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**Sequence of administration and methylation of SOCS3 may govern
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conventional chemotherapy in patients with refractory or relapsed
acute myelogenous leukemia (AML)**

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Abstract:

Introduction: In older patients suffering from AML, aggressive chemotherapy is accompanied with high treatment-related morbidity and mortality. Gemtuzumab ozogamicin (GO), a humanized monoclonal anti-CD33 antibody, represents a well tolerated treatment option, but optimal treatment schedules are still unknown. Additionally, *Suppressor of cytokine signaling 3 (SOCS3)* inhibits the CD33-induced block on cytokine-induced proliferation. Consequently, a variable response of AML cells to anti-CD33-targeted therapy may be caused by modulation of *SOCS3* expression.

Patients and Methods: 24 patients with refractory or relapsed CD33-positive AML received GO as a single agent before or after conventional chemotherapy. The methylation status of the *SOCS3* CpG island was assessed by methylation-specific polymerase chain reaction. **Results:** Response (RR) and overall survival (OS) were significantly higher in 16 patients receiving chemotherapy before GO (RR 81%, OS 14.8 months) compared to 3 patients who received GO single agent therapy (RR 33%, OS 7.2 months) or 16 with GO before chemotherapy (RR 0% OS 2.2 months, $p=0.01$ for RR and $p<0.001$ for OS). Methylation of the *SOCS3* CpG island was found in 8/24 patients. There was a trend towards a higher RR and longer OS in patients with *SOCS3* hypermethylation (RR 86%, OS 25.1 months) compared to unmethylated *SOCS3* (RR 56%, OS 10.3 months, $p = 0.09$). **Conclusion:** Administration of GO a few days after chemotherapy seems to provide better response and survival compared to administration of GO directly prior to chemotherapy. The potential role of *SOCS3* hypermethylation as a biomarker should be further investigated in patients undergoing GO containing therapies.

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1. Introduction

Acute myelogenous leukemia (AML) is the most frequent acute leukemia in adults with a median age at diagnosis of 68 years ¹⁻³. In younger patients, 5-year survival varies between 15% and 70% depending in particular on chromosomal aberrations ⁴. For patients older than 60 years, prognosis is poor, because advanced age is associated with adverse cytogenetics, preceding myelodysplastic syndrome (MDS) and elevated activity of multidrug resistance genes ⁵. Additionally, owing to severe comorbidities, aggressive chemotherapy is often accompanied with high treatment-related morbidity and mortality ².

Gemtuzumab ozogamicin (GO) is a humanized monoclonal murine Ig4 antibody which is linked to the potent antitumor antibiotic calicheamicin and targets the CD33 antigen ⁶⁻¹¹. After internalization, calicheamicin is cleaved from the antigen-antibody complex and induces DNA double-strand breaks and apoptosis by sequence specific binding to DNA ^{6,9,11}. CD33 is expressed on 90-99% of AML blasts, myeloid precursors cells, macrophages, monocytes and dendritic cells, but not on hematopoietic stem cells ^{9,11}. CD33 is member of the *sialic acid-binding immunoglobulin-like lectin (Siglec)* family of inhibitor receptors, and its activation induces apoptosis and inhibition of proliferation in AML cells. GO represents a treatment option for AML and was approved by the Food and Drug Administration (FDA) as a single agent for relapsed AML in patients older than 60 years who are not eligible to receive cytotoxic chemotherapy.

DNA methylation of cytosines within CpG dinucleotides and post-translational acetylation of histones are the most common epigenetic modifications with an impact on chromatin structure and transcriptional activity. Hypermethylation of CpG islands in the promoter region of genes is a well-characterized epigenetic modification associated with transcriptional silencing of cancer-related genes and plays a crucial role in

carcinogenesis¹². *Suppressor of cytokine signaling 3 (SOCS3)* was shown to inhibit the CD33-induced block on cytokine-induced proliferation by leading to its degradation because of forming complexes with CD33¹³. Consequently, Ball et al. suggested that a variable response of AML cells to anti-CD33-antibodies could be caused by differential levels or modulation of *SOCS3* expression¹⁴.

We here report our retrospective analysis of characteristics and outcome of 24 AML patients, treated with GO as a single agent or in combination with chemotherapy in our institution. Additionally, the methylation status of the promoter-associated CpG island of *SOCS3* was analyzed, because we hypothesized that patients with epigenetic dysregulation of *SOCS3* may have a better response to GO owing to a decreased degradation of CD33.

2. Patients and methods

2.1. Patients

Between December 2000 and March 2009, overall 156 patients have been treated for AML in our institution, including 81 patients with relapsed or refractory AML. At the individual physician's discretion, after consideration of CD33 expression, performance status (PS) and risk factors, 24 patients received a GO-containing therapy. If patients repeatedly received GO, we here report on the first GO containing treatment cycle only. All patients had an Eastern Cooperative Oncology Group performance status ≤ 2 and no severe hepatic (bilirubin < 2.0 mg/dl; aspartate aminotransferase < 100 U/l) or renal (creatinine < 2.0 mg/dl) dysfunction before starting treatment. Flow cytometric analysis of myeloblasts revealed positivity for CD33 in all 24 patients. Cytogenetic analyses were performed during routine clinical assessment and risk groups were classified according to Grimwade et al.¹⁵.

2.2. Matched pair analysis

To compare treatment related toxicity and OS in our patient cohort treated with a GO containing therapy, we performed a matched pair analysis with patients treated without GO. Patients were matched for cytogenetic risk group, stage of disease (refractory, 1st relapse, $\geq 2^{\text{nd}}$ relapse) and age. Table I shows the characteristics of the patient cohort and matched control group.

2.3. Treatment schedules

GO was administered either as a single agent (“GO”) or prior to (“GO→chx”) or after (“chx→GO”) various regimens with conventional chemotherapy. Table I shows the patient characteristics of the three treatment groups as well as the matched controls, table II the protocols and doses applied.

2.4. Response evaluation

Response was assessed as proposed by Cheson et al.¹⁶. Patients with all CR criteria but persistent thrombocytopenia (below 100 x 10⁹/L) were considered to have a CRp. Patients with recovery of peripheral blood (PB) values but at least 50% decrease of bone marrow (BM) blasts to 5-25% were reported to have a partial remission (PR). If less than 5% blasts were found in the BM without adequate recovery of peripheral blood counts, patients morphologic leukemia-free state (LFS) was considered. Response was considered in patients achieving a CR, CRp or PR¹⁶. The National Cancer Institute Common Toxicity Criteria version 3.0 (NCI-CTC 3.0) was used for toxicity assessment.

2.5. Methylation specific PCR (MSP)

From all patients with AML, BM and/or PB is routinely collected for analysis of genetic and epigenetic changes. The protocol was approved by the University hospital ethics committee and patients gave informed consent in accordance with the Declaration of Helsinki.

Genomic DNA was isolated from patient samples using standard methods. Approximately 1 µg of DNA was sodium bisulfite-modified and subjected to MSP as described previously¹⁷. MSP primers that specifically recognized the unmethylated *SOCS3* sequence were 5'-GTT GGA GGG TTT TGG GTA TTT AAT GT-3' (sense) and 5'-TAA ATA ACC ATA ACA CAC AAA ACC AAC A-3' (antisense); primers specific for the methylated *SOCS3* sequence were 5'-TGG AGG GTT TCG GGT ATT TAA CGC-3' (sense) and 5'-ATA ACC ATA ACG CAC GAA ACC AAC G-3' (antisense). Reactions were hot-started at 95°C for 5 min and held at 80°C before addition of 0.625 U of Taq polymerase. Temperature conditions for thermocycling were as follows: 35 cycles of 95°C for 30 sec, 58°C for 30 sec and 72°C for 30 sec, followed by 1 cycle of 72°C for 5 min. Normal DNA from PB was treated *in vitro* with *SssI* methyltransferase in order to generate *in vitro* methylated DNA (IVD) that served as a universally positive control for methylated alleles¹⁸. PCR products were separated on 2.5 % agarose gels and visualized by ethidium bromide staining.

2.6. Statistical analysis

Correlations between categorical variables were tested using the chi-square test or Fisher's exact test where appropriate. Overall survival (OS) was calculated from the first day of treatment until death or last follow-up. Relapse-free survival (RFS) was calculated from the day achieving LFS until death or last follow-up. OS and RFS were analyzed using the Kaplan-Meier method and compared using the log rank test.

All statistical tests are two-sided with a level of significance at $p < 0.05$. Statistical analyses were performed using the SAS software package version 9.1.3 (SAS Institute Inc., Cary, NC, USA).

3. Results

3.1. Response

Response evaluation was limited to 23 patients, since one patient received allogeneic stem cell transplantation during myelosuppression following GO containing therapy. Responses (including CR, CRp and PR) to the GO containing therapy were observed in 14 out of 23 patients (61%). Twelve patients achieved CR (52 %), one patient CRp (4%) and one patient PR (4%).

The response rate was significantly higher in patients receiving chemotherapy before GO (81% response) as compared to patients who received GO single agent therapy (33% response) or GO before chemotherapy (0% response, $p = 0.01$, chi-square test). Since no CR was achieved in the latter two groups, relapse-free survival could not be compared between these groups (table II). Noticeably, eight out of nine patients with primary refractory disease achieved a CR using a second chemotherapy followed by GO.

The median OS of all patients was 10.9 months with a RFS of 7.4 months among the 13 patients who achieved CR/CRp. Survival was significantly associated with different treatment schedules: Median survival was only 2.2 months for patients receiving GO followed by chemotherapy, whereas patients with single agent GO therapy survived 7.2 months. In patients who received chemotherapy followed by GO, the longest median survival of 14.8 months was observed ($p < 0.001$, log rank test, figure 1).

Compared to our matched control group, OS of all patients treated with GO was 10.9 months, whereas the median OS of the matched control group was only 3.4 months (figure 2). This difference, however, did not reach statistical significance ($p=0.11$, log rank test).

3.2. Methylation status of SOCS3

We performed MSP analysis to assess the methylation status of *SOCS3* in BM samples of our patient cohort treated with a GO containing therapy, because we hypothesized that response to GO may be modulated by epigenetic dysregulation of *SOCS3*. Hypermethylation of the *SOCS3* CpG island was found in 8/24 patients (33%, Figure 3). The response rate was slightly higher in patients with *SOCS3* methylation (86% response) compared to unmethylated *SOCS3* (56% response, $p=0.17$, chi-square test). Consequently, overall survival was longer in methylated cases (25.1 vs. 10.3 months, hazard ratio 0.29, 95% confidence interval 0.06 to 1.32, figure 4). However, the difference did not reach statistical significance ($p=0.09$, log rank test).

3.3. Toxicity

All patients treated with GO experienced severe myelosuppression (grade 4 thrombocytopenia and neutropenia). The median time until recovery of blood platelets exceeding $50 \times 10^9/L$ in patients with CR was 27 days (range 19-35 days). Recovery of ANC above $0.5 \times 10^9/L$ was also observed on day 27 (range 11-59) in these patients. Neutropenic fever emerged in 21 of all patients (88%), and one patient developed invasive aspergillosis after GO single agent therapy. However, sepsis with detection of pathogens in blood cultures seemed to be more frequent in the “GO”-group (67%) and “GO→chx”-group (60%), compared to the “chx→GO”-group (19%). Additionally, grade 3/4 elevation of bilirubin occurred in one patient (4%), and in six patients (25%),

grade 3/4 elevation of liver enzymes (aspartate aminotransferase = AST, alanine aminotransferase = ALT) was observed. Statistical analysis revealed no significant differences in the frequency of side effects between the three treatment groups. Overall, we observed four early deaths (17%) owing to infectious complications during severe myelosuppression.

In our matched control group, 23 out of 24 patients experienced grade 4 thrombocytopenia and all patients grade 4 neutropenia. Neutropenic fever emerged in 15 patients. Three patients died early due to infectious complications during severe myelosuppression. One additional patient died early owing to cerebral bleeding. We observed grade 3/4 elevation of bilirubin in 3 patients without grade 3/4 elevation of AST or ALT. Differences between the frequency of side effects in patients receiving GO and the matched control group were not statistically significant (all chi-square test).

3.4. Subsequent treatment

Following initial GO based chemotherapy, most patients later received additional treatment in the course of the disease: In three patients younger than 60 years, allogeneic stem cell transplantation (aSCT) was performed after achieving response to GO based treatment. One of these patients relapsed after 14 months, the remaining two patients are in continuous CR for 7 and 14 months, respectively. One additional patient, who did not respond to GO based therapy, underwent aSCT and subsequently died on day +68. Noticeably, although four patients underwent aSCT after a GO containing therapy, no veno-occlusive disease (VOD) was observed.

Five patients who achieved CR after GO based therapy received some GO based maintenance therapy.

4. Discussion

AML is mainly a disease of advanced age, since more than half of the patients are older than 60¹⁹ and one third older than 75 years². In western countries, the incidence rate in patients older than 65 years is about 15 per 100.000/year^{20,21}. In young patients, successful treatment with aggressive chemotherapy is possible in many cases, but the outcome of older patients or in relapsed/refractory cases is extremely poor. Despite the possibility of achieving a CR in 40-60% in these cases using conventional chemotherapy^{2,19,22-24}, the rate of DFS after 3 years is less than 10-20%^{19,22,24,25} with a median survival of only 5-12 months^{7,23}. Owing to multiple comorbidities and reduced PS, aggressive chemotherapy continues to remain a challenge in older patients². Additionally, time of hospitalization and therapy-related complications have to be considered. Consequently, targeted therapies have been developed to overcome the limitations of chemotherapy.

GO is a monoclonal antibody targeting the CD33 antigen linked to an antitumor antibiotic and has been approved for treatment of AML in the elderly. In the literature, many studies on GO-based therapies have been reported. Using GO as a single agent therapy in patients with refractory or relapsed AML, CR was achieved in 13 to 26%, leading to a median OS of 4.9 to 12 months^{10,21,25-27}. As a single agent, GO more likely induced a second CR than high dose araC especially in older patients with early relapse²⁸.

In studies, in which GO was administered within a few days before chemotherapy, CR was achieved in 9.5 to 55% with a median OS of 2 to 8.2 months^{8,11,29-32}. Chemotherapy followed by GO resulted in 10 to 70% CR and a median OS of 2.3 to 11 months^{22,33-36}. Doses of GO ranged from 3 mg/m² to 9 mg/m²^{25,37}. Thus, literature

provides a wealth of options for GO-containing therapies, but optimal patient selection, dose, schedule and combination partners remain still unclear.

After approval of GO in the United States and as proposed by the early literature^{10,29,30,37}, we started to use GO as a single agent or administered GO within a few days prior to conventional chemotherapy. Unfortunately, response and survival of our patient cohort until then was rather poor. Chevallier et al.³⁴ reported in 2005 promising results using GO 4 days after chemotherapy. Consequently, we changed our schedule to chemotherapy followed by GO. In our patients treated with single agent GO or GO followed by chemotherapy, we achieved an overall response in only 33% and 0%, respectively. Noticeably, no CR was observed. After changing our protocol to chemotherapy followed by GO, the response rate significantly increased to 81% (all CR/CRp). This marked increase translated into prolonged overall survival of 14.8 months, compared to 7.2 and 2.2 months in the “GO” or “GO→chx” group. Additionally, compared to a matched control group, a GO based treatment seemed to have a slight, but not statistical significant survival advantage in our patients comprising mainly refractory and relapsed AML.

Morris et al. showed an additive effect of GO and araC and etoposide in cell culture experiments. Thus, for combination of GO with chemotherapy, a better response compared to single agent GO should be expected³⁸. However, the mechanism of the different response according to the sequence of GO and chemotherapy is not clear. Internalization and hydrolytic release of the toxic calicheamicin moiety causes DNA damage and cell cycle arrest³⁹. Because most chemotherapeutic agents mainly target dividing cells, administration of GO before chemotherapy may lead to reduced efficacy of the following chemotherapy. The cytotoxic moiety of GO directly interacts with double-helical DNA in the minor groove and thereby causes site-specific double-stranded cleavage⁴⁰. This effect seems to be independent of DNA replication, thus,

prior administration of chemotherapy causing cell cycle arrest may not interfere with the effectiveness of GO. Additionally, a high CD33 antigen load in the peripheral blood simply consumes GO and thereby limits its penetration into the bone marrow. Prior chemotherapy may rapidly reduce CD33 antigen load and thus lead to a better efficacy of GO ⁴¹.

Recently, Ball et al. suggested a possible interaction between CD33, the target of GO, and *SOCS3*, since intracellular *SOCS3* binds to phosphorylated CD33 leading to proteosomal degradation of complexed *SOCS3* and CD33 ¹⁴. Consequently, we hypothesized that patients with epigenetic dysregulation of *SOCS3* may have a better response owing to decreased degradation of CD33. In our patient cohort, we found promoter methylation of *SOCS3* in 8 out of 24 cases (33%). The response rate in these cases was 86%, compared to 56% in cases without methylation of *SOCS3*. Overall survival was 25.1 months compared to 10.3 months, respectively. Statistical significance was not reached, possibly due to the relatively low patient number. However, the role of *SOCS3* hypermethylation as a biomarker should further be investigated in a larger number of patients undergoing GO containing treatment.

Elevation of liver enzymes is a frequent side effect of GO, and grade 3/4 hepatic toxicity was reported in 30% after GO in combination with idarubicin and araC ²⁹. Quite in accordance with this, we observed 25% grade 3/4 hepatic toxicity in our patient cohort. VOD, a severe complication with thrombosis of small liver veins, was reported with an incidence of 7-20% in studies using GO before chemotherapy ^{8,29,32,37}. However, in studies using GO after chemotherapy, incidence of VOD seems to be lower ^{22,33,36}. In our patient cohort, mainly including patients who received chemotherapy first followed by GO, no VOD was observed. Wadleigh et al. found the risk of VOD in patients who underwent myeloablative aSCT to be increased after prior GO exposure ⁴².

However, four of our patients received aSCT after GO without evidence of VOD.

All of our patients treated with GO experienced severe myelosuppression with grade 4 neutropenia and thrombocytopenia. Both, the median time to recovery of ANC $>0.5 \times 10^9/L$ and blood platelets $> 50 \times 10^9/L$ was 27 days and thus comparable to literature data^{10,19,26}. During myelosuppression, in almost all patients (88%) neutropenic fever emerged and four patients (17%) subsequently died due to infectious complications. Treatment-related mortality was reported to be 5-13% in published studies using GO after chemotherapy^{22,33,36}, whereas mortality in studies using GO before chemotherapy seemed to be higher (19-57%)^{8,11,37,43}.

Of note, multiple responses were observed in a single patient treated repeatedly with GO. One of our patients first received GO in 3rd relapse, comprising GO after araC and thioguanine leading to a CR, but this patient again relapsed after a few months. Subsequently, this patient was successfully treated with GO-containing regimens for seven times during the following 44 months.

Although the significance of our results is limited by the small number of patients, we propose that GO, given after chemotherapy, presents an efficient and safe therapeutic option for patients with refractory or relapsed AML, especially in advanced age. GO may even be administered, if a subsequent aSCT is planned. Administration of GO a few days after chemotherapy seems to provide better response and survival as well as fewer side effects compared to administration of GO directly prior to chemotherapy. Additionally, the methylation status of *SOCS3* may modulate response to GO. In order to optimize patient selection and treatment schedules containing GO and to further elucidate the impact of *SOCS3* hypermethylation on response and outcome, larger prospective studies are warranted.

5. Literature

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6. Figure legends

Figure 1: Estimated overall survival according to treatment schedule. OS was 2.2 months (“GO→chx”), 7.2 months (“GO”) and 14.8 months (chx→GO), respectively (p<0.001, log rank test)

Figure 2: Survival of all patients receiving a GO containing therapy (OS 10.9 months) compared to a matched control group treated without GO (OS 3.4 months, p=0.11, log rank test)

Figure 3: Representative MSP analysis of *SOCS3* in AML patient samples. Normal peripheral blood (PB) and. *in vitro* methylated DNA (IVD) and water served as controls. **Lane U**, amplified product with primers recognizing unmethylated *SOCS3* sequence. **Lane M**, amplified product recognizing methylated *SOCS3* sequence.

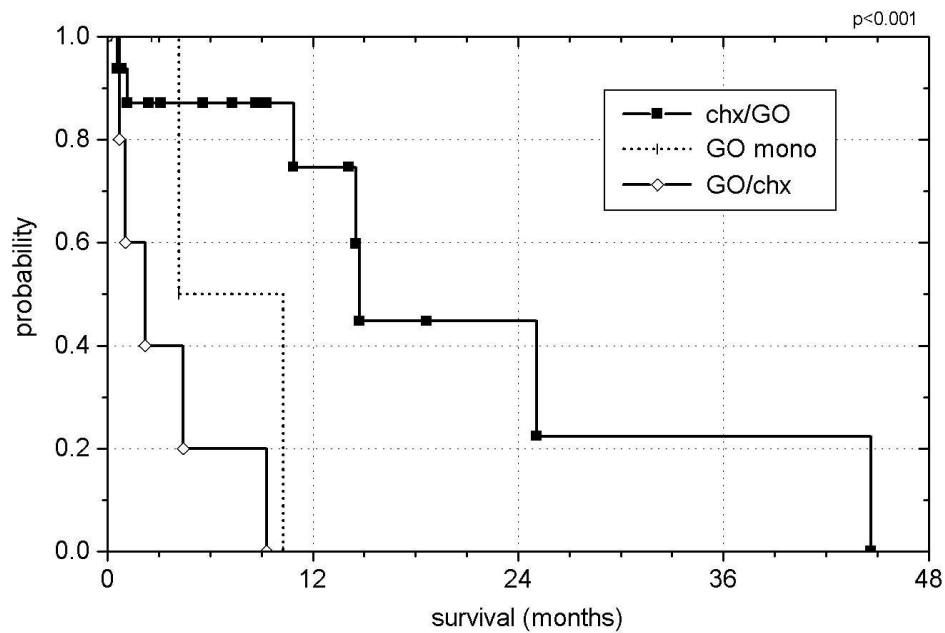
Figure 4: Survival according to the methylation status of *SOCS3* (25.1 months vs. 10.3 months, p=0.09, log rank test)

Table I: Patient characteristics and response; CR = complete remission, PR = partial remission, TF = treatment failure, OR = overall response

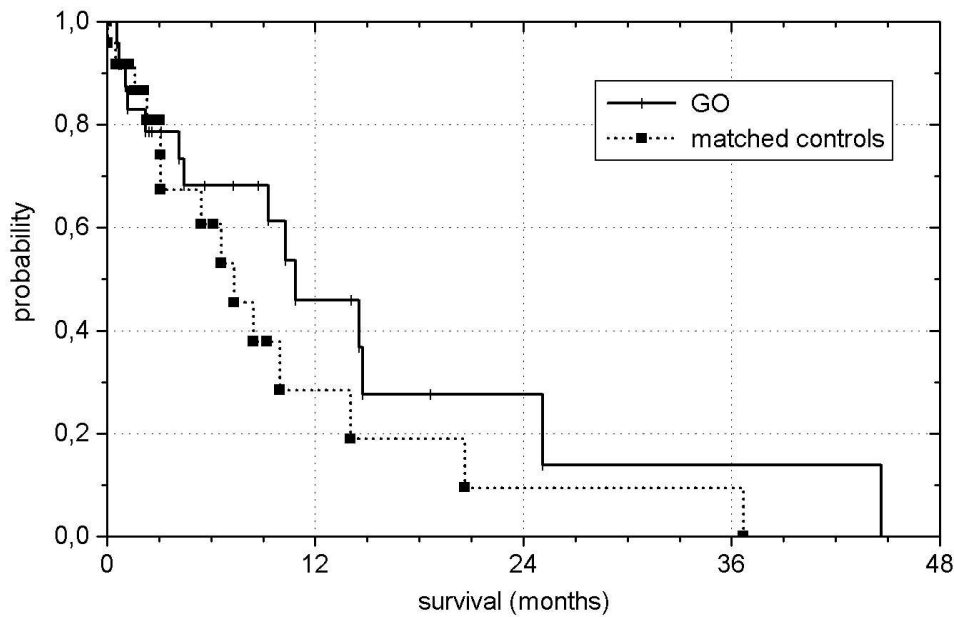
| | all patients | GO | GO→chx | chx→GO | p-value (among Pts with GO) | matched controls |
|------------------------|--------------|------------|------------|------------|-----------------------------------|---------------------|
| <u>no. of patients</u> | 24 | 3 | 5 | 16 | | 24 |
| <u>age (years)</u> | | | | | n.s. | |
| mean (range) | 66 (43-82) | 76 (70-82) | 62 (43-74) | 65 (43-76) | | 61 (38-84) |
| <u>disease status</u> | | | | | n.s. | |
| refractory | 12 | 2 | 1 | 9 | | 12 |
| 1. relapse | 8 | 1 | 2 | 5 | | 11 |
| ≥ 2. relapse | 4 | 0 | 2 | 2 | | 1 |
| <u>risk group</u> | | | | | n.s. | |
| favorable | 0 | 0 | 0 | 0 | | 0 |
| intermediate | 19 | 3 | 4 | 12 | | 20 |
| poor | 2 | 0 | 0 | 2 | | 2 |
| not available | 3 | 0 | 1 | 2 | | 2 |
| <u>response</u> | | | | | | |
| CR/CRp | 13 | 0 | 0 | 13 | | 8 |
| PR | 1 | 1 | 0 | 0 | | 1 |
| TF | 9 | 2 | 5 | 2 | | 14 |
| OR (CR(p)/PR) | 14 (61%) | 1 (33%) | 0 (0%) | 13 (81%) | 0.01 | 9 (38%) |
| not available | 1 | 0 | 0 | 1 | | 1 |
| early deaths | 4 (17%) | 0 (0%) | 2 (40%) | 2 (13%) | n.s. | 4 (17%) |

Table II: Treatment schedules; GO = gemtuzumab ozogamicin, araC = cytosine arabinoside, PO = orally, SC = subcutaneously

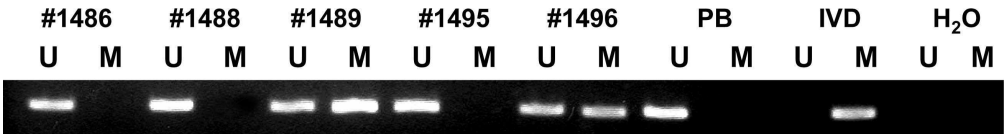
| patient number | protocol |
|-----------------|---|
| GO | |
| 2 | GO 9 mg/m ² day 1 |
| 1 | GO 9 mg/m ² day 1 and day 20 |
| GO→Chemo | |
| 1 | GO 9 mg/m ² d1, idarubicin 12 mg/m ² qd day 11-12, araC 100 mg/m ² qd day 11-15 |
| 3 | GO 9 mg/m ² d1, mitoxantrone 10 mg/m ² qd day 2-4, etoposide 100 mg/m ² qd day 2-4 |
| 1 | GO 9 mg/m ² d1, idarubicin 12 mg/m ² qd day 2-4, topotecan 1.25 mg/m ² qd day 2-4, araC 1000 mg/m ² qd day 2-6 |
| Chemo→GO | |
| 5 | araC 3 mg/kg sc q12h day 1-7, thioguanine 2.5 mg/kg PO qd, day 1-7, GO 9 mg/m ² once (between day 7 and 17) |
| 9 | mitoxantrone 10 mg/m ² qd day 1-3, etoposide 100 mg/m ² qd day 1-3, GO 9 mg/m ² once (between day 5 and 16) |
| 1 | fludarabin 30 mg/m ² qd day 1-5; araC 2000 mg/m ² qd day 1-5; idarubicin 10 mg/m ² qd day 1-3, GO 9 mg/m ² day 14 |
| 1 | idarubicin 12 mg/m ² qd day 1-2, araC 100 mg/m ² qd day 1-5, GO 9 mg/m ² day 15 |



Estimated overall survival according to treatment schedule. OS was 2.2 months ("GO→chx"), 7.2 months ("GO") and 14.8 months (chx→GO), respectively ($p < 0.001$, log rank test)
296x201mm (120 x 120 DPI)

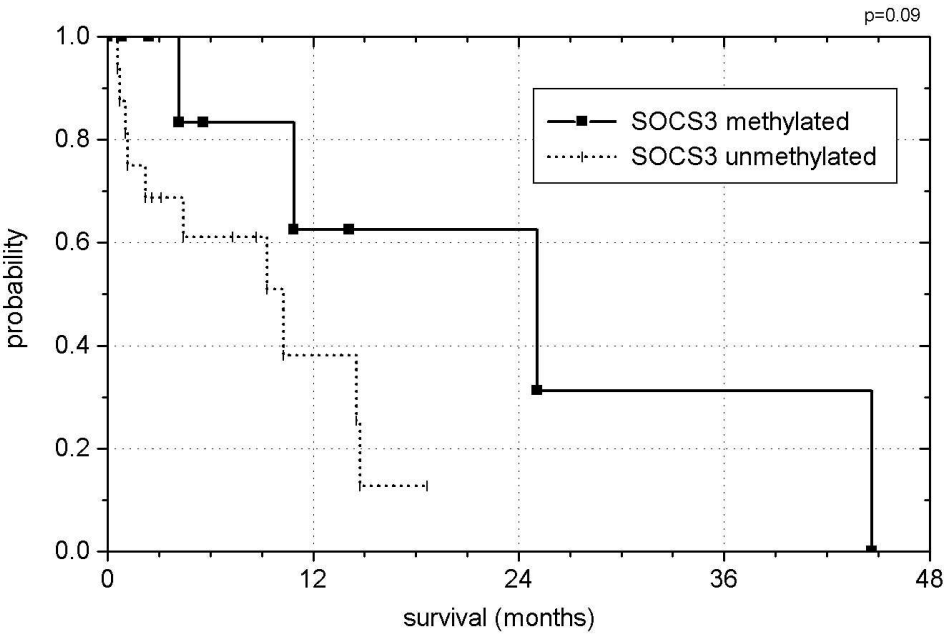


Survival of all patients receiving a GO containing therapy (OS 10.9 months) compared to a matched control group treated without GO (OS 3.4 months, $p=0.11$, log rank test)
296x201mm (120 x 120 DPI)



Representative MSP analysis of SOCS3 in AML patient samples. Normal peripheral blood (PB) and. in vitro methylated DNA (IVD) and water served as controls. Lane U, amplified product with primers recognizing unmethylated SOCS3 sequence. Lane M, amplified product recognizing methylated SOCS3 sequence.

690x88mm (96 x 96 DPI)



Survival according to the methylation status of SOCS3 (25.1 months vs. 10.3 months, $p=0.09$, log rank test)
296x201mm (120 x 120 DPI)