Chemical properties of fatty acid derivatives as inhibitors of DNA polymerases

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In this study, the chemical properties of organic acids as DNA polymerase inhibitors were examined. In total, we assayed the inhibitory activities of 23 compounds. We found that the DNA synthesis activity of DNA polymerase was usually reduced to less than 50% in the presence of 100 μ M monoprotic acids, which have a Clog P value greater than 7.0 and a p K_a value less than 5.4. With a minor modification these chemical properties applied to several organic fatty acids previously reported as DNA polymerase inhibitors. Moreover, we also examined the inhibitory activities of perfluorooctadecanoic acid (PFOdA) and perfluorooctanesulfonic acid (PFOS) against DNA polymerase β in detail. These compounds inhibited the polymerase activity of pol β competitively with template–primer DNA, and non-competitively with dNTPs. In addition, the 8 kDa domain-defective pol β was also sensitive to these compounds. Our results suggest that the inhibitory mode of action of PFOdA and PFOS is different from that mediated by the classic fatty acid inhibitors against DNA polymerase β .

Introduction

DNA polymerases are essential enzymes for DNA replication, repair and recombination. Eukaryotic cells contain three replicative DNA polymerases (pol α , δ and ϵ), a mitochondrial DNA polymerase (pol γ) and at least 10 repair-types of other DNA polymerases (pol β , δ , ϵ , ζ , η , θ , κ , λ , μ and ν). PR Recently, DNA polymerase inhibitors have been a major focus of research as candidate anti-cancer agents. In particular, repair-type DNA polymerases, including pol β , could be considered both as markers of tumorigenesis and as potential drug targets for chemotherapy.

We previously reported that long-chain fatty acids, such as stearic acid (1) and nervonic acid (2), inhibit the activities of a mammalian pol α and pol β *in vitro*, but that the fatty acids did not influence the activities of prokaryotic DNA polymerases or other DNA metabolic enzymes at a concentration below 100 μ M.⁷⁻¹³ Lineweaver–Burk plots of the inhibition of pol α by fatty acids indicate non-competitive inhibition with both the template DNA and the dNTPs.^{8,9} By contrast, inhibition of pol β was competitive with both the template DNA and the dNTPs.^{8,9} Moreover, our previous NMR study and docking simulation found that nervonic

acid (2) would form a 1 : 1 complex with the 8 kDa domain of pol β through interaction with four amino acids of pol β (Leu11, Lys35, His51, and Thr79). The hydrophobic region, which contains Leu11, His51 and Thr79, might interact with the side-chain of 2, and the hydrophilic Lys35, might interact with the carboxylic group of 2. The structure–activity relationships of fatty acids for inhibition of DNA polymerases revealed three major findings: (1) fatty acids with C18 or more carbon atoms inhibit the polymerases; (2) the carboxylic acid group is important for inhibition; (3) unsaturated fatty acids with the *cis*-configuration at the double bond display a stronger inhibition than those with the *trans*-configuration.

During the course of our investigation of DNA polymerase inhibitors, we became interested in the unique physical, chemical and biological properties of perfluorinated alkyl substances (PFAS). PFAS have a variety of applications, including acting as surfactants or additives for polymers and plastics. ¹⁴⁻¹⁷ The amphipathic, thermal, photochemical and chemical properties of PFAS, combined with their biological inertness, account for the versatile uses of these compounds for over 50 years. ¹⁴ However, PFAS are environmentally persistent and bioaccumulative. ¹⁴⁻¹⁶ In previous reports, PFOS was shown to display biological toxicity by targeting specific functions such as peroxisome proliferation and lipid metabolism. ^{14,17}

In this study, the inhibition assay was carried out with 23 organic acids to examine the structure–activity relationships in detail. On the basis of these results, we investigated the correlation between the chemical properties of the organic acids and the inhibitory activities against mammalian polymerases. We focused on the calculated $\log P$ value (partition coefficients for octanol–water) and pK_a of the organic acid as the chemical properties. Moreover, we examined the enzymatic inhibition of PFOdA and PFOS against DNA polymerases by kinetic analysis. A model for the inhibition by these compounds is also discussed.

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Materials and methods

General (chemistry)

¹H and ¹³C NMR spectra were recorded on a Bruker 400 MHz spectrometer (Avance DRX-400), using CDCl₃ solutions, unless otherwise noted. Chemical shifts were expressed in δ (ppm) relative to Me₄Si or residual solvent resonance, and coupling constants (J) were expressed in Hz. Melting points were determined with Yanaco MP-3S and were uncorrected. Infrared (IR) spectra were recorded on a Jasco FT-IR-410 spectrometer using NaCl (neat) or KBr pellets (solid), and were reported as wavenumbers (cm⁻¹). Mass spectra (MS) were obtained on an Applied Biosystems mass spectrometer (API QSTAR pulsar i) under high resolution conditions. Column chromatography was carried out on Fuji Silisia PSQ100B. Analytical thin-layer chromatography (TLC) was performed on precoated Merck silica gel 60 F₂₅₄ plates, and compounds were visualized by UV illumination (254 nm) or heating at 150 °C after spraying with phosphomolybdic acid in ethanol. HPLC was performed on an Agilent 1100 Series using a Mightysil Si 60 column (KANTO Inc.) with a 30 min linear gradient elution from 100:0 to 85:15 (hexane–i-PrOH with 0.1% trifluoroacetic acid) at a flow rate of 1.0 mL min⁻¹ with UV detection at 254 nm. THF was distilled from sodium-benzophenone. CH₂Cl₂ was distilled from P₂O₅. DMSO was distilled from CaH₂. All other solvent and reagents were obtained from commercial sources and used without further purification. Organic extracts were dried over Na₂SO₄, filtered, and concentrated using a rotary evaporator. Involatile oils and solids were vacuum dried.

Organic acids

Compounds 1, 2, 3, 5, 6, 13, 14, 15, 16, 17, 22, and 23 were obtained from commercial sources and used without further purification. Compounds 4, ¹⁸ 7, ¹⁹ 8, ²⁰ 9, ²¹ 10, ²² 12, ²³ 18, ²⁴ 19, ²⁴ 20²⁵ and 21²⁵ were prepared by literature procedures with slight modifications and characterized by NMR spectra. Compound 11 was prepared from the known compound 24²⁶ (Scheme 1).

Synthesis of 5-*tert*-butyldimethylsilyloxy-4-oxo-4*H*-pyran-2-carbaldehyde (25)

To a solution of **24** (114.7 mg, 0.50 mmol) and Et_3N (0.5 mL, 3.6 mmol) in CH_2Cl_2 –DMSO (4:1,5 mL) was added SO_3 -pyridine (363.6 mg, 2.3 mmol) at 0 °C. The mixture was stirred at rt for 10.5 h. Then the mixture was quenched by the addition of sat.

NaHCO₃ aq. and diluted with EtOAc. The layers were separated. Then the organic layer was washed with $\rm H_2O$ and brine, dried (Na₂SO₄), and evaporated *in vacuo*. The residue was purified by silica gel chromatography (hexane–EtOAc = 4 : 1) to yield **25** (31.4 mg, 28%) and the recovered **24** (68.8 mg, 61%). **25**: ¹H NMR (400 MHz) δ 0.25 (6H, s), 0.97 (9H, s), 7.00 (1H, s), 7.80 (1H, s), 9.67 (1H, s).

Synthesis of *E*- and *Z*-5-tert-butyldimethylsilyloxy-2-(1-heptadecenyl)-4-oxo-4*H*-pyran (26)

To a solution of the known phosphonium salt, hexadecyltriphosphenylphosphonium bromide, 27 (40.2 mg, 0.071 mmol) in THF (2 mL) was added n-BuLi (125 μ L of 1.56 M solution in hexanes, 0.20 mmol) at 0 °C. The mixture was stirred at rt for 10 min. Then a solution of 25 (31.4 mg, 0.12 mmol) in THF (1.5 mL) was added to the mixture at 0 °C and the mixture was stirred at rt for 3 h. The mixture was quenched by the addition of sat. NH₄Cl aq. and diluted with EtOAc. The layers were separated. Then the organic layer was washed with H₂O and brine, dried (Na₂SO₄), and evaporated in vacuo. The residue was purified by silica gel chromatography (hexane-EtOAc = 20:1) to yield 26 (15.1 mg, 46%) as a 1.4:1 mixture of *E*- and *Z*-isomers. *E*-isomer: ¹H NMR (400 MHz) δ 0.22 (6H, s), 0.88 (3H, t, J = 7.0 Hz), 0.95 (9H, s), 1.26 (24H, brs), 1.46 (2H, m), 2.21 (2H, q, J = 7.0 Hz), 6.02 (1H, d, J = 15.8 Hz), 6.16 (1H, s), 6.55 (1H, dt, J = 15.8 Hz)7.0 Hz), 7.59 (1H, s). Z-Isomer: 1 H NMR (400 MHz) δ 0.24 (6H, s), 0.88 (3H, t, J = 7.0 Hz), 0.96 (9H, s), 1.26 (24H, brs), 1.46 (2H, brs)m), 2.44 (2H, m), 5.92–5.93 (2H, m), 6.24 (1H, s), 7.64 (1H, s).

Synthesis of 2-heptadecyl-5-hydroxy-4*H*-pyran-4-one (11)

A solution of **26** (15.1 mg, 0.033 mmol) and Pd(OH)₂–C (4.4 mg) in THF (2 mL) was stirred under a H₂ atmosphere for 6 h. Then the solution was filtered through Celite and washed with THF. The filtrate was treated with 0.5 mL of 1 N HCl aq. and the mixture was stirred for 10 min. The mixture was quenched by the addition of sat. NaHCO₃ aq. and diluted with EtOAc. The layers were separated. Then the organic layer was washed with H₂O and brine, dried (Na₂SO₄), and evaporated *in vacuo*. The residue was purified by silica gel chromatography (hexane–EtOAc = 20 : 1) to yield **11** (8.5 mg, 74%) as a white solid. Mp = 85 °C. ¹H NMR (400 MHz, CDCl₃) δ 0.88 (3H, t, J = 6.8 Hz), 1.25 (26H, brs), 1.30 (2H, m), 1.64 (2H, m), 2.53 (2H, t, J = 7.6 Hz), 6.27 (1H, s), 7.79 (1H, s). ¹³C NMR (100 MHz, CDCl₃) δ 14.1, 22.7, 27.0, 28.9, 29.2, 29.35, 29.42, 29.56, 29.61, 29.65 (×3), 29.68 (×3), 31.9, 33.9, 110.3,

OTBS
$$Et_3N$$
 OTBS Et_3N OTBS $CH_3(CH_2)_{15}PPh_3Br$ n -BuLi n -BuLi

Scheme 1 Synthesis of 11.

137.4, 145.1, 170.4, 174.2. IR (KBr) 3224, 2919, 2849, 1734, 1702, 1653, 1620, 1585, 1461, 1378, 1226, 1186, 1149, 917, 892, 835, 757, 722, 701 cm⁻¹. ESIMS: calcd for $C_{22}H_{38}O_3Na$ ([M + Na]⁺) 373.2713, found 373.2711. HPLC analysis: $t_R = 9.9$ min. Anal. calcd for $C_{22}H_{38}O_3 \cdot H_2O$: C, 71.70; H, 10.94. Found: C, 72.05; H, 10.93%.

Materials for DNA polymerase assay

Calf thymus DNA, deoxynucleotide triphosphates and synthetic polynucleotides such as poly(dA), oligo(dT)₁₂₋₁₈, and [³H]-2′-deoxythymidine 5′-triphosphate (dTTP) (43 Ci mmol⁻¹) were purchased from GE Healthcare Bio-Sciences Corp (Piscataway, NJ 08855, USA). All other reagents were of analytical grade and were purchased from Wako Pure Chemical Industries, Ltd. (Osaka, Japan).

Enzymes

DNA polymerase α (pol α) was purified from the calf thymus by immuno-affinity column chromatography as described previously. Recombinant rat DNA polymerase β (pol β) and human terminal deoxynucleotidyl transferase (TdT) were purified from *Escherichia coli* BL21 (DE3) as described previously. The 31 kDa domain of pol β (residues 88–335) was cloned into the pET28a expression vector (Novagen), and purified by the same method as for the whole pol β .

DNA polymerase assays

The DNA polymerase activities were measured based on the methods described previously.³⁰ Briefly, the reaction mixture (25 µl) contained 50 mM Tris-HCl (pH 7.5), 5 mM MgCl₂, 1 mM dithiothreitol, 10% (v/v) glycerol, 10% (v/v) dimethyl sulfoxide, 50 μg ml⁻¹ poly(dA)-oligo(dT)₁₂₋₁₈, 0.5 μM dTTP, 0.1 μM [³H]dTTP, 4.1 fmol of pol α , β or TdT, and at least five different concentrations of each compound. When activated calf thymus DNA (100 μg ml⁻¹) was used as the DNA template, the reaction mixture was supplemented with 100 mM each of dATP, dGTP and dCTP, and dTTP was removed from it. After incubation at 37 °C for an hour, the reaction mixture was spotted onto a Whatmann DE81 ion-exchange filter, air-dried, and washed 3 times with 5% (w/v) Na₂HPO₄, 3 times with distilled water, and twice with ethanol. The radioactivity of each dried filter was measured in a toluene-based scintillator. The GraphPad Prism program (GraphPad Software Inc., San Diego, CA) was used to calculate the IC₅₀ value for each compound, and to determine the inhibitory mechanisms of compounds 3 and 17 by global fitting models.

Results and discussion

Enzymatic inhibitory activity of organic acids against DNA polymerases

The structures of all the compounds tested in this study are shown in Fig. 1. The calculated log P (Clog P), pK_a , and the IC₅₀ values of each compound are summarized in Table 1. Most of the Clog P and pK_a values of the organic acids were obtained from the calculated properties in SciFinder Scholar, which were originally

calculated using Advanced Chemistry Development (ACD/Lab) Software V8.14 for Solaris (ACD/Labs).

We previously reported that compound 1 inhibited the activity of pol α and pol β with IC₅₀ values of 9 μ M and 91 μ M, respectively. In order to examine the effect of the alkyl moiety on the inhibitory activity of fatty acids against DNA polymerase, we first changed the alkyl group of 1. Replacement of the octadecyl group in 1 with a perfluorooctadecyl moiety (3), thereby increasing the overall hydrophobicity and acidity of the molecule (Clog P =16.64, p $K_a = 0.37$), gave more potent inhibitory activity. The IC₅₀ values of perfluorooctadecanoic acid (PFOdA, 3) against pol α, pol β and TdT were 0.71 μ M, 0.63 μ M and 5.4 μ M, respectively. Since the CMC of 3 is reported to be 80 µM, 3 would not form a micelle at the concentration at which 3 inhibited DNA polymerases.³¹ Replacement of the alkyl group with pentaethylene glycol (4) increased the hydrophilicity (Clog P = -2.45), but 100 μM of compound 4 had no effect on DNA polymerase activity. Compound 5, which has a Clog P value of 2.9, also lacked inhibitory activity. By contrast, compound 6, which is considerably more hydrophobic (Clog P = 7.75), inhibited pol α and TdT with IC₅₀ values of 56 μM and 62.5 μM, respectively. These results suggested that the hydrophobicity of the alkyl moiety of the fatty acids was closely related to the inhibitory activity against DNA polymerases. This proposal is supported by our previous NMR study, in which the hydrophobic interaction between the hydrophobic region (Leu11, His51 and Thr79) in pol β and the side-chain in 2 had an important role in the binding between pol β and 2.10,13

The branched chain fatty acid (7) was a slightly weaker inhibitor of pol α activity (IC₅₀ = 63.4 μ M) than the straight chain fatty acid (1). However, the IC₅₀ value for 7 against pol β (82.2 μ M) was similar to that of 1. Because compound 7 has the same level of hydrophobicity (Clog P=8.03) and acidity (p $K_a=4.82$) as that of 1, any differences in inhibitory activity against pol α can be attributed to molecular shape. Compound 8 exhibited more potent inhibition than 7 against pol α , β and TdT with IC₅₀ values of 19.6 μ M, 36.1 μ M and 33.6 μ M, respectively. Indeed compound 8 displayed similar activities to compound 9. Thus the α -carboxyl substituents increased the inhibitory activity against the DNA polymerases.

We next turned our attention to the carboxylic group of stearic acid (1). Compounds 10–12, having similar p K_a values, are bioisosteres of 1. The IC₅₀ value of 10 against pol α (8.6 μ M) is similar to that of 1, whereas the IC₅₀ value of 10 against pol β (7 μ M) is much lower than that of 1. Although compound 11 is a weaker inhibitor of pol α than 1, inhibition of pol β by 11 was significantly more potent than that by 1 (IC₅₀ values of 11: IC₅₀ = 31.1 μ M against pol α ; IC₅₀ = 26.3 μ M against pol β). Both 10 and 11 did not affect the activity of TdT. Interestingly, the diprotic sulfonamide 12 showed no inhibitory activity against any DNA polymerase at a concentration less than 100 μ M. These results indicate that the acidic functional group affected not only inhibitory activity, but also selectivity against the enzymes.

The inhibitory activities of compounds 13–17, all of which have pK_a values much lower than that of 1, were also examined. The diprotic phosphoric acid 13 lacked inhibitory activity against pol α , pol β and TdT. Murofushi *et al.* reported that PHYLPA, a monoprotic lysophosphatidic acid, specifically inhibited pol α . ^{32,33} Thus, the type of protic acid might be a crucial factor for inhibition

Fig. 1 Chemical structures of compounds tested as DNA polymerase inhibitors.

against DNA polymerases. Octadecyl sulfate (14) and dodecyl sulfate (15) are alkyl sulfonates, which have the same p K_a value of -3.29. Although 14 efficiently inhibited all three polymerases (IC₅₀ = 5.8 μ M against pol α ; IC₅₀ = 12.8 μ M against pol β ; IC₅₀ = 18.3 μ M against TdT), 15 did not display any inhibitory activity even at a concentration of 100 μ M. Octadecyl sulfonic acid 16 inhibited the α and β polymerase activities with a similar dose to that of 14 (IC₅₀ values of 16: IC₅₀ = 12.9 μ M against pol α ; IC₅₀ = 21 μ M against pol β). PFOS (17) inhibited the activity of these polymerases in the submicromolar range (IC₅₀ = 24.5 μ M against pol α ; IC₅₀ = 46.4 μ M against pol β ; IC₅₀ = 31.3 μ M against TdT).

Since it is reported that the CMC of 17 ranges from 0.88 mM to 1.7 mM, 17 would not form micelles at the concentration used in this study. 34,35 These results suggest that (1) the type of protic acid as well as the acidic functional group is important for inhibition against polymerases; (2) the hydrophobicity of the side-chain moiety (Clog P) is important for the inhibitory activity against DNA polymerases.

Compounds **18–23** have greater pK_a values than that of **1**. The phenolic compounds (**18** and **19**), β -ketoester (**20**), ketone (**21**), amide (**22**) and alcohol (**23**) used in this study showed no inhibitory activity against pol α , pol β and TdT. These results

Table 1 The Clog P value and calculated p K_a value of organic acids 1–23, and the IC₅₀ and logIC₅₀ values for enzymatic inhibition of DNA polymerase α and β , and TdT by the compounds 1–23. The IC₅₀ and log IC₅₀ values were determined by using the GraphPad Prism program. The IC₅₀ values were obtained by taking the antilog of the log IC₅₀ values. Unless otherwise noted, both the Clog P values and the pKa values of organic acids were obtained from the calculated properties in SciFinder Scholar, which were originally calculated using Advanced Chemistry Development (ACD/Lab) Software V8.14 for Solaris (ACD/Labs)

Compounds	Clog P	pK_a	$IC_{50}/10^{-6} M$			Log IC ₅₀ /M		
			Pol α	Pol β	TdT	Pol α	Pol β	TdT
1	8.21	4.78	9	91	ND			
2	10.9	4.78	2	8	ND			
3	16.64	0.37	0.71	0.63	5.4	-6.15 ± 0.05	-6.20 ± 0.07	-5.27 ± 0.07
4	-2.45	3.39	>100	>100	>100			
5	2.9	4.78	>100	>100	>100			
6	7.75	0.35	56	>100	62.5	-4.25 ± 0.02		-4.20 ± 0.02
7	8.03	4.82	63.4	82.2	>100	-4.20 ± 0.05	-4.08 ± 0.01	
8	8.45	3.66	19.6	36.1	33.6	-4.71 ± 0.03	-4.44 ± 0.01	-4.47 ± 0.01
9	7.58	2.07	22.4	57.7	54.3	-4.65 ± 0.02	-4.24 ± 0.01	-4.27 ± 0.02
10	7.42	5.35	8.6	7	>100	-5.07 ± 0.01	-5.16 ± 0.02	
11	5.15 ^a	5.15^{b}	31.1	26.3	>100	-4.51 ± 0.02	-4.58 ± 0.03	
12	7.75	10.81 -5.27	>100	>100	>100			
13	7.01	2.42	>100	>100	>100			
14	8.58	-3.29	5.8	12.8	18.3	-5.23 ± 0.02	-5.89 ± 0.07	-4.74 ± 0.02
15	5.4	-3.29	>100	>100	>100			
16	7.14	1.84	12.9	21	>100	-4.89 ± 0.09	-4.68 ± 0.05	
17	7.03	-3.27	24.5	46.4	31.3	-4.61 ± 0.05	-4.33 ± 0.03	-4.51 ± 0.02
18	10.44	10.14	>100	>100	>100			
19	10.44	10.06	>100	>100	>100			
20	9.22	10.67	>100	>100	>100			
21	8.34	>15	>100	>100	>100			
22	7.27	16.61 -0.49	>100	>100	>100			
23	8.31	15.19	>100	>100	>100			

Errors mean \pm S.E. The Clog P values were obtained using CS ChemDraw version 6.0 software (Cambridge Soft, USA). The calculated p K_a value of compound 11 is substituted for that of 2-butyl-5-hydroxy-4*H*-pyran-4-one.

indicate that the acidic functional group is essential for inhibition.

After conducting a series of enzymatic inhibitory assays we conclude that the most potent DNA polymerase inhibitors possess the following properties. Monoprotic acids, which have a Clog P value greater than 7.0 and a p K_a value less than 5.4, have a tendency to inhibit the activity of one or more DNA polymerases. Although the mechanism of DNA polymerase inhibition for these inhibitors is not fully examined, both hydrophobicity and acidity are essential for inhibitory activity against mammalian polymerases. Our previous NMR study showed that the carboxylic group of 2 interacted with the Lys35 hydrophilic region and the alkyl side chain of 2 interacted with the Leu11–His51 hydrophobic region. 10,13 Thus, the acidic functional group and hydrophobic moiety of the inhibitor are required for binding to the basic amino acid residues and hydrophobic regions of the DNA polymerase, respectively.

Most of the previously identified fatty acid inhibitors showed chemical properties similar to those of inhibitors found in the present studies. The known DNA polymerase inhibitors, 8,36-44 which have a Clog P value greater than 7.0 and a p K_a value less than 5.4, are summarized in Fig. 2. Actually some phenolic^{45–49} and glycolipid^{50–52} inhibitors of DNA polymerases do not possess the anticipated chemical properties. However, our predictions have wide applicability to lipid compounds, assuming that the Clog P value is defined as more than 5.3 for steroidal inhibitors. For instance, fomitellic acid A^{53,54} (Clog P 5.3, pK_a 4.1), lithocholic acid (Clog P 6.7, pK_a 4.8),^{55–57} vitamin A acid⁵⁸ (Clog P 6.8, pK_a 4.7), tormentic acid⁵⁹ (Clog P 6.2, pK_a 4.5) and euscaphic acid⁵⁹ (Clog P 6.2, pK_a 4.5) inhibit DNA polymerases. We also found that the shape of some lipid inhibitors influences the selective inhibition of pol α .⁴⁴ By contrast, these chemical property–activity predictions do not apply to nucleotide analogue based inhibitors, presumably reflecting a different mode of inhibition.

Inhibition of DNA polymerase activity by PFOdA and PFOS

Since PFOdA (3) and PFOS (17) have unique physical, chemical and biological properties, we investigated the mechanism of these inhibitors in detail.

The mode of inhibition of pol β was examined by kinetic analysis (Fig. 3).89 In these experiments, poly(dA)-oligo(dT)₁₂₋₁₈ and dTTPs were utilized as the DNA template-primer and nucleotide substrate, respectively. The V_{max} and K_{m} values of the polymerase reactions with and without 3 and 17 were obtained by fitting the data to the Michaelis-Menten equation with Graphpad Prism4 software (Table 2). Considering the template–primer DNA as substrate, the Michaelis constant (K_m) increased with increasing concentrations of 3 and 17, but the apparent maximum velocity $(V_{\rm max})$ was almost unaltered. Thus the inhibition of pol β by 3 and 17 is considered to be competitive with DNA template-primer. The Lineweaver–Burk plot analysis also supports this suggestion (Fig. 3, A and E). In contrast, measuring the activity over a range of dTTP concentrations as substrate, $K_{\rm m}$ was independent of the

Linoleic acid (Clog P 7.2, pKa 4.78)

3-Geranyl-4-hydroxy-5-(3'-methyl-2'-butenyl) -benzoic acid (Clog P 7.6, pKa 4.7)

Hippospongic acid A (Clog P 9.8, pKa 3.6)

Elenic acid (Clog P 11.7, pKa 4.4)

Cholesterol derivative (Clog P 9.7, pKa 3.4)

Ursolic acid (Clog P 9.0, pKa 4.7)

SO₃H HO HO O HO O C₁₇H₃₅CO₂ O₂CC₁₇H₃₅

α-SQDG (Clog P 15.3, pKa 1.4)

Manzamenone A (Clog P 17.2, pKa 3.93)

Stigmasterol derivative (Clog P 10.0, pKa 3.4)

Cholestanol derivative (Clog P 10.0, pKa 3.5)

Kohamaic acid (Clog P 9.8, pKa 5.10)

Fig. 2 Known DNA polymerase inhibitors that have a Clog P value greater than 7.0 and a pK_a value less than 5.4.

concentration of **3** and **17**, although V_{max} was reduced in the presence of these compounds, suggesting that **3** and **17** inhibit the DNA synthesis by pol β non-competitively with a nucleotide substrate. This result was also consistent with that obtained from the Lineweaver–Burk plot (Fig. 3, B and F).

The melting temperature of double-stranded DNA was unchanged in the presence of 100 μM 3 or 17 (data not shown). Thus, compounds 3 and 17 must inhibit the enzymatic activity by

interacting with the enzyme in the reaction mixture. In order to analyze the mode of inhibition further, the inhibitory activity of 3 and 17 against the 31 kDa domain of pol β was examined (Table 3). Tryptic digestion of pol β liberates an N-terminal domain (8 kDa), which is responsible for single-strand DNA binding activity, and a C-terminal domain (31 kDa), which catalyzes DNA synthesis. ^{60,61} Compounds 3 and 17 inhibit the activity of the 31 kDa domain of pol β with an IC₅₀ value of 4.4 μ M and 51.1 μ M, respectively.

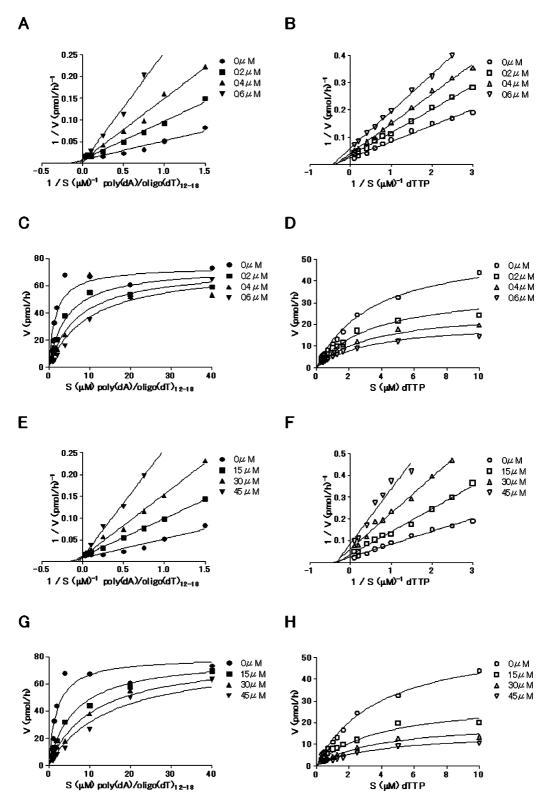


Fig. 3 Kinetic analysis of the inhibitory effects of compounds 3 (A–D) and 17 (E–H) on the activities of DNA polymerase β, as a function of the DNA template–primer dose and the nucleotide substrate concentration. (A) and (C): Pol β activity was measured in the absence or presence of the indicated concentrations of compound 3 using varying concentrations of the DNA template–primer (A: Lineweaver–Burk plots, C: Michaelis–Menten curves). (B) and (D): Pol β activity was measured in the absence or presence of the indicated concentrations of compound 3 using varying concentrations of the substrate dTTPs (B: Lineweaver–Burk plots, D: Michaelis–Menten curves). (E) and (G): Pol β activity was measured in the absence or presence of the indicated concentrations of compound 17 using varying concentrations of the DNA template–primer (E: Lineweaver–Burk plots, G: Michaelis–Menten curves). (F) and (H): Pol β activity was measured in the absence or presence of the indicated concentrations of compound 17 using varying concentrations of the substrate dTTPs (F: Lineweaver–Burk plots, H: Michaelis–Menten curves).

Table 2 Kinetic analysis of the inhibitory effects of compounds 3 and 17 on the activity of DNA polymerase β as a function of DNA template–primer dose and nucleotide substrate concentration. These data were obtained from global fitting models using the GraphPad Prism program. Errors mean \pm S.E

Substrate	Concentration $/\mu M$	$K_{\rm m}/\mu{ m M}$	$V_{\rm max}/{\rm pmol~min^{-1}~ng^{-1}}$	$K_{\rm i}/\mu{ m M}$	Inhibitory mode
Compound 3					
DNA Template–primer ^a	0	1.777 ± 0.563	0.795 ± 0.070	0.12 ± 0.03	Competitive
	0.2	3.970 ± 0.793	0.704 ± 0.045		•
	0.4	5.415 ± 3.021	0.725 ± 0.136		
	0.6	18.24 ± 1.713	0.994 ± 0.043		
Nucleotide substrate ^b	0	4.432 ± 0.3391	0.655 ± 0.026	0.37 ± 0.02	Non-competitive
	0.2	2.509 ± 0.230	0.323 ± 0.013		_
	0.4	2.737 ± 0.220	0.268 ± 0.010		
	0.6	2.462 ± 0.064	0.185 ± 0.002		
Compound 17					
DNA Template-primer ^a	0	1.777 ± 0.563	0.795 ± 0.070	6.8 ± 1.2	Competitive
• •	15	6.812 ± 0.658	0.825 ± 0.028		•
	30	14.79 ± 1.26	0.991 ± 0.037		
	45	27.75 ± 5.27	1.141 ± 0.115		
Nucleotide substrate ^b	0	4.432 ± 0.3391	0.655 ± 0.026	15.9 ± 0.7	Non-competitive
	15	2.652 ± 0.406	0.277 ± 0.019		Î
	30	3.193 ± 0.5181	0.195 ± 0.015		
	45	4.333 ± 0.8586	0.162 ± 0.016		

Table 3 The IC_{50} and log IC_{50} values of compounds 3 and 17 for enzymatic inhibition of the 31 kDa domain of DNA polymerase β. The IC_{50} and log IC_{50} values were determined by using the GraphPad Prism program. Errors mean \pm S.E

Compound	$IC_{50}/10^{-6} M$	Log IC ₅₀ /M
3	3.9	-5.41 ± 0.05
17	51.1	-4.29 ± 0.04

By contrast, 1 and 2 had almost no effect on the activity of the 31 kDa domain. We therefore conclude that the perfluorinated chain of 3 and 17 enhances the hydrophobic interaction between these compounds and the 31 kDa domain of pol β . However, the sensitivity of the 31 kDa domain to 3 was reduced by approximately 6-fold, whereas 17 inhibited the activity of both the 31 kDa domain and the full-length pol β equally. These results indicate that compound 17 interferes competitively with the interaction between the template–primer DNA and the active site of the 31 kDa domain of pol β , but non-competitively with incoming dNTPs. By contrast, compound 3 appears to display moderate affinity to the 8 kDa domain in addition to binding to the 31 kDa domain of pol β .

Conclusion

We have investigated the chemical properties of organic acids as DNA polymerase inhibitors. Monoprotic acids, which have a Clog P value greater than 7.0 and a p K_a value less than 5.4, were generally found to inhibit the activity of one or more DNA polymerases at concentrations lower than 100 μ M. By contrast, compounds with a Clog P value less than 7.0 and with a p K_a value greater than 5.4 generally had little influence on the polymerase activity. With a minor modification these chemical properties of DNA polymerase inhibitors also apply to several lipid compounds. Moreover, the molecular shape was suggested to be important for both inhibition and selectivity. In the present

study, we found that DNA polymerases are sensitive to PFOdA and PFOS. Furthermore, our kinetic analysis suggests that the perfluorinated chain might enhance the hydrophobic interaction between these compounds and the 31 kDa domain of pol β . These observations will provide insights not only for identifying new DNA polymerase inhibitors from natural or synthetic compounds, but for providing information on inhibitor–enzyme interactions in order to develop inhibitors of therapeutic utility.

Abbreviations

PFAS, perfluorinated alkyl substance; PFOdA, perfluorooctadecanoic acid; PFOS, perfluorooctanesulfonic acid; pol, DNA polymerase; TdT, terminal deoxynucleotide transferase; Clog *P*, calculated log *P*; IC₅₀, 50% inhibition concentration; dNTP, deoxynucleotide 5'-triphosphate; CMC, critical micelle concentration.

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