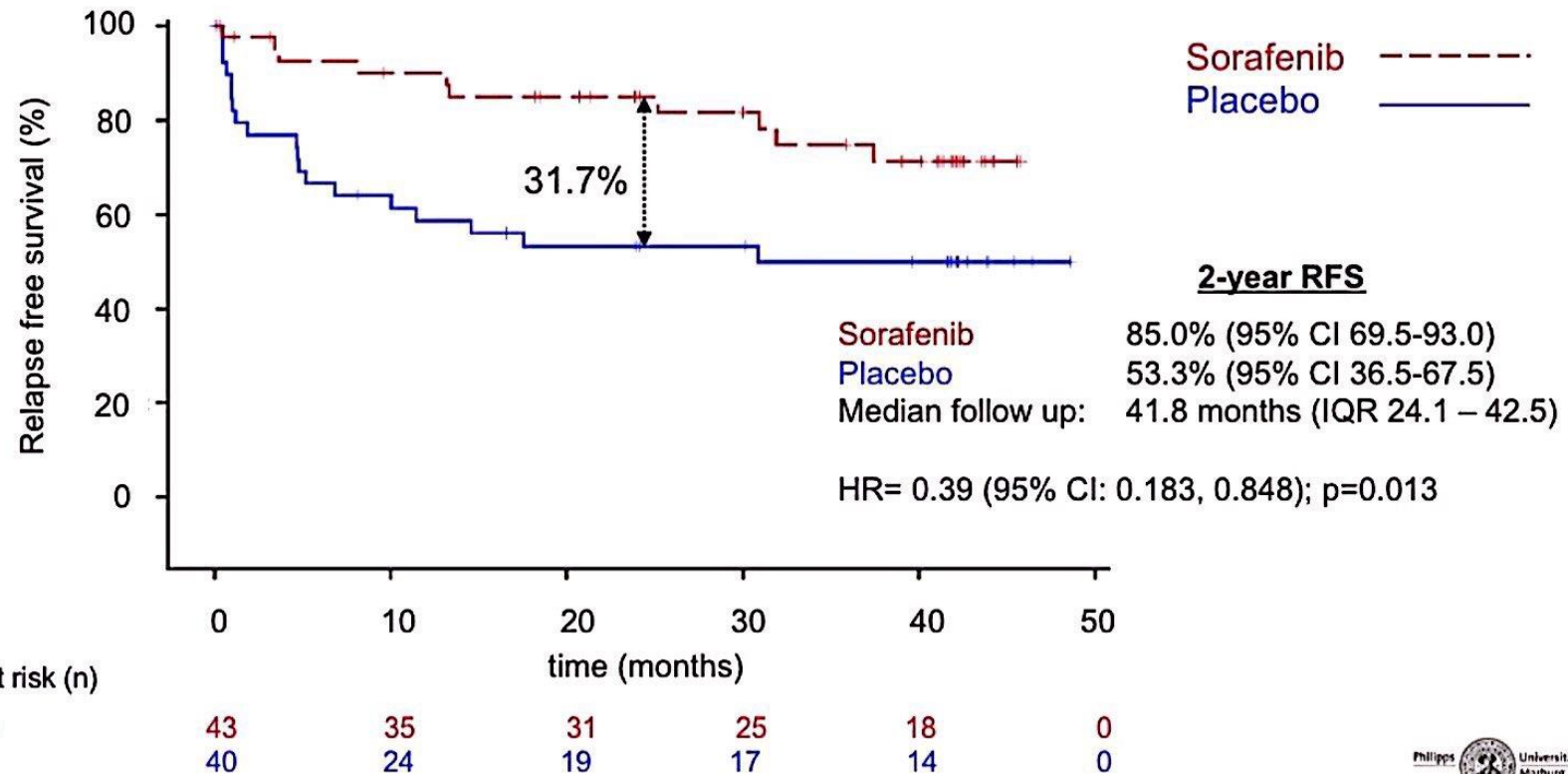
allo-HSCT后Quizartinib维持治疗可带来生存获益

Figure 2. Kaplan-Meier Plot of OS in Pts Who Achieved a Best Overall Response of CRc with Q Who Underwent Allo-HSCT with or without Q Resumption



Sorafenib维持治疗可明显延长RFS

SORMAIN – Results: Relapse free survival

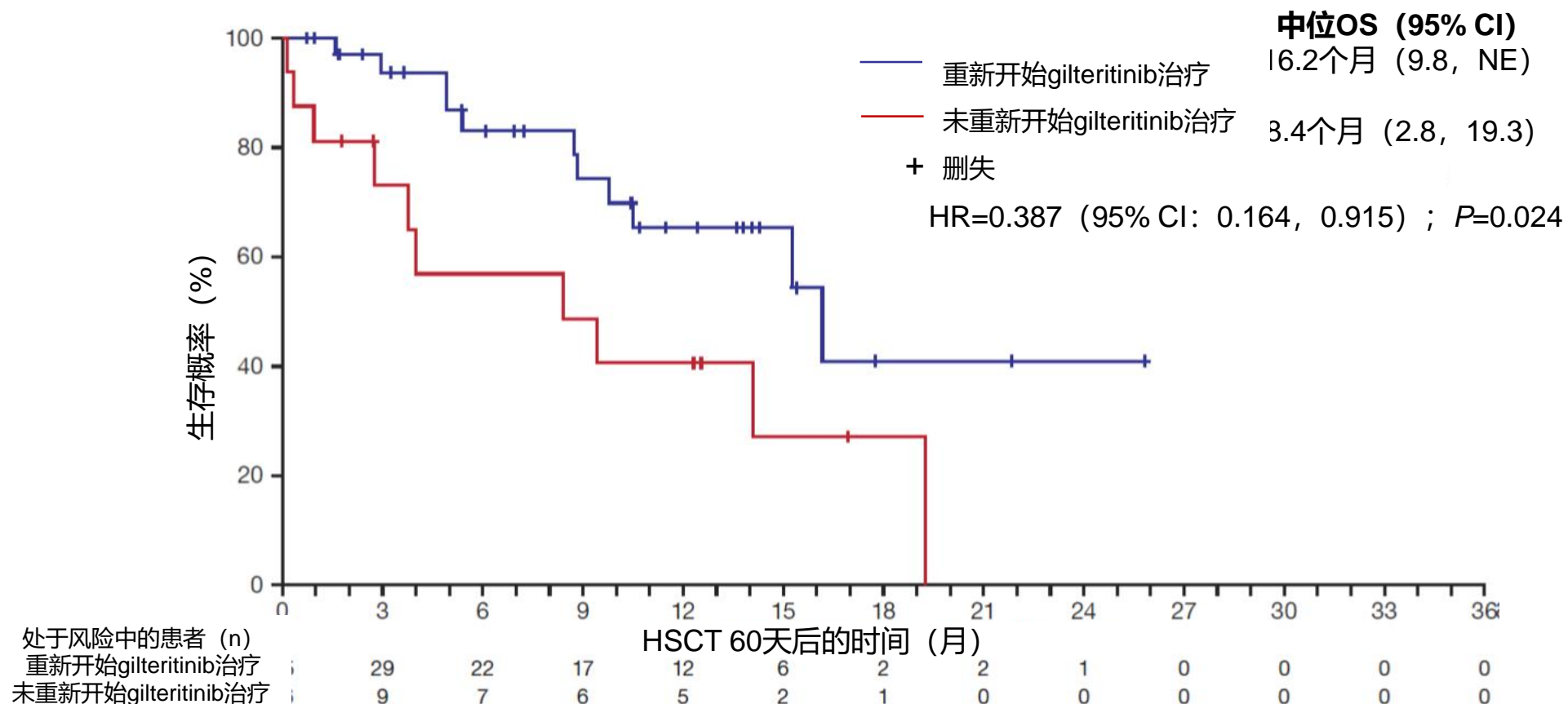


Abbreviations: CI, confidence interval; HR, hazard ratio; IQR, interquartile range; RFS, relapse free survival; EOT, end of treatment; EOS, end of study.



HSCT后Gilteritinib维持治疗可延长OS

Gilteritinib组的HSCT后生存期：维持治疗的效果
(自HSCT后第60天起的界标分析；n=51)



根据对数秩检验确定双侧P值；使用Kaplan-Meier方法结合Greenwood公式确定总生存期和相应的95%置信区间。
缩略语：CI，置信区间；HR，风险比；HSCT，造血干细胞移植；ITT，意向治疗；NE，不可估计；OS，总生存期。

总结

- AML的治疗越来越多的按照遗传学进行分层
- 靶向治疗越来越多的应用于一线、年轻患者
 - 更新的药物及治疗方法仍在研究中

New drugs influencing clinical practice in AML

Fit for intensive chemo

| | 1 st line therapy | Maintenance |
|---------------------------------|-------------------------------|-------------|
| <i>FLT3</i> ^{mut} | Intensive chemo + Midostaurin | Midostaurin |
| <i>CBF</i> | Intensive chemo + GO | CC-486 |
| tAML, sAML, AML MRC | CPX-351 | |
| Alternative non-targeted option | Intensive chemo ± GO | |

≥75 or co-morbidities

| | |
|---------------------------------|---|
| <i>FLT3</i> -ITD | AZA ± <i>FLT3</i> i |
| <i>IDH1</i> ^{mut} | AZA and/or Ivosidenib |
| <i>IDH2</i> ^{mut} | AZA and/or Enasidenib |
| <i>NPM1</i> ^{mut} | HMA or LDAC + Venetoclax |
| Alternative non-targeted option | HMA or LDAC + Venetoclax LDAC + Glasdegib Gemtuzumab ozogomycin |

| Relapsed/refractory AML | |
|----------------------------|--------------|
| <i>IDH1</i> ^{mut} | Ivosidenib |
| <i>IDH2</i> ^{mut} | Enasidenib |
| <i>FLT3</i> ^{mut} | Gilteritinib |
| <i>Other</i> | Chemo ± GO |