

# Salicylic acid metabolites and derivatives inhibit CDK activity: Novel insights into aspirin's chemopreventive effects against colorectal cancer

RAKESH DACHINENI<sup>1</sup>, D. RAMESH KUMAR<sup>1,3</sup>, EDUARDO CALLEGARI<sup>2</sup>, SIDDHARTH S. KESHARWANI<sup>1</sup>, RANJINI SANKARANARAYANAN<sup>1</sup>, TERESA SEEFELDT<sup>1</sup>, HEMACHAND TUMMALA<sup>1</sup> and G. JAYARAMA BHAT<sup>1</sup>

<sup>1</sup>Department of Pharmaceutical Sciences and Translational Cancer Research Center,

South Dakota State University College of Pharmacy and Allied Health Professions, Brookings, SD 57007;

<sup>2</sup>SD-BRIN Proteomic Facility, University of South Dakota School of Medicine, Vermillion, SD 57069, USA

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**Abstract.** Aspirin's potential as a drug continues to be evaluated for the prevention of colorectal cancer (CRC). Although multiple targets for aspirin and its metabolite, salicylic acid, have been identified, no unifying mechanism has been proposed to clearly explain its chemopreventive effects. Our goal here was to investigate the ability of salicylic acid metabolites, known to be generated through cytochrome P450 (CYP450) enzymes, and its derivatives as cyclin dependent kinase (CDK) inhibitors to gain new insights into aspirin's chemopreventive actions. Using *in vitro* kinase assays, for the first time, we demonstrate that salicylic acid metabolites, 2,3-dihydroxybenzoic acid (2,3-DHBA) and 2,5-dihydroxybenzoic acid (2,5-DHBA), as well as derivatives 2,4-dihydroxybenzoic acid (2,4-DHBA), 2,6-dihydroxybenzoic acid (2,6-DHBA), inhibited CDK1 enzyme activity. 2,3-DHBA and 2,6-DHBA did not inhibit CDK2 and 4; however, both inhibited CDK-6 activity. Interestingly, another derivative, 2,4,6-trihydroxybenzoic acid (2,4,6-THBA) was highly effective in inhibiting CDK1, 2, 4 and 6 activity. Molecular docking studies showed that these compounds potentially interact with CDK1. Immunoblotting experiments showed that aspirin acetylated CDK1, and pre-incubation with salicylic acid and its derivatives prevented aspirin-mediated CDK1 acetylation, which supported the data obtained from molecular docking studies. We suggest that

intracellularly generated salicylic acid metabolites through CYP450 enzymes within the colonic epithelial cells, or the salicylic acid metabolites generated by gut microflora may significantly contribute to the preferential chemopreventive effect of aspirin against CRC through inhibition of CDKs. This novel hypothesis and mechanism of action in aspirin's chemopreventive effects opens a new area for future research. In addition, structural modification to salicylic acid derivatives may prove useful in the development of novel CDK inhibitors in cancer prevention and treatment.

## Introduction

Increasing evidence from epidemiological studies and randomized clinical trials have suggested that aspirin decreases the incidence of colorectal cancers (CRC) (1-4). Most of aspirin's chemopreventive effects came to light during the course of its use as a cardio-protective and/or analgesic agent. It was observed that aspirin at doses ranging from 75 to 325 mg, when taken regularly over 5-10 years decreases the incidence and mortality associated with CRC (1,4). The low-dose 75 mg aspirin was just as effective as the high-dose in preventing CRC. Studies also revealed that aspirin is more effective against CRC as compared to the cancers of other tissues such as breast, prostate, liver, lung, and skin (5-7).

Aspirin is mainly absorbed in the acidic environment of the stomach and upper intestine. The oral bioavailability of aspirin is 40-50% (8); it undergoes hydrolysis in the intestine, liver, and plasma by esterases (9). In contrast to the short half-life of intact aspirin (20 min), salicylic acid, depending upon the dose, has a half-life varying between 4 to 6 h, and remains much longer in the plasma. In the liver, salicylic acid undergoes further metabolism through glucuronide formation to produce ester and ether glucuronides (1-42%), or conjugation with glycine to produce salicylic acid (20-65%). In addition, the cytochrome P450 (CYP450) enzymes in the liver metabolize salicylic acid to 2,5-dihydroxybenzoic acid (2,5-DHBA, gentisic acid) and 2,3-dihydroxybenzoic acid (2,3-DHBA), which accounts for less than 1% of the salicylic acid metabolites. The 2,5-DHBA can undergo conjugation with glycine to

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*Correspondence to:* Dr G. Jayarama Bhat, Department of Pharmaceutical Sciences, South Dakota State University College of Pharmacy and Allied Health Professions, Avera Health and Sciences Center, 1055 Campanile Avenue, Brookings, SD 57007, USA  
E-mail: jayarama.gunaje@sdsstate.edu

*Present address:* <sup>3</sup>University of Kentucky, Lexington, KY40546, USA

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form gentisuric acid. These metabolites are eliminated from the body either through the kidneys or bile (10-12).

Our laboratory has been investigating the hypothesis that aspirin and salicylic acid may target cell cycle regulatory proteins to exert their chemopreventive effects. Cyclin dependent kinases (CDKs) are key regulators of the cell cycle, and are activated by binding with temporally expressed cyclins during various stages of the cell cycle. The important CDKs in mammalian cell cycle regulation are: CDK1, CDK2, CDK4 and CDK6. CDK4 and CDK6, through binding to cyclin D, facilitate the progression through G1, whereas, CDK2 via binding to cyclin E helps to transition from G1 to S phase. CDK1 and CDK2 get activated via binding to cyclins A and B, which facilitate the progression through G2 and M phases. In many cancers, CDKs are deregulated and cyclins are over expressed (13). Due to their important regulatory role in cell cycle progression, CDKs have been attractive targets in cancer treatment (14-16), in this regard only two CDK inhibitors (palbociclib and ribociclib) specific to CDK4 and 6 have been approved for cancer treatment, which are classified as ATP-competitive inhibitors (17,18).

In our previous study, we demonstrated that exposure of colon cancer cells, to aspirin and salicylic acid caused down-regulation of cyclin A2, B1 and D3, and CDKs 1, 2, 4 and 6 (19). In that study, we also demonstrated that salicylic acid potentially binds to CDK2 via interactions with Asp145, and Lys33, both of which are found in the active site of the enzyme. Despite these potential interactions, salicylic acid failed to inhibit the CDK2 enzyme activity in an *in vitro* kinase assays. In this study, we considered the possibility that the two salicylic acid metabolites 2,3-DHBA and 2,5-DHBA, which are known to be produced through CYP450 catalyzed reactions may inhibit CDK enzyme activity. The goal of the present study was to determine the effect of salicylic acid metabolites (2,3-DHBA and 2,5-DHBA) and also derivatives (2,4-DHBA, 2,6-DHBA and 2,4,6-trihydroxybenzoic acid) on CDK activity *in vitro*, and determine the potential sites of interactions with CDK-1.

Herein, we report that salicylic acid metabolites 2,3-DHBA and 2,5-DHBA, as well as the salicylic acid derivatives 2,4-DHBA and 2,6-DHBA, inhibited CDK1 enzyme activity. The trihydroxy derivative of salicylic acid, 2,4,6-THBA, was also highly effective. While 2,3-DHBA and 2,5-DHBA are produced from the oxidation of salicylic acid through CYP450 metabolism, they are also reported to be naturally present in several fruits and medicinal herbs. This is the first report demonstrating the ability of salicylic acid metabolites, and naturally occurring salicylic acid derivatives to inhibit CDK activity, suggesting their potential role in chemoprevention.

## Materials and methods

**Cell lines.** HCT-116 human colorectal carcinoma cell line was purchased from American Type Culture Collection (ATCC). The cells were cultured in McCoy's 5A medium containing 10% FBS for 24 h before treatment with specified compounds for indicated times (20). Authentication of cell lines was done by ATCC through their DNA-STR profile.

**Reagents.** Aspirin, salicylic acid, 2,4,6-THBA, 3,4,5-THBA and trypsin-EDTA solution were purchased from Sigma (St. Louis,

MO, USA); Immobilon membranes, H1 Histones from EMD Millipore (Billerica, MA, USA);  $^{32}\text{P}$   $\gamma$ -ATP were from MP Biochemical; Super Signal™ West Pico Chemiluminescent Substrate, 2,3-DHBA, 2,4-DHBA, 2,5-DHBA, 2,6-DHBA, 3,4-DHBA, 3,5-DHBA, 2,4,6-THBA and protease inhibitor tablets and all other chemicals were obtained from Thermo Fisher Scientific, Inc. (Waltham, MA, USA).

**Recombinant proteins and antibodies.** Anti-CDK1 and anti- $\beta$  actin antibodies were purchased from Cell Signaling Technology (Danvers, MA, USA); Anti-acetyl lysine conjugated agarose beads from Immune Chem (Burnaby, BC, Canada); goat anti-rabbit and goat anti-mouse antibodies were obtained from Bio-Rad (Hercules, CA, USA). CDK1/cyclin B1 active enzyme from New England Biolabs (NEB) (Ipswich, MA, USA), CDK1/cyclin B1, CDK2/cyclin A2, CDK4/cyclin D1, CDK6/cyclin D1, Retinoblastoma (C-term) and kinase buffer were purchased from SignalChem (Richmond, BC, Canada).

**Cell lysate preparation and western blotting.** After treatment with aspirin at the indicated concentrations, cells were washed with phosphate buffered saline and scraped into lysis buffer and protein extracts were prepared as previously described (21). Samples containing 50  $\mu\text{g}$  of protein were separated by 8% polyacrylamide gel electrophoresis (PAGE) and immunoblotted with respective antibodies. For immunoprecipitations, 500  $\mu\text{g}$  of the protein was immunoprecipitated with anti-acetyl lysine conjugated agarose beads overnight, immunocomplexes were collected, and immunoblotted with anti-CDK1 antibody. Intensities of the bands were quantified using NIH ImageJ software.

**In vitro CDK assay.** *In vitro* CDK assays were performed as described by the protocols from NEB and SignalChem. Briefly, purified kinase was aliquoted into the reaction buffer provided by the respective kits and incubated with indicated compounds at various concentrations for 10 min at room temperature. Kinase reaction were performed by incubating the enzyme with a kinase buffer containing 15  $\mu\text{M}$  ATP, 2  $\mu\text{Ci}$  of [ $\gamma$ - $^{32}\text{P}$ ]ATP, 5  $\mu\text{g}$  of H1 Histone or retinoblastoma, at 30°C for 20 min. The final volume of the reaction was 50  $\mu\text{l}$ , the reactions were stopped by adding EDTA to a final concentration of 20 mM and 4X loading buffer. The samples were boiled for 10 min, loaded on to a 10% SDS-PAGE. The gel was stained using Coomassie Brilliant blue (R), dried and exposed to X-ray film. Intensities of the bands were quantified using NIH ImageJ software.

**MTT Assay.** Cytotoxicity was determined by MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assay. Briefly, HCT-116 cells were seeded in 24-well plates overnight at a density of 20,000 cells/well and treated with various compounds for 72 h. MTT assays were carried out as previously described (22).

**Molecular docking studies.** The crystallographic three dimensional structures of selected target proteins of CDK1 (4y72 A chain) and cyclin b1 (PDB ID: 2b9r) were retrieved from the Protein Data Bank (PDB) <http://www.pdb.org>. The

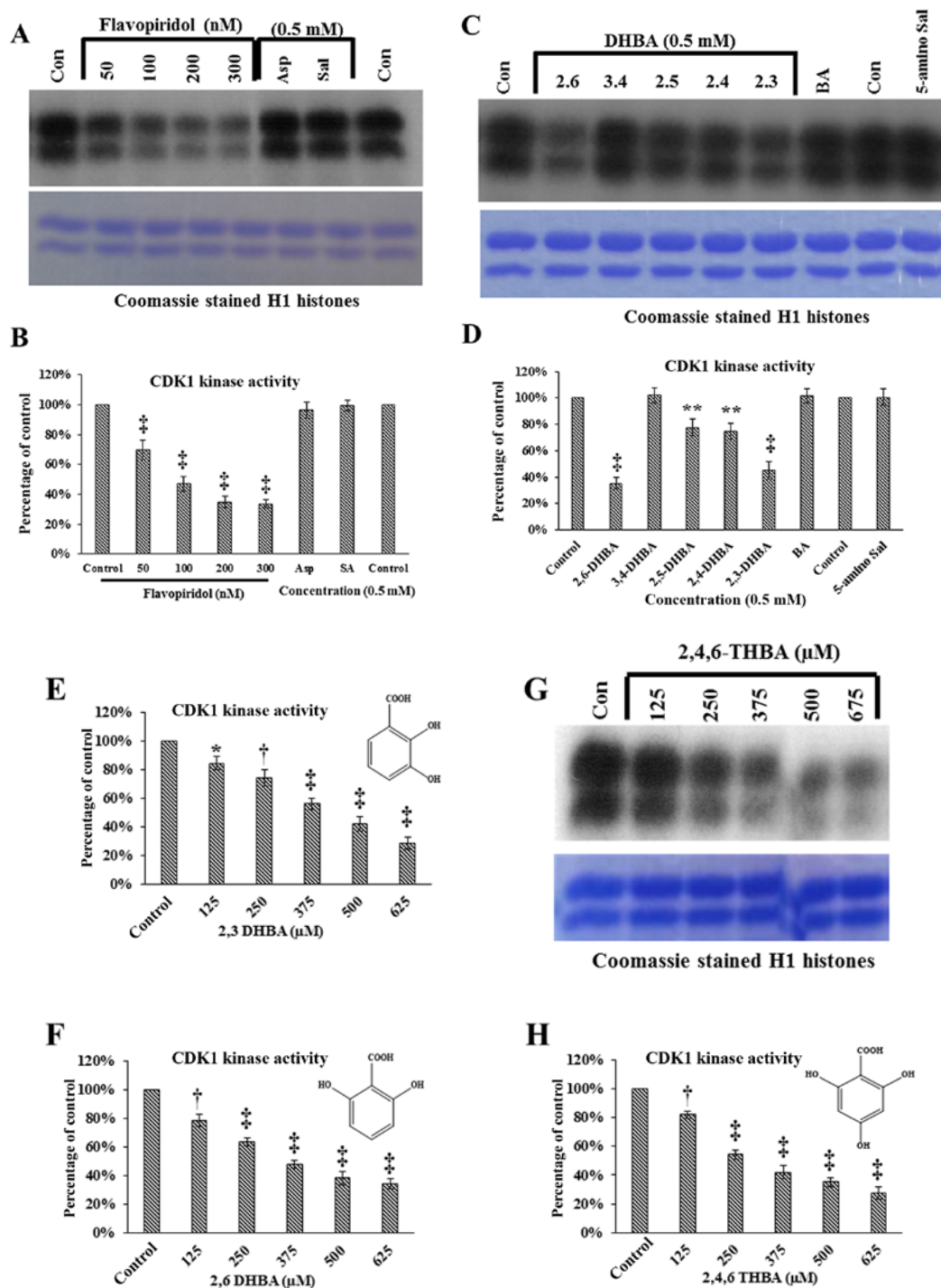


Figure 1. *In vitro* kinase assays showing the effect of aspirin, salicylic acid metabolites and derivatives on CDK1 enzyme activity. (A) Effect of aspirin (Asp), salicylic acid (Sal) and flavopiridol on CDK1 enzyme activity. (B) Quantification of the blot in (A). (C) Effect of salicylic acid metabolites 2,3-DHBA, 2,5-DHBA and derivatives 2,4-DHBA, 2,6-DHBA, 3,4-DHBA, 5-amino salicylic acid (5-amino Sal), and benzoic acid (BA) on CDK1 enzyme activity. The lower panels below (A and C) shows coomassie stained H1 histones. (D) quantification of the blot in (C). (E and F) The dose-dependent effect of 2,3-DHBA and 2,6-DHBA on CDK1 enzyme activity. (G) Dose-dependent effect of 2,4,6-THBA on CDK1. The lower panel of (G), shows coomassie stained pattern of histone (H). (H) Quantification of the blot in (G). The intensities of bands in various blots were quantified and expressed as percentage of control. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .

human CDK1, Cyclin b1 and CDK1\_cyclin b1 complex (PDB ID: 2b9r and 4y72) were selected for energy minimization using Gromacs 3.3.1 package with the GROMOS96 force field (23). These molecules were used as the receptor for virtual small molecule docking with ligand aspirin, salicylic acid, 2,3-DHBA, 2,5-DHBA, 2,6-DHAB and 2,4,6 THBA

using AutoDockVina. The molecular results were visualized by PYMOL molecular graphics system version 1.3.

**Statistical analysis.** All experiments were repeated 3-6 times independently of each other. One-way ANOVA followed by Tukey's post-hoc multiple comparison tests were adopted to

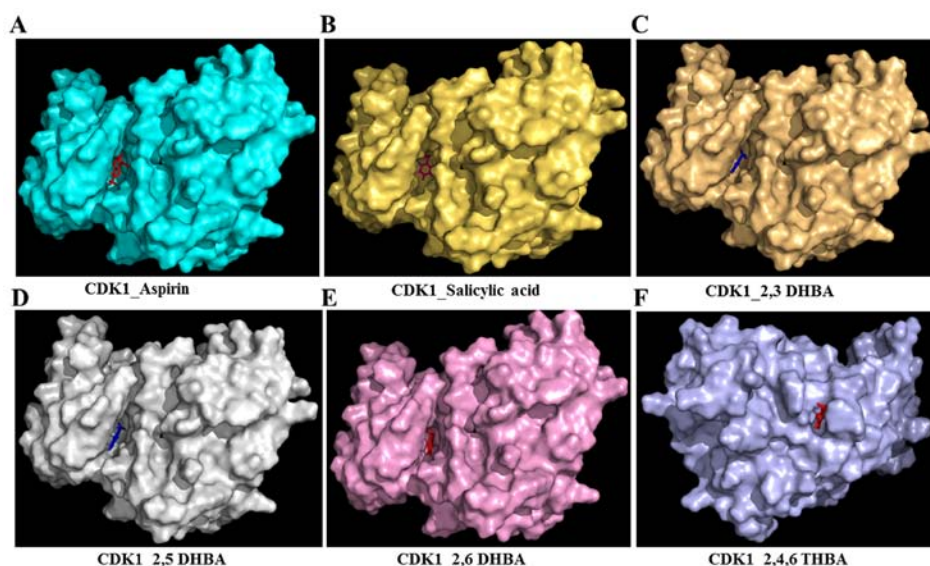


Figure 2. Molecular docking studies (space-filling model) showing the potential binding pockets in CDK1 for aspirin (A), salicylic acid (B), 2,3-DHBA (C), 2,5-DHBA (D), 2,6-DHBA (E) and 2,4,6-THBA (F).

compare group differences to control, and significance was defined at  $P < 0.05$ .

## Results

*Effect of aspirin, salicylic acid and salicylic acid metabolites and derivatives on CDK1 kinase activity in vitro.* In a previous study, we demonstrated that aspirin potentially binds to CDK2 through Lys33 and salicylic acid via Asp145 and Lys33. Despite the strong evidence of interactions of these drugs with CDK2, both failed to inhibit CDK2 enzyme activity in *in vitro* kinase assays (19). In the current study, we initially extended these observations to determine the effect of aspirin and salicylic acid on CDK1 enzyme activity *in vitro*. This was tested using commercially obtained CDK1/cyclin B1 assay kits, and by providing  $[\gamma\text{-}^{32}\text{P}]\text{ATP}$  and H1 histones as substrates. As a positive control, we used flavopiridol, a known inhibitor of CDK1 (24). Fig. 1A demonstrates that neither aspirin nor salicylic acid inhibited CDK1 activity as measured by the ability to phosphorylate H1 histones; however, flavopiridol dose-dependently inhibited CDK1 activity (upper panel). These samples contained similar levels of H1 histones (lower panel, Fig. 1A). Quantification of the bands in Fig. 1A (upper panel) are shown in Fig. 1B. These results show that aspirin and salicylic acid do not inhibit CDK1, consistent with the results previously obtained with CDK2 (19).

We next determined the ability of salicylic acid metabolites (2,3-DHBA and 2,5-DHBA) as well as salicylic acid derivatives (2,4-DHBA, 2,6-DHBA and 3,4-DHBA and 5-aminosalicylic acid) and benzoic acid to inhibit CDK1 enzyme activity. We observed that, of the seven different compounds tested, all DHBA compounds with a common -OH group at the 2nd carbon inhibited CDK1 enzyme activity to varying degrees (Fig. 1C, upper panel). It is interesting to note that the salicylic acid metabolites, 2,3-DHBA and 2,5-DHBA, both showed inhibitory effects on CDK1 activity. Quantification of the intensities of the bands showed that, four compounds showed varied levels of inhibition (2,6-DHBA,

65%; 2,5-DHBA, 22%; 2,4-DHBA, 25% and 2,3-DHBA, 55%) (Fig. 1D). Three compounds, 3,4-DHBA, 5-aminosalicylic acid and benzoic acid did not show any inhibition. In addition, 3,5-DHBA failed to inhibit the CDK1 enzyme activity (data not shown).

*Dose-dependent inhibition of CDK1 enzyme activity by 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA.* In experiments described in Fig. 1C, the salicylic acid metabolite 2,3-DHBA, and the salicylic acid derivative 2,6-DHBA, showed greater inhibition on CDK1 activity as compared to other compounds tested. Therefore, we focused initially on these two compounds to study their dose-dependent effects on CDK1 activity. Fig. 1E and F show that both 2,3-DHBA and 2,6-DHBA dose-dependently inhibit CDK1 activity with an  $\text{IC}_{50}$  of 386  $\mu\text{M}$  and 365  $\mu\text{M}$ , respectively. The ability of 2,3-DHBA and 2,6-DHBA to inhibit CDK1 enzyme activity suggested that the hydroxyl group at the 2nd and 6th carbon in these compounds may be important, and that a compound having these two -OH groups may more potently inhibit CDK1 activity. Therefore, in *in vitro* kinase assays, we determined the effect of 2,4,6-THBA on CDK1 activity at various concentrations. Fig. 1G (upper panel) demonstrates that 2,4,6-THBA inhibited CDK1 activity with an  $\text{IC}_{50}$  of 226  $\mu\text{M}$ . Quantification of the bands are shown in Fig. 1H. Of note, 3,4,5-THBA (gallic acid) did not show any inhibitory effect (data not shown). This again suggests that the -OH group at the 2nd and 6th carbon may contribute to greater inhibition of CDK1 enzyme activity.

*Molecular docking studies show potential interactions of aspirin, salicylic acid, salicylic acid metabolites and derivatives with CDK1 and cyclin B1.* We used AutoDockVina to understand the interactions between aspirin, salicylic acid, 2,3-DHBA, 2,5-DHBA, 2,6-DHBA and 2,4,6-THBA with CDK1 alone, cyclin B1 alone, and the CDK1/cyclin B1 complex. The binding free energy and hydrogen bond lengths were also determined. The results of the docking studies are shown in Table I, Fig. 2A-F (space-filling model) and Figs. 3-5.

Table I. Molecular docking studies of CDK1/Cyclin B1 with metabolites/derivatives of aspirin.

S. no.	Ligand	Receptor	PDB ID	Interacting amino acids	No. of H-bonds	Measurement (Å <sup>0</sup> )	Energy value (kcal/mol)	Interactions
1	Aspirin	CDK1	4y72 (A Chain)	Asn133, Gln132 and Tyr15	3	2.7, 2.1 and 2.6	-7.4	-COOH - Asn133, Gln132 and Tyr15
2	Salicylic acid	CDK1	4y72 (A chain)	Asp146 and Lys 33	2	1.8 and 2.5	-8.6	-OH - Lys33; -COOH - Asp146
3	2,3-DHBA	CDK1	4y72 (A Chain)	Asp146	1	2.0	-6.8	-COOH - Asp146
4	2,5-DHBA	CDK1	4y72 (A Chain)	Asp146	1	2.1	-6.8	-COOH - Asp146
5	2,6-DHBA	CDK1	4y72 (A Chain)	Asp146	1	2.0	-7.1	6th -OH - Asp146
6	2,4,6-THBA	CDK1	4y72 (A Chain)	Arg123, Arg151 and Gly154	3	2.5, 2.0 and 2.4	-5.8	2nd -OH - Arg151 and Gly154; -COOH -Arg123
7	Aspirin	Cyclin B1	2b9r	Arg135	2	1.9 and 2.5	-6.8	-COOH Arg135
8	Salicylic acid	Cyclin B1	2b9r	Arg135 and Tyr60	4	2.2, 2.4, 2.7 and 2.3 Å	-7.4	-OH - Arg135; -COOH - Tyr60, Arg135
9	2,3-DHBA	Cyclin B1	2b9r	Arg135, Arg38 and Asp67	6	2.1, 2.2, 2.2, 2.2, 2.4 and 2.3	-6.8	2nd -OH - Asp67; 3rd - OH -COOH - Arg38; Arg135
10	2,5-DHBA	Cyclin B1	2b9r	Arg135	2	2.0 and 2.4	-6.7	2nd -OH - Arg135
11	2,6-DHBA	Cyclin B1	2b9r	Arg135	2	2.1 and 2.5	-5.4	6th -OH - Arg135
12	2,4,6-THBA	Cyclin B1	2b9r	Arg135, Phe175, Asp67, Arg38, Thr166	6	2.2, 2.4, 2.2, 2.2, 2.3 and 2.1	-6.8	2nd -OH - Asp67; 4th -OH - Arg38, Thr166; 6th -OH - Asp67; -COOH - Arg135
13	Aspirin	CDK1_ Cyclin B1	4y72 (A, B chain)	Tyr15 (CDK1)	1	2.5	-7.0	-COOH - Tyr15 (CDK1)
14	Salicylic acid	CDK1_ Cyclin B1	4y72 (A, B chain)	Asp146 and Lys33 (CDK1)	3	1.9, 2.5 and 2.6	-9.6	-OH - Asp146; -COOH - Lys33, Asp146 (CDK1)
15	2,3-DHBA	CDK1_ Cyclin B1	4y72 (A, B chain)	Thr329, Val336, Phe338 (Cyclin B1)	2	2.4, 2.1 and 2.4	-6.1	2nd and 3rd -OH - Phe338; 3rd -OH Val336; -COOH - Thr329 (cyclin B1)
16	2,5-DHBA	CDK1_ Cyclin B1	4y72 (A, B chain)	Arg123 (3), Leu125, and Arg151 (CDK1)	5	2.2, 2.2, 2.4, 2.4 and 2.4	-6.0	5th -OH - Leu125 and Arg151; -COOH - Arg123 (CDK1)
17	2,6-DHBA	CDK1_ Cyclin B1	4y72 (A, B chain)	Val336 (Cyclin B1)	1	2.6	-5.6	6th -OH - Val336 (cyclin B1)
18	2,4,6-THBA	CDK1_ Cyclin B1	4y72 (A, B chain)	Tyr223, Asp 230, and Arg201 (Cyclin B1)	2	2.3, 2.5 and 2.5	-5.4	2nd -OH - Asp230; 4th -OH - Tyr223; -COOH - Arg201 (cyclin B1)

Free energy binding values and hydrogen bond lengths for the interaction of aspirin, salicylic acid, 2,3-DHBA, 2,5-DHBA, 2,6-DHBA and 2,4,6-THBA with CDK1, cyclin B1 and CDK1/cyclin B1 complex.

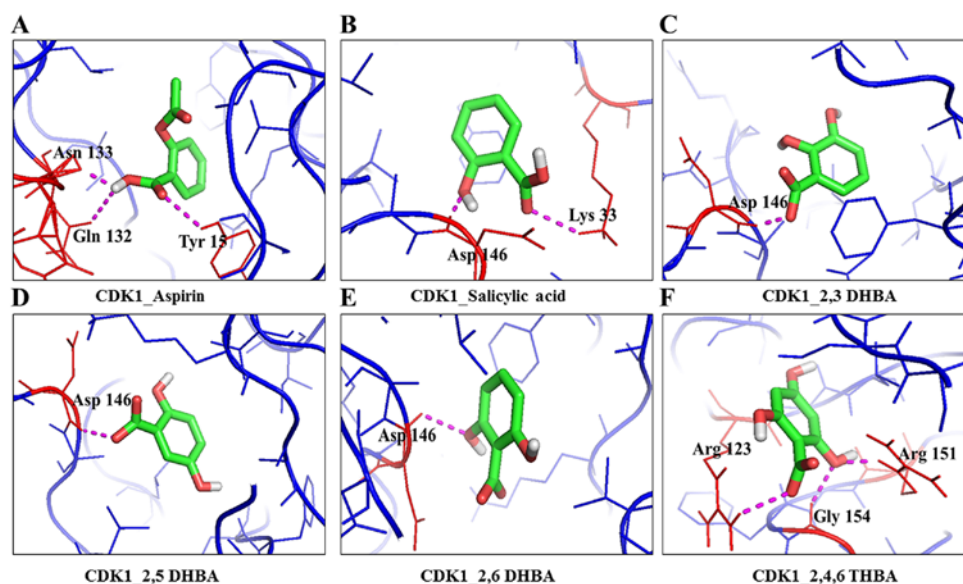


Figure 3. Molecular docking studies showing the potential interactions of aspirin (A), salicylic acid (B), 2,3-DHBA (C), 2,5-DHBA (D), 2,6-DHBA (E) and 2,4,6-THBA (F) with CDK1.

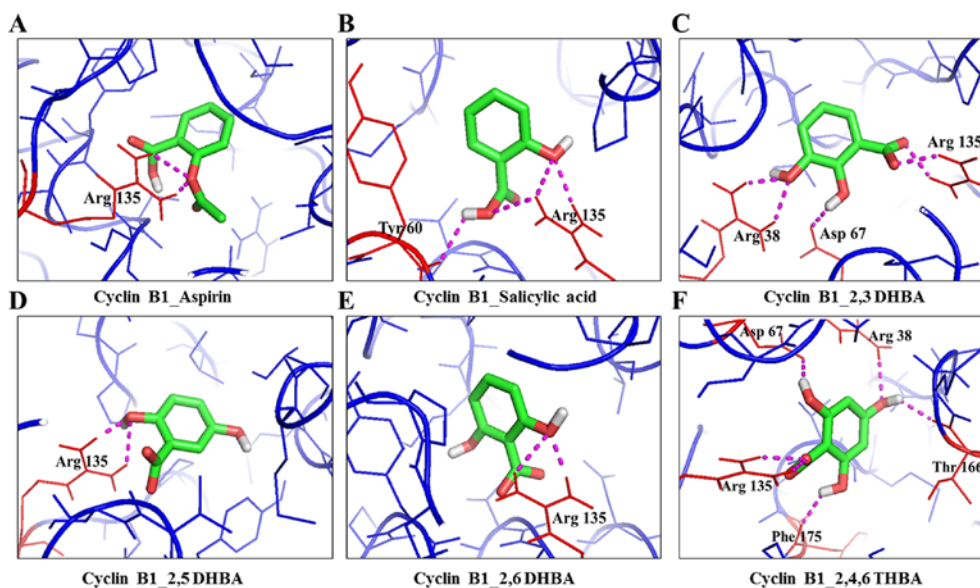


Figure 4. Molecular docking studies showing the potential interactions of aspirin (A), salicylic acid (B), 2,3-DHBA (C), 2,5-DHBA (D), 2,6-DHBA (E) and 2,4,6-THBA (F) with cyclin B1.

Among the potential interactions shown in Table I, salicylic acid interactions with CDK1 through Asp146 and Lys33 is significant as these two amino acids are part of the active site of the enzyme and are conserved among different CDK family members (25). Noteworthy, salicylic acid also interacts with CDK1 through Asp146 and Lys33 when it is present as part of the cyclin B1/CDK1 complex (Table I). Docking studies also revealed that aspirin can potentially interact with Tyr-15 in the CDK1 monomer, or in the CDK1/cyclin B1 dimer. It is interesting to note that Tyr15 is conserved in various CDKs, and its phosphorylation by Wee1/Mik1 kinase family inactivates the enzyme activity (26). Despite these predicted interactions of aspirin and salicylic acid with key amino acid residues in CDK1, both failed to inhibit CDK1 enzyme activity (Fig. 1).

Table I also shows that all three DHBA compounds (2,3-DHBA, 2,5-DHBA and 2,6-DHBA) showed potential interactions with CDK1 monomer through Asp 146, whereas 2,4,6-THBA interacted via Arg123, Arg151 and Gly154. Fig. 2A-F (space-filling model) shows that 2,3-DHBA, 2,5-DHBA and 2,6-DHBA bind to the same pocket in CDK1 (same as aspirin and salicylic acid); however, 2,4,6-THBA binds at a different site. Interestingly, all four compounds (DHBA/THBA) inhibited CDK1 enzyme activity (Fig. 1C-H).

*Aspirin acetylates recombinant CDK1, and pre-incubation with salicylic acid, 2,3-DHBA, 2,6-DHBA or 2,4,6-THBA inhibits aspirin's ability to acetylate CDK1.* We performed experiments to provide a second line of evidence for the

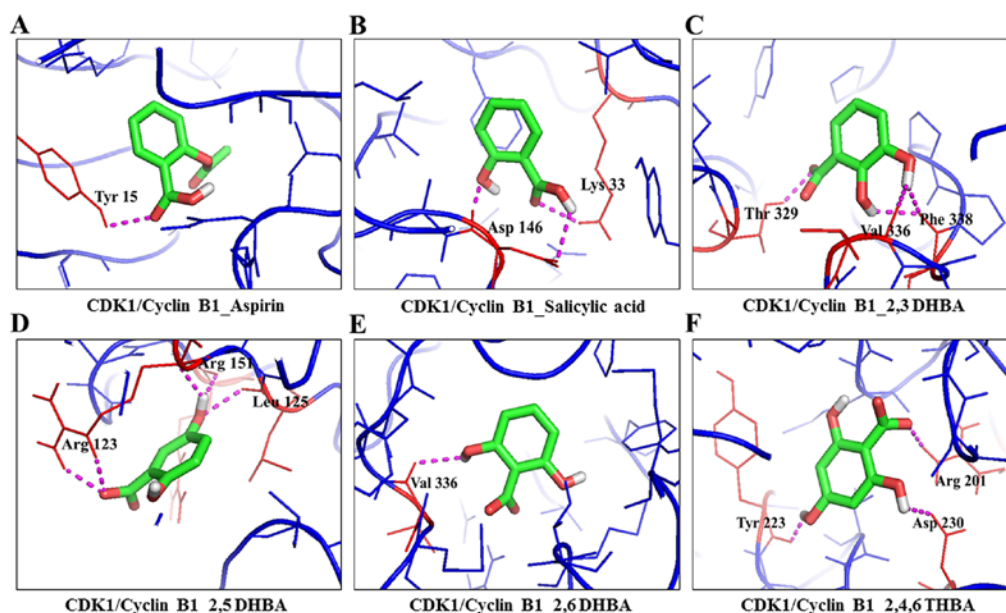


Figure 5. Molecular docking studies showing the potential interactions of aspirin (A), salicylic acid (B), 2,3-DHBA (C), 2,5-DHBA (D), 2,6-DHBA (E) and 2,4,6-THBA (F) with CDK1/cyclin B1 complex.

physical interactions of aspirin, salicylic acid, 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA with CDK1. We hypothesized that if aspirin binds to CDK1, it should be able to acetylate CDK1 and this could be detected by immunoblotting with anti-acetyl lysine antibody. If this is true, then pre-incubation of CDK1 with salicylic acid, 2,3-DHBA, 2,6-DHBA or 2,4,6-THBA should prevent aspirin-induced CDK1 acetylation. Therefore, first, we tested the ability of aspirin at various concentrations to acetylate recombinant CDK1 (Prospec) *in vitro*. For this, recombinant CDK1 (300 ng) was incubated with aspirin at various concentrations for 12 h, and then samples were immunoblotted with anti-acetyl lysine antibody. Fig. 6A demonstrates that aspirin dose-dependently acetylated recombinant CDK1 beginning at 0.25 mM (upper panel). Stripping and reprobing the blot showed similar amounts of CDK1 protein in all lanes (Fig. 6A, lower panel).

The ability of aspirin to acetylate CDK1 as shown in Fig. 6A, supports the results obtained from molecular docking studies which suggested that aspirin potentially binds to CDK1 (Table I). Since salicylic acid also potentially binds to CDK1, we next determined if pre-incubation of CDK1 with salicylic acid would prevent aspirin's ability to acetylate CDK1. For this, recombinant CDK1 was left untreated, or treated with aspirin alone (1.5 mM), or first pre-incubated with different concentrations of salicylic acid for 10 min (0.01-0.25 mM) and then treated with aspirin (1.5 mM) for 12 h. The samples were then immunoblotted with anti-acetyl lysine antibody. Fig. 6B demonstrates that aspirin caused acetylation of recombinant CDK1; however, this was dose-dependently prevented by pre-incubation with salicylic acid (upper panel). Reprobing the blot in Fig. 6B showed that all lanes contained equal amounts of CDK1 protein (lower panel).

In similar experiments as mentioned above, we next determined whether pre-incubation of recombinant CDK1 with 2,3-DHBA and 2,6-DHBA would prevent aspirin-mediated acetylation of CDK1. Recombinant CDK1 was pre-incubated

with 0.5 mM 2,3-DHBA and 2,6-DHBA for 10 min; then 1.5 mM, aspirin was added and the samples were incubated for 12 h at room temperature. The samples were analyzed by immunoblotting with anti-acetyl lysine antibodies. Fig. 6D (upper panel), demonstrates that pre-incubation of CDK1 with 2,3-DHBA or 2,6-DHBA significantly prevented aspirin-mediated acetylation of CDK1. In separate experiments, we also observed that pre-incubation of recombinant CDK1 with 2,4,6-THBA prevented aspirin-mediated acetylation of CDK1 (Fig. 6E, upper panel). These results collectively demonstrate that aspirin, salicylic acid, 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA can all directly bind to CDK1, and support the data obtained from molecular docking studies (Table I).

*Aspirin acetylates cellular CDK1.* We also performed experiments to determine if exposure of HCT-116 cells to aspirin would acetylate cellular CDK1. Cells were treated with aspirin at various concentrations (0.25-2.5 mM) for 12 h, and lysates prepared and immunoprecipitated with anti-acetyl lysine antibody agarose conjugates and immunoblotted with anti-CDK-1 antibody. Fig. 6F, upper panel, demonstrates that CDK1 was acetylated in aspirin treated samples in a dose-dependent fashion. We also analyzed equal amounts of proteins representing these samples by immunoblotting with anti-CDK1 antibody. Fig. 6F, lower panel, shows that these samples contained similar amounts of CDK1 protein.

*Effect of aspirin, salicylic acid, 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA on CDK2, CDK4 and CDK6 enzyme activity.* Having demonstrated the ability of 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA to inhibit CDK1 enzyme activity *in vitro*, we sought to determine if these compounds would also inhibit CDK2, 4 and 6 enzyme activity. Commercially available CDK2 enzymes were pre-incubated with these compounds for 10 min, and then *in vitro* kinase assays were carried out by providing H1 histones as substrates. Fig. 7A and B demonstrates that aspirin, salicylic

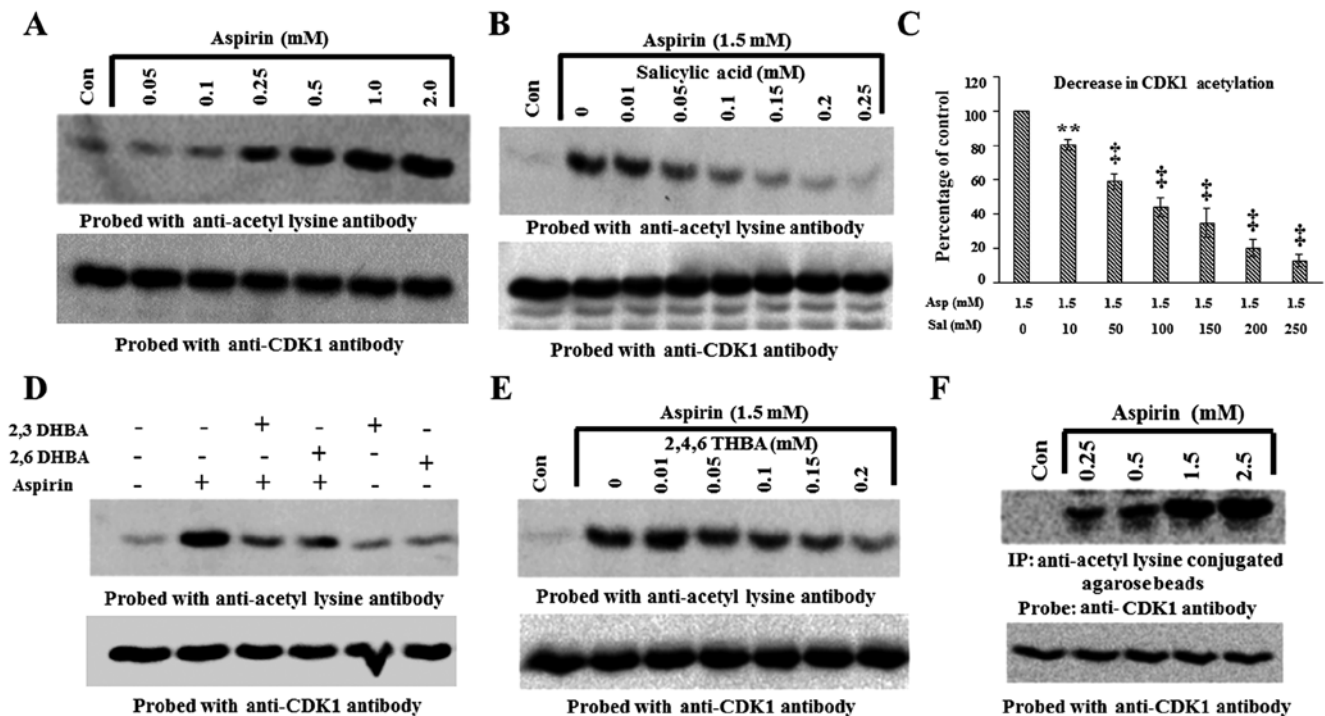


Figure 6. Aspirin acetylates recombinant CDK1 *in vitro* and cellular CDK1 in HCT116 cells. (A) Upper panel, dose-dependent effect of aspirin on recombinant CDK1 acetylation. (B) Salicylic acid pre-incubation prevents aspirin-induced acetylation (upper panel). (C) Quantification of blot in (B), the decrease in CDK1 acetylation is expressed as a percentage of the control. (D) Pre-incubation of recombinant CDK1 with 2,3-DHBA and 2,6-DHBA at 0.5 mM prevents aspirin induced acetylation of CDK1 (upper panel). (E) Pre-incubation of recombinant CDK1 with 2,4,6-THBA prevents aspirin induced acetylation of CDK1 (upper panel). (F) Aspirin acetylates cellular CDK1 in HCT-116 cells (upper panel). (F) Lower panel, equal amounts of proteins was immunoblotted with anti-CDK1 antibody. The lower panels in all other blots (A, B, D and E), the upper panel blots were stripped and reprobbed with anti-CDK1 antibody. \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .

acid, 2,3-DHBA and 2,6-DHBA did not inhibit CDK2 activity. However, 2,4,6-THBA showed dose-dependent inhibition of CDK2 enzyme activity (Fig. 7C and D). The  $IC_{50}$  for the inhibition of CDK2 by 2,4,6-THBA was  $\sim 300 \mu M$  (Fig. 7D).

We next performed experiments to determine the effect of aspirin, salicylic acid, 2,3-DHBA, 2,6-DHBA and 2,4,6-DHBA on CDK4 and CDK6 enzyme activity in *in vitro* kinase assays similar to the experiments performed in Fig. 7A except that retinoblastoma (Rb) protein was provided as substrate for phosphorylation. Fig. 7E and F demonstrates that aspirin, salicylic acid, 2,3-DHBA and 2,6-DHBA, all failed to inhibit CDK4; however, 2,4,6-DHBA inhibited CDK-4 activity (45%). Fig. 7G and H demonstrates that aspirin and salicylic acid did not affect CDK6 activity; interestingly, 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA significantly inhibited the CDK6 activity by 28, 37 and 40%, respectively.

## Discussion

Since the discovery of aspirin in 1897, many of its therapeutic properties and mechanisms of actions have been identified. It is used for a variety of illnesses including inflammation, fever, pain, and for its anti-platelet properties. These actions of aspirin have been attributed mainly to acetylation-mediated inhibition of cyclooxygenases (COX). There are two COX enzymes: COX-1 and COX-2. Aspirin's anti-platelet effect occurs through inhibition of COX-1 ( $IC_{50}$ - $1.67 \mu M$ ); whereas, its analgesic and anti-inflammatory effects occur primarily

through inhibition of COX-2 ( $IC_{50}$ - $278 \mu M$ ) (27). In recent years, evidence has emerged establishing aspirin's ability to decrease cancer incidence, igniting a renewed interest in its research and use. Despite extensive studies, aspirin's mode of action in cancer prevention, as well as how it primarily prevents CRC as compared to other cancers, is not clearly established. Although multiple targets and signaling pathways have been proposed, a unifying mechanism has not been identified till date, suggesting that different mechanisms may be responsible in different cancers or that the actual mechanism is yet to be discovered.

One of the hypotheses proposed for aspirin's anticancer effects is acetylation-mediated inactivation of COX, as upregulation of COX-2 is observed in nearly 80-90% of CRC; in contrast, COX-1 expression is unaffected (3). The observation that low dose aspirin, used for its cardio-protective effect, is also effective in the prevention of CRC, led to the hypothesis that a common pathway involving COX-1 inhibition in platelets may be central to both effects (28-30). Since low dose aspirin is insufficient to achieve complete inhibition of COX-2, it is argued that aspirin-mediated inhibition of platelet COX-1 plays a role in the prevention of CRC, and that aspirin's effects may occur through sequential steps involving COX-1 and COX-2. It is hypothesized that low dose aspirin by inhibiting COX-1 in platelets prevents their activation and the release of growth factors and lipid mediators, required for COX-2 expression and tumorigenesis in the colonic tissue. While this is a tenable hypothesis, it is yet to be confirmed.



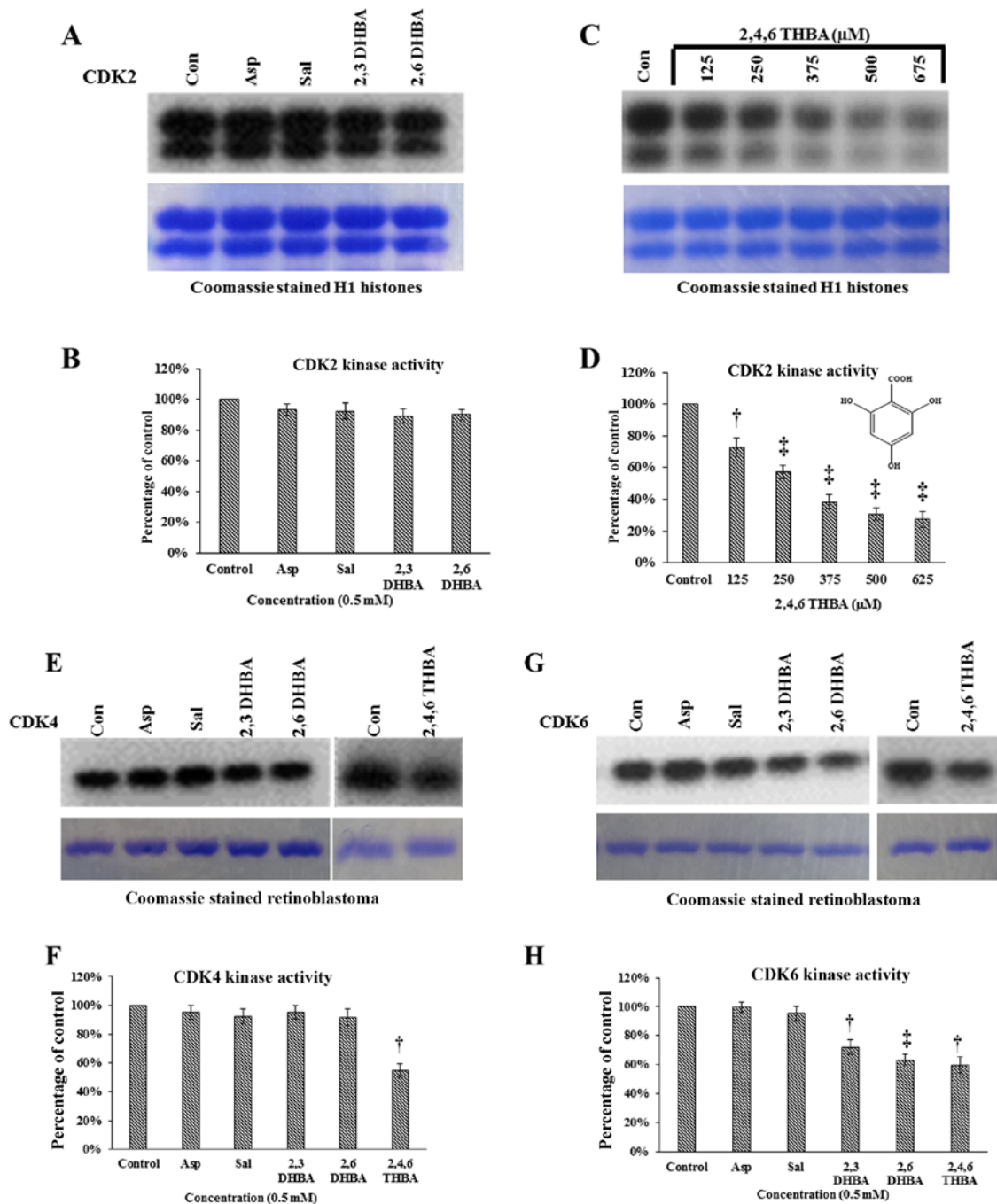


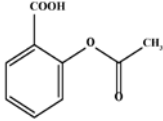
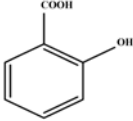
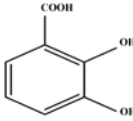
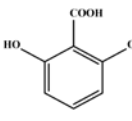
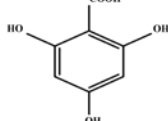
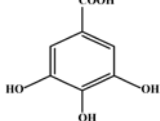
Figure 7. Effect of aspirin, salicylic acid, 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA (at 0.5 mM) on CDK2, 4 and 6 enzyme activity. (A) Upper panel represents *in vitro* kinase assays showing the effect of aspirin, salicylic acid, 2,3-DHBA and 2,6-DHBA on CDK2 activity. Lower panel shows coomassie stained H1 histones. (B) Quantification of the blot in (A). (C) Upper panel represents *in vitro* kinase assays showing the dose-dependent effect of 2,4,6-THBA on CDK2 activity. Lower panel shows coomassie stained H1 histones. (D) Quantification of the blot in (C). (E) Upper panel represents the effect of aspirin, salicylic acid, 2,3-DHBA and 2,6-DHBA and 2,4,6-THBA on CDK4 activity. Lower panel shows coomassie stained retinoblastoma protein. (F) Quantification of the blot in (E). (G) Upper panel represents the effect of aspirin, salicylic acid, 2,3-DHBA and 2,6-DHBA (0.5 mM) and 2,4,6-THBA on CDK6 enzyme activity. Lower panel shows coomassie stained retinoblastoma protein. (H) Quantification of the blot in (G). \*P<0.05, \*\*P<0.01, †P<0.001, ‡P<0.001.

A recent study demonstrated in human intestinal mucosal cells that low-dose aspirin caused acetylation of COX-1 and inhibited PGE2 synthesis leading to decreased levels of S6 kinase, implicating this in the prevention of early colorectal carcinogenesis (31). In addition, salicylic acid, the hydrolyzed product of aspirin, has also been implicated in aspirin's chemopreventive effects. Salicylic acid was shown to bind to a number of cellular proteins (Salicylic Acid Binding Proteins

or SABPs) such as I $\kappa$ B kinase (IKK), a component of the NF- $\kappa$ B complex (32), AMP activated protein kinase (33), High Mobility Group Box 1 proteins and GAPDH (34), and CDK2 (19), affecting their levels and/or functional activity.

Although specific protein targets have been identified for both aspirin and salicylic acid, no studies have reported on the cellular targets for salicylic acid metabolites. One of the most important findings of this study is the ability of salicylic acid

Table II. Effect of aspirin, salicylic acid, 2,3-DHBA, 2,6-DHBA, 2,4,6-THBA and 3,4,5-THBA on CDK-1, 2, 4 and 6.

						
CDKs	Aspirin	Salicylic acid	2,3-DHBA	2,6-DHBA	2,4,6-THBA	3,4,5-THBA
1	-	-	+	+	+	-
2	-	-	-	-	+	-
4	-	-	-	-	+	-
6	-	-	+	+	+	-

Effect of aspirin, salicylic acid 2,3-DHBA, 2,6-DHBA, 2,4,6-THBA and 3,4,5-THBA on CDK-1, 2, 4 and 6. Inhibitory effects are shown by plus sign; lack of inhibitory effect is shown by minus sign.

Table III. *In vitro* IC<sub>50</sub> determination for 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA using purified CDK1 enzyme and MTT assay in HCT-116 cells.

Compounds	CDK1 IC <sub>50</sub> ( $\mu$ M)	MTT assay HCT-116 ( $\mu$ M) 72 h
2,3-DHBA	386.1	1778.28
2,6-DHBA	365.2	1949.84
2,4,6-THBA	226.0	785.236

IC<sub>50</sub> values for CDK1 inhibition *in vitro* and cytotoxicity assays in HCT-116 cell.

metabolites, 2,3-DHBA and 2,5-DHBA, as well as other derivatives such as 2,4-DHBA, and 2,6-DHBA to inhibit CDK1 enzyme activity *in vitro* (Fig. 1). Among these, 2,3-DHBA (IC<sub>50</sub> 386  $\mu$ M) and 2,6-DHBA (IC<sub>50</sub> 365  $\mu$ M) showed greater degree of inhibition as compared to 2,4-DHBA and 2,5-DHBA. The ability of 2,3-DHBA and 2,5-DHBA to inhibit CDK1 activity is highly significant, because they are known to be generated from salicylic acid metabolism through CYP450-mediated reactions (11). Additionally, in this study, we also screened two THBA derivatives, 3,4,5-THBA and 2,4,6-THBAs, for their ability to inhibit CDK1. We observed that 3,4,5-THBA failed to inhibit; however, 2,4,6-THBA (IC<sub>50</sub>-226  $\mu$ M) significantly inhibited CDK1 activity in a dose-dependent fashion. In fact, the degree of inhibition observed with 2,4,6-THBA was greater than that of 2,3-DHBA and 2,6-DHBA (Fig. 1). Not all compounds tested inhibited CDK1 activity *in vitro*. For example, aspirin, salicylic acid, benzoic acid, 3,4-DHBA 3,5-DHBA, and 5-amino-salicylic acid failed to inhibit CDK1 activity. These results collectively suggest that the DHBAs or THBAs with a common -OH group at the 2nd carbon is likely important for the inhibitory effect on CDK1.

We also performed limited studies to determine the effect of 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA on CDK2, 4 and 6 enzyme activity. We observed that 2,3-DHBA and 2,6-DHBA had either no or marginal effect on CDK2 and 4 (at 500  $\mu$ M) activity (Figs. 7A and 4E); however, both inhibited CDK6

activity (Fig. 7G). 2,4,6-THBA was highly effective and inhibited all CDKs examined (CDK1, 2, 4 and 6) (Figs. 1G, 7C, E and G). This suggests that although CDK members share significant homology, the binding pockets for these compounds in CDKs may differ, perhaps accounting for their differential effects. A compilation of the inhibitory effect of 2,3-DHBA, 2,6-DHBA and 2,4,6-THBA on CDK-1, 2, 4 and 6 are shown in Table II. Cytotoxicity assays in HCT-116 cells showed that the IC<sub>50</sub> for these compounds differ. For 2,3-DHBA, the IC<sub>50</sub> was ~1.8 mM; for 2,6-DHBA, it was 1.9 mM and for 2,4,6-THBA, it was 0.8 mM (Table III). The IC<sub>50</sub> for compound's cytotoxicity in HCT-116 cells is significantly higher than observed for *in vitro* CDK1 inhibition (e.g., Fig. 1E-H); this may be related to a poor uptake of polyphenolic compounds by cancer cells (35). Our pilot studies using HPLC showed that the cellular uptake of dihydroxy and trihydroxy-benzoic acid are lower as compared to the mono-hydroxy benzoic acid (salicylic acid) (data not shown), possibly reflecting the presence of multiple OH groups making them more hydrophilic and less absorbable. The poor uptake of HBAs may also be related to the significantly lower expression of monocarboxylate transporters (MCTs, e.g., SLC5A8 and SLC5A12) in metastatic cancer cells (36). Therefore, the effect of these compounds on cellular functions, such as cell cycle regulation, can be better studied after formulation with compounds that can enhance cellular uptake, or in cell lines expressing MCTs. Further work is required to determine, how DHBAs and THBAs affect cell cycle regulation and functions.

Molecular docking studies revealed that aspirin, salicylic acid, 2,3-DHBA, 2,5-DHBA and 2,6-DHBA potentially bind at the same pocket in CDK1; in contrast, 2,4,6-THBA appears to bind to a different site (Fig. 2). Despite the common binding pocket, aspirin and salicylic acid failed to inhibit CDK1; however, 2,3-DHBA, 2,5-DHBA and 2,6-DHBA inhibited CDK1 activity (Fig. 1A-D). Inhibition was also observed with 2,4,6-THBA (Fig. 1G and H). The reason for the inability of aspirin and salicylic acid to inhibit CDK1 activity is not clear at this stage and requires further investigations. All DHBAs potentially showed interaction with Asp146 in CDK1, either through -COOH or -OH groups. 2,3-DHBA and 2,5-DHBA use -COOH group; while, 2,6-DHBA uses the -OH group at the 6th carbon to interact with CDK1. Salicylic acid, which

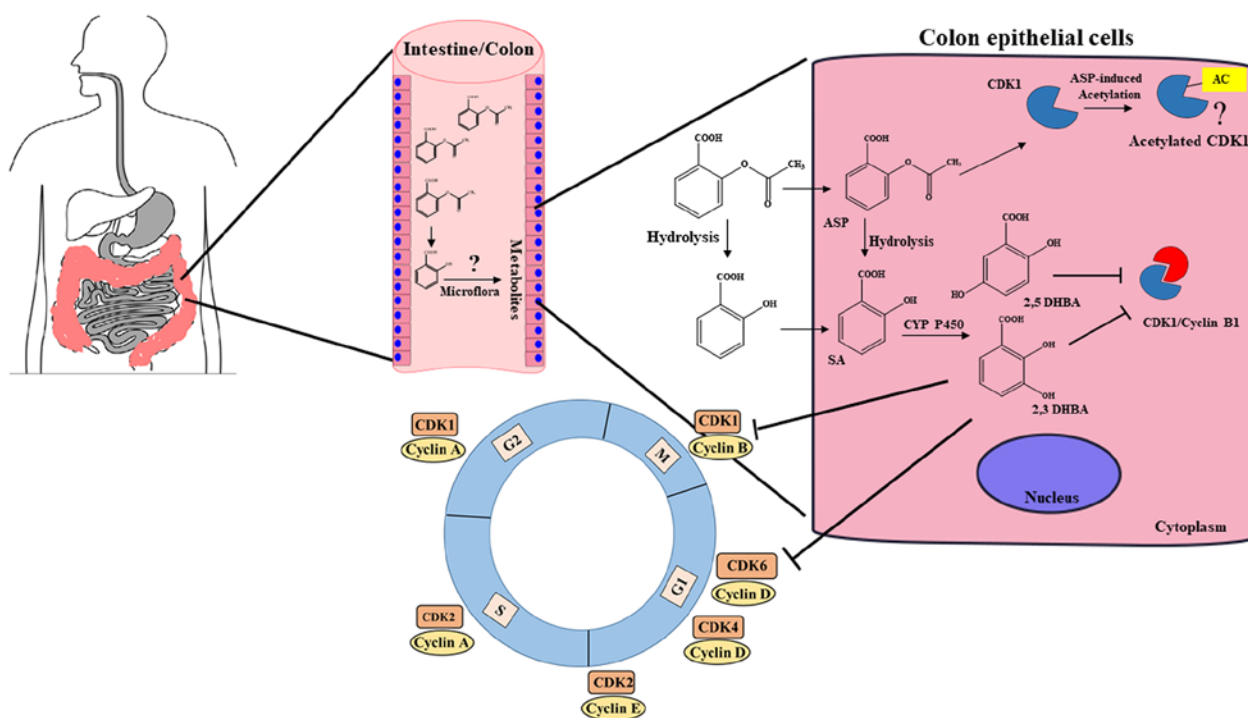


Figure 8. A model depicting how aspirin may preferentially act on colonic tissue to protect against CRC. We suggest that unabsorbed aspirin/salicylic acid in the stomach and upper intestine is passed on to the colon, taken up by the colonic epithelial cells, and metabolized by CYP450s to produce 2,3-DHBA and 2,5-DHBAs. These hydrophilic DHBAs may not easily cross the basolateral membrane of the epithelial cells and within these cells, they may accumulate to pharmacologically relevant concentrations, leading to the inhibition of CDK1 and CDK6, and exert anticancer effects. Colonic epithelial cells may also get exposed to salicylic acid metabolites generated by the gut microflora, and uptake of these HBAs by cells may also cause CDK inhibition. This additional layer of protection against tumor development may be unique to the colonic tissue allowing aspirin as a more effective drug against CRC. (?), Aspirin acetylates CDK1; however, how this modification affects its function is not clear at this stage.

did not show any inhibition of CDK1 appears to interact with Asp146 via the -COOH group, and with Lys33 via the -OH group at the 2nd carbon. It is also interesting to note that 2,4,6-TBHA which showed significant inhibition of CDK1 appears to interact with Arg123 via the -COOH group, and with Arg151 and Gly154 via the -OH group at the 6th carbon (Table I, Fig. 2). These data suggest that the ability of these compounds to inhibit CDK1 is likely determined by the interacting functional groups, as well as their orientation at the binding pocket (Fig. 2), which may affect CDK1 conformation and activity.

In this study, using immunoblotting, we also demonstrated that incubation of recombinant CDK1 (Prospec) monomer with aspirin for 12 h dose-dependently acetylated CDK1 (Fig. 6A), which was also observed in HCT-116 cells (Fig. 6F). An initial mass spectrometry analysis of aspirin-acetylated CDK1 showed that Lys34 is targeted for acetylation (data not shown). It is not clear at this stage whether this modification affects CDK1 activity; however, the Lys34 is adjacent to the amino acid Lys33 (in the active site). Our *in vitro* kinase assays using CDK1/cyclin B1 (NEB) were designed to address how the binding of aspirin affects CDK1 activity by preincubating with aspirin for 10 min, before assaying for H1 histone phosphorylation. The effect of aspirin-mediated acetylation on CDK1 activity could not be assessed, as the short incubation time (10 min) of CDK1/cyclin B1 complex with aspirin was insufficient to cause acetylation; longer incubation (12 h) caused degradation (unpublished data). The ability of aspirin to acetylate recombinant CDK1 monomer (Prospec) was taken advantage of to demonstrate

the binding of salicylic acid, and its derivatives 2,3-DHBA, 2,6-DHBA and 2,4,6-TBHA to CDK1, in competition experiments. Pre-incubation with all three compounds prevented aspirin-mediated CDK1 acetylation (Fig. 6A-E), which supported the data from molecular docking studies that all compounds potentially bind to CDK1 (Table I). These experiments also show that CDK1 is a salicylic acid binding protein.

It is well established that aspirin is more effective in preventing CRC as compared to other cancers, but a clear explanation for its preferential protective effect in colorectal tissues has not emerged in any reports. Our observation that salicylic acid metabolites 2,3-DHBA and 2,5-DHBA inhibit CDK1 activity provides an important insight into aspirin's chemopreventive actions. This is particularly relevant to the prevention of colon tumorigenesis, as expression of several CYP450 members, including CYP3A4/A5 implicated in aspirin/salicylic acid metabolism in the liver, are also reported to be present in healthy and cancerous colonic tissues (37-40). Although the stomach and upper intestine are the major sites of aspirin absorption, it is likely that some aspirin and its hydrolyzed product, salicylic acid, is passed on to the lower intestine and colon. Moreover, availability of aspirin in the intestine and colon will be much higher after ingestion of enteric coated tablets, as they are designed to resist dissolution in the acidic environment of the stomach. Aspirin and salicylic acid absorbed by the intestinal and colonic epithelial cells may be metabolized by local CYP450s to produce 2,3-DHBA and 2,5-DHBA. Although 2,3-DHBA and 2,5-DHBA are the minor metabolites generated by CYP450 metabolism in the

liver, the extent to which they are generated in epithelial cells of the GI are unknown, and it is possible that these metabolites may be produced at much higher levels in GI epithelial cells. These intracellularly produced hydroxyl-derivatives of salicylic acid are probably too hydrophilic to cross the basolateral side of the epithelial cell membrane, and within these cells, they may accumulate to pharmacologically relevant concentrations sufficient to inhibit CDKs and act locally to exert anticancer effects. Another source of salicylic acid derivatives for epithelial cell uptake may be generated through GI microflora. The epithelial cells of the intestine/colon are likely to be exposed to higher concentrations of the salicylic acid metabolites and derivatives (HBAs) as opposed to other tissues resulting in the inhibition of CDKs. This additional layer of protection against tumor development may be unique to the colorectal tissue as compared to other tissues such as breast, lung, ovary and skin, allowing aspirin to preferentially act against CRC. A model depicting new insights into aspirin's effect against CRC is shown in Fig. 8. This model envisions that the ability of aspirin to inhibit tumor formation in the intestinal/colonic mucosa is a local effect via locally generated salicylic acid metabolites/derivatives leading to inhibition of CDKs, and may not require systemic absorption into the blood.

Salicylic acid metabolites 2,3-DHBA and 2,5-DHBA were also reported to be present in the plasma of individuals even when there has been no intake of aspirin, indicating a dietary source (41). It is suggested that diet containing fruits and vegetables may also provide a rich source of HBAs. 2,3-DHBA is present in avocados, batoko plum and in medicinal herbs such as Madagascar rosy periwinkle. 2,5-DHBA is also widely present in fruits including kiwi fruit, aloe vera, mushrooms, apple, bitter melon, blackberries, grapes and pears; it is also present in wine (42). It is important to note that 2,4,6-THBA was demonstrated as one of the *in vitro* degradation product of catechins by the intestinal bacterium, *Acinetobacter calcoaceticus* (43,44). In addition, 2,4-DHBA and 2,4,6-THBA were reported to be produced through pH-dependent degradation of anthocyanins (45). The exposure of the colonic epithelial cells to these pharmacologically active CDK inhibitors may provide a link between diet rich in salicylic acid/DHBAs/catechins/anthocyanins and decreased occurrences of CRC (46-48).

We believe that our observations have opened a new frontier in aspirin research by identifying CDKs as novel cellular targets for salicylic acid metabolites and derivatives. The potential contributions of locally generated HBAs either in the colon or within the colonic epithelial cells, to aspirin's chemopreventive actions against CRC via CDK inhibition represent an important area for future research. Against the backdrop of our *in vitro* observation that salicylic acid metabolites inhibit CDKs, the question is raised whether aspirin through acetylation of COX or other proteins (7,49), or salicylic acid through binding to cellular targets (SABPs) (19,34), or salicylic acid metabolites through inhibition of CDKs, are responsible for the observed chemopreventive effects of aspirin. We suggest that although aspirin's ability to prevent CRC may occur through multiple mechanisms, salicylic acid metabolites formed locally in the colon, or within the intestinal epithelial cells, may contribute significantly to its protective effects. Other interesting questions that need to be addressed include: are there other cellular protein targets for

salicylic acid metabolites besides CDKs? Will daily supplements of DHBAs/THBAs capable of inhibiting CDKs provide protection against CRCs?  $IC_{50}$  for the inhibition of CDKs by DHBAs and THBAs are although in micromolar range (300  $\mu$ M), it could be argued that these concentrations are still physiologically relevant. Given the abundance of the occurrences of DHBAs and THBAs in fruits and vegetables (42), it is possible that CDKs have been evolved to be less sensitive to inhibition for these natural compounds avoiding cytotoxicity.

In recent times, attention is increasingly focused on developing CDK inhibitors to arrest cell cycle as a therapeutic intervention to treat cancer. Although several potential CDK inhibitors are undergoing clinical trials, palbociclib and ribociclib are the only FDA approved drugs used in cancer treatment (17). We are hopeful that structural modification to DHBAs and THBAs may potentially lead to the development of novel class of CDK inhibitors for cancer prevention and treatment.

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