

# GPX3 methylation is associated with hematologic improvement in low-risk myelodysplastic syndrome patients treated with Pai-Neng-Da

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## Abstract

**Objective:** The aim of this prospective randomized controlled clinical trial was to explore the relationship between *GPX3* methylation and Pai-Neng-Da (PND) in the treatment of patients with low-risk myelodysplastic syndrome (MDS).

**Methods:** There were 82 low-risk MDS patients who were randomly divided into the following two groups: androl, thalidomide, and PND capsule (ATP group,  $n = 41$ ); or androl and thalidomide (AT group,  $n = 41$ ). Hemoglobin and neutrophil and platelet counts and changes in *GPX3* methylation level were assessed.

**Results:** The plasma hemoglobin level increased in both groups after treatment. However, the platelet count increased only in the ATP group. Patients in the ATP group had a better platelet response than the AT group, and *GPX3* methylation markedly decreased after treatment with ATP but not after treatment with AT. Moreover, male patients had a significantly lower *GPX3* methylation level than female patients, while platelet counts from male patients increased dramatically after the ATP regimens compared with female patients. *GPX3* methylation changes were negatively correlated with platelet changes in ATP group.

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**Conclusion:** PND can improve hematological parameters and decrease the *GPX3* methylation level. Decreasing *GPX3* methylation is associated with the hematologic response that includes platelet in *GPX3* methylation.

**China Clinical Trial Bureau (ChiCTR; <http://www.chictr.org.cn/>) registration number:** ChiCTR-IOR-15006635.

## Keywords

Low-risk myelodysplastic syndrome, *GPX3*, DNA methylation, Pai-Neng-Da, hematological response, platelet

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## Introduction

Myelodysplastic syndrome (MDS) is a heterogeneous clonal myeloid disease that is characterized by ineffective hematopoietic function, which is accompanied by abnormal maturation and developmental abnormalities in one or more blood cell lineages.<sup>1</sup> MDS mainly affects elderly patients. The annual incidence of MDS in the elderly population is 12 to 50 per 100,000 people, which is much higher than 3.5 to 10 per 100,000 people for the general population.<sup>2,3</sup> MDS is associated with genetic mutations, immune disorders, chromosomal abnormalities, epigenetic abnormalities, and other factors.<sup>4,5</sup> Therapeutic strategies for low-risk MDS include red blood cell (RBC) infusion and iron chelation therapy, the combined use of hematopoietic growth factors and immunosuppressive agents, and androgen-assisted therapy, but the response to these treatments is unsatisfactory.<sup>5-7</sup>

Numerous studies have shown that glutathione peroxidase is involved in the development and progression of many tumors.<sup>8,9</sup> Glutathione peroxidase 3 (*GPX3*) is the most widely studied member of the *GPX* family.<sup>10</sup> *GPX3* reduces accumulation of hydrogen peroxide in the body during cellular oxidative metabolism.<sup>11,12</sup> The primary biochemical role of hydrogen peroxide is to modulate the characteristics of cancer

cells, including proliferation, invasion, migration, angiogenesis, and apoptosis.<sup>13</sup> Abnormal inactivation or low *GPX3* expression may result in the accumulation of excess reactive oxygen species (ROS), including hydrogen peroxide, leading to tumorigenesis.<sup>13,14</sup>

Abnormal DNA methylation at the CpG island of the promoter in tumor-associated genes is a common event in human cancer,<sup>14</sup> which has also been shown to be involved in the development and progression of MDS.<sup>14</sup> It is a biomarker for the prognosis, treatment, and survival of patients with MDS.<sup>15-19</sup> Recently, hypomethylated drugs, such as azacitidine and decitabine, have been shown to have clinical effects in the treatment of MDS.<sup>20,21</sup>

*GPX3* hypermethylation is associated with the development and progression of cancer, including gastric cancer, colorectal cancer,<sup>21</sup> chronic myeloid leukemia,<sup>22</sup> and acute promyelocytic leukemia.<sup>23</sup> *GPX3* methylation in the bone marrow can also predict poor prognosis of MDS and leukemia transformation.<sup>15</sup> CpG island methylation in the *GPX3* gene promoter region has a significant effect on the regulation of *GPX3* transcription and expression.<sup>24,25</sup>

The biologically active component in ginseng extract, the panaxadiol saponin component, was isolated from total saponins of ginsenosides using a hematopoiesis

biological activity assay, and it was formulated into capsules that are called Pai-Neng-Da (panaxadiol saponins component, PND), which is a class-five new Chinese patent medicine. The composition and content of PND have been analyzed and defined as five monomers of panaxadiol saponins. The PND patent for treating a variety of pancytopenia has been authorized by the State Intellectual Property Office of China. PND was used subsequently for a series of pharmacodynamics, toxicological, and clinical studies. All research data for the 23 items were submitted to the China Food and Drug Administration (CFDA), and two certificates for the new class-five Chinese patent medicine were authorized and granted by CFDA in 2010, including both PND capsule (approval No. 2010L00856) and panaxadiol saponins component (approval No. 2010L00857). This was then successfully transferred to Ningbo Tianzhen Pharmaceutical Co. Ltd. for clinical trials and commercial production. PND was demonstrated to be safe, and six to ten tablets daily was recommended as a safe dose in a phase I clinical trial.<sup>1</sup> A phase II clinical trial was performed in seven hospitals with a professional advantage for treating primary immune thrombocytopenic purpura (ITP) and chronic angiogenic leukocytopenia (neutropenia).<sup>26</sup> The clinical results confirmed that PND was effective and had no adverse side effects.<sup>27</sup> Consistent with previous studies and as supported by this project, this study used PND+andriol+thalidomide or andriol+thalidomide to investigate the efficacy of PND in the treatment of low-risk MDS, and to test the association between *GPX3* methylation and PND for the adjuvant treatment of patients with MDS.

## Methods

This study is a subanalysis of a prospective clinical trial that was registered with

the China Clinical Trial Bureau (ChiCTR; <http://www.chictr.org.cn/>; registration number ChiCTR-IOR-15006635).

## Inclusion and exclusion criteria

Low-risk MDS patients were diagnosed in accordance with the Vienna diagnostic criteria, including refractory anemia (RA), refractory anemia with multilineage dysplasia (RAEB-1), and refractory cytopenia with multilineage dysplasia (RCMD) type. On the basis of the International Prognostic Scoring System (IPSS) score, patients in the low-risk group had a score of 0 to 1.

The following patients are excluded, including pregnant women or women who planned to become pregnant, and breastfeeding women; patients with severe cardiovascular, hepatic, renal system or mental disease; patients with the complication of severe infection; patients with some types of MDS including refractory anemia with increased blasts, 5q-syndrome, MDS that were not classified, or IPSS score  $\geq 1.5$  points; or MDS patients who needed more active treatment such as demethylation chemotherapy, chemotherapy, or bone marrow transplantation.

## Drugs

PND capsule was provided by Ningbo Tianzhen Pharmaceutical Co. Ltd. (Ningbo, China; batch No. 20120101). Specifications were as follows: size 4 gelatin capsules and 40 mg per capsule. Each capsule contained five panaxadiol saponins monomers with a purity of 92.44% as analyzed and defined by high-performance liquid chromatography (HPLC) using specific monomers of ginsenosides as the reference standards. Androl was purchased from Catalent France Beenheim S.A. (Beenheim, France), and thalidomide was purchased from Changzhou Pharmaceutical Factory Co., Ltd. (Changzhou, China).

### Study design

Low-risk MDS patients from Ningbo First Hospital (from October 2015 to March 2019) were included in this prospective randomized controlled clinical trial. Patients were randomly divided into two groups. The ATP group comprised patients who received oral androl, thalidomide and PND (PND capsule, ginsengdiol group saponin extract; 40 mg, three capsules at a time twice per day). Andriol Testocaps capsule (80 mg) were taken twice daily, and thalidomide tablets 100 mg were taken before going to bed. Patients in the AT group received oral androl and thalidomide, as follows: Andriol Testocaps capsule 80 mg twice daily; and thalidomide tablets 100 mg before going to bed for a 4-week course of treatment, and there were five courses of medication. The neutrophil count, hemoglobin level, platelet count, hepatic function, and renal function of patients were assessed at the beginning of the experiment and at follow-up. Bone marrow samples from the two groups were collected before and after treatment for 3 months and stored at  $-80^{\circ}\text{C}$  until methylation detection. This study was approved by the Human Research Ethics Committees of Ningbo First Hospital (Ethical approval number, 2015-R001, Approval Date, 23 January 2015). All patients provided written informed consent for the bone marrow collection and immunophenotyping research.

### Efficacy evaluation

Therapeutic efficacy was evaluated in accordance with the efficacy standard of the MDS International Working Group.<sup>28</sup> In this study, the hematologic improvement including platelet response (HI-P), neutrophil response (HI-N), and erythroid response (HI-E) was our focus, especially HI-P. The major HI-P response was defined as follows: for patients with a pretreatment platelet count that was less than  $100,000/\text{mm}^3$ , it

was defined as an absolute increase of  $30,000/\text{mm}^3$  or more; and for platelet transfusion-dependent patients, it was defined as stabilization of the platelet count and platelet transfusion independence. The minor HI-P response was defined as follows: for patients with a pretreatment platelet count less than  $100,000/\text{mm}^3$ , it was defined as a 50% or more increase in platelet count with a net increase greater than  $10,000/\text{mm}^3$  but less than  $30,000/\text{mm}^3$ . The improvements must last at least 2 months in the absence of ongoing cytotoxic therapy.

### DNA methylation detection for GPX3

The Omega DNA Mini Kit (Omega Bio-Tek, Norcross, GA, USA) was used to extract genomic DNA from bone marrow and blood. The bisulfite conversion process of the DNA template was as described in our previous work.<sup>29</sup> The quantitative methylation-specific polymerase chain reaction (PCR) (qMSP) technology, which was based on fluorescent dye (SYBR-Green), was used to quantitatively detect the *GPX3* methylation level.<sup>30</sup> The qMSP procedure was performed as follows:  $95^{\circ}\text{C}$  for 10 minutes, then 45 cycles for 30 s at  $95^{\circ}\text{C}$ , 30 s at  $58^{\circ}\text{C}$ , and 30 s at  $72^{\circ}\text{C}$ . Melting curve analysis was performed at  $95^{\circ}\text{C}$  for 15 s and  $60^{\circ}\text{C}$  for 1 minute. The primer sequences for *GPX3* were 5'-TGTTTATGTTATTGTCGTTTC G-3' (forward primer) and 5'-CCTTACAA CCAATCGCTAA-3' (reverse primer).  $\beta$ -actin was used as an internal reference, and the  $\beta$ -actin sequences were 5'-TGGTG ATGGAGGAGGTTTAG TAAGT-3' (forward primer) and 5'-AACCAATAAAACC TACTCCTCCCTTAA-3' (reverse primer). The percentage of methylation reference (PMR) for *GPX3* was calculated as previously described.<sup>30</sup>

### Statistical analysis

SPSS version 17.0 software (SPSS Inc., Chicago IL, USA) was used for statistical

analysis. Measurement data were presented as the mean  $\pm$  standard deviation ( $x \pm s$ ). Paired-sample *t*-tests were used for statistical analysis. Count data were analyzed using the Chi-square test.  $P < 0.05$  was considered to be statistically different.

## Results

There were 101 MDS patients included in the main study, and 19 were excluded here because of incomplete data. Thus, 82 MDS patients were included in this subanalysis. There were 26 men and 56 women who were enrolled into this study. Among the enrolled patients, 41 were enrolled into the ATP group and 41 were enrolled into the AT group. The average age of patients in the ATP and AT groups was  $53.66 \pm 10.86$  years and  $52.63 \pm 13.46$  years, respectively. The clinical pathology and biochemical data from patients in the ATP and AT groups were matched for parameters such as age, gender, neutrophils, hemoglobin, and platelets (Table 1).

### *ATP promotes hematological improvement in patients with MDS*

After 3 months of treatment, 62.64% (21 of 33) of patients showed a hematologic improvement (platelet response, HI-P) in

patients receiving ATP compared with 36.67% (11 of 30) of patients in the AT group. As shown in Figure 1, platelet and hemoglobin increased significantly after 3 months of ATP treatment compared with before treatment ( $n = 41$ ; platelets:  $73.24 \pm 54.54 \times 10^9/L$  vs.  $98.76 \pm 75.46 \times 10^9/L$ ,  $P = 0.014$ ; hemoglobin:  $122.07 \pm 24.55$  g/L vs.  $128.76 \pm 26.61$  g/L,  $P = 0.012$ ). In addition, platelet and hemoglobin levels were significantly higher in the ATP group after 5 months of treatment compared with before treatment ( $n = 41$ ; platelets:  $73.24 \pm 54.54 \times 10^9/L$  vs.  $103.9 \pm 67.53 \times 10^9/L$ ,  $P = 0.001$ ; hemoglobin:  $122.07 \pm 24.55$  g/L vs.  $134.07 \pm 28.48$  g/L,  $P < 0.001$ ) (Figure 1a and 1b). However, in the AT group, only hemoglobin showed a significant increase after treatment compared with before treatment ( $n = 41$ ; 3 months:  $117.29 \pm 25.79$  g/L vs.  $124.64 \pm 25.28$  g/L,  $P = 0.001$ ; or  $n = 41$ , 5 months:  $117.29 \pm 25.79$  g/L vs.  $125.85 \pm 26.38$  g/L,  $P = 0.004$ ) (Figure 1c).

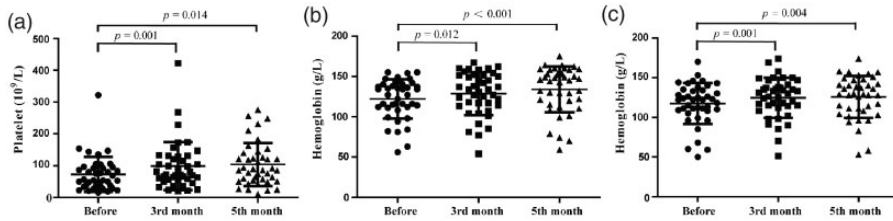
### *GPX3 demethylation induced by PND in patients with MDS was related to gender and could promote the recovery of PLT hematopoietic function*

As shown in Figure 2, we selected three CpG sites from the CpG island in the

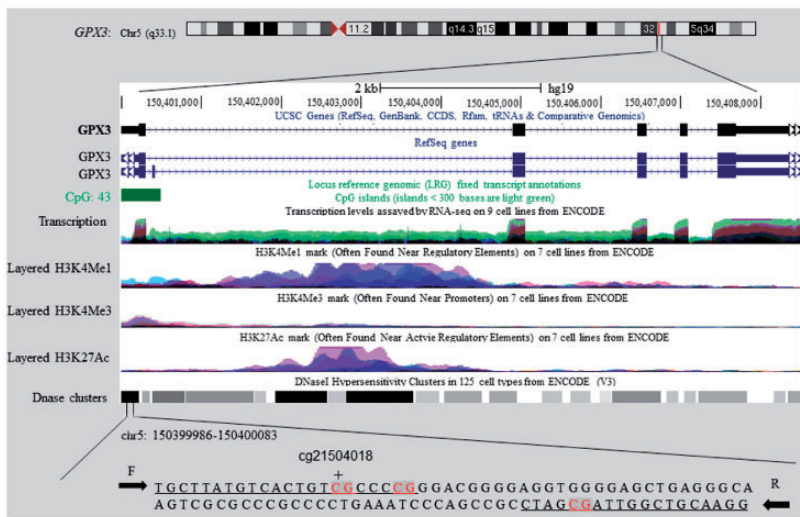
**Table 1.** Basic clinical characteristics of patients.

Variable	Treatment group (n = 41)	Control (n = 41)	p
Sex (n)			
Male	12	14	N.S.
Female	29	27	
Age (year)	$53.66 \pm 10.86$	$52.63 \pm 13.46$	N.S.
Baseline before treatment			
Neutrophils ( $\times 10^9/L$ )	$2.49 \pm 2.16$	$2.35 \pm 1.41$	N.S.
Hemoglobin (g/L)	$122.07 \pm 24.55$	$117.29 \pm 25.79$	N.S.
Platelets ( $\times 10^9/L$ )	$73.24 \pm 54.54$	$77.37 \pm 46.77$	N.S.
Blasts in bone marrow (%)	$0.62 \pm 0.813$	$0.73 \pm 0.832$	N.S.
Karyotype			
Normal	36	37	N.S.
Abnormal	5	4	

N.S., not significant.



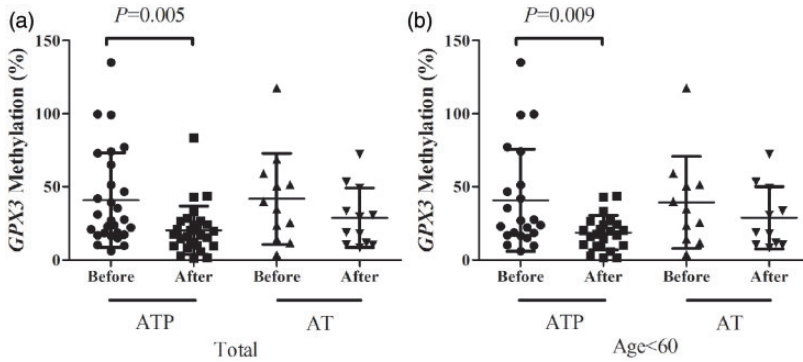
**Figure 1.** Routine blood test results for MDS patients changed significantly after 3 months and 5 months of treatment compared with before treatment in the ATP and AT groups. A. Change in the platelet level before and after treatment. B and C. Change in hemoglobin level before and after treatment in the two groups. MDS, myelodysplastic syndrome; ATP, group that received oral androl, thalidomide, and PND capsule; AT, group that received oral androl and thalidomide; PND, Pai-Neng-Da.



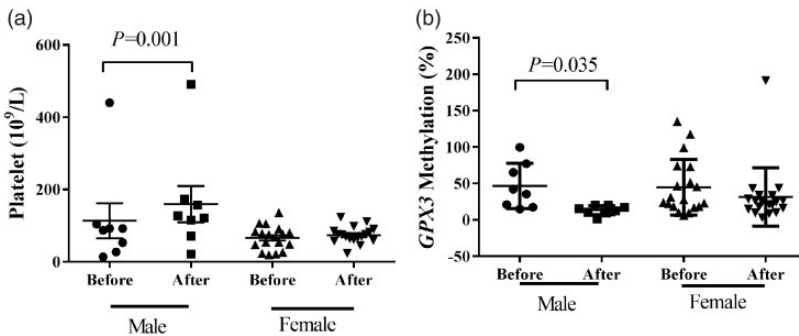
**Figure 2.** Drawing of the amplified fragment (chr5: 150399986-150400083) at *GPX3* CpG island that was obtained from the UCSC genome browser (GRCh37/hg19). The qMSP primers were underlined, and three CpG sites were in red. The targeted region was located at the CpG island of *GPX3*. F, forward primer; R, reverse primer; UCSC, University of California Santa Cruz; qMSP, quantitative methylation specific PCR; PCR, polymerase chain reaction.

*GPX3* promoter region to represent *GPX3* methylation.<sup>30</sup> We examined the bone marrow *GPX3* methylation level before and after treatment in 28 ATP-treated patients and 12 AT-treated patients. Our data confirmed that the *GPX3* methylation level was significantly reduced in MDS patients after 3 months of ATP treatment compared with before treatment (n = 28; 40.81% ± 32.36% vs. 22.31% ± 16.51%,

P = 0.005, Figure 3a). However, there was no difference in *GPX3* methylation in patients before or after AT treatment (n = 12; 41.76% ± 31.18% vs. 28.82% ± 20.33%, Figure 3a). In addition, in patients <60 years old, the *GPX3* methylation level was significantly reduced after 3 months of ATP treatment compared with before treatment (n = 22; 40.73% ± 34.93% versus 18.68% ± 11.74%, P = 0.009, Figure 3b).



**Figure 3.** Comparison of the difference in GPX3 methylation between the ATP and AT groups before and after treatment. a. After 3 months of treatment, GPX3 methylation was significantly reduced in the ATP group compared with before treatment, but there was no difference in the AT group; b. GPX3 methylation in patients who were younger than 60 years in the ATP and AT groups was investigated. GPX3 methylation was significantly lower in the ATP group after compared with before treatment. There was no difference in the AT group after 3 months of treatment compared with before treatment. Before, before 3 months of treatment; After, after 3 months of treatment; ATP, group that received oral androl, thalidomide, and PND capsule; AT, group that received oral androl and thalidomide; PND, Pai-Neng-Da.



**Figure 4.** In patients with low-risk MDS, GPX3 demethylation by ATP may be related to gender, and platelet elevation was more pronounced in male patients with stronger GPX3 demethylation. a. GPX3 demethylation was more pronounced in male MDS patients compared with female MDS patients after 3 months of ATP treatment. b. The platelet count increased significantly in low-risk male compared with low-risk female MDS patients with more significant GPX3 demethylation. Before, before 3 months of treatment; After, after 3 months of treatment; MDS, myelodysplastic syndrome; ATP, group that received oral androl, thalidomide, and PND capsule; PND, Pai-Neng-Da.

However, *GPX3* methylation level was not significantly different in patients receiving AT treatment before and after 3 months of treatment ( $n = 11$ ;  $39.32\% \pm 31.47\%$  vs.  $28.77\% \pm 21.32\%$ , Figure 3b).

In addition, the *GPX3* methylation level and platelet elevation after 3 months of

ATP treatment was different when compared by gender. In male patients, the *GPX3* methylation level was lower than that of female patients ( $n = 8$ ;  $46.58\% \pm 31.07\%$  vs.  $13.28\% \pm 6.23\%$ ,  $P = 0.035$ , Figure 4a). Conversely, platelet counts were significantly increased in male

compared with female MDS patients ( $n = 8$ ;  $113.58 \pm 135.98$  vs.  $159.5 \pm 142.23$ ,  $P = 0.001$ , Figure 4b). There was no notable change in the *GPX3* methylation level or in the platelet count in female MDS patients who were treated with ATP. Therefore, demethylation of ATP to *GPX3* may be associated with overall HI-P in MDS patients.

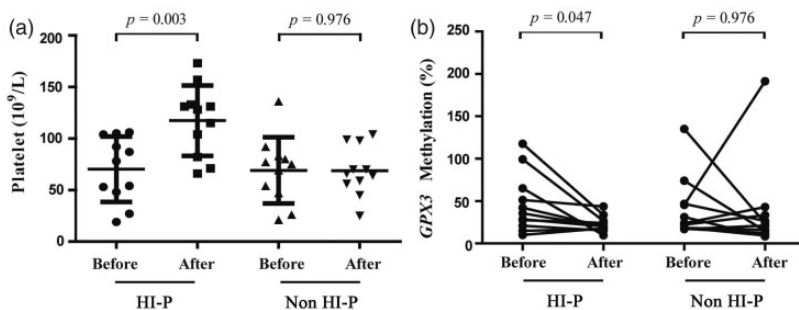
### The demethylation function of PND on *GPX3* might be related to the overall HI-P of patients with low-risk MDS

As described above, in the ATP group, *GPX3* methylation was significantly reduced in male patients, and platelet counts were significantly increased compared with female patients. To further investigate the relationship between demethylation of *GPX3* that is induced by PND and HI-P, we divided bone marrow specimens from the ATP group into a hematologic response group (HI-P,  $n = 11$ ) and a non-hematologic response group (non-HI-P,  $n = 11$ ). After 3 months of ATP treatment, platelet counts were significantly increased in HI-P patients compared with before treatment ( $n = 11$ ;  $70.27 \pm 31.61$

vs.  $117.36 \pm 34.09$ ,  $P = 0.003$ , Figure 5a) while non-HI-P patients showed no change ( $n = 11$ ). In addition, the *GPX3* methylation level was significantly lower in HI-P patients after compared with before ATP treatment ( $n = 11$ ;  $46.63\% \pm 34.61\%$  vs.  $22.58\% \pm 9.37\%$ ,  $P = 0.047$ ). However, there was no significant change in *GPX3* methylation in the non-HI-P group after compared with before ATP treatment ( $n = 11$ ;  $40.9\% \pm 35.91\%$  vs.  $36.44\% \pm 52.47\%$ , Figure 5b). Moreover, *GPX3* methylation was inversely correlated with changes in platelet count in patients with low-risk MDS ( $r = -0.352$ ,  $P = 0.048$ , Figure 6). Our results suggest that PND increases the platelet count and promotes hematopoiesis by decreasing the *GPX3* methylation level.

## Discussion

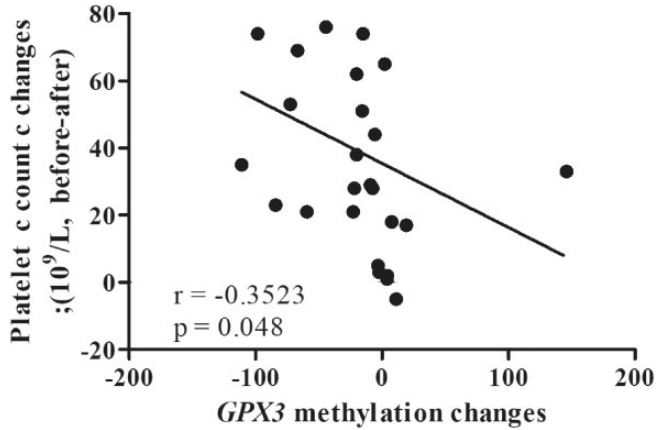
A series of preclinical studies of PND<sup>31–33</sup> were performed including pharmacodynamic studies in animal models with hemocytopenia, cell biology, molecular biology experiments, and toxicological studies to test the biologically active panaxadiol saponins component. These studies demonstrated



**Figure 5.** Effect of the change in *GPX3* methylation on the hematological response to low-risk MDS patients with ATP treatment. A. Compared with the Non HI-P group, platelet counts of patients in the HI-P group increased significantly after treatment. B. *GPX3* methylation decreased significantly in patients with a hematological improvement (HI-P), and those without hematological improvement (Non HI-P) had the negative results after treatment compared with before treatment.

Before, before 3 months of treatment; After, after 3 months of treatment; MDS, myelodysplastic syndrome; ATP, ATP group that received oral androl, thalidomide, and PND capsule; PND, Pai-Neng-Dal HI-P, hematological improvement in platelet response.





**Figure 6.** Correlation between the changes in *GPX3* methylation and PLT counts before and after ATP treatment. PLT, platelet; ATP, ATP group that received oral androl, thalidomide, and PND capsule; PND, Pai-Neng-Da.

that PND possessed dual activity, i.e., promoting proliferation and differentiation of hematopoietic progenitor cells and regulating the immune function. PND reduced peripheral blood CD8<sup>+</sup> cells, and increased CD4<sup>+</sup> cells, returning the CD4<sup>+</sup>-to-CD8<sup>+</sup> cell ratio to normal. It reduced the myelosuppression of bone marrow that was caused by chemotherapy, accelerated bone marrow recovery, and increased the peripheral WBC and platelet count, as shown by mice treated with hemocytopenia that was caused by cyclophosphamide chemotherapy. PND was effective in promoting proliferation for the three lineages of hematopoietic progenitor cells, increasing erythroid, myeloid, and megakaryocytic progenitor cell colony formation in bone marrow culture. A pharmacodynamic study in CD34<sup>+</sup> hematopoietic stem/progenitor cells suggested that PND was effective in promoting hematopoiesis and inducing differentiation, which suggests that its activity is similar to that of hematopoietic growth factors.<sup>34</sup> PND promotes hematopoiesis using an intracellular signal pathway, and it induced the up-regulation of multiple transcription factors such as GATA-1, GATA-2, c-Fos, c-Jun, and

NF-E2, and their protein expression level, phosphorylation status, and DNA binding activities were increased dramatically by PND treatment in hematopoietic cells, thereby promoting hematopoiesis and blood cell formation.<sup>26,35–37</sup> However, in this study, PND improved hematopoietic function in low-risk MDS patients by promoting *GPX3* demethylation.

In this study, after 3 months of treatment with ATP, the platelet count was significantly elevated in patients with low-risk MDS, and patients achieving HI-P. However, the *GPX3* methylation level was significantly reduced in the ATP group patients and in HI-P patients, and the *GPX3* methylation level was significantly lower in male patients in the ATP group after compared with before treatment. The platelet counts were also significantly increased in male compared with female patients. These findings suggest that *GPX3* demethylation of PND may be associated with improving the hematopoietic function. In patients with low-risk MDS, there may be an inverse correlation between platelet hematology improvement and *GPX3* methylation levels.

In our study, the hemoglobin level was significantly increased in both groups. After 3 months of treatment, 62.64% of ATP patients and 36.67% of AT patients achieved HI-P, indicating that PND as an auxiliary drug provided a better clinical outcome in low-risk MDS. Thrombocytopenia is common in patients with MDS and the estimated prevalence is 40% to 65%.<sup>38</sup> The average platelet component mean platelet volume is a useful screening marker for MDS.<sup>39</sup> MDS patients with thrombocytopenia have a poor prognosis.<sup>40</sup> Bleeding complications are a major cause of morbidity and mortality, and thrombocytopenia is also an independent factor in reducing survival. This has been included in the updated prognostic scoring system.<sup>41</sup> Platelet transfusion is a necessary support for clinically significant thrombocytopenia.<sup>42,43</sup> MDS patients with severe thrombocytopenia may benefit from the recovery of platelet activity and further allo-hematopoietic stem cell transplantation (HSCT) after decitabine treatment.<sup>44</sup> In this study, the platelet count was significantly increased after 3 months of treatment in the ATP group compared with before treatment, indicating that PND capsule has a positive effect on enhancing hematopoietic function and improving symptoms in low-risk MDS patients.

Promoter DNA hypermethylation and down-regulation of the *GPX3* gene occur in a variety of solid tumors.<sup>30,45-47</sup> The *GPX3* promoter is methylated in chronic myelogenous leukemia, and *GPX3* hypermethylation is an independent prognostic biomarker in non-M3 acute myeloid leukemia.<sup>23</sup> The preliminary study also reported that *GPX3* methylation in the bone marrow was associated with poor prognosis of MDS and leukemia transformation.<sup>15</sup> *GPX3* may be a predictor of NKTCL and NESM response chemotherapy.<sup>48</sup> Additionally, *GPX3* hypermethylation in gastric cancer (GC) predicted a shorter

tumor recurrence time in patients aged >60 years.<sup>30</sup> In the current study, we observed a significant decrease in *GPX3* methylation levels in MDS patients after a 3-month ATP regimen compared with before treatment. Similarly, the methylation level of patients in the ATP group was significantly lower in patients under, compared with over, 60 years of age.<sup>48</sup> We also found a significant decrease in the *GPX3* methylation level in MDS patients who received HI-P after a 3-month ATP regimen compared with before treatment, but we found no significant difference of *GPX3* methylation in the non-HI-P MDS patients before or after treatment. Additionally, after a 3-month ATP regimen, the *GPX3* methylation level in male patients was lower than that of female patients. HI-P or an increase in the platelet count in male MDS patients was more pronounced than that in female MDS patients. Because of the increased ROS production that was associated with platelet activation and apoptosis, *GPX3* can reduce platelet apoptosis while enhancing their activity as a ROS scavenger to increase the platelet count.<sup>49</sup> However, a few results of the statistical analysis were not meaningful owing to the moderate sample size. We have reason to hypothesize that *GPX3* expression in low-risk MDS patients is regulated by epigenetic regulation of DNA methylation and that PND may reduce the *GPX3* methylation level, leading to up-regulation of *GPX3* expression, which thereby increases the platelet count and improves hematopoietic function.

In this study, we enrolled 82 patients who had low-risk MDS, which did not reach the expected number of cases, and this may affect the statistical significance of the results. However, in conclusion, the study shows, to a certain extent, that PND capsule is able to increase the platelet count to obtain HI-P in low-risk MDS patients, which showed better results than the group

without PND. Decreasing the *GPX3* methylation level may be involved in the hematological improvement process, and the effect of PND on the *GPX3* methylation level provides clues to the drug action mechanism of PND capsules.

### Declaration of conflicting interest

The authors declare that there is no conflict of interest.

### Funding

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