Research Report

Evaluation of the Effects of Opium on the Expression of *SOX2* and *OCT4* in Wistar Rat Bladder

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Abstract.

BACKGROUND: Bladder cancer is a malignancy greatly affected by behavioral habits. The aim of this study was to examine the effect of opium on changes in the expression of *OCT4* and *SOX2* in the bladder tissue of rats.

METHOD: Thirty six rats were divided into six groups: 24 rats in the addicted group received morphine and opium for 4 months with 12 rats in the control group. Blood testing was done for the evaluation of CBC, MDA, and TAC. The bladder tissue was removed and checked by histopathological examination. All total RNA was extracted, then cDNAs were synthesized and the OCT4 and *SOX2* gene expressions were evaluated by Real-time PCR.

RESULTS: The *OCT4* mRNA expression level in the opium group of rats was significantly increased compared to the control group (13.5 and 6.8 fold in males and females respectively). Also, in the morphine group, similar augmentation was detected (3.8 and 6.7 fold in males and females respectively). The *SOX2* mRNA over-expression level was seen in the morphine group of both genders as compared to the control group (3.7 and 4.2 fold in male and female respectively) but in the opium group, enhancement of mRNA level was seen only in males (6.6 fold). Opium increases both *OCT4* and *SOX2* expression more than morphine in male rats, but in female rats, *SOX2* is increased more by morphine.

CONCLUSION: Over expression of *OCT4* and *SOX2* was observed in rats treated with opium and morphine. Increased *OCT4* and *SOX2* expression was seen in opium-treated male rats, but in female rats, *SOX2* was increased more by morphine.

Keywords: Opium, morphine, SOX2, OCT4, bladder

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Abbreviations

BAX	Bcl-2-associated X protein
BCl2	B-cell lymphoma 2
CBC	complete blood count
cDNAs	complementary deoxyribonucleic acid
H & E	hematoxylin and eosin

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HCT	hematocrit
Hgb	hemoglobin
MCHC	mean corpuscular hemoglobin concentration
MDA	malondialdehyde
OCT4	octamer-binding transcription factor 4
PLT	platelets
RBC	red blood cell
RNA	ribonucleic acid
SOX2	SRY-Box Transcription Factor 2
TAC	total antioxidant capacity
WDC	1 1 1 1 .

WBC white blood count

INTRODUCTION

Bladder cancer is the most prevalent genitourinarysystem malignancy, and plays a primary role in more than 7% of neoplasms, especially among men [1]. Yearly reports show a further 12 million cases of bladder cancer, putting it in the eleven top-ranked cancers, by incidence, in the world [2] It is the third most common cancer in Iranian men [3] and it occupies the ninth place in prevalence among Iranian women [4]. The risk factors which can lead to bladder cancer include high body mass index [5], genetic factors [6], occupational hazards [7, 8], environment and nutrition [9], and behavioral factors [8, 10-12]. In previous studies, behavioral factors such as smoking and opium addiction were counted as the most important risk factors for several cancers, including bladder cancer [13-15]. Several new treatment strategies are suggested for bladder cancer [16–19]. Meta-analysis agrees that opium consumption is potentially responsible for the development of bladder cancer [20]. The prevalence of opioid consumption ranges from 0.1% to 2% globally [21], and a high prevalence of opioid use has been correlated to Iranians for several hundred years [22]. Although the use of opium is prohibited in Iran, it is the most commonly used drug, and up to 14% of the Iranian population are regular users of opium [23]. Opioids such as morphine, heroin, and opium are the most common form of drugs used in Iran. They can have a negative impact on the immune system [24–26], and have been reported to augment the susceptibility of animals to bacterial and viral infections [27]. They also cause increased metastasis and - as this leads to apoptosis - cell death, and can decrease survival in tumor-bearing animals [28]. Research has also shown apoptosis in the brain and liver of opium-addicted rats [29].

Uncontrolled self-renewal is seen as an essential mechanism in carcinogenesis [30]. By referring to

the cancer stem cell (CSC) hypothesis, a batch of cancer cells with stem-cell-like features can sustain a tumor; these batches should be self-renewing and pluripotent [31-33]. The CSCs have an indistinct proliferation, and can also lead to tumorigenesis [34]. The hypothesis is that the single pathway that manages self-renewal in normal stem cells is also employed to manage CSCs in tumors [35]. It is also believed that a member of the POU family, called Octamer 3/4 (Oct 3/4), is a crucial stem-cell marker and an essential transcription factor through human embryogenesis [36]. Recent reports have contained information regarding the presence of Oct 3/4 among differentiated benign and malignant human cells [37]. Sox2 is responsible for encoding transcription factors with a single HMG DNA-binding domain. Another function of this member of the SOX (SRY-Related HMG box) gene family is controlling neural progenitor cells by prohibiting their ability to differentiate [38]. Sox2 expression is also present in various malignant tissues [39].

This study aims to examine the effect of opium on changes in the expression of *OCT4* and *SOX2*, the main controlling genes of the self-renewal pathway, in the bladder tissue of rats.

MATERIALS AND METHODS

This research was carried out in the Animal Laboratory of the Urology Research Center of Tehran Medical University, Tehran, Iran, between October 2021 and September 2022, and approved by the Ethics Research Committee of Tehran University of Medical Sciences (IR.TUMS.SINAHOSPITAL.REC.1399.026). Ethical principles of working with rats were considered in all phases of the research.

Animal care

In this study, 38 4-week old male and female adult Wistar rats weighing between 200 and 220 g each, were purchased from the Pharmacology College of Tehran University, Iran. Four rats were placed in each Plexiglas cage, and given *ad libitum* access to food and water. The temperature of the laboratory was set at $22 \pm 2^{\circ}$ C, relative humidity was $45.00 \pm 2.00\%$ and the light/dark cycle was 12/12 h. Before commencing the experiments, the rats were allowed to adapt to the lab's environment for seven days. Maintenance of all animals was in accordance with institutional guidelines for animal care and use and all the experimental protocols were approved by the National Institutes of Health guidance for the care and use of laboratory animals [40].

Exposure of rats to morphine sulfate and powdered opium

In Iran, opium is generally used in one of three different forms; Teriak (crude opium), Sukhteh (remnants of smoked opium or dross), and Shireh (opium juice, an opium product usually made by boiling Teriak or Sukhteh in water, filtering the mixture several times and then evaporating the filtrate) [41].

Morphine sulfate and opium powder (Teriak) (Temad Manufactory, Tehran, Iran), and Naloxone hydrochloride (Tolid Daru company, Tehran, Iran) were used in this study. The drugs were dissolved in sterile 0.9% saline and injected intraperitoneally (i.p.) at a volume of 1 ml/kg.

Two rats were dissected at the beginning of the study. The 36 remaining rats were divided into six groups: 12 rats in the control group (6 females and 6 males) received a single i.p. injection of saline for four months and were treated in the same conditions as the other groups. Rats in the addicted group (6 males and 6 females in each group of morphine and opium) received an initial dose of morphine (2 mg/kg) and opium (2 mg/kg).The morphine dose was then increased every fifteen days with a 4, 8, 10, 15, 20, 25 & 40 mg/kg dose respectively, and the opium dose every 30 days with 10, 20, 40 mg/kg doses respectively, via i.p. injection (a total of 120 days) [40, 42, 43].

To confirm the addiction of the rats, one month after the first injection of morphine and opium, 2 rats in each group received Naloxone randomly (2.5 mg/kg, to confirm addiction) [40, 44] and were placed in a transparent glass box to observe any withdrawal signs such as diarrhea, genital licking, wet-dog shakes, yawing, and sweeping tail movements.

Investigation of serum parameters

Blood samples (5 ml) were taken from rats exposed to opium and morphine from each group before starting the study and immediately after the completion of the project. Nihon Kohden MEK-1305 Celltac α + cell counter device (automated hematology and Erythrocyte sedimentation rate)ESR(analyzer was used to conduct a complete blood count (CBC) examination. Malondialdehyde (MDA) content was determined by colorimetric method, using a kit purchased from Eagle Biosciences, Inc., USA, according to the method of Satoh [45]. Total Antioxidant Capacity (TAC) was determined using the ELISA technique, using a kit purchased from Cell Biolabs, Inc, San Diego, USA, according to Campbell et al. [46].

Bladder pathology examination

Bladder tissue was removed from the Wistar rats and placed in 10% neutral-buffered formalin. After bladder tissue fixation, the bladder specimens were sent to the comparative histopathology laboratory in the pathology department of the Tehran Medical University, Urology Research Center. According to the standard method, paraffin blocks were processed after molding, and five-micron microscopic incisions from diverse sections were prepared and stained according to the H & E (combination of two histological stains containing hematoxylin and eosin) standard protocol. Observations were performed by Olympus light microscope with \times 100, \times 200 and \times 400 magnifications.

RNA isolation and real-time PCR

All total Ribonucleic acid (RNA) of Wistar rat Bladder tissues was extracted using a High Pure RNA Isolation Kit (Cat. No. 11 828 665 001) and then synthesizing the complementary Deoxyribonucleic acid (cDNAs) (Cat. #RR014A/B). The relative expression of genes was then evaluated using the $2^{-\Delta\Delta CT}$ method used for calculating relative expression. The primers used for the Rotor gene real-time PCR cycler of OCT4-F mRNA were as follows: forward primer, CCGTGTGAG-GTGGAACCTG and reverse primer: CGGTTACA-GAACCACACTCG, SOX2 mRNA were as follows: forward primer, CGAGTAGGACATGCTGTAGG and reverse primer: ACATGAACGGCTGGAG-CAAC. The forward primer sequence for GAPDH mRNA quantification as housekeeping gene was GGCAAGTTCAACGGCACAG, the reverse primer sequence was CGCCAGTAGACTCCACGAC. The performance of Triplicate qPCR reactions was applied to the samples one by one. We describe the actual time of qPCR condition as 95°C for 10 min in the role of the pre-denature step, after that 40 PCR cycles at 95°C for 30s, 60°C for 30s, and 72°C for 30 s. Melting curve analysis was performed, to measure the specificity of the real-time RT-PCR.

Statistical analysis

All experiments were performed three times and showed similar results. At the end of the study, the data was achieved from the graph pad prism 9. Statistical significance was defined at *P < 0.05, **P < 0.01, and ***P < 0.001. one way, two way ANOVA, and *T* test were performed to evaluate the significance of differences.

RESULTS

Expression of the OCT4 and SOX2 genes

The different mRNA expression levels of the main controlling genes of the self-renewal pathway (*OCT4* and *SOX2*) before and after the administration of opium and morphine were detected by RT-PCR.

As shown in Fig. 1, the OCT4 mRNA expression level in opium-group rats was significantly increased compared to the control group (13.5 and 6.8 fold in males and females respectively). Similar augmentation was also detected in the morphine group (3.8 and 6.7 fold in males and females respectively).

The relative mRNA expression levels of the *OCT4* gene were also evaluated by real-time PCR. As shown in Fig. 2, the mRNA expression of *OCT4* is detectable for all of the groups studied. Although the mRNA quantities are changing, a comparison of the relative gene-expression levels indicated significant differences among the studied groups (p < 0.05) except in males of the morphine group (p = 0.0596).

Increase of the SOX2 mRNA expression level was seen in the morphine group of rats of both genders as compared to the control group (3.7 and 4.2 fold in male and female respectively) but in the opium group, enhancement of mRNA level was seen only in males (6.6 fold) (Fig. 3).

The relative mRNA expression levels of the *SOX2* gene are shown in Fig. 4. As shown in Fig. 4, the mRNA expression of *SOX2* is detectable for all of the studied groups. A comparison of the relative gene-expression levels indicated significant differences among the morphine groups. (p < 0.05).

Figure 5 demonstrates that opium increases *OCT4* and *SOX2* expression more than morphine in male rats but in female rats, *SOX2* is increased more by morphine.

Histopathological findings

In order to confirm our results, various tissues such as the kidney and liver were harvested from the Wistar rats and analyzed with Hematoxylin and Eosin (H&E) staining. We required a fixative in which to soak the tissue sections, and we chose 10% neutralbuffered formalin for this purpose. Tissues were fixed, mounted in paraffin wax, and the slides were prepared before staining with H&E. Staining was carried out on all tissues. A light microscope was then utilized for monitoring the slides in a pathology laboratory, slide observation was also performed using an OPTIKA, C-P2504 microscope.

The results of staining on bladder tissue showed urothelial atrophy in the morphine group (star in Fig. 6) and urothelial cell hyperplasia in the opium group (Arrowhead at Fig. 6). Also, hypertrophy of smooth muscle of the vascular wall and hyperemia were seen in the morphine group as compared with those in the control group (arrows in Fig. 6). Pathol-

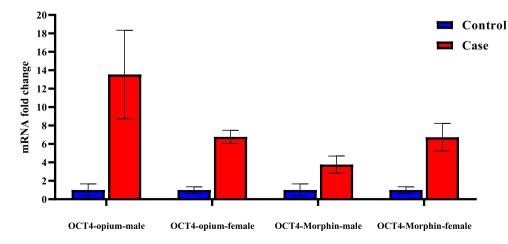


Fig. 1. OCT4 mRNA fold changes were significantly increased in all groups compared to the control group.

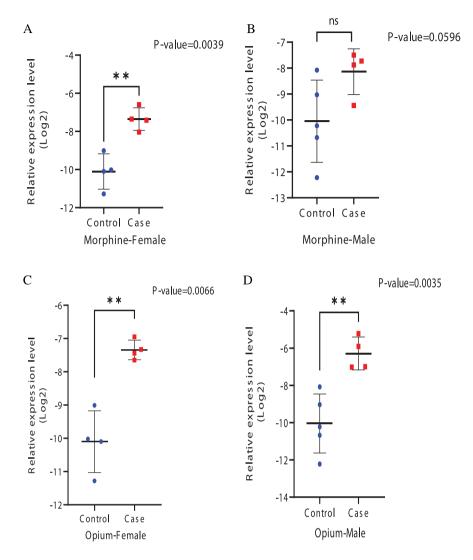


Fig. 2. Relative expression levels of *OCT4* gene in different groups A) Morphine-Female. B) Morphine-Male. C) Opium-Female. D) Opium-male. There was no significant difference in males of the morphine group, but a noticeable difference was observed in other groups.

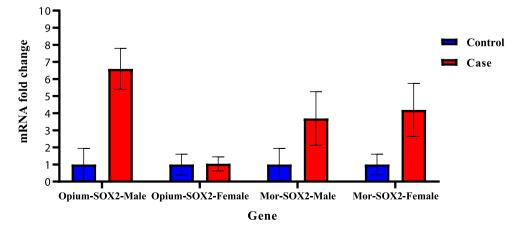


Fig. 3. SOX2 mRNA fold change was increased in all groups except females in the opium group.

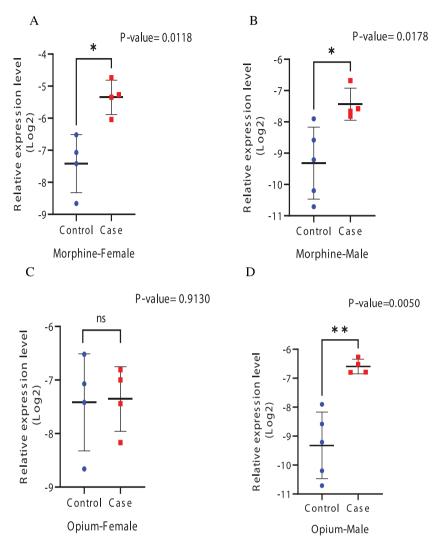


Fig. 4. Relative expression levels of *SOX2* gene in case and control groups A) Morphine-Female. B) Morphine-Male. C) Opium-Female. D) Opium-male.

ogy findings for urinary bladder of all rats are also illustrated in Supplementary Table 1.

Serological findings

The mean values of the serological parameters for rats from each of the six groups were evaluated (Table 1). Values included are: CBC, total antioxidant capacity (TAC), and Malondialdehyde (MDA). A comparison was made of these parameters in the various groups (Fig. 7).

As regards male rats, observations indicated that opium treatment led to a reduction in white blood cells (WBC), hemoglobin (Hgb), hematocrit (HCT), and platelets (PLT), Increases in red blood cells (RBC), mean cell hemoglobin concentration (MCHC), MDA, and TAC were observed as compared to the control group. However, opium treatment caused a decrease in HCT and MDA levels and an increase in TAC levels in female rats as compared to the control group.

By comparing opium and morphine in detail, it was established that opium raises RBC and MCHC and causes a decline in Hgb, HCT, PLT, and MDA more than morphine in male rats, but that in female rats, only Hgb, HCT, and MDA are decreased more by opium.

DISCUSSION

Bladder cancer is the most prevalent malignancy of the genitourinary system [1]. Behavioral factors

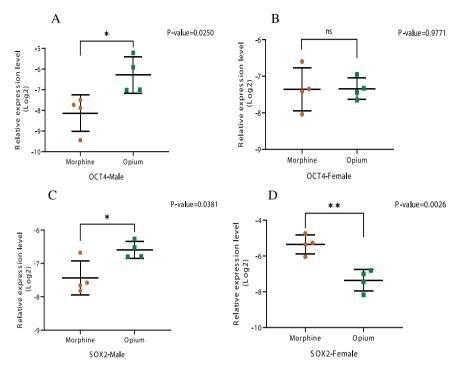


Fig. 5. relative expression levels of genes between opium and morphine A) OCT4-Male B) OCT4-Female C) SOX2-Male D) SOX2-Female.

such as smoking and opium addiction are counted as the most important risk factors for several cancers, including bladder cancer [13, 14]. The prevalence of opioid consumption ranges from 0.1% to 2% of population globally [21]. In the Middle East and Asia, illicit opioid use exists across a spectrum between heroin and opium. Among 204 opioid-use disorder patients, in 56 (29.3%) opium was the primary substance of choice [47]. Based on a national survey conducted among Iranian workers, opium proved to be the most popular substance, and 2.7% of workers were using it [48], and it is estimated that about 20% of the adult population of Iran consumes it [49]. In recent years, there have also been reports of the presence of OCT4 and SOX2 expression in a number of malignant tissues. In this study, we found that morphine and opium significantly increased OCT4 expression in both male and female rats. SOX2 was expressed more in morphine cases as compared to the control group, but in the opium group, this increase was only observed in male rats. In male rats, opium increases OCT4 and SOX2 expression more than morphine. In female rats however, morphine increases SOX2 more than opium.

There are other studies that confirm our results. An increase observed in the mRNA levels of *OCT4*, *SOX2*, and Nanog in both MCF-7 and BT549 cells was caused by morphine, as discussed by Niu et al..

They reported that the mRNA levels of OCT4, SOX2, and Nanog were enhanced as compared to untreated controls by:13.08, 10.57, and 19.18 fold in MCF-7 cells, while the initial values reported were: 6.15, 10.37, and 14.92 fold in BT549 cells. We observed a similar effect; the growth in OCT4 expression was reported as 3.8 and 6.7 fold in males and females, and SOX2 expression was increased by 3.7 and 4.2 fold in both males and females in our study. Increases in OCT4 and SOX2 expression were greater in the Niu study as compared to our study. They also confirmed their findings by western blot, showing that the increase in protein levels of OCT4, SOX2, and Nanog in MCF-7 and BT549 cells is dependent on morphine dose, so they recommend that tuning OCT4, SOX2, and Nanog by means of morphine may promote cancer stem cell properties [50].

SOX2, *OCT4*, and Nanog are stem-cell markers that develop cancer and increase drug resistance [51]. It is believed that stem-cell markers are significantly increased by morphine, so reports agree that the enrichment of stem-cell properties and drug resistance may be caused by long-time morphine use [50].

Gonzalez et al. also agree with these results. In their report, it was observed that *SOX2*, *OCT* 4, and Nanog were highly expressed in zebra-fish embryos treated with morphine, as qPCR indicated [52].

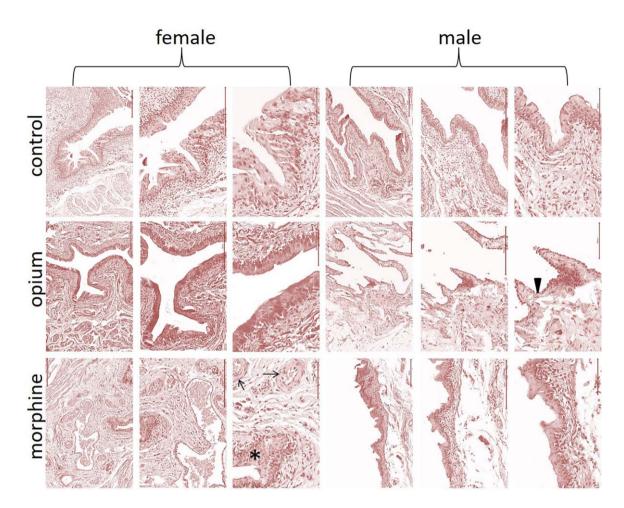


Fig. 6. Histopathological sections of urinary bladder. Female and male rats, H&E staining. Arrows indicate the urothelial atrophy and star indicates the urothelial cell hyperplasia. Arrows indicate hypertrophy of smooth muscle of vascular wall and hyperemia.

SOX2 and OCT4 are crucial for the maintenance of neural stem cells, and expression of these transcription factors is associated with p300 [53, 54]. In transcriptional complexes, single participants can induce the expression and activation of others, and these complexes are performed by the essential transcription factors SOX2 and OCT4 [55]. Regulation of SOX2 is associated with the expression of the other two transcription factors. They suggest that morphine effectively promotes numerous mechanisms whereby the members of this transcriptional complex are up-regulated, and this state was concluded from observations. The observations showed that during several stages of zebra-fish development, morphine treatment exerted a progressive up-regulatory effect on the mRNA levels of these genes. SOX2 and OCT4 can be effectively regulated with the increase of p300, induced by morphine [54].

Various mechanisms explain the variation of transcription-factor expression caused by morphine, and the effect and feedback regulation established between the transcription factors themselves can be part of the mechanisms mentioned (Table 2). The over expression of the pluripotency transcription factors *SOX2/OCT4*/Nanog and p300 is one of the reactions for which morphine facilitates the increase, as Gonzalez and their co-workers reported. They believe that there are several mechanisms for the morphine effect [52].

There are also a number of studies discussing the effect of synthetic opioids. Fentanyl is a very potent synthetic opioid, used as pain-relief medication and in anesthesia. Zhang et al. observe that fentanyl wasn't significantly effective in a decrease of the expression of *OCT4* and *SOX2*, though there was an occasional decrease. In comparison, the expres-

Variables		Control	Morphine	Opium	<i>p</i> -value	Post-hoc <i>p</i> -values		
			•	*		CvsM	CvsO	MvsO
WBC	Female	5.72 (0.42)	6.86 (1.82)	5.54 (0.37)	0.162	0.379	1	0.244
	Male	10.22 (0.80)	7.00 (1.63)	7.70 (0.10)	0.001	0.001	0.008	0.940
RBC	Female	7.48 (0.78)	7.78 (0.80)	6.78 (0.36)	0.096	1	0.386	0.114
	Male	8.76 (0.55)	8.18 (0.40)	9.60 (0.25)	0.001	0.149	0.025	0.001
HGB	Female	13.86 (0.82)	14.6 (0.55)	12.92 (0.44)	0.004	0.253	0.102	0.003
	Male	14.8 (1.09)	14.7 (0.47)	10.10 (0.16)	< 0.001	1	< 0.001	< 0.001
HCT	Female	37.50 (1.64)	39.80 (0.96)	34.88 (0.18)	< 0.001	0.019	0.008	< 0.001
	Male	40.88 (0.76)	39.86 (0.86)	27.54 (0.42)	< 0.001	0.124	< 0.001	< 0.001
MCV	Female	50.02 (3.08)	51.02 (1.06)	51.60 (0.94)	0.459	1	0.679	1
	Male	46.70 (7.79)	48.66 (1.72)	47.62 (1.62)	0.808	1	1	1
МСН	Female	18.48 (0.50)	17.94 (1.00)	19.1 (0.89)	0.128	0.969	0.779	0.141
	Male	16.90 (0.89)	17.98 (0.71)	17.6 (0.35)	0.079	0.088	0.404	1
MCHC	Female	36.90 (1.31)	36.92 (0.72)	37.00 (1.01)	0.987	1	1	1
	Male	36.30 (0.94)	36.94 (0.64)	45.70 (0.46)	< 0.001	0.537	< 0.001	< 0.001
PLT	Female	692.00 (31.91)	715.50 (58.23)	737.00 (61.03)	0.42	1	0.591	1
	Male	920.00 (18.50)	715.50 (35.42)	510.56 (0.53)	< 0.001	< 0.001	< 0.001	< 0.001
RDW	Female	10.50 (0.87)	11.60 (0.89)	11.02 (0.04)	0.092	0.097	0.826	0.679
	Male	12.20 (0.45)	12.84 (1.32)	11.96 (0.21)	0.250	0.714	1	0.340
TAC	Female	0.98 (0.28)	2.06 (0.68)	2.09 (0.49)	0.007	0.018	0.015	1
	Male	1.20 (0.45)	2.06 (0.58)	2.40 (0.23)	0.003	0.029	0.003	0.746
MDA	Female	8.54 (0.54)	9.14 (0.48)	5.96 (0.37)	< 0.001	0.195	< 0.001	< 0.001
	Male	6.06 (0.13)	9.34 (0.29)	8.21 (0.40)	< 0.001	< 0.001	< 0.001	< 0.001

Table 1 Mean values (SD) of CBC, TAC, and MDA of rats from each group

p-values from ANOVA, CvsM = control versus morphine; CvsO = control versus opium; MvsO = morphine versus opium.

sion of OCT4 and SOX2 remained unchanged in the remifentanil treatment groups [56]. Furthermore, Kocak and colleagues agree with the results, and also showed that the expression of BAX, BCl2, OCT4, SOX2, and Nanog genes in MCF7 cancer stem cells can be decreased with fentanyl treatment. All these results confirm the statement that fentanyl decreases the expression of stem-cell markers. Fentanyl caused a decrease in SOX2, Nanog, and Bax genes, however, there was an increase in the OCT4 gene, in HEK293 stem cells [57]. Acceleration of cell numbers in the G1 phase and the decrease of OCT4 and SOX2 gene expression can be caused by synthetic opioids such as fentanyl, as Zhang indicated. So, the observations support the idea that regulation of the cell cycle and maintenance of mESCs can be performed by fentanyl, which leads to its contribution to proliferation and differentiation. Clues extracted from this evidence are useful in rationalizing the mechanisms that explain the effects of fentanyl on mESCs [56].

Dynorphin-B is a natural agonist of the κ -opioid receptor that decreases the expression of pluripotency markers like *OCT4* in mESCs. This is another similar example of synthetic opioids, which confirms the previous paragraph [58].

Results from a couple of other studies do, however, contradict this. Yang et al. concluded that the expression of Nanog, *SOX2*, and *OCT4* genes detected by qPCR is increased by fentanyl. Western blot also confirmed these results. Utilizing immunohistochemistry led to the detection of the expression of *OCT4* in mouse xenograft tumor tissues and this expression of *OCT4* was observed to be significantly greater in the fentanyl treatment group as compared to the control group [59].

Further experimental (*in-vivo* and *in-vitro*) studies have utilized opium. A study performed on 70 Wister rats showed that consumption of high doses of opium tincture may lead to hepatotoxicity [60]. Opium decreased the amplitude of duodenal and ileum con-

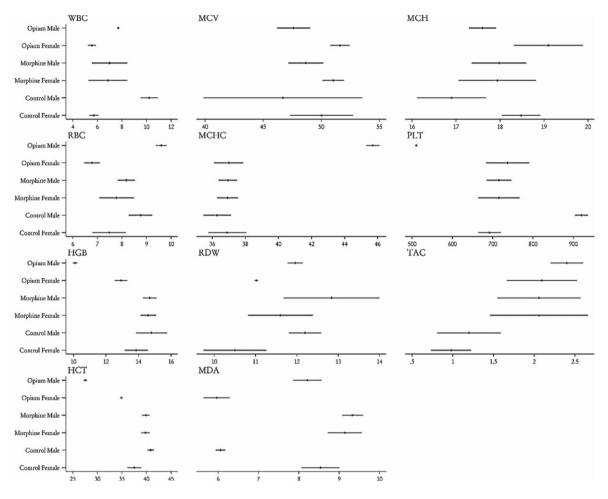


Fig. 7. Comparison of serological parameters in various groups; mean (95% CI).

Table 2					
Comparing the expression of OCT4, SOX2, and Nanog, affected by the various opioids					

Author	Opioid	Sample	Effect on OCT4 expression	Effect on SOX2 expression	Effect on Nanog expression
Our study	Morphine	Rat bladder	Increase	Increase	Wasn't evaluated
Our study	Opium	Rat bladder	Increase	Increase	Wasn't evaluated
Niu et al. [50]	Morphine	MCF-7 and BT549 cells	Increase	Increase	Increase
Gonzalez et al. [52]	Morphine	Zebra fish embryos	Increase	Increase	Increase
Zhang et al. [56]	Fentanyl	mESCs	Decrease	Decrease	Wasn't evaluated
Zhang et al. [56]	Remifentanil	mESCs	Unchanged	Unchanged	Wasn't evaluated
Kocak et al. [57]	Fentanyl	MCF7 cancer stem cells	Decrease	Decrease	Decrease
Šínová et al. [58]	Dynorphin-B	mESCs	Decrease	Wasn't evaluated	Wasn't evaluated
Yang et al. [59]	Fentanyl	Mouse xenograft tumor tissue	Increase	Increase	Increase

tractions, but increased the frequency of duodenal and mid-colon contractions. In another study, 66 male mice who received incrementally increased doses of opium [61]. Other experiments have also studied the effects of opium *in-vivo* and *in-vitro* [62–64].

The effects of opium and morphine on Nanog expression were not assessed in our study. Evaluating

the effect of opium on Nanog expression is suggested for future studies. Also, Due to the restrictions related to keeping rats, it was not possible to continue the study for a longer period. There is, therefore, a possibility that if the study were conducted over a longer time, we would observe the occurrence of cancer cells.

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CONCLUSION

We approached the direct effects of morphine and opium on *OCT4* and *SOX2*. In detail, an increase in *OCT4* and *SOX2* expression in rats treated with opium and morphine was confirmed by our results. Opium increased *OCT4* and *SOX2* expression more than morphine in male rats, but in female rats, *SOX2* was increased more by morphine. However, more clinical studies are required to extend our observations and explore the potential mechanism of the effects of opium and morphine on the expression of *SOX2* and *OCT4*.

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The authors report no funding.

AUTHOR CONTRIBUTIONS

SMKA is the principal investigator, IMO wrote the manuscript, LZ, PZ, GM, MN, and

AM collected data, AKH analyzed the data, and RM ran molecular testing.

CONFLICTS OF INTEREST

IMO, LZ, RM, PZ, AM, AK, MN, GM and SMKA have no conflict of interest to report.

DATA AVAILABILITY

All information, data, and photos are provided through the manuscript and additional will be provided if requested.

SUPPLEMENTARY MATERIAL

The supplementary material is available in the electronic version of this article: https://dx.doi.org/ 10.3233/BLC-230076.

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