A Novel Class of Tyrosyl-DNA Phosphodiesterase 1 Inhibitors That Contains the Octahydro-2H-chromene-4-ol Scaffold

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Abstract: Tyrosyl-DNA phosphodiesterase 1 (Tdp1) is a DNA repair enzyme that mends topoisomerase 1-mediated DNA damage. Tdp1 is a current inhibition target for the development of improved anticancer treatments, as its inhibition may enhance the therapeutic effect of topoisomerase 1 poisons. Here, we report a study on the development of a novel class of Tdp1 inhibitors that is based on the octahydro-2H-chromene scaffold. Inhibition and binding assays revealed that these compounds are potent inhibitors of Tdp1, with IC50 and Kd values in the low micromolar concentration range. Molecular modelling predicted plausible conformations of the active ligands, blocking access to the enzymatic machinery of Tdp1. Our results thus help establish a structural-activity relationship for octahydro-2H-chromene-based Tdp1 inhibitors, which will be useful for future Tdp1 inhibitor development work.

Keywords: anticancer agent; Tdp1 inhibitor; DNA repair enzyme; synthesis; biochemical assay; molecular modeling; chemical space; structural-activity relationships

1. Introduction

Tyrosyl-DNA phosphodiesterase 1 (Tdp1) is an enzyme that belongs to the phospholipase D superfamily [1,2]. Tdp1 functions as a DNA repair enzyme in cells, in which it removes stalled topoisomerase 1 (Top1)–DNA complexes by catalysing the hydrolysis of phosphodiester bonds between the tyrosine residue of Top1 and the 3′-phosphate of DNA [3]. Top1 is a current inhibition target for chemotherapeutic agents [4]. However, Tdp1 also repairs DNA damage that is caused by Top1 inhibitors. Thus, the inhibition of Tdp1 may enhance the efficacy of Top1 poisons, potentially reducing the side effects of chemotherapeutic treatments and enabling a lower dosage to be used.

Given the potential of Tdp1 inhibition in enhancing the efficacy of anticancer treatments, several types of Tdp1 inhibitors have been developed [5–14]. Most of these inhibitors show a good to moderate
inhibition potency, with IC_{50} values in the concentration range between 0.1 and 20 µM. These include 7-azaindeneisoquinoline derivatives 1 such as triple Top1/Tdp1/Tdp2 inhibitors [5]; achyrodimer F 2, which is a dimeric cyclobutane metabolite from an Australian fungus of the Cortinariaceae family [6]; 7-hydroxycoumarins derivatives 3 that contain aromatic or monoterpane substituents [7]; usnic acid derivatives, such as 4 with an enamine moiety [8]; compounds that contain the benzopentathiepine moiety 5 [9]; aminoadamantane-containing monoterpane-derived compound 6 [10]; and furamidine (NSC 305831), which is a commercial Tdp1 inhibitor [11] (Figure 1), as well as the other less potent inhibitors of various chemical classes [12–14].

![ Structures of reported Tdp1 inhibitors 1–6, commercial Tdp1 inhibitor Furamidine, and octahydro-2H-chromene-4-ol derivatives 7 that we investigated in this study.](image)

Octahydro-2H-chromene is an oxygen-containing heterocycle with a tetrahydropyran moiety. This structural scaffold occurs in many important natural products and biologically-active molecules [15,16]. For example, compounds that contain octahydro-2H-chromene-4-ol were found to exhibit antiviral and analgesic activities [17,18]. The synthesis of molecules that contain the octahydro-2H-chromene scaffold has been investigated extensively. In particular, Prins cyclisation was found to be a flexible method for the synthesis of octahydro-2H-chromenes derivatives. Typically, the reaction involves a homoallylic alcohol and a carbonyl compound being catalysed by Brønsted or Lewis acids. The most commonly used catalysts in this transformation are montmorillonite clays or Lewis acids such as Sc(OTf)₃, BF₃·Et₂O, or I₂ [19–22]. Montmorillonite clays were also found to be effective catalysts for the synthesis of octahydro-2H-chromenes with heterocyclic motifs [23–26]. Although aldehydes containing amino groups cannot be involved in such reactions because of catalyst poisoning [26], in principle, such products can be obtained through reactions with nitro-aromatic aldehydes, followed by the reduction of the nitro group. For example, we have successfully obtained nitro-containing octahydro-2H-chromene from 5-nitro-thiophene-2-carbaldehyde in our previous work [18]. Moreover, the electron-rich thiophene ring may facilitate the reduction of the nitro group into an amino group, which is especially important for compounds that are relatively labile, such as those that contain a tert-hydroxy group. Thus, we reasoned that, due to the versatility and ease of synthesis, thiophene-2-carbaldehyde is an ideal starting point for the synthesis and derivatisation of compounds that contain the octahydro-2H-chromene scaffold.

By using a range of octahydro-2H-chromene-4-ol derivatives with different amide motifs 7 (Figure 1), preliminary molecular modelling studies were performed. Our modelling results suggested that these derivatives may bind to Tdp1. We therefore reasoned that octahydro-2H-chromene-4-ol may be a useful scaffold for the development of new Tdp1 inhibitors. Herein, we report our work on the synthesis of a structural series of octahydro-2H-chromene-4-ol derivatives as potential Tdp1 inhibitors. Inhibition and binding assays were performed to evaluate their inhibition potency and binding affinity.
to Tdp1, respectively. Finally, molecular modelling was performed to predict the plausible binding
conformations of these compounds.

2. Results and Discussion

2.1. Synthesis of Octahydro-2H-chromen-4-ol Derivatives

Nitro compounds 8 with an octahydro-2H-chromen-4-ol scaffold were synthesised as a mixture of
diastereomers on the C-4 position in a reaction of (-)-isopulegol with 5-nitrothiophene-2-carbaldehyde
in the presence of montmorillonite K10 clay (Scheme 1).

Scheme 1. Synthesis of (4R)-9 and (4S)-9 compounds and proposed mechanism for the synthesis of 4R
and 4S, as proposed by Nazimova et al. [18].

The possible mechanism of the transformation depicted in Scheme 1 has been previously
proposed [18]. After purification using column chromatography, the individual products (4R)-8
and (4S)-8 were obtained with yields of 25% and 28%, respectively. In order to avoid dehydration of
the compounds, their reduction was carried out in a slightly alkaline aqueous alcohol solution with
sodium dithionite. This resulted in the production of primary amines (4R)-9 and (4S)-9 with yields of
60% and 80%, respectively.

The primary amines obtained were selectively acylated with anhydrides and acid chlorides
(Scheme 2). Acylation of amine groups solely occurred, even when the acylating reagent was in
excess. Using acid chlorides resulted in products of smaller yields compared to acid anhydrides.
The desired amide derivatives of octahydro-2H-chromen-4-ol 10–13 were purified using column
chromatography. In each case, except for the reaction with 1-adamantanecarbonyl chloride, the yield
of (4S)-diastereomer was higher than the one of (4R). We supposed that the isolation of (4R)-diastereomers
was less successful, likely due to their instability during column chromatography.

The structure of the compounds was determined by nuclear magnetic resonance (NMR)
spectroscopy, including $^1$H, $^{13}$C, $^1$H–$^1$H 2D homonuclear correlation spectroscopy (COSY), $^1$H–$^{13}$C
2D heteronuclear single quantum coherence (HSQC), and $^1$H–$^{13}$C 2D heteronuclear multiple bond
correlation (HMBC) experiments. As the hydroxyl group of (4R)-isomers 8–13 is in the axial position,
this causes the $^1$H resonances of neighbouring protons, H-1 and H-3, to shift upfield by ~0.3 ppm and
~0.4 ppm, respectively, when compared with (4S)-isomers. This phenomenon is likely caused by the
1,3-diaxial interaction, which was previously observed in our studies of similar compounds [17,18].
properties of all the chromenes. BHQ1 (Black Hole Quencher-1) at the 3′-end hence reflects the catalytic activity of the enzyme. Hexadecameric oligonucleotide was selected as a binding mode is predicted as the ligands block access to these key amino acid residues. It can therefore be stated that a plausible function and the binding pocket was defined there [1,7,9].

Interestingly, all our compounds appeared to bind Tdp1 to more than one binding pocket (i.e., -13). The docked configuration of the most active derivative (4S)-10, (4R)-10 95% 35% (4S)-11, (4R)-11 80% 30% displayed a hydrogen bonding interaction with His493 through its hydroxyl group, unlike (4R, 4S)-8, (4R, 4S)-9, and (4R)-10. Both isomers (4R, 4S)-12, (4R)-12 40% 30% (4S)-13, (4R)-13 14% 20% were found to be similar. This suggests that the different substituents on the adamantane moiety and the thiophene displayed a hydrogen bonding interaction with the amino acids.

2.2. Tdp1 Inhibition and Binding Assays

The inhibition potency of our synthesised compounds against Tdp1 was evaluated by using a real-time oligonucleotide biosensor-based assay [27]. In this assay, Tdp1 catalyses the removal of fluorophore quenchers from the 3′-end of a modified DNA substrate [27]. Fluorescence intensity hence reflects the catalytic activity of the enzyme. Hexadecameric oligonucleotide was selected as a Tdp1 substrate, as it contains 5(6)-carboxyfluorescein (FAM) at the 5′-end and fluorophore quencher BHQ1 (Black Hole Quencher-1) at the 3′-end [9]. This approach was used to determine the inhibitory properties of all the chromenes.

All our synthesised compounds except derivatives 8 and 9 were found to exhibit inhibitory activity against Tdp1 in the low micromolar range (Table 1). Compounds with IC50 > 15 µM were considered inactive. The (4S)-diastereomer with the most bulky substituent was found to be the most potent (13), with an IC50 value of 1.24 µM. In general, the IC50 values for the (4S) series are in the IC50 of 1.24 to 5.8 µM range. For (4R)-diastereomers, their IC50 values were found to be similar.

| Compound | R           | IC50, µM | Kd, µM |
|----------|-------------|----------|--------|
| 8        | NO2-        | >15      | >15    | 30.6 ± 9.3 23.3 ± 7.4 |
| 9        | NH2-        | >15      | >15    | n.d.  n.d. |
| 10       | CH3-        | 5.8 ± 3.0 | 2.9 ± 0.8 | n.d.  n.d. |
| 11       | CF3-        | 1.4 ± 0.3 | 4.0 ± 0.4 | 2.0 ± 1.2 19.8 ± 2.4 |
| 12       |             | 5.0 ± 1.5 | 3.3 ± 0.2 | 24.5 ± 5.7 n.d. |
| 13       |             | 1.24 ± 0.02 | 2.8 ± 0.6 | 17.9 ± 3.4 12.4 ± 7.5 |
| Fur *    |             | 1.2 ± 0.3 | n.d. |

* Furamidine was used as a positive control. n.d. = not determined.

Scheme 2. Synthesis of 10–13 amide derivatives of octahydro-2H-chromen-4-ol.
In addition to the inhibition assay, the binding of our compounds to Tdp1 was also evaluated. An intrinsic protein fluorescence-based assay, which we have previously applied to study the binding of esters combining monoterpeneoid and adamantane fragments, was applied [28]. In agreement with the inhibition data, all our compounds were relatively strong binders to Tdp1, with dissociation constants (K_D) in the low µM region (Table 1 and Supplementary Figures S1–S7). Interestingly, all our compounds appeared to bind Tdp1 to more than one binding pocket (i.e., specific and non-specific binding). This is possibly due to the large hydrophobic surface of Tdp1. The slight discrepancy between the measured K_D and IC_50 values is likely due to these non-specific binding events.

2.3. Molecular Modelling

Both isomers (4R, 4S) of the six ligands were docked into Tdp1 (PDB ID: 1MU7) [2,29] using GOLD and all the ligands fitted into the pocket in a similar way with a reasonable predicted affinity for the scoring functions used, as can be seen in Table 2. GoldScore (GS) and, in particular, Chem Piecewise Linear Potential (ChemPLP), scoring functions give higher scores for 10–13 (4R, 4S) compared to (4R, 4S)-8 and (4R, 4S)-9, which is consistent with their IC_50 values. A similar trend is also observed for ChemScore (CS) and Astex Statistical Potential (ASP), albeit not as pronounced as for GoldScore and ChemPLP. The correlation between the measured affinities and the predicted binding values is reassuring, particularly since ChemPLP is considered to be the most robust function used [30] and strengthens the overall applicability of the modelling.

The docked configuration of the most active derivative (4S)-13 is shown in Figure 2. Our docking model revealed that the ligand is in lipophilic contact (LC) with His263, Tyr204, Ala520, Ala521, and Pro461, and exhibited hydrogen bonding with Gly458 and Ser459 via the hydroxyl group. Also, the hydrophobic pocket in the binding site is occupied by the adamantane moiety and the thiophene group occupies a cleft exposed to the water environment, as shown in Figure 2A. Derivative (4S)-11, which shows a similar inhibition potency to (4S)-13, is predicted to have LC with His263 and Tyr204 in a similar manner to (4S)-13. However, (4S)-11 displayed a hydrogen bonding interaction with His493 through its hydroxyl group, unlike (4S)-13. This suggests that the different substituents on the amide group can have an important effect in orienting the ligands in the binding site of the enzyme. It is established that for Tdp1, both His263 and His493 play a key role in its biological function and the binding pocket was defined there [1,7,9]. It can therefore be stated that a plausible binding mode is predicted as the ligands block access to these key amino acid residues.

The calculated molecular descriptors (MW=molecular weight, log P octanol–water partition coefficient, HD=hydrogen bond donors, HA=hydrogen bond acceptors, PSA=polar surface area, and RB=rotatable bonds) for the six isomers (4S, 4R) are given in Table S1 (see Supplementary Material). The ligands are relatively average in size, with a molecular weight between 281.4 and 443.6 (4R, 4S), and nine (4R, 4S) derivatives lie in the lead-like space, while others are within the boundaries of the drug-like space (for definition of lead-like, drug-like, and Known Drug Space regions, see ref [31]). The log P values range from 2.3 to 5.1, with only compounds (4R, 4S)-8, (4R, 4S)-9, and (4R)-10 being slightly outside of the ideal drug-like space but within the Known Drug Space. Moreover, the Polar Surface Area (PSA) values also indicate that they lie in the drug-like space.

The SAR and modelling studies clearly indicate that lipophilic R-groups are potentially beneficial for improved binding of this class of compounds to Tdp1. To investigate this possibility, four derivatives with the R-groups α-naphthyl (14), β-naphthyl (15), isopropyl (16), and tert-butyl (17) were docked into the binding site and their scores are given in Table 2. Compared with the scores for the most active compounds 11 and 13, it can be seen from Table 2 that the α-naphthyl substitution gives favourable scores for both GS and ChemPLP, suggesting that this derivative may be a promising derivative to synthesise and test. The other substitutions gave similar or worse scores compared to 11 and 13. Similar binding poses were predicted for derivatives 14–17 as for their 8–13 counterparts. Finally, derivatives 14 and 15 have slightly higher Log P values than 13, but have essentially the same values for the other molecular descriptors (see Table S1 in the SI).
was dried with calcined Al2O3. A 30-m quartz column HP-5MS (copolymer 5%–diphenyl–95%–dimethylsiloxane) with an inner diameter of 0.25 mm and the stationary phase film thickness of 0.25 μm was used. Optical rotation parameters: polAAr 3005 spectrometer; CHCl3 soln. 1H and 13C NMR: Bruker DRX-500 apparatus at 500.13 MHz (1H) and 125.76 MHz (13C), 2D homonuclear correlation, J-modulated 13C NMR spectra (JMOD), and 13C–1H 2D heteronuclear correlation with one-bond and long-range spin-spin coupling constants (C–H COSY, 1H, 13C). HR-MS: DFS Thermo Scientific spectrometer in a full scan mode (15–200 m/z, 70 eV electron impact ionisation, direct sample administration).

### 3. Materials and Methods

#### 3.1. Chemistry

The purity of all target compounds reported in this paper exceeds 95%. From 0 to 100% EtOAc in hexane, ethanol. The purity of the target compounds was determined by analysing the 1H NMR spectra, including 1H–1H double resonance spectra and 1H–1H 2D homonuclear correlation, J-modulated 13C NMR spectra (JMOD), and 13C–1H 2D heteronuclear correlation with one-bond and long-range spin-spin coupling constants (C–H COSY, 1H, 13C). GC-MS methods. The purity of all target compounds reported in this paper exceeds 95%.

#### 3.1.1. General Materials and Methods

Chemicals were purchased from commercial sources (Sigma-Aldrich, St. Louis, MO, USA; Acros, Morris Plains, NJ, USA) and used with no further treatment. The K10 clay (Aldrich) was used as the catalyst of the reaction. The clay was dried at 105 °C for 3 h prior to use. Dichloromethane CH2Cl2 was dried with calcined Al2O3. For the products structure analysis, GC-MS was used, consisting of an Agilent 7890A gas chromatograph equipped with a quadrupole mass spectrometer Agilent 5975C as a detector. A 30-m quartz column HP-5MS (copolymer 5%–diphenyl–95%–dimethylsiloxane) with an inner diameter of 0.25 mm and the stationary phase film thickness of 0.25 μm was used. Optical rotation parameters: polAAr 3005 spectrometer; CHCl3 soln. 1H and 13C NMR: Bruker DRX-500 apparatus at 500.13 MHz (1H) and 125.76 MHz (13C), 2D homonuclear correlation, J-modulated 13C NMR spectra (JMOD), and 13C–1H 2D heteronuclear correlation with one-bond and long-range spin-spin coupling constants (C–H COSY, 1H, 13C). HR-MS: DFS Thermo Scientific spectrometer in a full scan mode (15–200 m/z, 70 eV electron impact ionisation, direct sample administration).

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#### Table 2. Results of the Scoring Functions (GS, CS, ChemPLP, and ASP) for Compounds 8–17 with Tdp1. The IC50 Values of Compounds 8–13 with Tdp1 were Also Included.

| Compound | GS   | CS   | ChemPLP | ASP  | IC50 (μM) |
|----------|------|------|---------|------|-----------|
| (4R)-8   | 40.1 | 26.0 | 47.9    | 27.4 | >15       |
| (4S)-8   | 41.0 | 25.3 | 45.0    | 26.6 | >15       |
| (4R)-9   | 41.5 | 25.0 | 41.1    | 27.2 | >15       |
| (4S)-9   | 42.2 | 25.5 | 45.5    | 29.0 | >15       |
| (4R)-10  | 42.3 | 25.9 | 50.7    | 28.9 | 2.9 ± 0.8 |
| (4S)-10  | 44.0 | 25.2 | 49.4    | 28.4 | 5.8 ± 3.0 |
| (4R)-11  | 39.6 | 23.8 | 51.8    | 34.4 | 4.0 ± 0.4 |
| (4S)-11  | 44.0 | 25.2 | 50.6    | 31.3 | 1.4 ± 0.3 |
| (4R)-12  | 55.8 | 30.3 | 57.2    | 29.1 | 3.3 ± 0.2 |
| (4S)-12  | 56.0 | 30.8 | 56.4    | 29.8 | 5.0 ± 1.5 |
| (4R)-13  | 51.5 | 32.3 | 57.1    | 30.5 | 2.8 ± 0.6 |
| (4S)-13  | 52.5 | 31.4 | 56.1    | 30.4 | 1.24 ± 0.02 |
| (4R)-14  | 60.9 | 30.9 | 60.4    | 31.3 | -         |
| (4S)-14  | 62.5 | 29.6 | 61.3    | 32.8 | -         |
| (4R)-15  | 50.2 | 30.5 | 64.8    | 32.6 | -         |
| (4S)-15  | 53.6 | 28.2 | 59.6    | 28.5 | -         |
| (4R)-16  | 48.7 | 23.2 | 51.6    | 25.9 | -         |
| (4S)-16  | 48.8 | 21.5 | 48.5    | 27.5 | -         |
| (4R)-17  | 49.2 | 21.9 | 49.3    | 25.1 | -         |
| (4S)-17  | 51.2 | 22.2 | 47.9    | 27.7 | -         |

**Figure 2.** Docked configurations of derivative (4S)-13 to the substrate binding sites predicted by the ChemPLP algorithm. The protein surface is rendered where partial positive, negative charges, and neutral regions are colored blue, red, and grey, respectively (A). The hydrogen bond interactions are depicted as green lines between compound (4S)-13 and the amino acids residues; Ser459 and Gly458 (B). Furthermore, lipophilic contacts (LC) are shown as purple dashed lines with His263, Tyr204, Ala520, Ala521, and Pro461.

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#### 3.1. Chemistry

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Chemicals were purchased from commercial sources (Sigma-Aldrich, St. Louis, MO, USA; Acros, Morris Plains, NJ, USA) and used with no further treatment. The K10 clay (Aldrich) was used as the catalyst of the reaction. The clay was dried at 105 °C for 3 h prior to use. Dichloromethane CH2Cl2 was dried with calcined Al2O3. For the products structure analysis, GC-MS was used, consisting of an Agilent 7890A gas chromatograph equipped with a quadrupole mass spectrometer Agilent 5975C as a detector. A 30-m quartz column HP-5MS (copolymer 5%–diphenyl–95%–dimethylsiloxane) with an inner diameter of 0.25 mm and the stationary phase film thickness of 0.25 μm was used. Optical rotation parameters: polAAr 3005 spectrometer; CHCl3 soln. 1H and 13C NMR: Bruker DRX-500 apparatus at 500.13 MHz (1H) and 125.76 MHz (13C), 2D homonuclear correlation, J-modulated 13C NMR spectra (JMOD), and 13C–1H 2D heteronuclear correlation with one-bond and long-range spin-spin coupling constants (C–H COSY, 1H, 13C). HR-MS: DFS Thermo Scientific spectrometer in a full scan mode (15–200 m/z, 70 eV electron impact ionisation, direct sample administration).
inner diameter of 0.25 mm and the stationary phase film thickness of 0.25 µm was used. Optical rotation parameters: polAAr 30005 spectrometer; CHCl₃ soln. ¹H and ¹³C NMR: Bruker DRX-500 apparatus at 500.13 MHz (¹H) and 125.76 MHz (¹³C), J in Hz; structure was determined by analysing the ¹H NMR spectra, including ¹H–¹H double resonance spectra and ¹H–¹H 2D homonuclear correlation, J-modulated ¹³C NMR spectra (JMOD), and ¹³C–¹H 2D heteronuclear correlation with one-bond and long-range spin-spin coupling constants (C–H COSY). ¹J(C,H) = 160 Hz, COLOC, ²J(C,H) = 10 Hz. HR-MS: DFS Thermo Scientific spectrometer in a full scan mode (15-500 m/z, 70 eV electron impact ionisation, direct sample administration).

All product yields are given as pure compounds obtained from the column chromatography. Column chromatography: silica gel (SiO₂; 60-200 µ; Macherey-Nagel); hexane, solution containing from 0 to 100% EtOAc in hexane, ethanol. The purity of the target compounds was determined by GC-MS methods. The purity of all target compounds reported in this paper exceeds 95%.

Please note that the numeration of the atoms of products is performed as shown in Figure 3.

3.1.2. Chemical Syntheses

The syntheses and characterisations of (2R,4aR,7R,8aS)-4,7-Dimethyl-2-(5-nitrothiophen-2-yl)octahydro-2H-chromen-4-ols, (4S)-8, and (4R)-8, were reported previously [18].

(2R,4R,4aR,7R,8aS)-4,7-Dimethyl-2-(5-aminothiophen-2-yl)octahydro-2H-chromen-4-ol, (4R)-9. To the solution of nitro compound (4R)-8 (0.436 g) in EtOH (20 mL), Na₂S₂O₄ (2.300 g) and K₂CO₃ (1.900 g) in water (10 mL) were added, and the mixture was stirred at r.t. for 2 h. After the completion of the reaction, 10 mL of a saturated aq. NaCl solution was added to the reaction mixture and extracted with ethyl acetate (3 x 10 mL). The organic layer was washed with brine (2 x 10 mL) and dried over anhydrous sodium sulphate. The organic layer was concentrated in a rotary evaporator and separated using an SiO₂ column. Purification resulted in the yield of (4R)-9 (0.234 g, 60%). ¹H-NMR (CDCl₃) (ppm): δ: 0.88–0.97 (m, 1H, Ha-8); 0.92 (d, J(16, 9a) = 6.5 Hz, 3H, H-16); 1.05 (ddd, J(10a, 10e) = J(10a, 9a) = 12.1 Hz, J(10a, 1a) = 11.2 Hz, 1H, Ha-10); 1.09–1.20 (m, 2H, Ha-6, Ha-7); 1.24 (s, 3H, H-15); 1.41–1.52 (m, 1H, Ha-9); 1.67 (dd, J(4a, 4e) = 13.4 Hz, J(4a, 3a) = 11.6 Hz, 1H, Ha-4); 1.72 (dm, J(8e, 8a) = 13.1 Hz, other J < 3.5 Hz, 1H, He-8); 1.77–1.83 (m, 1H, He-7); 1.95 (dd, J(4e, 4a) = 13.4 Hz, J(4e, 3a) = 2.4 Hz, 1H, He-4); 1.97 (dm, J(10e, 10a) = 12.1 Hz, other J < 4.5 Hz, 1H, He-10); 3.21 (ddd, J(1a, 10a) = 11.2 Hz, J(1a, 6a) = 9.7 Hz, J(1a,10e) = 4.2 Hz, 1H, Ha-1); 4.49 (ddd, J(3a, 4a) = 11.6 Hz, J(3a, 4e) = 2.4 Hz, J(3a, 14) = 1.0 Hz, 1H, Ha-3); 5.99 (d, J(14, 13) = 3.5 Hz, 1H, H-14); 6.53 (d, J(13, 14) = 3.5 Hz, 1H, H-13). ¹³C-NMR (CDCl₃) (ppm): δ: 72.13 (d, C-1); 70.63 (d, C-3); 47.25 (t, C-4); 69.99 (s, C-5); 49.22 (d, C-6); 22.28 (t, C-7); 34.14 (t, C-8); 31.08 (d, C-9); 40.83 (t, C-10); 150.56 (s, C-11); 132.81 (s, C-12); 122.36 (d, C-13); 107.64 (d, C-14); 27.92 (q, C-15); 22.01 (q, C-16). HR-MS: 281.1449 (M⁺, C₁₅H₂₂O₂NS⁺; calc. 281.1444). [α]D²⁶.₀ = +7.2 (c 0.334, EtOH).

(2R,4S,4aR,7R,8aS)-4,7-Dimethyl-2-(5-aminobenzo-2-yl)octahydro-2H-chromen-4-ol, (4S)-9. To the solution of nitro compound (4S)-8 (0.223 g) in EtOH (20 mL), Na₂S₂O₄ (1.300 g) and K₂CO₃ (1.000 g) in water (10 mL) was added, and the mixture was stirred at r.t. for 2 h. After the completion of
the reaction, 10 mL of a saturated aq. NaCl solution was added to the reaction mixture and extracted with ethyl acetate (3 × 10 mL). The organic layer was washed with brine (2 × 10 mL) and dried over anhydrous sodium sulphate. The organic layer was concentrated in a rotary evaporator and separated on an SiO₂ column. Purification resulted in the yield of (4S)-9 (0.167 g, 80%). 1H-NMR (CDCl₃) (ppm) δ: 0.88–0.89 (m, 1H, Ha-8); 0.93 (d, J(16, 9a) = 6.5 Hz, 3H, H-16); 0.94 (ddd, J(10a, 10e) = J(10a, 9a) = 12.1 Hz, J(10a, 1a) = 11.2 Hz, 1H, Ha-10); 1.03–1.12 (m, 2H, Ha-6, Ha-7); 1.14 (s, 3H, H-15); 1.33–1.43 (m, 1H, Ha-9); 1.63 (ddd, J(4a, 4e) = 13.4 Hz, J(4a, 3a) = 11.6 Hz, 1H, Ha-4); 1.70–1.89 (m, 4H, He-7, He-8, He-4, He-10); 2.03 (s, 3H, H-18); 3.50 (ddd, J(1a, 10a) = 11.2 Hz, J(1a, 6a) = 9.7 Hz, J(1a, 10e) = 4.2 Hz, 1H, Ha-1); 3.89 (q, C-16); 167.39 (s, C-17); 22.03 (q, C-18). HR MS: 323.1547 (M + 1.63 (dd, J(1a, 10e) = 4.2 Hz, 1H, Ha-1); 4.85 (ddd, J(3a, 4a) = 11.6 Hz, J(3a, 4e) = 2.4 Hz, 1H, Ha-3); 6.39 (d, J(14, 13) = 3.8 Hz, 3H, H-14); 6.62 (d, J(13, 14) = 3.8 Hz, 1H, H-13). 13C-NMR (CDCl₃) δ: 77.51 (d, C-3); 72.51 (d, C-3); 49.32 (t, C-4); 70.26 (s, C-5); 51.50 (d, C-6); 22.83 (t, C-7); 34.16 (t, C-8); 31.08 (d, C-9); 40.97 (t, C-10); 138.77 (s, C-11); 138.64 (s, C-12); 120.60 (d, C-14); 22.37 (q, C-15); 20.60 (q, C-16); 167.39 (s, C-17); 21.95 (q, C-18). HR MS: 323.1547 (M⁺, C₁₅H₂₁O₂NS⁺; calc. 323.1550). [α]D²⁰ = +20.5 (c 0.16, EtOH).

General Procedure for the Preparation of Compounds 10–11. To the solution of amine (R)-4S (-)-28.0 = +20.5 (c 0.16, EtOH).

N-(5-((2R,4R,4aR,7R,8aR)-4-Hydroxy-4,7-dimethyloctahydro-2H-chromen-2-yl)thiophen-2-yl)acetamide, (4R)-10. The reaction of amine (4R)-9 (0.100 g) and acetic anhydride (0.037 g) in the presence of 4-DMAP (0.01 mmol) was added dropwise, and the mixture was stirred at r.t. for 2 h. After the completion of the reaction, 10 mL was added. The reaction mixture was extracted with ethyl acetate (3 × 10 mL). The organic layer was washed with brine (2 × 10 mL) and dried over anhydrous sodium sulphate. The organic layer was concentrated in a rotary evaporator and separated using an SiO₂ column.

N-(5-((2R,4R,4aR,7R,8aR)-4-Hydroxy-4,7-dimethyloctahydro-2H-chromen-2-yl)thiophen-2-yl)acetamide, (4S)-10. The reaction of amine (4S)-9 (0.040 g) and acetic anhydride (0.015 g) in the presence of 4-DMAP (0.01 mmol) was added dropwise, and the mixture was stirred at r.t. for 2 h. After the completion of the reaction, 10 mL was added. The reaction mixture was extracted with ethyl acetate (3 × 10 mL). The organic layer was washed with brine (2 × 10 mL) and dried over anhydrous sodium sulphate. The organic layer was concentrated in a rotary evaporator and separated using an SiO₂ column.
2,2,2-Trifluoro-N-(5-((2R,4R,4aR,7R,8aR)-4-hydroxy-4,7-dimethyloctahydro-2H-chromen-2-yl)thiophen-2-yl)acetamide, (4R)-11. The reaction of amine (4R)-9 (0.110 g) and trifluoroacetic anhydride (0.082 g) in the presence of 4-DMAP (0.001 g) for 2 h resulted in the yield of (4R)-11 (0.043 g, 30%). [α]D 28 = +22.38 (c 0.143, EtOH).

General Procedure for the Preparation of Compounds 12–13. To a solution of amine 9 (1.0 mmol) in EtOAc, chloroform (1.0 mmol) in the presence of NEt3 (1 mmol) was added dropwise and the mixture was stirred at r.t. for 2 h. After the completion of the reaction, 10 mL of a saturated aq. NaHCO₃ solution was added to the reaction mixture and extracted with ethyl acetate (3 × 10 mL). The organic layer was washed with brine (2 × 10 mL) and dried over anhydrous sodium sulphate. The organic layer was concentrated in a rotary evaporator and separated using an SiO₂ column.

N-(5-((2R,4R,4aR,7R,8aR)-4-Hydroxy-4,7-dimethyloctahydro-2H-chromen-2-yl)thiophen-2-yl)benzamide, (4R)-12. The reaction of amine (4R)-9 (0.130 g) and benzoyl chloride (0.064 g) in the presence of NEt₃ (1 mmol) was stirred at r.t. for 2 h. After the completion of the reaction, 10 mL of a saturated aq. NaHCO₃ solution was added to the reaction mixture and extracted with ethyl acetate (3 × 10 mL). The organic layer was washed with brine (2 × 10 mL) and dried over anhydrous sodium sulphate. The organic layer was concentrated in a rotary evaporator and separated using an SiO₂ column.
(0.015 g) for 2 h resulted in the yield of (4S)-12 (0.020 g, 40%). $^1$H-NMR (CDCl$_3$) (ppm) δ: 0.88–0.97 (m, 1H, Ha-8); 0.92 (d, J(16, 9a) = 6.5 Hz, 3H, H-16); 1.05 (ddd, J(10a, 10e) = J(10a, 9a) = 12.1 Hz, J(10a, 1a) = 11.2 Hz, 1H, Ha-10); 1.09–1.20 (m, 2H, Ha-6, Ha-7); 1.24 (s, 3H, H-15); 1.41–1.52 (m, 1H, Ha-9); 1.67 (dd, J(4a, 4e) = 13.4 Hz, J(4a, 3a) = 11.6 Hz, 1H, Ha-4); 1.72 (dm, J(8e, 8a) = 13.1 Hz, other J < 3.5 Hz, 1H, He-8); 1.77–1.83 (m, 1H, He-7); 1.95 (dd, J(4e, 4a) = 13.4 Hz, J(4e, 3a) = 2.4 Hz, 1H, He-4); 1.97 (ddm, J(10e, 10a) = 12.1 Hz, other J < 4.5 Hz, 1H, He-10); 3.57 (ddd, J(1a, 10a) = 11.2 Hz, J(1a, 6a) = 9.7 Hz, J(1a, 10e) = 4.2 Hz, 1H, Ha-1); 4.96 (ddd, J(3a, 4a) = 11.6 Hz, J(3a, 4e) = 2.4 Hz, J(3a, 14) = 1.0 Hz, 1H, Ha-3); 6.60 (dd, J(14, 13) = 4.2 Hz, J(14, 3a) = 1.0 Hz, 1H, H-14); 6.73 (d, J(13, 14) = 4.2 Hz, 1H, H-13); 7.42–7.46 (m, 2H, H-20, H-22); 7.78–7.86 (m, 1H, H-21); 8.13–8.14 (m, 1H, H-24); 8.52 (d, J(1a, 1a) = 11.2 Hz, J(1a, 6a) = 9.7 Hz, J(1a, 10e) = 4.2 Hz, 1H, Ha-1); 4.57 (d, J(3a, 4a) = 11.6 Hz, 1H, Ha-3); 6.45 (d, J(14, 13) = 4.2 Hz, 1H, H-14); 6.68 (d, J(13, 14) = 4.2 Hz, 1H, H-13). $^{13}$C-NMR (CDCl$_3$) (ppm) δ: 77.37 (d, C-1); 72.48 (d, C-3); 49.50 (t, C-4); 70.71 (s, C-5); 51.89 (d, C-6); 22.91 (t, C-7); 34.23 (t, C-8); 37.31 (d, C-9); 41.14 (t, C-10); 136.97 (s, C-11); 132.56 (s, C-12); 112.06 (d, C-13); 121.19 (d, C-14); 21.08 (q, C-15); 22.05 (q, C-16); 174.08 (s, C-17); 40.69 (s, C-18); 38.99 (t, C-19, C-25, C-26); 27.89 (d, C-20, C-22, C-24); 36.23 (t, C-21, C-23, C-27). HR MS: 443.2481 (M$^+$, C$_{26}$H$_{37}$O$_3$NS$^+$; calc. 443.2489). [α]$_D^{25.0}$ = +19.3 (c 0.114, EtOH).

N-(5-((2R,4R,4aR,7R,8aS)-4-Hydroxy-4,7-dimethyloctahydro-2H-chromen-2-yl)thiophen-2-yl)adamantan-1-carboxamide, (4S)-13. The reaction of amine (4R)-9 (0.055 g) and 1-adamantanecarboxyl chloride (0.040 g) in the presence of NEt$_3$ (0.030 g) for 2 h resulted in the yield of (4R)-13 (0.016 g, 20%). $^1$H-NMR (CDCl$_3$) (ppm) δ: 0.87–0.96 (m, 1H, Ha-8); 0.89 (d, J(16, 9a) = 6.5 Hz, 3H, H-16); 0.98–1.09 (m, 1H, Ha-10); 1.09–1.16 (m, 2H, Ha-6, Ha-7); 1.21 (s, 3H, H-15); 1.40–1.49 (m, 1H, Ha-9); 1.67–1.77 (m, 7H, Ha-4, 2H-21, 2H-23, 2H-27); 1.84–1.97 (m, 2H, He-8, He-7); 1.91 (bs, 3H, 6H, 2H-19, 2H-25, 2H-26); 1.98–2.01 (m, 2H, He-4, He-10); 2.06 (bs, 3H, H-20, H-22, H-24); 2.33 (ddd, J(1a, 1a) = 11.2 Hz, J(1a, 6a) = 9.7 Hz, J(1a, 10e) = 4.2 Hz, 1H, Ha-1); 4.57 (d, J(3a, 4a) = 11.6 Hz, 1H, Ha-3); 6.45 (d, J(14, 13) = 4.2 Hz, 1H, H-14); 6.68 (d, J(13, 14) = 4.2 Hz, 1H, H-13). $^{13}$C-NMR (CDCl$_3$) (ppm) δ: 77.37 (d, C-1); 72.48 (d, C-3); 49.50 (t, C-4); 70.71 (s, C-5); 51.89 (d, C-6); 22.91 (t, C-7); 34.23 (t, C-8); 37.31 (d, C-9); 41.14 (t, C-10); 136.97 (s, C-11); 132.56 (s, C-12); 112.06 (d, C-13); 121.19 (d, C-14); 21.08 (q, C-15); 22.05 (q, C-16); 174.08 (s, C-17); 40.69 (s, C-18); 38.99 (t, C-19, C-25, C-26); 27.89 (d, C-20, C-22, C-24); 36.23 (t, C-21, C-23, C-27). HR MS: 443.2481 (M$^+$, C$_{26}$H$_{37}$O$_3$NS$^+$; calc. 443.2489). [α]$_D^{25.0}$ = +7.9 (c 0.304, EtOH).

3.2. Tdp1 Assay

The pET 16B-Tdp1 was kindly provided by Dr. K.W. Caldecott (University of Sussex, United Kingdom). Recombinant Tdp1 was expressed in Escherichia coli BL21(DE3) and was purified by chromatography on Ni-chelating resin and phosphocellulose P11, as described in [29].

Real-time Tdp1 activity measurements were carried out as described in [9]. In brief, Tdp1-biosensor (50 nM) was incubated in a buffer that contains 50 mM Tris-HCl pH 8.0, 50 mM NaCl, and 7 mM β-mercaptoethanol (200 µL). The mixture was then supplemented with purified Tdp1 (1.3 nM) and various
concentrations of the inhibitor. Fluorescence measurements (Ex$_{485}$/Em$_{520}$ nm) were carried out during the linear phase of reaction (from 0 to 8 min) every 55 s. The reactions were incubated at a constant temperature of 26 °C using a POLARstar OPTIMA fluorimeter (BMG LABTECH). The influence of the compounds was evaluated by comparing the fluorescence increase rate in their presence to that of 1.5% DMSO control wells. The data were imported into the MARS Data Analysis 2.0 program (BMG LABTECH), and IC$_{50}$ values were calculated by non-linear curve fitting. Tdp1-biosensor $5'$-(5,6 FAM-aac gtc gtc ttc c-BHQ1)-3' was synthesised in the Laboratory of Biomedical Chemistry, Institute of Chemical Biology and Fundamental Medicine, Novosibirsk, Russia.

3.3. Binding Studies

Synthetic DNA encoding human Tdp1 (residues 149-608 [2]) was cloned into pET-28a(+) (GenScript), which was then transformed into *Escherichia coli* BL21(DE3) for recombinant protein production. Protein production was induced with 1 mM IPTG at 28 °C with overnight incubation. Tdp1 was purified using affinity and size exclusion chromatography. Fluorescence was measured using a PerkinElmer EnSpire Multimode Reader. Tdp1 concentration was 10 µM and compound concentrations were 1 µM, 2.5 µM, 5 µM, 10 µM, 15 µM, 20 µM, 30 µM, 50 µM, 100 µM, 200 µM, 350 µM, 500 µM, and 750 µM. Buffer was 20 mM Tris and 250 mM NaCl (pH 8). Excitation wavelength was 280 nm and intrinsic fluorescence was measured between 300 and 450 nm. Control experiments with Tdp1 or compound on its own were also conducted. Background fluorescence arising from the compounds was subtracted from the final spectrum. The total volume per well was 30 µL. Dissociation constant (K$_D$) values were calculated using the following formula (1), which takes non-specific binding into account [32].

$$I = \frac{I_{max} \times [LT]}{K_D + [LT]} + N_s [LT]$$

$I$ indicates changes in fluorescence intensity from the titrations, $I_{max}$ indicates the maximum fluorescence intensity change, [LT] is the titrated ligand concentration, and $N_s$ is the non-specific term. Non-linear curve fitting was conducted using SigmaPlot 13.0 (Systat Software, San Jose, CA, USA). Experiments were conducted in triplicate and the errors shown are standard derivations.

3.4. Molecular Modelling Methods

Molecular modelling and the generation of molecular descriptors of Tdp1 inhibitors were conducted as described in [10]. Briefly, the crystal structure of Tdp1 was used for docking (PDB ID: 1MU7) [33] with the GOLD v5.4 software and the molecular descriptors were generated using QikProp v5.2 [34,35].

4. Conclusions

In this study, we reported a new class of Tdp1 inhibitors that are based on the (-)-isopulegol-derived octahydro-2H-chromene scaffold. By using a three-step reaction, a series of chiral heterocyclic compounds that contain the octahydro-2H-chromene scaffold and different amide substituents were synthesised. These structural series of compounds were found to be good inhibitors and binders of Tdp1, with IC$_{50}$ and K$_D$ values in the low micromolar concentration range. In particular, the octahydro-2H-chromen-4-ol derivative with 1-adamantane moiety (4S)-13 appeared to be the most potent inhibitor of Tdp1 among this series, with an IC$_{50}$ value of 1.24 µM. The plausible binding modes of these compounds to Tdp1 were also predicted, which found that these compounds may block the access of substrates to the enzyme active site. Overall, our results helped establish a structural-activity relationship for these novel octahydro-2H-chromene-based Tdp1 inhibitors. This information will be useful for future inhibitor development work for Tdp1.

**Supplementary Materials:** The Supplementary Materials are available online.
Author Contributions: Designing the experiments, performing chemistry experiment, writing—original draft preparation, N.S.L.-Z.; performing Tdp1 assay, A.L.Z. and A.A.C.; supervision, project administration, writing—review and editing, I.K.H.L., J.P., K.P.V., N.F.S., and O.I.L.; molecular modelling, J.R. and A.Z.; binding studies J.P. and I.K.H.L. All authors read and approved the final manuscript.

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**Sample Availability:** Samples of the compounds are available from the authors.
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