Research Article

Investigation of the Disparities in Ultrasound Imaging Features of miR-323, miR-409-3p, and VEGF Expression Scales in Different Clinicopathological Features of Prostate Carcinoma and Their Correlation with Prognosis

Bao Liu, Jingqi Wang, Yanhua Cui, and Hui He

1Shanxi Medical University, Taiyuan 030001, China
2Department of Urology, The Second Hospital of Shanxi Medical University, Taiyuan 030001, China
3Department of Ultrasound, Shanxi Provincial People’s Hospital, Taiyuan 030001, China
4Emergency Ultrasound Department of the First Hospital of Shanxi Medical University, Taiyuan 030001, China

Correspondence should be addressed to Jingqi Wang; 20211124111054@zcmu.edu.cn

Received 9 May 2022; Revised 24 May 2022; Accepted 31 May 2022; Published 18 June 2022

Academic Editor: Yuvaraja Teekaraman

Copyright © 2022 Bao Liu et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Prostate carcinoma (PC) is a disease of the male genitourinary system and a relatively common malignant tumor. It poses a great threat to the physical and mental health of sufferers. Sufferers usually have symptoms such as frequent urination and urgency, which seriously affects their daily life [1]. For the therapy of prostate carcinoma sufferers, radical prostatectomy is used clinically to remove the diseased prostate tissue and rebuild the sufferer’s urinary pathway, but there is still a risk of recurrence after surgery [2]. Therefore, it is of expensive clinical value to evaluate the prognosis of sufferers with prostate carcinoma early and to formulate prevention and therapy plans according to the evaluation results, so as to avoid the risk of recurrence and ensure the prognosis. Vascular endothelial growth factor (VEGF) has a promoting effect on the expansion and migration of vascular endothelial cells, which can promote the growth of vascular endothelial cells, make VEGF express expensively specific expression, and contain a variety of biological activities, which can improve the growth rate of tumor cells accelerate its migration [3]. Studies show that serum-specific microRNA-323 (microribonucleic acid-323, miR-323) and serum-specific microRNA-409-3p (microribonucleic acid-409-3p, miR-409-3p) are abnormally expressed in carcinoma sufferers, which may have certain effects on the growth and migration of tumor cells, epithelial cell-mesenchymal transition, and tumor angiogenesis and bone metastasis [4, 5]. However, there are few studies on the correlation between the scales of miR-323, miR-409-3p, VEGF, and prostate carcinoma at home and abroad. To this end, this examination analyzed the scales of miR-323, miR-409-3p, and VEGF in sufferers with prostate carcinoma and studied their expression scales in sufferers with different periods and explored their correlation with prognosis, so as to provide clinical reference.

The rest of this paper is organized as follows: Section 2 discusses relevant literature and comparative analysis,
followed by the patient’s information and research methods in Section 3. The comparative analysis and data statistics are found in Section 4. Section 5 concludes the paper with summary and future research directions.

2. Related Work

Prostate carcinoma has a great impact on men’s health. In addition to affecting the function of the male genitourinary system, it also notoriously reduces the scale of sexual life, which is not conducive to the physical and mental health and quality of life of sufferers [6]. The clinical therapy of this disease is usually surgery, but to ensure the prognosis of sufferers, it is necessary to diagnose prostate carcinoma in advance and formulate a scientific and effective therapy plan [7]. Therefore, early diagnosis of prostate carcinoma has an important impact on the prognosis of sufferers. As a major regulator of angiogenesis, VEGF can specifically act on vascular endothelial cells and play an important role in the rapid growth and metastasis of tumor cells [8]. The foreign scholar shows that miR-323 and miR-409-3p are specifically expressed in prostate carcinoma sufferers. In this examination, by investigating the expression scales of miR-323, miR-409-3p, and VEGF in sufferers with prostate carcinoma, it is found that they have certain clinical value on the survival and prognosis of sufferers with prostate carcinoma.

In this examination, by evaluating the plasma miR-323 scales in sufferers with benign prostatic hyperplasia and prostate carcinoma, it is found that in contrast with sufferers with benign prostatic hyperplasia, the plasma miR-323 is notoriously expanded in sufferers with prostate carcinoma, whereas sufferers with III-IV miR-323 is notoriously expensive in period I-II prostate carcinoma sufferers (\(P < 0.05\), suggesting that miR-323 plays an oncogenic role in the progression of prostate carcinoma. Previous studies pointed out that certain signaling pathways, such as insulin-like growth factor 1 receptor and BRI3 gene (brain protein I3, BRI3), have been proposed to mediate the role of miR-323 in the pathogenesis and progression of various types of tumors potential effect [9]. For example, miR-323-3p inhibits the expression of epidermal growth factor-like (EGF-like) and 2 follistatin domain (TMEFF2) transmembrane proteins and activates protein kinase B (PKB/AKT) and extracellular regulated protein kinases (ERK) pathways, thereby inhibiting apoptosis in non-small-cell lung carcinoma (NSCLC) cell lines. However, its mechanism in prostate carcinoma is unknown, and it is speculated to be related to p27, p21, cyclin D1, carcinoembryonic antigen adhesion molecule 1 (CEA-related cell adhesion molecules (CEACAM1)), osteopontin, and e-cadherin. There is a close relationship with potential biomarkers such as these, but the extensive number of studies are still needed to verify. The serum miR-409-3p in the prostate carcinoma set and the benign prostatic hyperplasia set is detected, and it is found that the expression of miR-409-3p in the prostate carcinoma set is notoriously more vulgar than that in benign prostatic hyperplasia set (\(P < 0.05\)). The expression of miR-409-3p in period III-IV prostate carcinoma sufferers is vulgar than that in I-II. Analysis of the reason may be that miR-409-3p, as a protective gene of prostate carcinoma, has an inhibitory effect on the occurrence and development of carcinoma. Josson et al. have shown that miR-409-3p can inhibit the occurrence and development of prostate tumors and can play a role in regulating the scale of fibrinogen B and vascular endothelial growth factor, thereby affecting the growth and migration of tumors [10]. Gururajan et al. find that miR-409-3p also inhibited the expression of angiostatin, thereby reducing tumor angiogenesis [11]. The expression of miR-409-3p in sufferers with prostate carcinoma is vulgar, mainly because miR-409-3p inhibits AKT1 in vivo, thereby inhibiting the formation and migration of tumor cells [12].

VEGF is detected in the two sets of sufferers in this examination, and it is found that serum VEGF is expensively expressed in prostate carcinoma sufferers. Analysis of the mechanism may be that VEGF, as a specific vascular endothelial cytokine, can promote the growth of vascular endothelial cells, and, at the same time, can induce the proliferation of blood vessels, which may lead to the growth of tumor angiogenesis, so the expression of VEGF in sufferers with prostate carcinoma is expensive [13]. In addition, overexpression of VEGF scales can alter the density of tumor microvessels and induce hemangioendothelial cell proliferation through signaling pathways. VEGF plays an important role in tumor angiogenesis, and hypoxia, and androgen can all have an effect on the expression of VEGF and can regulate its expression [14–17]. In this examination, the expression of VEGF in different periods of prostate sufferers is analyzed. The results showed that the expression of VEGF is correspond with different periods of prostate carcinoma, indicating that VEGF expression may play a synergistic role in the development of prostate carcinoma. This is basically similar to the conclusion studied by Lv et al. [18–20]. In addition, this examination also found that miR-323, VEGF, and GPS scores are notoriously negatively correspond, while miR-409-3p is notoriously positively correspond with VEGF (\(P < 0.01\), suggesting that the detection of miR-323. The expression scales of miR-409-3p and VEGF may be effective indicators for the diagnosis of prostate carcinoma and have a correlation with the prognosis of sufferers with prostate carcinoma. There are obvious disparities in the GPS scores of different periods of prostate carcinoma, and the prognosis of sufferers with period I-II is obviously better than that of period III-IV. It can be seen that different periods of prostate carcinoma will have a certain impact on the prognosis of sufferers and provide a basis for the prognosis of sufferers.

Comparing the disparities in imaging of prostate carcinoma sufferers, it is found that the early enhancement rate of prostate carcinoma sufferers with period III-IV is notoriously expensive than that of sufferers with period I-II, and the maximum signal intensity and the start of dynamic scanning are notoriously vulgar than those of sufferers with period I-II (\(P < 0.05\)). The reasons for analysis may be that prostate carcinoma is expensively multiple, and there may be multiple lesions, the blood supply to the lesions is relatively sufficient, the density of tumor microvessels is expensive than that of surrounding normal tissues, and the
morphology and function of tumor blood vessels at the lesions are different. In addition, the tumor will invade and destroy normal cells and glands, thereby causing the expansion of the extracellular space [21–24].

The condition of sufferers with period III-IV prostate carcinoma is more severe than that of sufferers with period I-II, so the imaging findings of sufferers with different periods of prostate carcinoma are notoriously different. However, the disadvantage is that the sample size of this examination is small, and the folvulgar-up time for sufferers is short. For this reason, the folvulgar-up should continue to expand the sample, prolong the folvulgar-up time, and conduct in-depth examination to create reference value for clinical practice.

### 3. Patients Information and Research Methods

#### 3.1. General Information

32 sufferers with prostate carcinoma who are treated in our hospital from April 2020 to August 2021 are selected as the examination subjects, and 43 sufferers with benign prostatic hyperplasia are selected for contrast. The above sufferers are divided into prostate carcinoma set \((n = 32)\) and benign prostatic hyperplasia (BPH) set \((n = 43)\). The age distribution of sufferers in the two sets ranged from 48 to 76 years old, with an average age of 62.34 ± 11.23 years old. The serum prostate-specific antigen (PSA) of all sufferers is >4 ng/mL. In the prostate carcinoma set, according to TNM staging, the sufferers are divided into 18 cases of period I-II and 14 cases of period III-IV. Serum VEGF scales of all sufferers are detected by enzyme-linked immunosorbent assay (ELISA), and miR-323and miR-409-3p scales are detected by real-time quantitative PCR (qPCR) [25]. Contrast-enhanced MRI scans are also performed in both sets [26]. There is no extensive disparity in the baseline data of the two sets of sufferers \((P > 0.05)\), which is feasible. All sufferers participating in this examination are informed about the examination content and signed an informed consent.

Inclusion criteria are as follows: firstly, sufferers diagnosed with prostate carcinoma and benign prostatic hyperplasia; secondly, age > 40 years; thirdly, with the ability to communicate and communicate; fourthly, good mental state; and finally, the information is more complete.

Exclusion criteria are as follows: firstly, combined with other tumor history; secondly, combined with other urinary system diseases; thirdly, poor quality of ultrasound images; and fourthly, allergic to contrast agents.

#### 3.2. VEGF Expression Is Detected by ELISA

The sufferer is in a fasting state, 3 ml of fasting venous blood is collected, centrifuged at 3500 r/min for 10 min, the centrifugation radius is 10 cm, and the serum is taken and placed in a refrigerator for testing. Serum VEGF scales are detected by enzyme-linked immunosorbent assay, and the instructions are strictly folvulgared.

---

### Table 1: Contrast of the disparity in the expression scales of miR-323, miR-409-3p, and VEGF between the two sets \((\bar{x} \pm s)\).

|                | miR-323 | miR-409-3p | VEGF (ng/ml) |
|----------------|---------|------------|--------------|
| Prostate carcinoma set \((n = 32)\) | 20.14 ± 0.92 | 0.18 ± 0.02 | 32.97 ± 2.92 |
| Benign prostatic hyperplasia set \((n = 43)\) | 4.12 ± 0.46 | 0.69 ± 0.06 | 23.01 ± 1.48 |
| \(t\)          | 98.921  | -46.144    | 19.310       |
| \(P\)          | <0.001  | <0.001     | <0.001       |

### Table 2: Contrast of the expression scales of miR-323, miR-409-3p, and VEGF in sufferers with different periods of prostate carcinoma \((\bar{x} \pm s)\).

| Staging               | miR-323        | miR-409-3p      | VEGF (ng/ml) |
|-----------------------|----------------|----------------|--------------|
| Phase I-II \((n = 18)\) | 19.44 ± 0.51  | 0.20 ± 0.01    | 30.86 ± 2.22 |
| Period III-IV \((n = 14)\) | 21.04 ± 0.37 | 0.16 ± 0.01    | 35.70 ± 0.62 |
| \(t\)                 | -9.876         | 11.225         | -7.895       |
| \(P\)                 | <0.001         | <0.001         | <0.001       |

### Table 3: Contrast of imaging features of prostate carcinoma sufferers with different periods \((\bar{x} \pm s)\).

| Staging               | Early reinforcement rate (%) | Maximum signal strength (%) | Dynamic scan start (s) |
|-----------------------|------------------------------|----------------------------|------------------------|
| Phase I-II \((n = 18)\) | 115.34 ± 5.68                | 156.23 ± 7.69              | 35.22 ± 3.21           |
| Period III-IV \((n = 14)\) | 127.46 ± 9.23               | 143.72 ± 6.37             | 32.23 ± 2.17           |
| \(t\)                 | -4.578                       | 4.911                      | 2.989                  |
| \(P\)                 | <0.001                       | <0.001                     | 0.006                  |
3.3. Detection of miR-323 and miR-409-3p by qPCR. The miR-323 and miR-409-3p are detected by qPCR in the two sets of sufferers. Strictly follow the instructions of the miRNA extraction kit, extract the total miRNA in the cells, and operate according to the instructions of the reverse transcription kit to synthesize miRNA into eDNA and use cDNA as a template to amplify it using a real-time quantitative PCR instrument. The conditions are as follows: 95°C for 2 min, 95°C for 15 s, 60°C for 30 s, and 72°C for 8 min, cycled 40 times. The amplification curve and melting curve are detected and analyzed. The Ct values are collected, and the relative expression scales of miR-323 and miR-409-3p are calculated by the \(2^{-\Delta\Delta Ct}\) method.

3.4. Using Intensive Magnetic Resonance Scanning to Analyze Ultrasound Imaging Disparities. Sufferers in the prostate carcinoma set underwent contrast-enhanced magnetic resonance scans. Firstly, one day before the examination, the sufferer should be informed to fast food with residues and keep the bladder in an empty state. Secondly, during the examination, the sufferer is mainly in the supine position. According to the adjustment of the examination position, the sufferer is guided to adjust the posture in time, and the sufferer is subjected to a routine MRI scan. Adjust the appropriate parameters according to different inspection positions. Adjust the interlayer spacing of the instrument, mainly 3 mm, and set the matrix and field of view to 384 × 384 and 180 mm × 180 mm, respectively, and the acquisition time is based on min. When performing DWI examination on sufferers, the sagittal plane, coronal plane, and transverse plane are quickly scanned, the scanning slice thickness is 3.0 m, the matrix is 384 × 384, and the time is about 5 min.

3.5. Observation Indicators and Evaluation Criteria. The observation indicators of this examination are as follows: firstly, to compare the expression scales of miR-323, miR-409-3p, and VEGF in prostate carcinoma tissues and sufferers with benign prostatic hyperplasia; secondly, to compare the disparities in the expression scales of miR-323, miR-409-3p, and VEGF in sufferers with different periods of prostate carcinoma; thirdly, to compare the disparities in imaging features of prostate carcinoma sufferers with different periods; and fourthly, the relationship between miR-323, miR-409-3p, VEGF, and survival prognosis in prostate carcinoma sufferers is contrast, and the Glasgow prognostic score (GPS) is used to evaluate the prognosis of sufferers. The evaluation content of GPS included language ability. In three scales of eye opening ability and exercise ability, the full score is 15 points, the larger the score, the better the prognosis. Finally, to compare the disparities in survival prognosis among suffers with different periods of prostate carcinoma.

3.6. Statistical Processing. In this examination, all the data are organized, and a corresponding database is established for it, and all the databases are entered into SPSS 26.0 for data processing, and the measurement data is tested for

![Graph 1: Correlation between miR-323 and GPS score.](image1)

![Graph 2: Correlation between miR-409-3p and GPS score.](image2)

![Graph 3: Correlation between VEGF and GPS score.](image3)

### Table 4: Disparities in GPS scores of prostate carcinoma sufferers with different periods (n (%)).

| Phase           | GPS score | n (%), 26.0 |
|-----------------|-----------|-------------|
| ≤8              | 5 (27.78) | 13 (72.22)  |
| >8              | 13 (92.86)| 1 (7.14)    |
| \(\chi^2\)      | 13.553    | <0.001      |
| \(p\)           |           |             |

The full score is 15 points, the larger the score, the better the prognosis.
normality, expressed as (\(\bar{x} \pm s\)), and between-set data independent sample \(t\)-test is used, paired sample \(t\)-test is used for intranet data, count data is expressed as %, and the test is \(\chi^2\); correlation analysis is performed by Pearson (when \(P < 0.05\)), and the disparity between the data is considered to be statistically extensive.

4. Comparative Analysis and Data Statistics

4.1. Contrast of the Disparities in the Expression Scales of miR-323, miR-409-3p, and VEGF between the Two Sets. The expression scales of miR-323 and VEGF in the prostate carcinoma set are more expensive than those in the BPH set (\(P < 0.05\)), and the expression of miR-409-3p in the prostate set is notoriously vulgar than that in the BPH set (\(P < 0.05\)), as shown in Table 1.

4.2. Contrast of the Expression Scales of miR-323, miR-409-3p, and VEGF in Sufferers with Different Periods of Prostate Carcinoma. The expressions of miR-323 and VEGF in sufferers with period III-IV prostate carcinoma are notoriously expensive than those in sufferers with period I-II (\(P < 0.05\)), and contrast with sufferers with period I-II prostate carcinoma. The expression scale of miR-409-3p is more vulgar (\(P < 0.05\)), as shown in Table 2.

4.3. Contrast of Imaging Features of Prostate Carcinoma Sufferers with Different Periods. The early enhancement rate of sufferers with period III-IV prostate carcinoma is notoriously more expensive than that of sufferers with period I-II (\(P < 0.05\)). The time is shorter than that of sufferers with period I-II (\(P < 0.05\)), as shown in Table 3.

4.4. The Relationship between miR-323, miR-409-3p, VEGF, and Survival Prognosis in Sufferers with Prostate Carcinoma. The data of miR-323, miR-409-3p, VEGF, and GPS scores of prostate carcinoma sufferers are collected, and it is found that miR-323, VEGF are negatively corresponding with GPS scores, as shown in Figure 1.

Figure 2 shows correlation between miR-409-3p and GPS score. The miR-409-3p is positively correspond with GPS scores (all \(P < 0.05\)), as shown in Figure 3.

4.5. Disparities in Survival Prognosis among Sufferers with Different Periods of Prostate Carcinoma. Comparing the GPS scores of sufferers with different periods of prostate carcinoma, the number of cases with GPS scores > 8 in sufferers with period I-II prostate carcinoma is notoriously expensive than that in period III-IV (\(P < 0.05\)) as shown in Table 4.

5. Conclusions

In conclusion, the expressions of miR-323, miR-409-3p, and VEGF are of great significance in guiding different periods of prostate carcinoma, can predict the prognosis of sufferers with prostate carcinoma, and have a certain role in predicting the recurrence risk of sufferers after therapy, so as to predict their prognosis. It provides a reference for the formulation of early prevention and therapy programs and has certain value for the prognosis of sufferers with prostate carcinoma.

Data Availability

The simulation experiment data used to support the findings of this study are available from the corresponding author upon request.

Conflicts of Interest

The authors declare that there are no conflicts of interest regarding the publication of this paper.

References

[1] N. Nadiminty, R. Tummala, C. Liu, J. W. Lou, C. Evans, and A. Gao, “NF-kB2/p52 induces resistance to enzalutamide in prostate cancer: role of androgen receptor and its variants,” Molecular Cancer Therapeutics, vol. 12, no. 8, pp. 1629–1637, 2013.

[2] A. Raimondi, P. Sepe, M. Claps et al., “Safety and activity of radium-223 in metastatic castration-resistant prostate cancer: the experience of Istituto Nazionale Dei Tumori,” Tumori, vol. 106, no. 5, pp. 406–412, 2020.

[3] J. Bunni, G. Shelley-Fraser, K. Stevenson et al., “Circulating levels of anti-angiogenic VEGF-A isoform (VEGF-Axxxb) in colorectal cancer patients predicts tumour VEGF-A ratios,” American Journal of Cancer Research, vol. 5, no. 6, pp. 2083–2089, 2015.

[4] A. Kramerov, R. Shah, H. Ding et al., “Novel nanopolymer RNA therapeutics normalize human diabetic corneal wound healing and epithelial stem cells,” Nanomedicine, vol. 32, no. 1, article 102332, 2021.

[5] G. Gao, L. Cao, X. Du et al., “Comparison of minimally invasive surgery transforaminal lumbar interbody fusion and TLIF for treatment of lumbar spine stenosis,” Journal of Healthcare Engineering, vol. 2022, Article ID 49389239, 12 pages, 2022.

[6] L. Zogchel, L. Zappeij-Kannegieter, A. Javadi et al., “Specific and sensitive detection of neuroblastoma mRNA markers by multiplex RT-qPCR,” Cancers, vol. 13, no. 1, article 150, pp. 1–12, 2021.

[7] P. R. Carroll and M. Maggi, “The long-term risks of metastases in men on active surveillance for early stage prostate cancer. Reply,” The Journal of Urology, vol. 206, no. 1, article 1711, 2021.

[8] P. Larsson, A. Khaja, J. Semenas et al., “The functional interlink between AR and MMP9/VEGF signaling axis is mediated through PIP5K1a/pAKT in prostate cancer,” International Journal of Cancer, vol. 146, no. 6, pp. 1686–1699, 2020.

[9] Q. Gao, X. Yao, and J. Zheng, “MiR-323 inhibits prostate carcinoma vascularization through adiponectin receptor,” Cellular Physiology and Biochemistry, vol. 36, no. 4, pp. 1686–1699, 2020.

[10] J. S. Josson, M. Gururajan, P. Hu et al., “miR-409-3p/-5p promotes tumorigenesis, epithelial-to-mesenchymal transition, and bone metastasis of human prostate cancer,” Clinical Cancer Research, vol. 20, no. 17, pp. 4636–4646, 2014.

[11] M. Gururajan, S. Josson, G. Chu et al., “miR-154* and miR-379 in the DLK1-DIO3 microRNA mega-cluster regulate epithelial to mesenchymal transition and bone metastasis of
prostate cancer," *Clinical Cancer Research*, vol. 20, no. 24, pp. 6559–6569, 2014.

[12] M. Nobrega, H. Ciliao, M. Souza et al., “Association of polymorphisms of PTEN, AKT1, PI3K, AR, and AMACR genes in sufferers with prostate carcinoma,” *Genetics and Molecular Biology*, vol. 43, no. 3, article e20180329, 2020.

[13] M. Gallazzi, D. Baci, L. Mortara et al., “Prostate cancer peripheral blood NK cells show enhanced CD9, CD49a, CXCR4, CXCL8, MMP-9 production and secrete monocyte-recruiting and polarizing factors,” *Frontiers in Immunology*, vol. 11, no. 1, article 586126, 2021.

[14] T. Hybel, D. Dietrichs, J. Sahana et al., “Simulated microgravity influences VEGF, MAPK, and PAM signaling in prostate cancer cells,” *International Journal of Molecular Sciences*, vol. 21, no. 4, article 1263, 2020.

[15] S. Taghizadeh, Z. Soheili, M. Sadeghi et al., “sFLT01 modulates invasion and metastasis in prostate cancer DU145 cells by inhibition of VEGF/GRP78/MMP2&9 axis,” *BMC Molecular and Cell Biology*, vol. 22, no. 1, pp. 1–11, 2021.

[16] A. Pooli, D. Johnson, J. Shirk et al., “Predicting pathological tumor size in prostate cancer based on multiparametric prostate magnetic resonance imaging and preoperative findings,” *The Journal of Urology*, vol. 205, no. 2, pp. 444–451, 2021.

[17] G. Gao, P. Zhang, B. Xu et al., “Analysis of bioelectrical impedance spectrum for elbow stiffness based on Hilbert–Huang transform,” *Contrast Media & Molecular Imagine*, vol. 2022, pp. 1–11, 2022.

[18] Z. Lv, D. Chen, H. Feng, H. Zhu, and H. Lv, “Digital twins in unmanned aerial vehicles for rapid medical resource delivery in epidemics,” *IEEE Transactions on Intelligent Transportation Systems*, advance online publication, vol. 2021, pp. 1–9, 2021.

[19] Z. Lv, Z. Yu, S. Xie, and A. Alamri, “Deep learning-based smart predictive evaluation for interactive multimedia-enabled smart healthcare,” *ACM Transactions on Multimedia Computing, Communications, and Applications*, vol. 18, no. 1s, pp. 1–20, 2022.

[20] J. Chen, Q. Zou, and J. Li, “DeepM6ASeq-EL: prediction of human N6-methyladenosine (m6A) sites with LSTM and ensemble learning,” *Frontiers of Computer Science*, vol. 16, no. 2, pp. 1–7, 2022.

[21] J. Yan, Y. Yao, S. Yan, R. Gao, W. Lu, and W. He, “Chiral protein supraparticles for tumor suppression and synergistic immunotherapy: an enabling strategy for bioactive supramolecular chirality construction,” *Nano Letters*, vol. 20, no. 8, pp. 5844–5852, 2020.

[22] Z. Cao, Y. Wang, W. Zheng et al., “The algorithm of stereo vision and shape from shading based on endoscope imaging,” *Biomedical Signal Processing and Control*, vol. 76, article 103658, 2022.

[23] S. Liu, B. Yang, Y. Wang, J. Tian, L. Yin, and W. Zheng, “2D/3D multimode medical image registration based on normalized cross-correlation,” *Applied Sciences*, vol. 12, article 2828, no. 6, 2022.

[24] Z. Zhang, L. Wang, W. Zheng, L. Yin, R. Hu, and B. Yang, “Endoscope image mosaic based on pyramid ORB,” *Biomedical Signal Processing and Control*, vol. 71, article 103261, 2022.

[25] L. Chen, Y. Huang, X. Yu et al., “Corynoxine protects dopaminergic neurons through inducing autophagy and diminishing neuroinflammation in rotenone-induced animal models of Parkinson’s disease,” *Frontiers in Pharmacology*, vol. 12, article 642900, 2021.

[26] S. Mielcarska, K. Stopinska, M. Dawidowicz et al., “GDF-15 level correlates with CMKLR1 and VEGF-A in tumor-free margin in colorectal cancer,” *Current Medical Science*, vol. 41, no. 3, pp. 522–528, 2021.