

Review of Poly (ADP-ribose) Polymerase (PARP) Mechanisms of Action and Rationale for Targeting in Cancer and Other Diseases

Julio C. Morales,¹ Longshan Li,¹ Farjana J. Fattah,¹ Ying Dong,¹ Erik A. Bey,² Malina Patel,¹ Jinming Gao,¹ & David A. Boothman^{1*}

¹Simmons Cancer Center, University of Texas Southwestern Medical Center, Dallas, TX 75399; ²Department of Basic Pharmaceutical Sciences & Mary Bapp Randolph, West Virginia University, Morgantown, WV 26506

* Address all correspondence to: David A. Boothman, University of Texas Southwestern, 6001 Forest Park Dr. North Campus ND2.210, Dallas, TX 75390; Tel.: 214-645-6371; Fax: 214-645-6347; david.boothman@utsouthwestern.edu

ABSTRACT: Poly (ADP-ribose) polymerases (PARPs) are a family of related enzymes that share the ability to catalyze the transfer of ADP-ribose to target proteins. PARPs play an important role in various cellular processes, including modulation of chromatin structure, transcription, replication, recombination, and DNA repair. The role of PARP proteins in DNA repair is of particular interest, in view of the finding that certain tumors defective in homologous recombination mechanisms, may rely on PARP-mediated DNA repair for survival, and are sensitive to its inhibition. PARP inhibitors may also increase tumor sensitivity to DNA-damaging agents. Clinical trials of PARP inhibitors are investigating the utility of these approaches in cancer. The hyperactivation of PARP has also been shown to result in a specific programmed cell death pathway involving NAD⁺/ATP depletion, mu-calpain activation, loss of mitochondrial membrane potential, and the release of apoptosis inducing factor. Hyperactivation of the PARP pathway may be exploited to selectively kill cancer cells. Other PARP forms, including tankyrase 1 (PARP 5a), which plays an important role in enhancing telomere elongation by telomerase, have been found to be potential targets in cancer therapy. The PARP pathway and its inhibition thus offers a number of opportunities for therapeutic intervention in both cancer and other disease states.

KEY WORDS: Parp1, DNA repair, metabolism, β -lap, DNA damage, hyperactivation, synthetic lethality, chromatin, programmed necrosis

I. WHAT IS PARP?

The poly (ADP-ribose) polymerases (PARPs) are an emerging family of enzymes that share the ability to catalyze the transfer of ADP-ribose to target proteins (poly ADP-ribosylation).^{1,2} There are at least 18 members of the PARP family that are encoded by different genes, and share homology in a conserved catalytic domain.¹ Although some isoforms including PARP1 and PARP2 are best known for their involvement in DNA repair processes, it is now clear that these and other PARPs have an important role in several cellular processes including cell proliferation and cell death.¹ A number of cellular substrates for PARP have been defined, and a majority of these proteins are nuclear proteins that are involved in nucleic acid metabolism, modulation of chromatin

structure, DNA synthesis, and DNA repair.² PARP also automodifies itself in the presence of DNA strand breaks, and is one of the main acceptors of poly ADP ribose *in vivo*.² PARP1 is the first and best characterized member of the PARP family. PARP2 is most closely related to PARP1 with 69% similarity in its catalytic domain, and was identified on the basis of the persistence of PARP activity in PARP1-deficient cells.^{1,3}

II. ROLE OF PARP IN DNA REPAIR

Evidence for an important role of PARP in DNA repair comes from the finding that DNA damaging agents and radiation-induced DNA damage causes increased PARP activity.⁴ The accumulation of DNA lesions has also been found to result in a significant

increase in PARP levels in cells.⁵ PARP is involved in base excision repair (BER) in response to single-stranded DNA breaks (SSBs) and is a component of the BER complex, which consists of DNA ligase III, DNA polymerase beta, and the XRCC1 protein.⁶ In cell-free systems, it has been shown that the unmodified PARP enzyme binds tightly to DNA strand breaks and following auto-poly ADP-ribosylation, is released and allows for repair enzyme access to the damaged DNA.^{2,7} Both PARP1 and PARP2 also interact, and share common partners in the SSB repair and BER pathways, although PARP2 also has unique partners, such as the telomeric protein TRF-2.^{1,8,9}

It has been shown that PARP1 plays a role in nucleotide excision repair (NER), since NER processes are reduced with PARP1 inhibition.¹⁰ Both BER and NER are key pathways that enable repair of DNA damage that can be caused by certain alkylating and chemotherapeutic agents.¹¹ PARP^{-/-} mice are therefore sensitive to alkylating agents such as N-methyl-N-nitrosourea (MNU) and gamma radiation exposure.¹² An increased capacity for DNA repair and/or DNA repair activity is frequently found in cells that acquire resistance to chemotherapy agents such as platinum.^{13,14} *In vitro* studies demonstrate that cells resistant to cisplatin display an increased ability to repair cisplatin-DNA damage.¹³ Upregulation of DNA repair mechanisms is therefore one of several mechanisms by which tumor cells can become resistant to chemotherapies.

III. INHIBITION OF PARP IN CHEMOTHERAPY

Because of its role in DNA repair, PARP inhibition results in genomic instability and accumulation of damaged cells in cell cycle arrest.¹⁵ This shows that ADP ribosylation reactions are needed following DNA damage and for cells to progress through G2 and M phases of the cell cycle.¹⁵ The inhibition of PARP activity using dominant negative mutant PARPs has also been shown to result in an increase in apoptosis, which arises in part due to a reduced DNA repair capacity.¹⁶ It has been suggested that PARP is a key component of the cell

cycle G2 checkpoint, which prevents a damaged cell with DNA strand breaks from being able to enter mitosis.¹⁶ Expression of a dominant negative DNA-binding domain of PARP thus sensitizes cells to SSB caused by alkylating agents.^{16,17} As noted earlier, PARP^{-/-} deficient mice are also extremely sensitive to gamma radiation, and DNA damaging agents cause rapid apoptosis in PARP^{-/-} cells.¹² These findings demonstrate the importance of PARP in post-DNA-damage repair.¹² The viability of PARP^{-/-} mice further suggests that PARP is relatively dispensable for normal activity, but is an essential survival factor for DNA damage.¹² These characteristics of PARP make it an attractive candidate for therapeutic inhibition in combination with cancer chemotherapy or radiotherapy.

There is evidence of upregulation of PARP activity in some cancer types. It has been shown that tumor tissue from hepatocellular carcinoma patients displayed significantly increased levels of ADP ribosylated PARP than did non-tumorous adjacent tissues.¹⁸ Recent results further indicate that PARP1 mRNA was upregulated in several tumor types with the most striking differences observed in primary tumors of the breast, endometrium, lung, ovary, and skin.¹⁹ In particular, a high expression of PARP1, but not PARP2, was found in triple-negative breast cancer (TNBC) tumors.¹⁹ The latter findings suggest that inhibition of PARP, either alone or in combination with DNA-damaging agents, could be a potential therapeutic approach in TNBC and other tumor types.¹⁹ This therapeutic approach is currently under investigation in several clinical development programs. Inhibition of PARP has potential for use in cancer treatment through at least two mechanisms, i.e., by increasing tumor sensitivity to chemotherapeutic agents that damage DNA, and also by inducing “synthetic lethality” in cells that are highly dependent on PARP, due to deficiency in HR, such as BRCA1 mutants (Fig. 1).

IV. INHIBITING PARP AND SYNTHETIC LETHALITY

The breast cancer-associated gene BRCA1 is

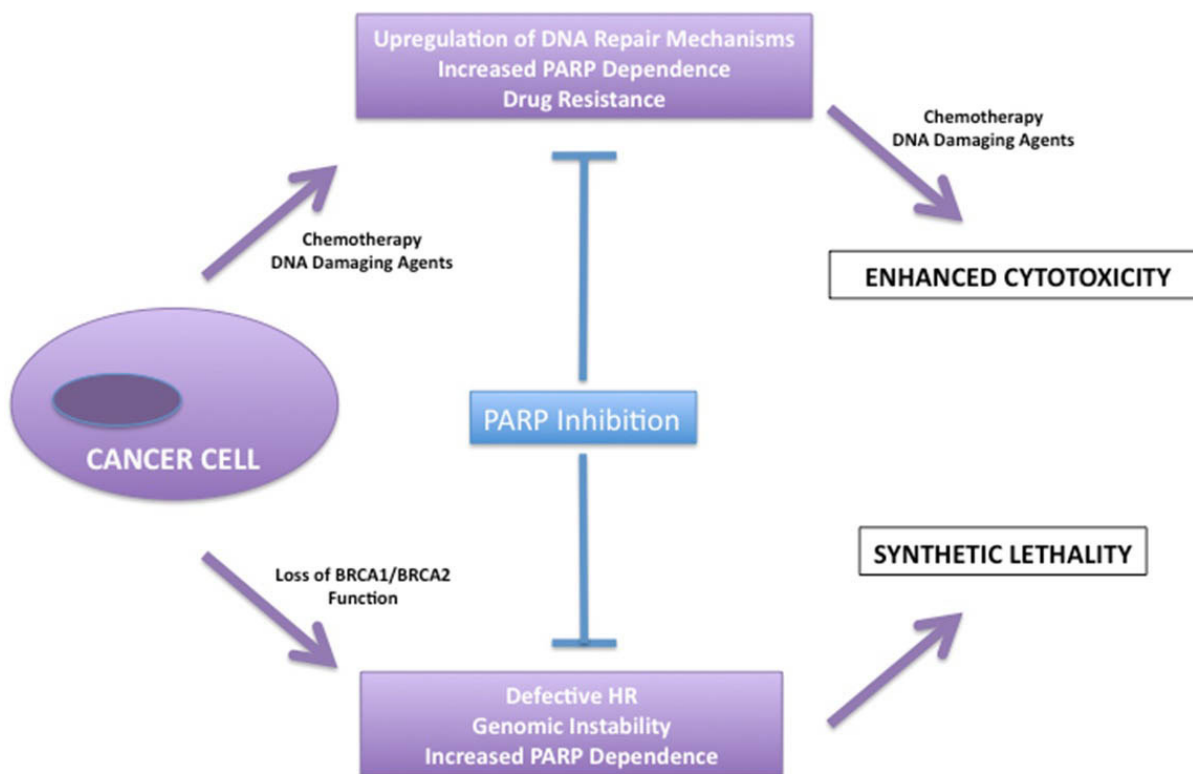


FIG. 1: Dual therapeutic potential for PARP inhibition in oncology

known to play an important role in repair of DS DNA breaks via homologous recombination (HR) because cells that are deficient in BRCA1 display impaired HR and an inability to repair defective chromosomes.²⁰ Similarly, BRCA2 interacts with the DNA repair protein RAD51 and has also been shown to play an important role in HR because cells deficient in RAD51-interacting regions of BRCA2 display hypersensitivity to DNA cross-links and chromosomal instability.²¹ It has been shown that defects in HR repair mechanisms, arising from deficiencies in key repair proteins such as RAD51, DSS1, RPA1, or CHK1, cause cells to be highly dependent on the activity of PARP and therefore highly sensitive to its inhibition.²² It has been postulated that PARP inhibition compromises SSB repair and BER, and, in cells lacking intact HR mechanisms (e.g., BRCA1 and BRCA2 mutants), these are then converted into double-

stranded breaks, resulting in cell lethality.²³ Accordingly, it has been shown that cells deficient in BRCA1 or BRCA2 are, respectively, 57- and 133-fold more sensitive than normal cells to PARP inhibition.²⁴ Thus, cells with defects in HR mechanisms may also be targeted with PARP inhibitors to further impair DNA repair mechanisms, resulting in “synthetic lethality.” The benefit of this approach is that PARP inhibition is likely to be highly selective for tumors, for example, that are BRCA2^{-/-} in patients with BRCA2 heterozygosity.^{23,25} Immunoreactivity for PARP1, for example, was found in over 80% of BRCA1 tumors, suggesting that PARP inhibition may be a useful intervention for patients with breast cancer harboring BRCA1 mutations.²⁶ Interestingly, however, it has been found that loss of 53BP1, a DNA-damage response factor, in BRCA1 tumors alleviated their recombination defect and reverted their hy-

persensitivity to DNA damage.^{27,28} Further, loss of 53BP1 appears to be common among both TNBC and tumors with BRCA1/2 mutations, suggesting that this may be an important biomarker for determining sensitivity to PARP inhibitors and/or DNA damaging agents in these tumors, or whether other therapeutic strategies might be necessary.^{27,28} An assay utilizing RAD51 focus formation has been developed to rapidly assess the HR state of tumors and their corresponding sensitivity to PARP inhibitors.²⁹

V. PARP FUNCTION IN CHROMATIN REMODELING

Because chromatin structure can affect several important processes such as DNA transcription, replication, and repair, its modulation by PARP is important.² Early studies demonstrated that poly ADP-ribosylation of DNA poly nucleosomes by PARP results in an overall relaxation of chromatin structure.³⁰ This relaxation of chromatin structure could be an important event that facilitates processes such as DNA replication, repair, and transcription of DNA.³⁰ Relaxation of chromatin is also likely to allow damaged DNA to become more accessible to repair enzymes.³⁰ More recent findings have shown that PARP1 functions as a recruiter and activator of proteins such as Alc1 that are involved in the remodeling of chromatin.³¹ In this study, the PARP1-mediated poly ADP-ribosylation of Alc1 both directed its recruitment to chromatin as well as activated its ATPase and chromatin remodeling activity.³¹ PARP2 also appears to function as a chromatin modulator, and it has been suggested, based on gene targeting studies in mice, that PARP1 and PARP2 share complementary, but not fully overlapping functions; double-mutant PARP1 and PARP-2-deficient mice are not viable.¹

VI. PARP EFFECTS ON TRANSCRIPTION FACTORS AND EPIGENETIC FACTORS

Early studies also suggested that PARP plays an important role in the regulation of DNA transcription,

since PARP activity was found to be more abundant on regions of the chromatin that were transcriptionally active (euchromatin) and less associated with transcriptionally inert regions.³² Similarly, it has been shown that PARP serves as a transcriptional cofactor and could stimulate both activator-dependent as well as basal transcriptional activity mediated by RNA polymerase II.³³ A role for PARP in the regulation of a specific gene has also been demonstrated by virtue of its interaction with a transcription enhancer factor TEF1, at MCAT promoter elements in the promoter region of the cardiac troponin T (cTNT) gene.³⁴ In this study, PARP could ADP-ribosylate TEF1, and MCAT-dependent transcriptional activity could be blocked by inhibition of PARP enzymatic activity, suggesting that PARP has a role in the modulation of muscle-specific gene transcription.³⁴ Studies have also shown that PARP interacts with the transcription factor AP-2 and enhances its transcriptional activity as a coactivator in cell lines.³⁵ Conversely, PARP can bind to nuclear receptors and inhibit ligand-dependent transcriptional activity mediated by receptors such as the retinoid x receptor (RXR).³⁶ Taken together, these results suggest that poly ADP-ribosylation can serve to either stimulate or repress transcriptional activity and modulation of PARP may therefore have therapeutic implications for many disease states. Another more recently identified example of PARP1 function in transcription is its impact on the promoter regions of genes known to be regulated by PARP1.³⁷ In this study, PARP1 was shown to promote a permissive chromatin environment onto which RNA polymerase II transcriptional machinery could be efficiently loaded; depletion of PARP1 activity caused these processes to become inefficient.³⁷

VII. PARP1 HYPERACTIVATION AND PROGRAMMED NECROSIS

PARP1 hyperactivation has also been shown to result in a specific programmed cell death pathway that is characterized by depletion of NAD⁺/ATP, calcium imbalance, mu-calpain activation, loss of mitochondrial membrane potential, rapid

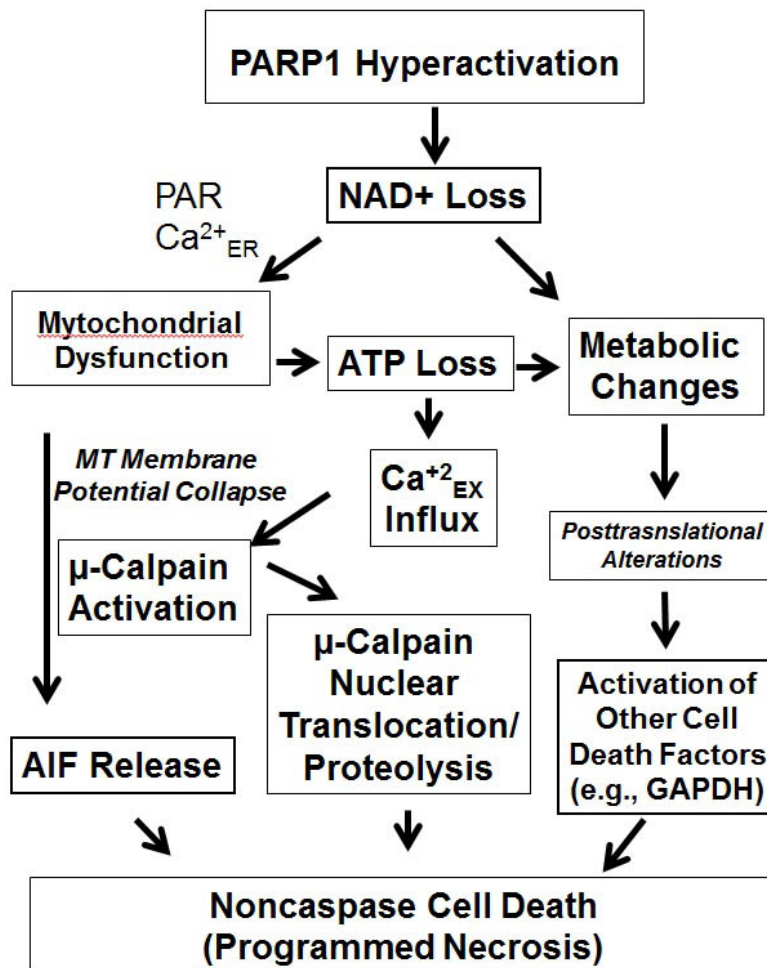


FIG. 2: PARP-mediated cell death mechanisms

mitochondrial dysfunction, and release of apoptosis-inducing factor (AIF) resulting in cell death (Fig. 2).

VIII. “ENERGY CRISIS”: NAD⁺/ATP DEPLETION/MITOCHONDRIAL ENERGY FAILURE

Early studies showed that PARP was the key molecule that resulted in NAD⁺ depletion following exposure to DNA damaging agents and gamma radiation.⁴ As originally proposed by Berger as the “suicide hypothesis,” in cells with extensive DNA damage or damage that is not repaired, PARP remains activated, leading to continued NAD⁺ de-

pletion and further ATP consumption in order to resynthesize NAD⁺.³⁸ This process escalates the energy crisis in the cell resulting in programmed necrosis.³⁹ It has been shown that depletion of NAD⁺ and mitochondrial permeability transition are important sequential steps involved in PARP1-mediated cell death.⁴⁰ Furthermore, the restoration of NAD⁺ leads to recovery of mitochondrial membrane potential, and blocks translocation of AIF from the mitochondria to the nucleus.⁴⁰ PARP1 activation has also been shown to induce a rapid mitochondrial dysfunction, which was followed by a collapse in mitochondrial potential, and release of AIF and cytochrome c.⁴¹ More recently, this mi-

tochondrial energy failure has been shown to be a direct consequence of PARP1 hyperactivation; Chiarugi and coworkers have found that the PARP product poly ADP-ribose becomes catabolized to AMP via the action of poly ADP-ribose glycohydrolase (PARG) and nucleoside diphosphate X (NUDIX) hydrolases.⁴² The accumulated AMP then serves to compete with ADP for binding to the adenine nucleotide transporter, abrogating energy production by the mitochondria and further contributing to the “energy crisis.”⁴²

IX. PAR FORMATION AND AIF RELEASE

Dawson and coworkers first connected PARP1 hyperactivation with AIF release from mitochondria, signaling a caspase-independent programmed cell death pathway.⁴³ Their studies showed that PARP1 was required for AIF release from the mitochondria, and that AIF was required for a PARP1-mediated cell death program that was caspase independent.⁴³ Consequently, it was shown that poly ADP-ribose polymer, or PAR, was a “death signal” that causes AIF release from mitochondria in response to PARP1 hyperactivation; e.g., cells with reduced AIF are resistant to PARP1 mediated death or PAR toxicity.¹⁴ AIF release from mitochondria thus plays a major role in cell death induced by PARP1 (which is also termed “parthenatos”), a process that is not dependent on calpain activation.⁴⁴ The translocation of AIF from the mitochondria to the nucleus has been shown to be required for cell death induced by agents such as arsenic trioxide.⁴⁵ This mechanism also involves production of ROS and PARP1 hyperactivation, and PARP1 inhibitors can attenuate this cell death mechanism.⁴⁵ Collectively, the data indicate that PARP1 activation triggered by ROS causes AIF release from mitochondria resulting in a caspase-independent cell death pathway.⁴⁵ The translocation of AIF from mitochondria to the nucleus causes chromatin condensation and DNA fragmentation, and AIF downregulation confers resistance to DNA alkylating agent-induced necrosis.^{46,47} These findings demonstrate the importance of AIF as a mediator of cell death in the PARP pathway.

X. POTENTIAL THERAPEUTIC ROLE(S) OF PAR/PARG MANIPULATION

PARP enzymes function to convert NAD⁺ into polymers of ADP-ribose, or PAR, which, as noted earlier, plays an essential role in PARP-mediated cell death, however, there is also evidence that PAR may act on mitochondria directly, independent of NAD⁺ depletion.^{47,48} PAR polymer has been shown to be directly cytotoxic to neurons, and manipulation of the PAR polymer signal may therefore be of value in disease states that involve cellular injury.⁴⁹ PARG is needed for the hydrolysis of PAR, and also has an important function in regulating cell death following DNA injury.⁵⁰ Analysis of PARG-deficient mice suggests that PARG plays an important role in the maintenance of PAR levels.⁵¹ Studies have shown that in undamaged cells, PARG depletion is not detrimental, and could protect cells from spontaneous SSB and telomeric abnormalities.⁵⁰ In contrast, PARG deficiency has been shown to increase the sensitivity of cells to radiation, by virtue of a defect in repair of SSB and DSB repair, leading to centrosome amplification and mitotic spindle defects; PARG could thus be another potential therapeutic target for radiotherapy, and PARG inhibition could be useful strategy to sensitize cells to DNA-damaging agents and chemotherapies.^{52–54} Inhibition of PARG may also be useful in physiologic conditions such as ischemia, inflammation, and stroke.⁵³

Tight junctions are important in structures such as the blood-brain barrier (BBB), and the tight junction protein occludin has been found downregulated following PARP1 hyperactivation in animal models of ischemia-reperfusion injury, with an accompanying increase in BBB permeability.⁵⁵ The PARP inhibitor PJ34 prevented the loss of occludin and preserved BBB function in this model, suggesting a therapeutic benefit of PARP inhibition in this disease state.⁵⁵ Similarly, in animal models of ischemic stroke, inhibition of PARP using 3-aminobenzamide (3-AB) significantly reduced brain and plasma levels of matrix metalloproteinase 9 (MMP 9), brain

nuclear factor KappaB (NF- κ B), and neutrophil infiltration as assessed by myeloperoxidase activity.^{56,57} Thus, in addition to its potential for use in cancer therapy, the modulation of the PARP pathway may have important implications for other disease states. Recent studies suggest that PARP knockout mice are highly susceptible to diet-induced obesity.⁵⁸ Compared to their wild-type counterparts, PARP knockout mice display fat accumulation, hyperleptinemia, insulin resistance, and glucose intolerance; this weight gain is due in part to decreased metabolic rate and total energy expenditure.⁵⁸ These findings argue for an important role of the PARP pathway in obesity.

XI. PARP1 AND TUMOR-SELECTIVE PROGRAMMED NECROSIS (BETA-LAPACHONE)

There is evidence that the PARP1 hyperactivation pathway itself may also be exploited to selectively kill certain types of cancer cells. Beta-lapachone (β -LAP) is a chemotherapeutic and radiosensitizer agent that causes the initiation and execution of apoptosis in cancer cells with characteristic DNA ladder formation.⁵⁹ In HL-60 cells, this apoptosis was also associated with activation of caspase 3 and proteolytic cleavage of PARP, and could be inhibited by expression of the apoptosis regulator Bcl2.⁵⁹ Findings in MCF7 cells, however, also showed that β -LAP, at higher concentrations, killed cells through cell-cycle-independent activation of a noncaspase proteolytic pathway.⁶⁰ Subsequent studies uncovered a novel mechanism of PARP1-mediated cell death in response to agents that generate ROS; in cells with elevated levels of the two-electron reductase NAD(P)H:quinone oxidoreductase 1 (NQO1), β -LAP caused a depletion of NAD⁺ and ATP that were at least partly mediated by PARP1, since inhibitors of PARP abrogated these effects and the ensuing apoptotic program.⁶¹ These findings established PARP1-mediated NAD⁺ and ATP depletion as essential upstream events in the β -LAP-mediated cell death pathway.⁶¹ Accordingly, it has also been shown that lung cancer cells expressing high levels of NQO1 are selectively killed by β -LAP treatment

in a manner that involved ROS formation, extensive DNA damage, and PARP1 hyperactivation, and this pathway could be blocked with PARP inhibition.⁶² By comparison, prostate cancer cells (LNCaP) that are deficient in NQO1 were resistant to killing by β -LAP, and these findings demonstrated that NQO1 activity was a key determinant of β -LAP toxicity in prostate cancer cells.⁶³ MCF7 breast cancer cells expressing NQO1 also undergo a proteolytic apoptosis pathway that involves activation of mu-calpain and is independent of caspase activation.⁶⁴

As noted above, PARP1 has been shown to be necessary for initiation of cell death by β -LAP and calcium has been found to be an important cofactor in this process.⁶¹ From a kinetic standpoint, cell death by β -LAP has been shown to involve reactive oxygen species (ROS) formation with extensive DNA damage, PARP1 hyperactivation, depletion of NAD⁺/ATP levels, proteolytic cleavage of p53/PARP1, mu-calpain activation, and apoptosis (Fig. 3).⁶² PARP1 hyperactivation, depletion of nucleotides, and apoptosis could be blocked with a calcium chelator or PARP inhibitor.⁶² β -LAP has also been shown to act as a potent radiosensitizing agent. β -LAP sensitized NQO1-expressing prostate cancer cells to ionizing radiation (IR) through a mechanism involving NQO1 bioactivation, induction of ROS, irreversible DNA damage via SSB, and PARP hyperactivation with NAD⁺/ATP depletion and mu-calpain-induced necrosis.⁶⁵ Furthermore, this radiosensitization by β -LAP was also blocked with PARP inhibition or the NQO1 inhibitor dicoumarol.⁶⁵ This synergistic combination of radiation with β -LAP was effective when β -LAP was administered between 0 and 10 h following irradiation, which served to upregulate NQO1 levels; β -LAP in turn sensitized the cells by inhibiting the repair of sublethal radiation damage.⁶⁶ β -LAP has also been found to synergize with IR in its antitumor effects in mouse xenograft models; these findings suggest that β -LAP may be an effective radiosensitizer in prostate cancers that overexpress NQO1.⁶⁵ Taken together, these findings demonstrate that the actions of β -LAP, which closely involve the PARP

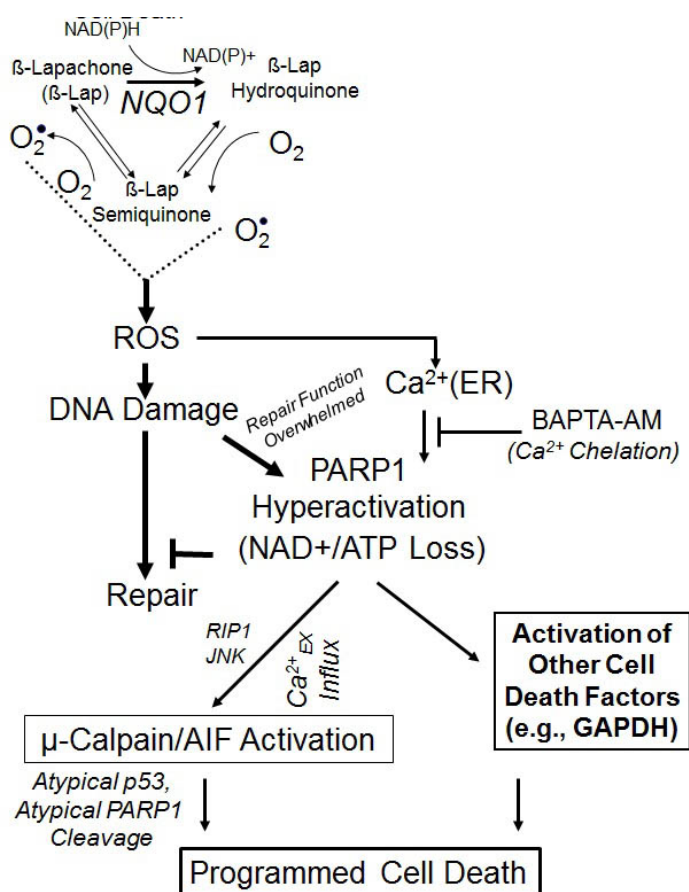


FIG. 3: Mechanisms of β -lapachone-induced toxicity in cancer cells

pathway, may be useful to selectively kill cells with NQO1 overexpression, and increase their sensitivity to radiotherapy.

XII. EXPLOITING PARP1 PROCESSES USING THERANOSTIC NANOMEDICINE

The first endeavor to enhance the solubility and bioavailability of β -LAP was undertaken using cyclodextrin inclusion complexes (alpha-, beta-, gamma-, and HPbeta-CD) by Gao and coworkers.⁶⁷ Their findings showed that the complexing of β -LAP with HPbeta-CD improved its solubility and bioavailability, with a TD50 of 2.1 microM observed in cytotoxicity studies.⁶⁷ Nano-delivery vehicles such as polymer micelles have also been used to enhance the solubility and biocompat-

ibility of β -LAP, and these β -LAP micelles have been shown to be effective in killing a variety of tumor cells expressing NQO1 with a similar cytotoxic effect involving PARP1 hyperactivation as the free drug.⁶⁸ More recently, these polymer micelles have been shown in mouse xenograft models to exhibit prolonged blood circulation, with a half-life of \sim 28 h, and increased accumulation in tumors, with an accompanying delay in tumor growth and increased survival.⁶⁹ Finally, β -LAP-encapsulated polymer implants (millirods) have been demonstrated to achieve tumor-specific therapeutic effects with site-specific release after direct implantation into prostate tumors.^{70,71} Implantation of β -LAP millirods into prostate tumor xenografts demonstrated significant tumor regression and enhanced survival. These findings offer

an attractive potential therapy for prostate tumors expressing NQO1 with minimal systemic toxicity observed.⁷¹

XIII. POTENTIAL UTILITY OF OTHER PARPS IN CANCER (TELOMERIC PARP)

Telomeres are DNA-protein complexes that protect chromosome ends in eukaryotic cells, and the “end replication” problem refers to the inability of the DNA replication machinery to replicate the very ends of DNA, resulting in a gradual loss of telomeres with each replication of DNA.⁷²⁻⁷⁴ Although new telomeric DNA can be synthesized by telomerase, the level of telomerase activity is generally insufficient to prevent progressive attrition of telomeres in most somatic cells; this results in replicative senescence.⁷³ In contrast, because the majority of cancer cells have high telomerase activity, they can continue to replicate indefinitely and this forms the basis for the use of telomerase inhibitors as cancer therapy.^{73,74} The disadvantage of telomerase inhibition, however, is the need for the continued cell division and consequently the repetition of the end replication problem that ultimately results in senescence and cell death.^{73,74} In addition, the long treatment periods required for sufficient telomere loss to occur can result in the emergence of resistance to telomerase inhibitors.^{73,74} It has been suggested that increasing the efficiency of telomere shortening and therefore hastening the onset of the “crisis” would be a more favorable approach for targeting the cancer cell.^{73,74} In this regard, TRF-1 interacting ankyrin-related ADP-ribose polymerase 1, or tankyrase 1 (also known as PARP 5a), has been shown to enhance access of telomeres to telomerase in a manner dependent on its ADP-ribosylating activity, thereby enhancing telomere elongation by telomerase.^{73,75} As such, increased expression and/or activity of tankyrase 1 can thereby promote telomere elongation in the face of telomerase inhibition. Consistent with this notion, tankyrase 1 has been found to be overexpressed in certain cancer cells including gastric cancer and breast cancer. Significantly higher expression of tankyrase 1

has been detected in human breast tumor tissues as compared to uninvolved normal breast tissue.⁷⁶ Expression of tankyrase 1 was also significantly elevated at both the mRNA and protein level in gastric cancers as compared to normal tissues, and this increased expression was further correlated with increased telomerase activity compared to normal tissue.⁷⁷ Moreover, the inhibition of tankyrase 1 using antisense RNA has been found to enhance the shortening of telomeres in gastric cancer cells without affecting telomerase activity, suggesting that tankyrase 1 may be a valid target for inhibition for this and other cancers.⁷⁸

Because the PARP activity of tankyrase 1 is required for its activity in promoting telomere elongation, the use of PARP inhibitors, in conjunction with telomerase inhibitors, has also been suggested as a means to synergize and enhance the activity of the latter in cancer cells.⁷³ Interestingly, recent findings from Ji and coworkers suggest that iniparib (BSI-201) was distinct from three other PARP inhibitors in that it did not show activity as an inhibitor of PARP1 or PARP2 in A375 tumor xenografts, or of PARP1 in BRCA-deficient MX-1 cells, MCF7 cells, or healthy donor peripheral blood mononuclear cells.⁷⁹ However, in this study, iniparib and another PARP inhibitor (ABT-888) caused dose- and time-dependent increases in phosphorylated histone H2AX (γ -H2AX), a marker of DNA damage, independent of PARP inhibition in MX-1 cells, suggesting that these agents both induced DNA damage in a BRCA1-negative background.⁷⁹ Both iniparib and ABT-888 increased telomeric γ -H2AX foci, however, only iniparib increased telomeric γ -H2AX foci, and these findings were consistent with differential gene expression and sequence analyses, which further suggested that iniparib suppressed genes in the telomere maintenance pathway, including DKC1 and PINX1.⁷⁹ These findings have important implications regarding the PARP specificity of these inhibitors and raise the interesting prospect that iniparib, unlike other PARP inhibitors, may act specifically on telomeric PARP.

As noted earlier, PARP1 is the first and best characterized member of the PARP family, and

has been the focus of most of the studies described herein. Many other members of the PARP family, however, have been characterized and their requirement for PARP enzymatic activity could further expand the therapeutic potential of PARP inhibitors.¹ PARP3, a newly characterized member of the PARP family, also appears to function in the cellular response to DSBs, and is important in the stabilization of the mitotic spindle and telomere integrity.⁸⁰ Interestingly, PARP3 has also been shown to interact with and regulate tankyrase 1, and these functions may be potentially important for cancer therapeutics. Vault PARP (VPA, PARP4) is a component of the large ribonucleoprotein complex, the vault particle, and, although initial gene targeting experiments failed to identify a function, subsequent work has shown that VPA-deficient mice are more susceptible to carcinogen-induced tumorigenesis.^{1,81,82} A second tankyrase, tankyrase 2 (PARP5b) has also been identified that has some distinct properties from tankyrase 1, however, gene targeting experiments suggest that this tankyrase is functionally redundant with tankyrase 1 during mouse development.^{83,84} Further study will be necessary to elucidate the molecular basis and implications of this redundancy, as well as the redundancy of other PARP homologs (e.g., PARP1 and PARP2).¹ As additional PARPs are identified, and the functions of the various PARP family members are elucidated, their potential utility as therapeutic targets in cancer and other disease processes is likely to expand even further.

XIV. SUMMARY AND CONCLUSIONS

PARP is a critical enzyme involved in DNA repair and many other cellular processes including transcription and modulation of chromatin structure. PARP plays a central role in NER and BER, and enables repair of DNA damage caused by alkylating agents and chemotherapeutic drugs. Studies have demonstrated that PARP is upregulated in certain cancers including TNBC and those that express BRCA1 mutations. Because certain tumor types (e.g. BRCA1 mutants) are more de-

pendent on PARP, they are also more sensitive to “synthetic lethality” with PARP inhibitors. PARP inhibition may also be useful in combination with chemotherapy to selectively sensitize cancer cells to DNA damaging agents and promote tumor cell death.

Other mechanisms by which PARP exerts its effects are via rapid mitochondrial dysfunction with membrane permeability transition, NAD⁺ depletion, and translocation of AIF from the mitochondria to the nucleus. Inhibition/manipulation of other components of the PARP pathway, including PAR and PARG, may be therefore be useful therapeutic interventions not only for cancer, but for other disease states. Hyperactivation of PARP1 may also be exploited to selectively kill cancer cells with agents such as β -LAP, and the combination of this drug with novel delivery systems may offer promising new therapeutic options for patients. Finally, other PARP family members such as telomeric PARPs (tankyrases) may offer another potential avenue for therapeutic intervention in cancer and possibly other disease states. The specificity of the different PARP inhibitors that are currently under development and/or undergoing clinical evaluation could further expand the utility of PARP and its inhibition.

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