

Yh2ax As A Marker Of DNA Damage In Oral Precancer And Oral Cancer: Implications For Drug Delivery, Therapeutic Monitoring, And Clinical Outcomes

Ramanathan Manikandhan¹, Saurabh Arya^{2*}, Vikram Khanna³, Dr Sureka V.⁴

¹meenakshi Ammal Dental College, Alapakkam Main Road, Madhuravoyal, Chennai, India
Meenakshi Academy Of Higher Education And Research (Maher)Drmaniramanathan@Gmail.Com

^{2*}meenakshi Academy Of Higher Education And Research(Maher)
Sourav.Arya@Gmail.Com

³affiliation Meenakshi Academy Of Higher Education And
Research (Maher)Drvikramkhanna@Gmail.Com

⁴affiliation Meenakshi Academy Of Higher Education And
Research Maher

***Corresponding Author:** Saurabh Arya

*Meenakshi Academy Of Higher Education And Research(Maher)
Sourav.Arya@Gmail.Com

Abstract

Oral squamous cell carcinoma (OSCC) is a multistage carcinogenesis that is initiated by oral potentially malignant disorders, as a result of cumulative DNA damage and genomic instability. To enhance the result of treatment, it is crucial to identify feasible biomarkers capable of linking the biology of the disease with the evaluation of its treatment to assess whether the disease was fixed or developed into a completely functional cell. γ H2AX, which is a phosphorylated form of histone generated in response to the activity of breaks on the DNA strands, has become a sensitive measure of cellular damage and repair processes. This review is a critical analysis of the role of γ H2AX in oral precancer and cancer with a focus on its growing use in drug delivery and pharmaceuticals. It has been shown that, the expression of γ H2AX rises gradually through normal mucosa, dysplasia and carcinoma as a prognostic of tumour aggressiveness and treatment outcome. Notably, γ H2AX is a pharmacodynamic biomarker of assessing the efficacy of chemotherapeutic agents, radiotherapy, and targeted therapies. Its incorporation into nanotechnology-based systems of drug delivery and controlled release formulations can serve as a quantifiable endpoint of the evaluation of drug performance and therapeutic optimization. It can also be used to develop pathway-based targeted therapies as this, in combination with the fact that it relates to DNA damage response pathways. As a whole, γ H2AX provides an opportunity in the connection of molecular diagnostics with sophisticated drug delivery methodology, which will improve accuracy of treatment in the treatment of oral cancer.

Keywords: γ H2AX; Drug Delivery, Oral Squamous Cell Carcinoma, Nanocarriers, Pharmacodynamic Biomarker

How To Cite This Article: Manikandhan R, Arya S, Khanna V, Sureka V. Yh2ax As A Marker Of DNA Damage In Oral Precancer And Oral Cancer: Implications For Drug Delivery, Therapeutic Monitoring, And Clinical Outcomes. Int J Drug Deliv Technol. 2026;16(25s):961-968. Doi: 10.25258/ijddt.16.25s.112

1. Introduction

OSCC is a major health challenge in the world since it is prevalent in locations where tobacco use, alcohol consumption and betel nut chewing are high and ends with the ultimate genetic alterations and unremitting damages to the DNA leading to malignant transformation¹.

Regardless of the additional diagnostic and treatment strategies, OSCC remains a low prognosis after late-stage representation and low efficacy of the conventional therapy, therefore, requiring the addition of molecular biomarkers and the advanced drug delivery options to enhance the instrument of earlier diagnosis, therapy localization, and patient outcomes².

In the last few years, Nanosystems of drug delivery through nanotechnology have significantly enhanced the targeted delivery of chemotherapeutic compounds, and have resulted in enhanced bioavailability, reduced

*Author for Correspondence: _sourav.arya@gmail.com

systemic poisoning and enhanced therapeutic effect of delivery of orally introduced chemotherapeutic agents³. Liposomes, polymeric nanoparticles and metallic nanoparticles are examples of nanocarriers that have shown potential as anticancer drugs delivery vectors that target specific tumor tissue, as well as had shown advantages by overcoming the negative side effects and potential toxicity limit of the traditional methods of delivery and administration⁴.

Within the emerging pharmaceuticals, the method of drug delivery and response to treatment has acquired increased importance of determining valid pharmacodynamic biomarkers, which are capable of measuring drug delivery and cause cell damage in real-time clinical application⁵.

H2AX, a phosphorylated form of the histone variant H2AX, γ H2AX at serine 139, is deemed an extremely sensitive DNA strand break repair data and an important

part of the cellular DNA damage response cascade, and therefore a convenient measurement of genomic instability and cytotoxic effects of treatments⁶.

Another evidence that is emerging to indicate that the level of γ H2AX expression increases gradually between normal mucosa of the mouth and dysplastic lesions and increases exponentially in OSCC represents the accretion of the DNA damage and its close relationship with the tumor progression, invasiveness, and the ability to metastasize⁷.

Further, γ H2AX can also interact with key molecule regulators, such as p53, ATM, CHK2 and BRCA1, indicating its primary position in the repair of DNA damage, and its potential use in the framework of the evaluation of the effectiveness of targeted drug delivery system which is aimed to induce selective destruction of DNA in cancer cells⁸.

It is interesting to note that γ H2AX has been explored to be a predictive and prognostic bi- marker in oral cancer in which the levels of expressions have been associated with treatment response particularly in radiotherapy and chemotherapy resulting in clinicians the ability to monitor treatment response and thus differentiating between treatment regimens⁹.

These qualities can inform its application of γ H2AX in drug delivery inquiry as a new way to bridge the molecular oncology and pharmaceutical disciplines by providing an appraisable end point of measuring drug activity, better-controlled and directed models of delivery and the approaches of precision drug development in the cure of oral cancer¹⁰. The objectives of this review are:

1. To assess γ H2AX as a biomarker of DNA damage in oral precancer and oral cancer
2. To evaluate the role of γ H2AX in drug delivery systems for monitoring therapeutic efficacy and optimizing treatment

2. Review Methodology

To integrate the existing evidence of the literature on the subject of the γ H2AX expression in oral precancerous lesions and oral squamous cell carcinoma (OSCC), in particular, its usage in drug delivery and therapeutic monitoring, a detailed narrative literature review was conducted. Electronic databases, including pubmed, scopus and web of science among others, were searched using keywords, including, but not limited to, the γ H2AX, DNA double-strand breaks, oral cancer, oral dysplasia, drug delivery, nanoparticles, and pharmacodynamic biomarkers, up to 2025.

The inclusion criteria were that the peer-reviewed article was an original research article or a review article that addresses γ H2AX expression in oral potentially malignant disorders (OPMDs) and OSCC; and studies

assessing responses to DNA damage caused by either chemotherapeutic agents, radiotherapy or advanced drug delivery systems. The study of γ H2AX as a biomarker to assess the effectiveness of treatments, controlled drug delivery and the delivery of drugs to specific target sites was given special attention.

Articles were filtered by them due to not being in English, not being peer reviewed, case report, and not being relevant to either oral carcinogenesis or drug delivery use. Narrative as opposed to statistical meta-analysis because the literature analysis was critical and synthesized to guarantee such consistency, since the review was exploratory and integrative. Such a strategy made it possible to integrate the expertise on the molecular, clinical, and pharmaceuticals with an additional role of the biomarker to optimize the diagnostics and optimization of drugs delivery technologies.

3. Molecular Basis of γ H2AX in DNA Damage and Repair

H2AX, a phosphorylated derivative of the histone protein H2AX, forms quickly and at DNA break sites, and is a sensitive and early sign of DNA damage and genomic instability in a range of disease pathologies, including cancer¹¹.

Such pharmacognosy must recognize and induce the activation of the DNA damage response by phosphorylation of H2AX at serine 139 that allows the identification of the presence of the damage to the DNA and signaling to ensure the preservation of genomic integrity.

Due to the broken repair pathways, continued expression of γ H2AX even with background defective DNA repair, has been attributed to overemphasized cell sensitivity to radiation, especially in cancer treatment patients receiving radiotherapy¹².

The γ H2AX foci are sites of recruitment of most DNA repair factors, including CHK2, BRCA1, and p53 to form repair complexes and activate downstream signaling pathways that have roles in cell cycle regulation and inhibiting apoptosis¹³.

Besides its effects in acute DNA damage response, it has been reported that γ H2AX is also engaged in a large number of physiological processes which implies that it is more broadly applicable in the maintenance of genomic stability in cell development and aging processes as shown in Table 1.

The microenvironmental factors of hypoxia also contribute to the regulation of the γ H2AX expression to remodel the effectiveness of DNA repair and to contribute to the heterogeneity of tumors, and this effect was also used on the responsiveness of therapy and resistance to cancer development¹⁵.

Table 1: Role of γ H2AX in Drug-Induced DNA Damage and Therapeutic Evaluation

Therapeutic Agent/System	Delivery Type	Mechanism of Action	γ H2AX Response	Clinical/Pharmaceutics Implication (Refs)
Radiotherapy	Localized radiation delivery	DNA double-strand breaks	Increased foci formation	Radiosensitivity assessment ¹⁶

Targeted therapy (PARP inhibitors)	Oral/systemic targeted delivery	Inhibition of DNA repair pathways	Enhanced γ H2AX accumulation	Therapy optimization ¹⁷
Chemotherapy (Cyclophosphamide + Veliparib)	Combination drug delivery	DNA damage + PARP inhibition	Elevated DNA damage signaling	Improved treatment efficacy ¹⁸
Flavonoid-based therapy (Apigenin)	Natural compound delivery	Induces apoptosis via DNA damage	Increased γ H2AX expression	Antitumor potential evaluation ¹⁹
Prognostic biomarker application	Diagnostic/monitoring system	DNA damage response signaling	Persistent γ H2AX levels	Survival prediction and prognosis ²⁰

Collectively, these molecular events reveal that γ H2AX plays the central role in DNA damage signaling and repair, and the genomic instability is linked to cancer progression and provides a mechanistic foundation to its application in the measurement of drug-induced DNA damage and optimization of the targeted therapeutic strategies.

4. γ H2AX Expression in Oral Precancer and Cancer

4.1 Expression in Oral Potentially Malignant Disorders (OPMDs)

It has been demonstrated that the expression of γ H2AX increases gradually with expression in normal oral mucosa and varying levels of epithelial dysplasia that are indicative of the accumulation of DNA damage during the early stages of carcinogenesis²¹. This biomarker has a very low level of expression in normal tissues, but is evident in basal and parabasal layers in case of mild dysplasia and this is an early sign of genomic instability. When dysplasia progresses to moderate and severe levels, the count and the localization of γ H2AX-positive nuclei go up in the layers in the epithelia indicating that the double-strand breaks in DNA and repair systems are exacerbated. To a large extent, these patterns of expression are inextricably linked to the risk of malignant transformation and γ H2AX is a useful device to forecast high-risk OPMDs.

Much more, it has been linked to other proteins of the DNA damage response, which further support its role in the initial molecular alterations preceding development of overt malignancy. By doing so, γ H2AX may serve as a diagnostic biomarker and as a tool to stratify patients by transformation risk.

4.2 Expression in Oral Squamous Cell Carcinoma (OSCC)

Our findings reveal that γ H2AX expression is very high in OSCC compared to the normal mucosa and the

dysplastic lesions and this is a sign of broad genomic instability and incessant DNA damage. The higher γ H2AX levels correlate with high tumor grade, high aggressive behavior and high invasive potential, meaning that it is very applicable in tumor progression. Moreover, both γ H2AX expression and lymph node metastasis have been observed to be significantly correlated and therefore, the γ H2AX expression is a useful prognostic biomarker in a clinical setting. Large foci of γ H2AX occur in tumor cells, and this suggests sustained activation of the DNA damage response, and it can be triggered by both internally acquired genetic modifications and externally administered therapeutic treatment. The findings demonstrate the importance of γ H2AX in tumor biology research, in addition to the evaluation of disease severity and progression²².

This implies that γ H2AX can be utilized as a treatment monitoring and prognosis biomarker in the case of OSCC and, therefore, it can be incorporated into a complex strategy of drug delivery and treatment assessment²³.

5. γ H2AX as a Pharmacodynamic Biomarker in Drug Delivery

5.1 Evaluation of Chemotherapeutic Drug Action

As a result of chemotherapeutic agents, including cisplatin, doxorubicin, and alkylating agents, DNA breaks in the form of double strands are induced resulting in the activation of cellular pathways of DNA damage response culminating in the eventual death of cancer cells²⁴, as shown in Figure 1. The γ H2AX pharmacodynamic biomarker is highly sensitive and enables the detection of such lesions at an early stage following the exposure to drugs. foci formation of the γ H2AX are directly correlated with the extent of the DNA damage, and therefore, good predictor of treatment success in cancer.

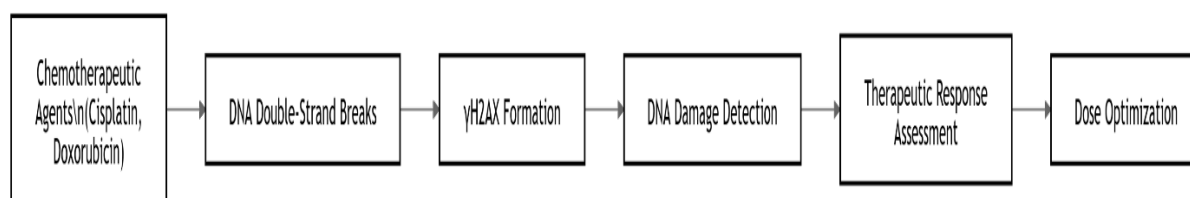


Figure 1: Schematic Representation of γ H2AX-Mediated Evaluation of Chemotherapeutic Drug-Induced DNA Damage and Therapeutic Response

This allows the clinicians and researchers to assess the workings of the drug at the molecular level and hence make the treatment plans better. Moreover, γ H2AX-based assays help to optimize the chemotherapeutic dosage regimes to kill tumor cells to the utmost, and reduce the toxicity to the minimum.

5.2 Monitoring Radiotherapy Response

Radiotherapy continues to be part of the foundations of cancer treatment and its therapeutic actions are largely based on the induction of DNA base strand breaks in cancer cells²⁵.

H2AX phosphorylation to γ H2AX is a rapid reaction to radiation and therefore it is a useful biomarker to determine radiation damages in DNA²⁶. The amount of DNA damage as well as the effectiveness of repair mechanisms in irradiated cells can be researched using the γ H2AX foci in a quantitative manner. This enables the clinicians to determine the radiosensitivity of the tumours and provide a clue on how a certain patient

would respond to a certain treatment. Consequently, γ H2AX surveillance can be useful in improving the personal radiotherapy planning by informing adaption in dose and improving delivery of treatment.

5.3 Role in Targeted Drug Delivery Systems

The objective of targeted drug delivery systems is to enhance tumor tissue therapeutic effects through specific targeting of anticancer drugs at tumor tissues but without causing systemic side effects. Their ability to induce sufficient damage to the DNA strands within the cancer cells that can be detected using gamma H2X as one such biomarker can be tested to determine the functionality of these systems^{27,28}, as also presented in Table 2. This especially applies in the development of nanocarrier and molecule-directed therapy. The γ H2AX-based model of drug delivery effectiveness and biological reaction is a highly useful instrument in the process of validation and optimization of targeted therapeutic modalities.

Table 2: γ H2AX in Drug Delivery and Therapeutic Evaluation

Therapy/System	Delivery Type	Target/Pathway	γ H2AX Role	Implication
Radiotherapy	Ionizing radiation	DNA DSBs	Foci formation	Damage quantification ²⁹
High-dose irradiation	External beam	Radiosensitivity	Residual foci	Response prediction ³⁰
DDR biomarker panel	Molecular profiling	PARP1/BRCA1	Expression levels	Survival prediction ³¹
DDR-targeted therapy	Targeted drugs	DNA repair pathways	Increased signal	Therapy advancement ³²
ATR/CHK1/2 inhibitors	Targeted therapy	Replication stress	γ H2AX activation	Precision targeting ³³

6. γ H2AX in Nanotechnology-Based Drug Delivery

The creation of nanotechnology-based drug delivery system has been a potential treatment in cancer treatment wherein therapeutic agents are transported to the tumor in the body through the delivery system with great precision and moreover, the bioavailability of drugs is also increased. These systems include liposomes, polymeric nanoparticles and hydrogel based carriers that allow the controlled release of drugs and targeted location of their release and this system is better as it enhances the efficiency of therapy and minimizes the systemic toxicity³⁴.

Some of the mechanisms that can result in the DNA damage of cancer cells by the Nanoparticles include oxidative stress, generation of reactive oxygen species, and disruption of DNA repair pathways. Such nanoparticle-induced genotoxicity can be conveniently assessed with the help of γ H2AX that is a sensitive assay of DNA double-strand breaks and cellular stress responses³⁵.

One of the tools that is applicable in studies of nanomedicine to identify the cytotoxic efficacy and pharmacologic efficacy of advanced drug delivery mechanisms is the application of γ H2AX analysis. Mapping the γ H2AX expression is provided by the DNA

damage caused by nanoparticle, which allows the researcher to optimize the system of delivery so as to target the protein in a more accurate location and to release it whenever it is required.

Thus, a significant role in the relationship between nanotechnology and pharmaceuticals is the γ H2AX which offers safer and effective targeted therapy in the management of oral cancer.

7. Role of γ H2AX in Controlled and Sustained Release Systems

Process Controlled and sustained delivery systems are used to ensure therapeutic levels of drugs, over a more extended duration, and increase treatment efficacy and reduce dosing frequency of oral cancer treatment. These systems ensure that the tumor cells have prolonged lives of the chemotherapeutic agents that augment the induction of DNA injuries and curative results³⁶.

γ H2AX is an appropriate pharmacodynamics biomarker that can be used in the measurement of the long-lasting DNA damage when the systems are subjected to a continuous exposure to a drug. Its expression reveals the preservation of the DNA interrupts of the two strands of the DNA strands, and permits the indirect determination

of the kinetics of the drug release and the intracellular drug activity over time³⁷.

The relationship between the level of γ H2AX and chronic exposure to therapy allows researchers to estimate the efficacy of controlled release preparations

and plan them to produce the most intense cytotoxic effect (Figure 2). This applies in particular to advanced drug delivery mechanisms where maintaining a constant amount of drug in the body is significant in the provision of the desired therapeutic effect.



Figure 2: γ H2AX-Guided Evaluation of Controlled and Sustained Drug Delivery Systems

Thus it can be restated that the γ H2AX analysis as part of the experiments of the controlled drug delivery is a robust solution to maximize the dosage regime or efficacy of the particular curing technique and improve the overall clinical outcome of the cancer treatment.

8. Integration with Molecular Pathways for Drug Targeting

Integration of γ H2AX and fundamental pathway in the DNA damage response presents a strong mechanistic foundation of targeted drug delivery methodologies in cancer treatment. Centrally regulated by the kinases ATM, ATR, and DNA-PK, the signalling events that control the detection of DNA double-strand breaks and initiate downstream signalling cascades, including CHK2 and p53, are all important to ensuring cell cycle arrest, repair or apoptosis³⁸.

The γ H2AX action in this pathway is that it is an initial signal to recruit repair factors, like BRCA1, to enhance this signal, which is entirely required between the identification of molecular damage and targeting of therapeutic actions.

This discovery of these pathways has led to new modes of treatment, especially the use of PARP blockers that exploit the genetic defects in the homologous recombination repair mechanism, which induces synthetic lethality in cancer cells with impaired DNA repair mechanisms³⁹.

At the same time, the γ H2AX with other biomarkers such as p53 and p-ATM can be utilized to offer a multidimensional measure of treatment response to improve precision in treatment delivery.

By so doing, integrating γ H2AX in the DNA damage repair networks will contribute to developing the complex drug delivery and combination therapy that will enhance their therapeutic effect and decrease drug resistance.

9. Clinical and Pharmaceutics Applications

The γ H2AX has become an important biomarker in the clinical oncology field as well as in drug development with reference to measuring drug efficacy and the optimal therapeutic regimens. γ H2AX is frequently applied in preclinical drug screening to determine the

DNA-damaging potential of new compounds which are useful in the case of anticancer drug development. The quantitative measure of its value is a valid pharmacodynamic endpoint to compare the various drug formulations and administration systems. γ H2AX has the potential to be used in personalized medicine to predict the specific response and repair ability of various people by showing changes in DNA damage response and repair. Also, the γ H2AX levels can be monitored in response to treatment to early identify therapeutic resistance, which can be used to adjust treatment programs in time. It is also useful in stratification of high-risk patients, which is the identification of patients with severe genomic instability. Moreover, the application of γ H2AX as a vital biomarker has been used in clinical trials that use advanced drug delivery systems to determine treatment response and confirm the effectiveness of the targeted and controlled drug delivery strategies.

10. Challenges in Translating γ H2AX into Drug Delivery Research

Despite this promising application, there are other obstacles to the overall application of γ H2AX in drug delivery research. This is because of the lack of standardized protocols of detection and quantification which are one of the most critical issues since they can lead to the variability of results in various studies and labs. The differences in sample preparation, specificity of antibodies and imaging also contribute to the differences in γ H2AX measurements. Also, γ H2AX is not entirely cancer specific DNA damage, as it can also result due to physiological stress or inflammation or environmental agent and it can be interpreted with difficulty in clinical practice. The other weakness is that, no scoring systems are universally accepted thus, findings could not be easily compared across studies. In addition, there is still a lot of clinical validation required to ascertain its reliability and repeatability in connection to being a pharmacodynamic biomarker. These issues should be overcome so as to integrate γ H2AX into the drug delivery and therapeutic monitoring models.

11. Future Perspectives

It is foreseen that the application of γ H2AX in additional therapeutic use and drug delivery would be prolonged in the future research. The combination of γ H2AX systems and nanomedicine platforms leaves the room to produce delivery systems that can be dynamically traced basing on the real-time analysis of the DNA damage. The use of the γ H2AX patterns can be further improved as a result of the new technologies, including artificial intelligence and machine learning, that would be created, allowing to predict the outcome of the treatment. Adaptive strategies that react to tumor-specific levels of DNA damage through the use of a system that responds to γ H2AX-guided delivery of therapeutic agents could be developed. Moreover, γ H2AX analysis along with some molecular treatments such as repair inhibitors may also lead to more specificity and efficacy of treatment. The technology also allows the use of innovations to monitor non-invasive and real-time γ H2AX in clinical practice through the use of biosensing technologies. All these tests indicate that γ H2AX has the potential to revolutionize the research in drug delivery and provide the new generation of targeted cancer therapy.

12. Conclusion

γ H2AX is a sensitive and dependable biomarker of DNA double-strand breaks, with important potential in the interface of molecular oncology and drug delivery technology. Its gradual onset during oral possibly malignant diseases up to oral squamous cell carcinoma illustrates its usefulness in identifying the presence of genomic imbalance and the disease progression. In addition to its diagnostic value, γ H2AX has also been found to be an important pharmacodynamic biomarker in assessing the presence of DNA damage caused by treatment and is, therefore, especially useful in determining the effectiveness of chemotherapeutic agents, radiotherapy, and targeted therapies. In the framework of pharmaceuticals, γ H2AX is an endpoint that can be quantified to measure the efficacy of new drug delivery systems, such as nanocarriers, controlled release formulations, and targeted therapeutic platforms. The effective reflection of real-time cellular responses to the exposure to drugs allows it to optimize dosing regimens and increases accuracy in the treatment planning. In addition, the combination of γ H2AX with molecular pathways including ATM -ATR signaling and DNA repair pathways enhances the design of pathway-based drug delivery strategies. Despite the persistent issues with the standardization and clinical validation, the integration of γ H2AX into the research of drug delivery has a great potential of enhancing the treatment outcomes. All in all, γ H2AX can be utilized as a useful tool to fulfill the goals of precision pharmaceuticals and the next generation of targeted therapies in the management of oral cancer.

References

1. Kattimani, V., Bhukya, N. K. N., Panga, G. S. K., Chakrabarty, A., & Lingamaneni, P. (2024). Nano-drug carriers for targeted therapeutic approaches in

oral cancer: A systematic review. *Journal of Maxillofacial and Oral Surgery*, 23(4), 763-771.

2. Herrada Céspedes, A., Reyes, M., & Morales, J. O. (2025). Advanced drug delivery systems for oral squamous cell carcinoma: A comprehensive review of nanotechnology-based and other innovative approaches. *Frontiers in Drug Delivery*, 5, 1596964.
3. Zhang, Y., Wu, Y., Du, H., Li, Z., Bai, X., Wu, Y., ... & Chen, X. (2023). Nano-drug delivery systems in oral cancer therapy: recent developments and prospective. *Pharmaceutics*, 16(1), 7.
4. Deivayanai, V. C., Thamarai, P., Karishma, S., Saravanan, A., Yaashikaa, P. R., Vickram, A. S., ... & Shruthi, S. (2025). Advances in nanoparticle-mediated cancer therapeutics: current research and future perspectives. *Cancer Pathogenesis and Therapy*, 3(04), 295-310.
5. Nicol, A. J., Ching, J. C., Tam, V. C., Liu, K. C., Leung, V. W., Cai, J., & Lee, S. W. (2023). Predictive factors for chemoradiation-induced oral mucositis and dysphagia in head and neck cancer: a scoping review. *Cancers*, 15(23), 5705.
6. Nikitakis, N. G., Rassidakis, G. Z., Tasoulas, J., Gkouveris, I., Kamperos, G., Daskalopoulos, A., & Sklavounou, A. (2018). Alterations in the expression of DNA damage response-related molecules in potentially preneoplastic oral epithelial lesions. *Oral surgery, oral medicine, oral pathology and oral radiology*, 125(6), 637-649.
7. Philouze, P., Gauthier, A., Lauret, A., Malesys, C., Muggioli, G., Sauvaigo, S., ... & Rodriguez-Lafresse, C. (2022). CD44, γ -H2AX, and p-ATM expressions in Short-Term ex vivo culture of tumour slices predict the treatment response in patients with oral squamous cell Carcinoma. *International journal of molecular sciences*, 23(2), 877.
8. Hosking, H. (2025). *Using DNA Double Strand Break Damage and Repair Capacity as an Indicator of Cancer Risk and Ageing Related Conditions* (Doctoral dissertation, CQUniversity).
9. Lyu, S. I., Fretter, C., Eckel, H. N. C., Knipper, K., Schultheis, A. M., Büttner, R., ... & Simon, A. G. (2025). High expression of H2AX/ γ -H2AX is associated with distinct biological pathway alterations and shorter survival in oropharyngeal squamous cell carcinoma. *Oral Oncology*, 161, 107171.
10. Prabhu, K. S., Kuttikrishnan, S., Ahmad, N., Habeeba, U., Mariyam, Z., Suleman, M., ... & Uddin, S. (2024). H2AX: A key player in DNA damage response and a promising target for cancer therapy. *Biomedicine & Pharmacotherapy*, 175, 116663.
11. Stope, M. B. (2021). Phosphorylation of histone H2A. X as a DNA-associated biomarker. *World Academy of Sciences Journal*, 3(3), 31.
12. Lobachevsky, P., Leong, T., Daly, P., Smith, J., Best, N., Tomaszewski, J., ... & Martin, O. A. (2016). Compromized DNA repair as a basis for identification of cancer radiotherapy patients with

- extreme radiosensitivity. *Cancer letters*, 383(2), 212-219.
13. Merighi, A., Gionchiglia, N., Granato, A., & Lossi, L. (2021). The phosphorylated form of the histone H2AX (γ H2AX) in the brain from embryonic life to old age. *Molecules*, 26(23), 7198.
 14. Kunachowicz, D., Tomecka, P., Sędzik, M., Kalinin, J., Kuźnicki, J., & Rembiałkowska, N. (2025). Influence of hypoxia on tumor heterogeneity, DNA repair, and cancer therapy: from molecular insights to therapeutic strategies. *Cells*, 14(14), 1057.
 15. Burger, K., Ketley, R. F., & Gullerova, M. (2019). Beyond the trinity of ATM, ATR, and DNA-PK: multiple kinases shape the DNA damage response in concert with RNA metabolism. *Frontiers in Molecular Biosciences*, 6, 61.
 16. Kawashima, S., Kawaguchi, N., Taniguchi, K., Tashiro, K., Komura, K., Tanaka, T., ... & Uchiyama, K. (2020). γ -H2AX as a potential indicator of radiosensitivity in colorectal cancer cells. *Oncology Letters*, 20(3), 2331-2337.
 17. de Miguel-Luken, M. J., Chaves-Conde, M., Quintana, B., Menoyo, A., Tirado, I., de Miguel-Luken, V., ... & Carnero, A. (2016). Phosphorylation of γ H2AX as a novel prognostic biomarker for laryngoesophageal dysplasia-free survival. *Oncotarget*, 7(22), 31723.
 18. Alhusaini, A., Cannon, A., Maher, S. G., Reynolds, J. V., & Lynam-Lennon, N. (2021). Therapeutic potential of PARP inhibitors in the treatment of gastrointestinal cancers. *Biomedicines*, 9(8), 1024.
 19. Kummar, S., Wade, J. L., Oza, A. M., Sullivan, D., Chen, A. P., Gandara, D. R., ... & Doroshow, J. H. (2016). Randomized phase II trial of cyclophosphamide and the oral poly (ADP-ribose) polymerase inhibitor veliparib in patients with recurrent, advanced triple-negative breast cancer. *Investigational new drugs*, 34(3), 355-363.
 20. Masuelli, L., Benvenuto, M., Mattera, R., Di Stefano, E., Zago, E., Taffera, G., ... & Bei, R. (2017). In vitro and in vivo anti-tumoral effects of the flavonoid apigenin in malignant mesothelioma. *Frontiers in pharmacology*, 8, 373.
 21. Leung, E. Y., McMahon, J. D., McLellan, D. R., Syed, N., McCarthy, C. E., Nixon, C., ... & Adams, P. D. (2017). DNA damage marker phosphorylated histone H2 AX is a potential predictive marker for progression of epithelial dysplasia of the oral cavity. *Histopathology*, 71(4), 522-528.
 22. Gocol, H., Zeng, J. H., Chang, S., Koh, B. Y., Nguyen, H., & Cirillo, N. (2023). A critical interpretive synthesis of the role of arecoline in oral carcinogenesis: is the local cholinergic axis a missing link in disease pathophysiology?. *Pharmaceuticals*, 16(12), 1684.
 23. Philouze, P., Gauthier, A., Lauret, A., Malesys, C., Muggioli, G., Sauvaigo, S., ... & Rodriguez-Lafrasse, C. (2022). CD44, γ -H2AX, and p-ATM expressions in Short-Term ex vivo culture of tumour slices predict the treatment response in patients with oral squamous cell Carcinoma. *International journal of molecular sciences*, 23(2), 877.
 24. Mitra, A., Coyne, G. H. O. S., Zlott, J., Kummar, S., Meehan, R., Rubinstein, L., ... & Chen, A. P. (2024). Pharmacodynamic effects of the PARP inhibitor talazoparib (MDV3800, BMN 673) in patients with BRCA-mutated advanced solid tumors. *Cancer chemotherapy and pharmacology*, 93(3), 177-189.
 25. Ni, J., Bucci, J., Chang, L., Malouf, D., Graham, P., & Li, Y. (2017). Targeting microRNAs in prostate cancer radiotherapy. *Theranostics*, 7(13), 3243.
 26. Rassamegevanon, T. (2020). *Translation and optimization of a gamma H2AX foci assay for the prediction of intrinsic radiation sensitivity* (Doctoral dissertation, Dissertation, Dresden, Technische Universität Dresden, 2020).
 27. Sharma, S. K., & Bagshawe, K. D. (2017). Translating antibody directed enzyme prodrug therapy (ADEPT) and prospects for combination. *Expert opinion on biological therapy*, 17(1), 1-13.
 28. Jurkovicova, D., Neophytou, C. M., Gašparović, A. Č., & Gonçalves, A. C. (2022). DNA damage response in cancer therapy and resistance: challenges and opportunities. *International journal of molecular sciences*, 23(23), 14672.
 29. Valente, D., Gentileschi, M. P., Guerrisi, A., Bruzzaniti, V., Morrone, A., Soddu, S., & Verdina, A. (2022). Factors to consider for the correct use of γ H2AX in the evaluation of DNA double-strand breaks damage caused by ionizing radiation. *Cancers*, 14(24), 6204.
 30. van Oorschot, B., Hovingh, S., Dekker, A., Stalpers, L. J., & Franken, N. A. (2016). Predicting radiosensitivity with gamma-H2AX foci assay after single high-dose-rate and pulsed dose-rate ionizing irradiation. *Radiation Research*, 185(2), 190-198.
 31. Kim, K. M., Moon, Y. J., Park, S. H., Park, H. J., Wang, S. I., Park, H. S., ... & Jang, K. Y. (2016). Individual and combined expression of DNA damage response molecules PARP1, γ H2AX, BRCA1, and BRCA2 predict shorter survival of soft tissue sarcoma patients. *PloS one*, 11(9), e0163193.
 32. Qian, J., Liao, G., Chen, M., Peng, R. W., Yan, X., Du, J., ... & Yang, Z. (2024). Advancing cancer therapy: new frontiers in targeting DNA damage response. *Frontiers in Pharmacology*, 15, 1474337.
 33. Sofianidi, A., Dumbrava, E. E., Syrigos, K. N., & Nasrazadani, A. (2024). Triple-negative breast cancer and emerging therapeutic strategies: ATR and CHK1/2 as promising targets. *Cancers*, 16(6), 1139.
 34. Madhyastha, H. (2024). Nanoparticles in Focus: Understanding Genotoxicity and Carcinogenicity. *Materials Research Foundations*, 171.
 35. Oliva, N., Conde, J., Wang, K., & Artzi, N. (2017). Designing hydrogels for on-demand therapy. *Accounts of chemical research*, 50(4), 669-679.
 36. Wilsker, D. F., Barrett, A. M., Dull, A. B., Lawrence, S. M., Hollingshead, M. G., Chen, A., ... & Kinders,

- R. J. (2019). Evaluation of pharmacodynamic responses to cancer therapeutic agents using DNA damage markers. *Clinical Cancer Research*, 25(10), 3084-3095.
37. Ambati, S. R., Shieh, J. H., Pera, B., Lopes, E. C., Chaudhry, A., Wong, E. W., ... & Moore, M. A. (2016). BO-1055, a novel DNA cross-linking agent with remarkable low myelotoxicity shows potent activity in sarcoma models. *Oncotarget*, 7(28), 43062.
38. Blackford, A. N., & Jackson, S. P. (2017). ATM, ATR, and DNA-PK: the trinity at the heart of the DNA damage response. *Molecular cell*, 66(6), 801-817.
39. Lord, C. J., & Ashworth, A. (2017). PARP inhibitors: Synthetic lethality in the clinic. *Science*, 355(6330), 1152-1158.