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DOI: 10.1016/j.jconrel.2024.03.036

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Document Version Publisher's PDF, also known as Version of record

Citation for published version (Harvard):

Wilson, JJ, Bennie, L, Eguaogie, O, Elkashif, A, Conlon, PF, Jena, L, McErlean, E, Buckley, N, Englert, K, Dunne, NJ, Tucker, JHR, Vyle, JS & McCarthy, HO 2024, 'Synthesis and characterisation of a nucleotide based pro-drug formulated with a peptide into a nano-chemotherapy for colorectal cancer', *Journal of Controlled Release*, vol. 369, pp. 63-74. https://doi.org/10.1016/j.jconrel.2024.03.036

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Synthesis and characterisation of a nucleotide based pro-drug formulated with a peptide into a nano-chemotherapy for colorectal cancer

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ARTICLE INFO

Keywords: Selenium RALA Dinucleoside tetraphosphate Anticancer Non-viral Nanoparticles dUTPase Mechanochemsitry

ABSTRACT

Recent studies in colorectal cancer patients (CRC) have shown that increased resistance to thymidylate synthase (TS) inhibitors such as 5-fluorouracil (5-FU), reduce the efficacy of standard of care (SoC) treatment regimens. The nucleotide pool cleanser dUTPase is highly expressed in CRC and is an attractive target for potentiating anticancer activity of chemotherapy. The purpose of the current work was to investigate the activity of P^1 , P^4 -di (2',5'-dideoxy-5'-selenouridinyl)-tetraphosphate (P4-SedU2), a selenium-modified symmetrically capped dinucleoside with prodrug capabilities that is specifically activated by dUTPase. Using mechanochemistry, P₄-SedU₂ and the corresponding selenothymidine analogue P_4 -SeT₂ were prepared with a yield of 19% and 30% respectively. tively. The phosphate functionality facilitated complexation with the amphipathic cell-penetrating peptide RALA to produce nanoparticles (NPs). These NPs were designed to deliver P4-SedU2 intracellularly and thereby maximise in vivo activity. The NPs demonstrated effective anti-cancer activity and selectivity in the HCT116 CRC cell line, a cell line that overexpresses dUTPase; compared to HT29 CRC cells and NCTC-929 fibroblast cells which have reduced levels of dUTPase expression. In vivo studies in BALB/c SCID mice revealed no significant toxicity with respect to weight or organ histology. Pharmacokinetic analysis of blood serum showed that RALA facilitates effective delivery and rapid internalisation into surrounding tissues with NPs eliciting lower plasma C_{max} than the equivalent injection of free P₄-SedU₂, translating the *in vitro* findings. Tumour growth delay studies have demonstrated significant inhibition of growth dynamics with the tumour doubling time extended by >2weeks. These studies demonstrate the functionality and action of a new pro-drug nucleotide for CRC.

1. Introduction

Colorectal cancer (CRC) is the 4th most common cancer with the 2nd highest mortality rate in the UK at 11% and ~117 new daily diagnoses [1]. Current CRC standard of care (SoC) includes FOLFOX (Folinic acid, 5-fluorouracil and oxaliplatin) and FOLFIRI (Folinic acid, 5-fluorouracil and irinotecan) both of which contain 5-FU, a uracil homologue that functions to terminate DNA synthesis, causing cancer cell apoptosis [2,3]. Other SoC includes Capecitabine, and Trifluridine which are nucleoside analogues derived from pyrimidines that compete with endogenous nucleosides to yield antimetabolic activity [4–6].

Despite advancements in systemic therapy, SoC prolongs survival by 5-years [7]. This poor prognosis is attributed to numerous 5-FU resistance mechanisms; from innate evolution of drug transporters to acquired mutation of metabolic enzymes [8]. 5-FU is readily metabolised *in vivo*, leading to rapid clearance of the drug, necessitating prolonged treatment regimens and high doses of the therapeutic [9]. Additionally, 5-FU has poor membrane permeability and has been shown to be exported from cells *via* ATP-binding cassette (ABC) transporters, leading to resistance [10,11]. This may lead to reduced efficacy due to the inability of the drug to exert its intracellular therapeutic effects. With 90% of metastatic patients becoming resistant [12]. This necessitates the

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https://doi.org/10.1016/j.jconrel.2024.03.036

Received 5 December 2023; Received in revised form 1 March 2024; Accepted 18 March 2024 Available online 23 March 2024 0168-3659/© 2024 The Authors. Published by Elsevier B.V. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/). need for new chemotherapies [13]. The enzyme, 2'-deoxyuridine 5'triphosphate nucleotidohydrolase (dUTPase) is overexpressed in CRC compared to normal/healthy tissues by a factor of 2–5 [14,15]. dUTPase protects the cell by preventing genotoxic incorporation of 2'-deoxyuridine (dU) into DNA by converting the 2'-deoxyuridine 5'-triphosphate (dUTP) into the corresponding monophosphate, enabling cancer proliferation (Fig. 1) [16,17]. Furthermore, dUTPase can cause resistance to 5-FU by preventing bioaccumulation of the active 5-FU metabolite; 5fluoro-2'-deoxyuridine 5'-triphosphate (5F-dUTP) [18].

Studies by Wilson [19] *et al* have demonstrated the function of wildtype and mutant p53 on the transcriptional modulation of dUTPase by the E2F-1 and Sp1 genes [20,21]. In HCT116 CRC cells, p53-mediated DNA-damage response caused by oxaliplatin treatment resulted in transcriptional downregulation of dUTPase, potentially sensitising cells to treatment with 5-FU [22]. Moreover, elevated dUTPase has been found in a range of CRC cancer cells including SW60 cells [23], which also display characteristic TS inhibition resistance attributed to the expression of dUTPase [24,25]. With 60% of CRC patients showing p53 dysregulation [26] the interplay between p53 and dUTPase expression, and the role in resistance, dUTPase is an attractive target for enhancing TS inhibitors yielding greater anticancer activity [27–30].

Kool [23] created a novel luminescence-based assay for the detection of dUTPase activity. This inspired the synthesis of a symmetric dinucleoside tetraphosphate designed to be specific to dUTPase and exploit its action. A symmetric tetraphosphate has the advantage of 'dual dosing', as the compound can be cleaved twice by dUTPase to yield two active moieties. However, phosphorylated compounds have limited bioavailability within cells and need to be delivered intracellularly for a therapeutic effect [31]. Intracellular barriers include crossing the cell membrane, escaping the endocytic cycle, efflux pumps, and non-specific phosphatases [32–34].

RALA is an amphipathic, fusogenic, cationic peptide that can penetrate through extra and intracellular barriers [35]. RALA is rich in arginine which confers positive charges at a physiological pH to encapsulate anionic cargoes, such as nucleic acids [36], and bisphosphonates [37] with efficient intracellular delivery of these cargos through fluorescent imaging. From these studies, it is clear that

RALA nanoparticles with a size of <200 nm and a positive charge >20mv readily enter cells and escape endosomes, irrespective of the cargo [38,39]. RALA NPs have been utilised for injectable therapeutic and prophylactic vaccines as well as biomaterial devices such as microneedle arrays [40], thermo-responsive hydrogels [41], and electro-spun wound healing patches delivering a wide range of molecules [42]. Indeed, RALA/miR26a formulations remained stable over 28 days and up to 40 °C with respect to zeta potential and z-average [43]. Nanoparticles also remain intact and functional following exposure to heat inactivated and non-heat inactivated serum [43]. RALA delivers cargo in vivo, resulting in upregulated protein expression (for nucleic acid delivery) and therapeutic effects (for small molecule delivery) [38,44]. Nanoparticle-mediated delivery can be especially beneficial in oncology, where a discontinuous epithelium, coupled with leaky vasculature and reduced lymphatic drainage can lead to accumulation of the therapeutic at tumour sites. This is referred to as the enhanced permeability and retention (EPR) effect [45].

RALA NPs enable passage through cell membranes, are subsequently encased by an endosome and subjected to an environment of pH \approx 4.5 by v-ATPase mediated proton influx [46,47]. Protonation of histidine instigates α -helicity of the peptide, aided by the glutamic acid and leucine residues. This concerted action causes endosomal membrane destabilisation, increases to the osmotic potential, and leads to release of the cargo from the endosome into the cytoplasm. Overall, the residues' combinatorial effects facilitate cargo in the cytosol and confer protection from degradation [36]. These effects have been demonstrated experimentally by performing transfections in the presence of chloroquine, a known endosomal disrupting agent, with no significant effect on transfection noted [38].

This study reports for the first time on the synthesis of a dinucleoside tetraphosphate prodrug that incorporates selenium which provides additional anticancer activity through a variety of proposed mechanisms including immune regulation, oxidative signaling and chemo preventive mechanisms [48–50]. The bioisostere modification changing the 5' oxygen to selenium is postulated to not interfere with recognition by dUTPase [23,51,52], allowing liberation of an unstable monophosphate that spontaneously degrades to form an active selenol species, that

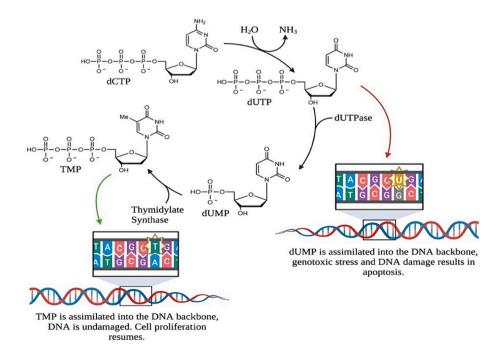


Fig. 1. Role of dUTPase. dUTP occurs naturally in the nucleotide pool by spontaneous deamination of dCTP. Assimilation of dUTP into the DNA chain yields genomic stresses and can result in apoptosis. The cell is protected by dUTPase by converting accumulated dUTP into dUMP which is sequentially converted to TMP by thymidylate synthase (TS).

exerts anticancer activity [48,50,53]. It is also hypothesised that complexation with RALA will result in improved bioavailability and efficacy as determined *via* i) the optimal mole ratio (MR) of RALA/P₄-SedU₂ in terms of physiochemical characteristics, cellular uptake, functionality and specificity for dUTPase; ii) the pharmacokinetic profile of RALA/P₄-SedU₂ NPs compared to P₄-SedU₂ alone; and iii) the *in vivo* safety profile, dosing profile and anti-tumour efficacy.

2. Materials & methods

Information regarding total synthesis of P₄-SedU₂ and P₄-SeT₂ including NMR, MS, HPLC spectra and methodologies, detailed nanoparticle formulation rationale, lyophilisation recipe, RT-qPCR primers and full *in vivo* protocols, can be found in the supplementary information. RALA (NH₂-WEARLARLARALARALARALARALARALARALARACEA-CO₂H) (RFM: 3327.98 g mol-1) was synthesised by solid state FMOC Chemistry and provided in the acetate salt form at \geq 95% purity as a lyophilised powder (Biomatik, USA) and stored at -20 °C. RALA when used is reconstituted to 1.0 µg/µL (determined using Nano-Drop 2000C spectrophotometer using pre-set protein A280 setting (Thermoscientific, USA) for use to formulate nanoparticles. For lyophilisation protocols, trehalose (Pfanstiehl, USA, No. T104–4) was dissolved in Ultrapure DNase / RNase free water (Invitrogen, UK) at 20% *w/v* was used as the cryoprotectant and lyoprotectant.

2.1. Formulation and characterisation of RALA nanoparticles

Nanoparticles (NP) were assembled by pipetting the required volume of P₄-SedU₂, or P₄-SeT₂ at 0.5 mg/mL to achieve 1 nmol in solution followed by an aliquot of H₂O and finally varying volumes of RALA at 1.0 mg/mL (to give stoichiometric ratios of 1:1 to 10:1 peptide/cargo) in a final volume of 50 µL. The solution was incubated at ambient temperature (~21 °C) for ~30 min before characterisation. Formulated NPs were characterised using a Malvern Nano-ZS Zetasizer (Malvern Instruments, UK). 50 µL of NP sample was placed in a 2 mL disposable microcuvette and was used to characterise the NP size using the Zaverage parameter measured in diameter nm (d.nm) by means of dynamic light scattering (DLS). The 50 μL NP sample was diluted to 1 mL and transferred to a 1 mL disposable folded-capillary zeta-cell cuvette and was used to characterise the NP charge potential using the zeta potential parameter and measured in mV by means of laser doppler velocimetry (LDV). Both analyses were carried out at 20 °C. For transmission electron microscopy (TEM), 2:1 mol ratio of RALA/P₄-SedU₂ NPs were produced, lyophilised, and loaded onto carbon reinforced 400 mesh copper grids (TAAB laboratories, UK) for 15 min before overnight drying, subsequent staining with Uranyless EM stain (EMS, USA) for 5 min at RT and dried overnight. The samples were imaged using a JEOL-JEM 1400 Plus TEM (JEOL, USA) at a 80 kV accelerating voltage.

2.2. Cell culture & maintenance

HCT116 CRC and HT29 CRC cell lines (ATCC, USA) were maintained in McCoy's 5 A serum; modified with L-Glutamine (Invitrogen, UK) and supplemented with 10% FCS (Invitrogen, UK). NCTC-929 fibroblast cell lines were maintained in Dulbecco's Modified Eagles Medium (Invitrogen, UK) and supplemented with 10% foetal calf serum (Invitrogen, UK). All cells were cultivated in T75 flasks and grown as monolayers in a humidified incubator at 37 °C and 5% CO₂ conditions. At *ca.* 80% confluency, cells are passaged using 10% *v*/v trypsin in phosphate buffered saline (PBS) to detach the monolayer and then neutralised with a corresponding volume of culture media, followed by centrifugation and removal of the supernatant. The cell pellet was resuspended in culture media and re-seeded in flasks to maintain exponential growth. Cell genotypes are authenticated by short-tandem repeat (STR) profiling by the suppliers and routine monthly testing ensures no *mycoplasma* contamination is present during *in vitro* research.

2.3. Cell viability assay

Cell viability was assessed using alamarBlue[™] (Invitrogen, UK). HCT116 & HT29 CRC cell lines were seeded onto 96-well plates at 0.5 imes 10^4 and 1.5×10^4 cells per well respectively and incubated overnight at 37 $^\circ C$ and 5% CO_2 to adhere to the well surface to form a monolayer. Culture media was replaced with OptiMEM (Invitrogen, UK) 2 h prior to treatments with lyophilised RALA NPs to achieve final exposure concentrations from 1×10^{-4} – 1×10^{-9} M respectively. Drug-only controls were also performed to the equivalent concentrations. Cells were incubated for 6 h with the treatments and then replaced with culture media and incubated for 1–3 days at 37 $^\circ \rm C$ and 5% $\rm CO_2$ and cell viability was assessed using alamarBlueTM (non-toxic cellular reduction of resazurin to resorufin) (Invitrogen, UK) at 24 h intervals. AlamarBlue™ was added to a final concentration in culture media of 10% and incubated (avoiding light) for 2–4 h, after which 50 µL of the 10% alamarBlue™ media was transferred to black 96-well plates and recorded using fluorescence with excitation 530 nm and emission of 590 nm using a FLUOstar Omega microplate reader with MARS data analysis software (BMG Labtech, UK). Cell viability data was normalised from the untreated cells as 100% to the null-cell blank as 0% and EC₅₀ was calculated as half-maximal response using GraphPad Prism v10.0.3 (GraphPad Software, USA) and using eq. $Y = 100/(1 + 10^{((LogEC50-X)*HillSlope))}$.

2.4. Clonogenic survival assay

HCT116 and HT29 CRC cell lines were plated onto 6-well plates at a density of 1.5×10^5 cells per well and incubated overnight at 37 °C and 5% CO₂ to adhere to the well surface to form a monolayer. Culture media was replaced with OptiMEM (Invitrogen, UK) 2 h before treatments. Cells were then treated for 6 h with uncomplexed dinucleoside tetraphosphate and NPs at a concentration respective to the cell line EC_{50} , with a trehalose negative control at the equivalent percentage concentration as the reconstituted lyophilised NPs and untreated well control. Following 6 h treatments, cells were washed with PBS and detached using $1 \times$ trypsin in PBS. Culture media is added to neutralise the trypsin and cells are centrifuged at 1500 rpm for 5 min. The cell pellet is resuspended in culture media and vortexed gently to homogeneity and counted using the Beckman-Coulter counter (Beckman-Coulter, IRE). Each cell experiment is plated at cell densities of 250, 500, 750 and 1000 cells with 3 replicates per density, cells are placed back in an incubator at 37 °C and 5% CO2 for 14 days. Culture media was decanted, and colonies fixed with 0.4% crystal violet in 70% MeOH for 15 min, decanted and washed with H₂O and dried for 24-48 h before manual colony counting. Plating efficiencies and surviving fractions were calculated as per Nature Protocol [54].

2.5. Spheroid culture

HCT116 CRC cell lines were seeded onto a 96-round-bottom well plates at 0.5×10^4 cells per well and incubated overnight at 37 °C and 5% CO₂. Spheres were allowed to grow for one week and monitored to establish spheroid growth profiles. After a week, culture media was carefully replaced with OptiMEM (Invitrogen, UK) 2 h before treatments with lyophilised 2:1 RALA/P₄-SedU₂ NPs to achieve final exposure concentrations from 1×10^{-5} – 1×10^{-9} M respectively. P₄-SedU₂ only controls were also added to the equivalent concentrations. Cells were incubated for 6 h with the treatments and then replaced with culture media and incubated for 1 day at 37 °C and 5% CO₂. Spheroid volume was assessed using the Cell3iMager X neo (SCREEN, JPN) cell imager and software to calculate approximate spheroid volumes.

2.6. mRNA extraction

HCT116 and HT29 cells were plated onto 6-well plates with appropriate culture media and grown until \sim 100% confluent. Culture media

was discarded and 1 mL TRIzol reagent (Invitrogen, UK) was added to each well and scraped to completely lyse cells and dissociate nucleic acid content. TRIzol / cell lysate mixture was transferred into 1.5 mL tubes, 0.2 mL of CHCl₃ was added, and vortexed for 5 s, and then incubated for 2 min. The mixture was centrifuged for 15 mins at 12,000 g at 4 °C and the supernatant was transferred into new 1.5 mL tubes. 0.5 mL of isopropanol was added to the extracted nucleic acid mixture, vortexed for 5 s and incubated overnight at 4 °C to facilitate complete mRNA precipitation and then centrifuged for 15 min at 12,000 g at 4 °C to form a mRNA pellet. The mRNA pellet was resuspended in 1 mL of EtOH and vortexed for 5 s and centrifuged for 5 min at 7500 g at 4 °C. The supernatant was discarded, and the mRNA pellet was partially dried before resuspension in ddH2O where yield and quality was determined using a Nano-Drop 2000C spectrophotometer (Thermoscientific, USA). An A260/A280 ratio \sim 1.8–2.0 and A260/A230 ratios \sim 2.0–2.2 are considered pure.

2.7. cDNA synthesis

HCT116 and HT29 mRNA samples were used to produce the corresponding cDNA using Transcriptor First Strand cDNA Synthesis kit (Roche, SUI). mRNA was analysed and diluted with ddH₂O to obtain a working concentration of 0.1 μ g/ μ L (10 μ L total volume required per reaction well). Each reaction well requires 1 μ L of Anchored-Oligo Primer and 2 μ L of Random Hexamer Primer (this is performed in duplicate to provide samples for reverse transcriptase (RT) negative reactions). The final reaction volume is 13 μ L and the sample is placed into a thermocycler at 65 °C for 10 min. After denaturation, each reaction well has 4 μ L of 5× Reaction Buffer, 0.5 μ L RNase Inhibitor, 2 μ L DNTP mix and 0.5 μ L RTase (equivalent volume of ddH₂O is added in place for the RT negative reaction wells). The total reaction volume is 20 μ L. Reaction wells were placed into the thermocycler, at 25 °C for 10 min – 55 °C for 30 min – 85 °C for 5 min and then stored on ice. cDNA was stored at –20 °C until further use in PCR.

2.8. RT-qPCR

RT-qPCR was performed using a LightCycler® 480 SYBR Green I Master (Roche, SUI). Theoretically, 1 µg of RNA in 20 µL yields 50 ng μ L⁻¹ of cDNA. The cDNA solution is diluted to 5 ng μ L⁻¹ for experimental use. To each reaction well, 5 µL of SYBR, 0.5 µL of forward and reverse primers (5 µM), and 4 µL of cDNA is added. The SYBR and primers are prepared separately as a master mix by adding the total volume required of both SYBR and the primers together before adding 6 µL of the master mix to the cDNA reaction wells. The PCR is run on a LightCycler 480 and the annealing step must be set to 2–3 °C lower than the lowest Tm of the primers used in the experiment.

2.9. Animals

BALB/c SCID mice were purchased from Charles River Laboratory (USA) and housed in an open pathogen controlled clean zone at 21 °C at 50% relative humidity with a supply of food and water *ad libitum*. All protocols and procedures conform to the UK scientific act of 1986 and are covered by the Department of Health, Social Services and Public Safety, Project License 2903 and Personal License 2128.

2.10. Statistical analysis

Statistical analyses were performed using GraphPad Prism v10.0.3 (GraphPad Software, USA). Statistically significant differences were calculated using a one-way ANOVA, two-way ANOVA, or unpaired *t*-test, preferring parametric analysis only if normality and lognormality tests pass (alpha = 0.05), otherwise nonparametric tests are used. *p*-value of \leq 0.05 considered significant, (*) $p \leq$ 0.05, (**) $p \leq$ 0.01, (***) $p \leq$ 0.001.

3. Results & discussion

3.1. Synthesis and characterisation of selenium-modified dinucleoside tetraphosphate analogues P_4 -SedU₂ (4a) and P_4 -SeT₂ (4b)

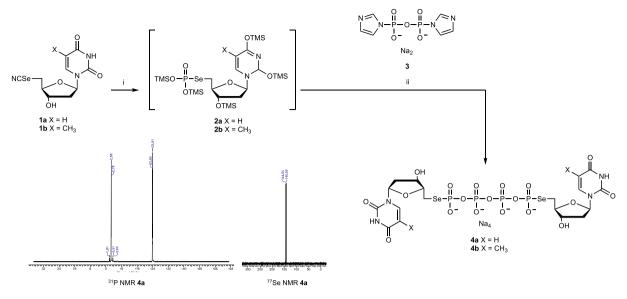
To introduce the selenium at the 5'-position of 2'-deoxyuridine (dU) or thymidine (T), b Selenocyanate-mediated displacement of a 5'-tosylate in refluxing anhydrous acetonitrile gave the corresponding nucleoside 5'-selenocyanates 1a or 1b (Scheme 1). Phosphate coupling using acid-promoted P—N bond activation under Khorana-type solution-phase conditions [55] is complicated by the sensitivity of the P—Se moiety in precursor phosphoroselenolate monoesters towards cleavage in the presence of aqueous acid [56]. In contrast, the corresponding persilylated monoesters such as the selenonucleoside derivatives 2a and 2b are relatively stable when stored under anhydrous conditions and can be readily accessed in near quantitative yields (by ³¹P NMR) following reaction of the corresponding selenocyanates (1a and 1b) with tris (trimethylsilylphosphite).

Our group has recently exploited mechanochemistry to provide access to dinucleoside pyrophosphate analogues containing P-Se-C^{5'} linkages [57] *via* intermediates such as 2b following *in situ* hydrolytic desilylation and coupling with AMP-morpholidate [58,59] in a vibration ball mill (VBM). More efficient phosphate coupling in a VBM was described subsequently using reactive NMP-imidazolide substrates generated *in situ* to access di-, tri- and tetraphosphate-linked dimers. [60,61] Following these latter reports, we attempted *in situ* hydrolytic desilylation and coupling of 2a or 2b with half an equivalent of the pyrophosphate bis-imidazolide coupling partner 3 [62] under mechanochemical activation.

Crude reaction mixtures were analysed by ³¹P NMR following VBM for 90 min. Under optimised conditions (1.5 equivalents MgCl₂.6H₂O and 13 equivalents H₂O) complete consumption of 3 (δ_P -20.94) was observed. The reaction mixtures were extracted from the vessel in buffer (pH 7.4) and the filtered suspension subjected to ion-pair, reversed-phase HPLC purification. Pure 5',5"-linked dinucleoside tetraphosphate analogues were isolated following desalting with a volatile buffer, co-evaporation with water and finally ion exchange chromatography. The sodium salts of P₄-SedU₂ (4a) and P₄-SeT₂ (4b) were fully characterised by multinuclear NMR (*e.g.*, ³¹P and ⁷⁷Se - Scheme 1) and mass spectrometry.

3.2. Formulation and characterisation of RALA NPs

RALA NPs were synthesised using mole ratios (RALA:P4-SedU2 and RALA: P₄-SeT₂) from 1:1 to 10:1 and then lyophilised to overcome coldchain storage limitations with trehalose as the lyoprotectant [63]. Previous work on the RALA peptide by Massey et al [64] revealed the capacity of RALA to complex phosphorous containing small molecules. Those formulations were based on a mass-mass ratio of cargo and RALA. The use of a mole-mole relationship was employed in this study as it better describes the quantity of RALA molecules necessary for successful encapsulation of a single molecule of cargo. Using mole ratios, which can be compared across molecules, P₄-SedU₂ and P₄-SeT₂ successfully form NPs upon mixing with RALA (Fig. 3), apart from mole ratio 1:1 with dTSep4SedT (shown in supplementary information). Ideal NPs were produced with a 2:1 mol ratio for RALA: P4-SedU2 NPs and 5:1 mol ratio for RALA: P₄-SeT₂ NPs. The NPs had a size <150 nm diameter (size and morphology of particles is confirmed with transmission electron microscopy (TEM)), a charge >10 mV and a relatively homogenous population with a PdI of <0.4 (Fig. 2); characteristics that are ideal for cellular uptake and stability [65,66]. P₄-SeT₂ NPs are larger, less homogenous and require more RALA to successfully form NPs compared to P₄-SedU₂. This may be attributed to steric hinderance by the C5 methyl, reducing electrostatic interactions between with RALA [67]. Lyophilisation of the 2:1 mol ratio RALA: P₄-SedU₂ NPs improved the PdI to <0.2, and had no significant impact on the distribution spectra of 5:1



Scheme 1. Synthesis of selenium modified dinucleoside tetraphosphates. i) BSA (13.1 eq.) TMSO₃P (1.1 eq.), CHCl₃, rt, 12 h (2a, 2b > 99%, 4b > 99% - by ³¹P NMR). ii) 3 (0.5 eq.), MgCl₂.6H₂O (1.5 eq.), H₂O (13.0 eq.), 30 Hz, 1.5 h (4a 19\%, 4b 30\%).

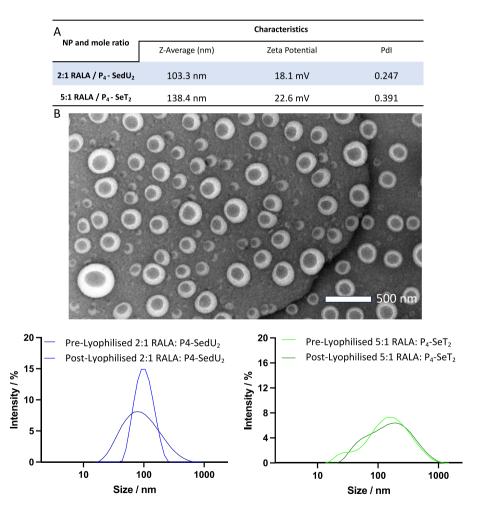


Fig. 2. Nanoparticle formulation and characterisation of lyophilised RALA NPs. A) Size, charge, and Polydispersity characteristics of lyophilised 2:1 RALA/P₄-SedU₂ NPs and lyophilised 5:1 RALA/ P₄-SeT₂ NPs. B) Transmission Electron Microscopic image of lyophilised 2:1 RALA/P₄-SedU₂ NPs (Scale bar = 500 nm). C) DLS spectrographs of pre and post lyophilised 2:1 RALA/ P₄-SedU₂ NPs, and pre and post lyophilised 5:1 RALA/ P₄-SeT₂ NPs.

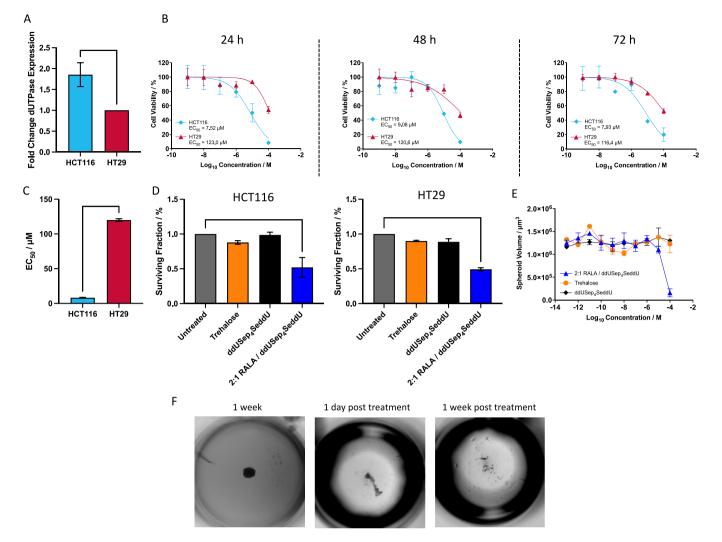


Fig. 3. dUTPase activity and anticancer efficacy in HCT116 and HT29 cells. A) Fold change difference in dUTPase expression by RT-qPCR in HCT116 cells compared to HT29 cells. Results are normalised to dUTPase expression in HT29 cells. B) Cell viability of HCT116 (light blue) and HT29 (red) cells with lyophilised 2:1 RALA/P₄-SedU₂ NPs over 3 days at 24 h intervals. C) Difference in average EC₅₀ value over 3 days between HCT116 and HT29 cells when treated with 2:1 RALA/P₄-SedU₂. D) Clonogenic survival assays on HCT116 and HT29 cells with treatments of trehalose, P₄-SedU₂ only and 2:1 RALA/P₄-SedU₂ at the corresponding observed EC₅₀ values of 2:1 RALA/P₄-SedU₂ treatments. E) Spheroid volume of HCT116 cells after 7 days after treatment with trehalose, P₄-SedU₂ only and 2:1 RALA/P₄-SedU₂. F) Photographs of spheroid of HCT116 cells, 1 week after plating, 1 and 7 days after treatment with 2:1 RALA/P₄-SedU₂.

mol ratio RALA: P_4 -SeT₂ NPs (Fig. 3). These findings prove that the formulated nanoparticles can be lyophilised, allowing ease of storage without compromising physicochemical characteristics.

3.3. Potentiating an anticancer response in vitro

The anti-cancer effects of the 2:1 RALA/P₄-SedU₂ were interrogated *via* cell viability, clonogenic assay, and 3D culture experiments on the HCT116 and HT29 cell lines; two colorectal cancer cell lines with reported high and low dUTPase expressions respectively [68–70]. This was confirmed by RT-qPCR revealing dUTPase expression is $1.9 \times$ fold higher in HCT116 cells compared to HT29 (Fig. 3 A). This was reflected in the viability assay where 2:1 RALA/P₄-SedU₂ (Fig. 3 B and C) had significantly lower anticancer activity ($p \leq 0.0001$) in HT29 cells with an EC₅₀ of >115 µM compared to >7.5 µM in the HCT116 cell across all time points. Despite a mere 1.9-fold difference in dUTPase there was a > 15 fold enhancement in activity which is indicative of significant potentiation. These results indicate that higher dUTPase levels present in HCT116 cells facilitate greater release of the active metabolite, leading to greater cytotoxicity.

Clonogenic survival studies (Fig. 3 D) were performed to assess

impacts on tumourigenicity, using corresponding 3-day EC_{50} dose of the 2:1 RALA/P₄-SedU₂ NP treatments of 8.2 µM on HCT116 and 360 µM on HT29 which resulted in 50% reduction in colony formation. Trehalose was also included as a control as carbohydrate can potentiate an independent m-TOR pathway to induce autophagy [71,72] thus promoting proliferative effects in HCT116 cells. However, the addition of trehalose controls had no significant effect on proliferation on either cell line. The 2:1 RALA/P₄-SedU₂ NP treatments outperformed free P₄-SedU₂, indicating that efficient intracellular trafficking of the drug is required to exert a cytotoxic effect.

The 3D culture yielded a reduction in anti-cancer activity when treated with an EC_{50} of 22.33 µM in HCT116 spheroids (Fig. 3 E), which could be attributed to the formation of an extra cellular matrix (ECM) from increased cell-cell interactions. The ECM can act as a barrier to drug delivery by trapping NPs within the collagen rich matrix. Additionally, penetration into the inner core is diminished in spheroid cultures. Although spheroids provide a pseudo-tumour environment, with ECM, hypoxia and angiogenic factors present, fundamentally they do not have the tortuous vasculature to support the transport of nanoparticles [73,74]. Other studies have found similar results with the EC_{50} increasing between 2D and 3D cultures depending on; the 3D scaffold

used, cell type, particle size, and the relative tissue densities within the 3D culture [75–77].

It was observed with the highest dose of 100 µM, that the cellular structures were abolished after 7 days, (Fig. 3 F) leaving scattered fragments indicating internal stresses on the cellular architecture. Selenol species' are highly reactive and can destabilise disulfide bridges [78-80], that are found in lysosomal proteins, lumen of the RER, mitochondrial intermembrane space, and collagen III [81]. In addition, the potential to trigger nucleocytoplasmic coagulation, a terminal cascade triggered by cellular-injury causes proteolytic destruction of the cellular contents into macromolecular aggregates. This then results in apoptosis to prevent uptake of misfolded proteins [82]. It has been shown in nucleic acid and small molecule delivery studies that problems typically arise due to the extra cellular effects in 3D models that are not found in 2D cultures. However, efficacy in a 3D model does provide confidence that a therapeutic effect would be observed in vivo [83-85]. Herein, the 3D model results indicate that the potency of the RALA/P₄-SedU₂ NPs should be retained in vivo.

3.4. Specificity of dUTPase in vitro

The 5:1 RALA/P₄-SeT₂ NPs were used on the HCT116 cell line to test the specificity of dUTPase to the uracil motif on the drug molecules. Work by Vertessy [52,86] used computational modelling to show that the dUTPase active site is robust enough to differentiate dUTP and dTTP and prevent mis binding. These findings were confirmed *in vitro* by Kool [23]. 5:1 RALA/P₄-SeT₂ NPs do not induce any anticancer activity (Fig. 4) supporting the specificity of P₄-SedU₂. Experimental data provides evidence to support the hypothesis that the dUTPase active site is specific to uridine, where scission between the α and β phosphates occurs to liberate a monophosphate and triphosphate species, to release the anti-cancer potential of the selenols (Fig. 4). Zeng observed a 2.2-fold increase in cellular apoptosis with 5 μ M treatments with methylselenol on HCT116 cells. Zeng also reported that selenol species interrupt protein-thiol moieties on redox responsive signaling and transcription factors (E2F) [19] ultimately leading to G₁ arrest, and diminution of MMP-2 and VEGF [49,87–89]. Further investigations into the downregulation of MMP-2 and VEGF would need to be performed to build a better understanding of the downstream mechanisms with P₄-SedU₂. Furthermore, the uncomplexed materials (Fig. 3 D & Fig. 4) also do not elicit anti-cancer activity, revealing the necessity of RALA, as negatively charged molecules are unable to penetrate the cell [90,91].

The evidence presented indicates that patients could be stratified based on dUTPase expression prior to treatment with RALA/P₄-SedU₂, and those with the highest dUTPase expression could be predicted to have an improved response as this would be the first dUTPase targeted chemotherapy. This personalised approach to treatment is advantageous compared to other chemotherapies such as cisplatin which are administered indiscriminately and target all genomic DNA in replicating cells [92–96].

3.5. Establishing safety and pharmacokinetics

To investigate potential anticancer effects *in vivo* on tumour models, safety studies were initially performed to establish potential cytotoxicity. Firstly, *in vitro* treatments with 2:1 RALA/P₄-SedU₂ NPs were

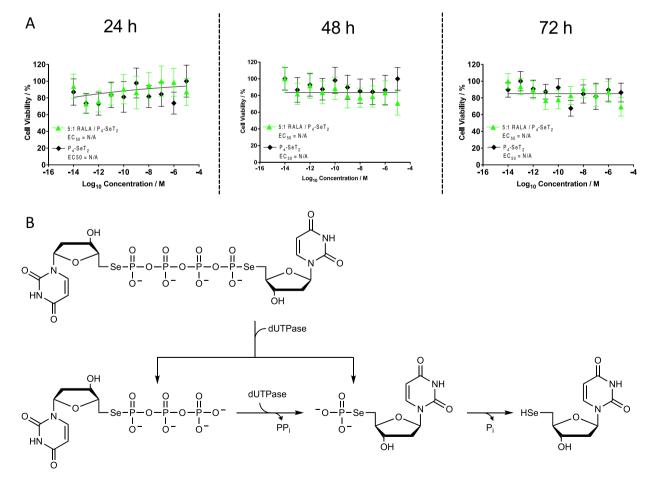


Fig. 4. Determination of 5:1 P_4 -SeT₂ NPs EC₅₀ on dUTPase positive HCT116 cells. A) Cell viability studies on HCT116 cells over 72 h following treatment with uncomplexed P_4 -SeT₂ or 5:1 RALA/P₄-SeT₂ NPs. B) Theoretical mechanism of action of dUTPase activity on the P_4 -SedU₂. Scission between the α and β phosphates may liberate an unstable monophosphate species which dephosphorylates the corresponding selenol, which could cause anticancer activity.

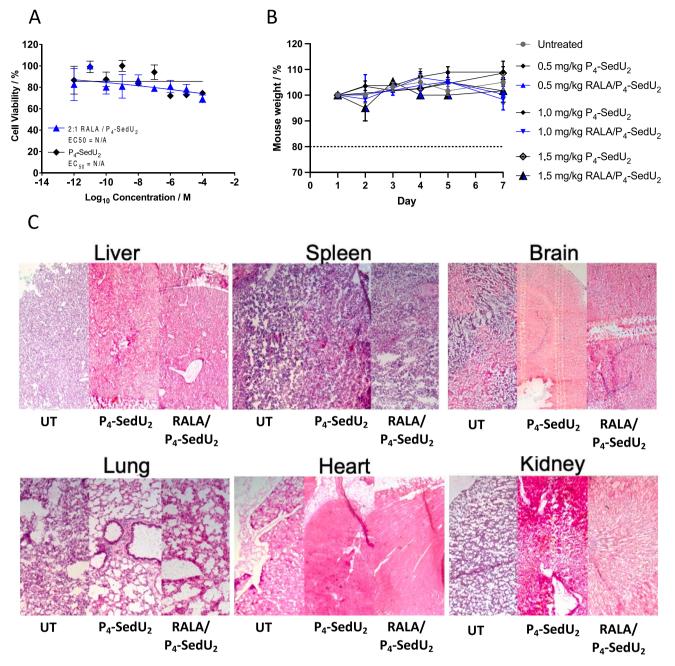


Fig. 5. Investigation into the safety profile of P_4 -SedU₂ and RALA NPs. A) Cell viability assay on NCTC-929 cells with 2:1 RALA/ P_4 -SedU₂ and P_4 -SedU₂ only after 24 h. B) Mouse weight measurements of MTD study. Untreated control, 0.5–1.5 mg/kg doses of P_4 -SedU₂ only and 2:1 RALA/ P_4 -SedU₂, NPs. C) Histology slides of H&E-stained mouse organ sections from MTD study, $10 \times$ magnification. Each slide represents the corresponding organ and directly comparing treatments. UT = untreated, P_4 -SedU₂ = 1.5 mg/kg 2:1 RALA/ P_4 -SedU₂, and RALA/ P_4 -SedU₂ = 1.5 mg/kg P_4 -SedU₂ only.

performed on the NCTC-929 murine fibroblast cell line to investigate the impacts on non-neoplastic tissues (Fig. 5 A). This revealed a non-cytotoxic response, as there is no significant impact on cell survival with the concentration used for an effective treatment on HCT116 cells, since a cell viability <70% is considered cytotoxic [97].

A minimal effective dose (MED) study was performed on BALB/c SCID mice; using doses of 0.5 mg/kg, 1.0 mg/kg and 1.5 mg/kg of either 2:1 RALA/P₄-SedU₂ NPs or P₄-SedU₂ only (Fig. 5 B). Results revealed no significant weight change throughout the week of experiment, and no mice weights dropping below the 20% weight-loss threshold. In addition, health checks 1 h, 3 h, 5 h, and 24 h post treatment showed no signs of distress or negative behavioural characteristics. This was supported by histology of vital organs (Fig. 5 C). Evaluations of the organ sections

of untreated and treated mice with 1.5 mg/kg (0.5 & 1.0 mg/kg not shown) of both P₄-SedU₂ only and 2:1 RALA/P₄-SedU₂ NPs displayed no signs of necrotic tissue within the gross structures.

Pharmacokinetic (PK) analysis was performed to investigate differences between P₄-SedU₂ only and 2:1 RALA/P₄-SedU₂ NPs with respect to drug exposure, bioavailability, clearance, metabolism, and excretion [98,99]. Mice were treated with 1.5 mg/kg P₄-SedU₂ only and 2:1 RALA/P₄-SedU₂ NPs, and sampled at 0.25 h, 0.5 h, 1 h, 5 h post treatment (Fig. 6). The P₄-SedU₂ only treatment is highly present in the blood plasma compared to the 2:1 RALA/P₄-SedU₂ NP treatment, with a C_{max} of 21.174 mg/L compared to 2.394 mg/L respectively. The AUC is 76.74 mg*h/L for the P₄-SedU₂ only, which was $4.4 \times$ higher than then 2:1 RALA/P₄-SedU₂ NPs treatment. Both T_{max} values were the same for both

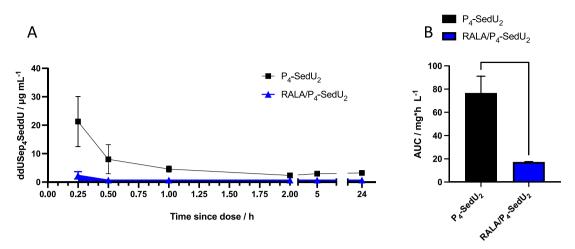


Fig. 6. Pharmacokinetic Data of 1.5 mg/kg P₄-SedU₂ and RALA/P₄-SedU₂ NP treatments. A) Recovered P₄-SedU₂ (μ g/mL) over time (from 0.25 to 24 h, N = 3 for each time point) Area under curve (AUC) highlighted. Concentration of P₄-SedU₂ was extrapolated from standard curve of peak area (R² = 0.9922) in mg/L. B) AUC plot and unpaired *t*-test statistics reveal significance ($p \le 0.05$) between the two treatments, AUC represents total exposure to analyte N = 3.

Table 1

 P_{K} parameters extrapolated from PK profile, Area Under the Curve (AUC), Maximum concentration (C_{max}), Time to maximum concentration (T_{max}) and putative half-life (T_{V}).

	AUC _[0-24 h] [mg*h/L]	C _{max} [mg/ L]	T _{max} [min]	T _½ [min]
P ₄ -SedU ₂	76.74	21.17	15	23.3
2:1 RALA/P ₄ -SedU ₂ NPs	17.36	2.39	15	17.1

treatments, showing highest concentrations with the first sample point. P₄-SedU₂ had a longer half-life compared to the 2:1 RALA/P₄-SedU₂ NPs treatment, with a $T_{1/2}$ value of 23.3 min compared to 17.1 min (Table 1).

P_K studies of prodrugs typically investigate the concentration of the active metabolite [100], however in this study only the parent molecule was observed. The putative metabolite ddUSeH was not readily available to build the standard curve by HPLC. The metabolite could not be extracted from an enzyme digest study, as during the synthesis of P₄-SedU₂. the ddUSeH species spontaneously forms the corresponding diselenide that would readily precipitate out of solution, causing issues with HPLC analysis. Such studies will be investigated in future to confirm biodistribution of the active metabolite. Herein, the results above represent the inverse effect. The AUC represents total exposure in plasma, and results indicate that the 2:1 RALA/P₄-SedU₂ NPs treatment was being delivered and retained within the exposed tissues/cells as it has a much lower concentration in plasma compared to P₄-SedU₂ which is unable to penetrate the intracellular environment and thereby is retained within the plasma and eliminated. Additionally, the rate of uptake of 2:1 RALA/P₄-SedU₂ NPs into tissues indicates a rapid process of cellular uptake as the C_{max} value was 8.8-fold less than the $P_4\mbox{-}SedU_2$ treatment, this effect is reflected with the shorter half-life time of 17.1 mins with the 2:1 RALA/P₄-SedU₂ NPs treatment.

These results correlate with other P_K analyses of prodrugs where DB289 (an anti-fungal prodrug) by Trendler exhibited similar P_K characteristics to the parent compound with a rapid T_{max} values followed by rapid elimination in Sprague-Dawley albino rats [98]. Furthermore, reduction in AUC is observed with PEGylated NPs of 5-FU and gemcitabine, while both treatments significantly improving anti-tumour efficacy in CRC and breast cancer xenograft studies respectively [101–103]. The results in the present study, indicate that RALA rapidly delivers of P_4 -SedU₂ into surrounding tissues thus preventing rapid elimination.

3.6. Tumour growth delay study

Male BALB/c SCID mice were subcutaneously implanted in the left rear flank with HCT116 CRC cells. Once tumours reached a volume of 100 mm³, mice were randomly allocated treatments of 1.5 mg/ kg of either P₄-SedU₂ only, 2:1 RALA/P₄-SedU₂ NPs or untreated. Treatments had no significant impact on the bodyweight of mice (Fig. 7 A). All Tumours developed visible dense vasculature after ~1 week of tumour size reaching 100 mm³, indicating angiogenic factors were stimulated. This supports findings by Sun with ~1.4-fold increase of VEGF-A secretion in HCT116 in hypoxic conditions ($\leq 1\%$ O₂) *in vitro* along with upregulation of CD31, CD34 and VE-cadherin [104] – endothelial factors that are associated with tumour vascular establishment [105–107]. Treatment regimens associated with 5-FU and associated adjuvants (*e.g.* Leucorovin) [108,109] consist of single intravenous treatment weekly. Herein a single dose of 1.5 mg/kg⁻¹ P₄-SedU₂ was administered weekly for three weeks.

There is evidence of effective tumour growth inhibition during treatment with 2:1 RALA/P₄-SedU₂ NPs from day 0 to ~25 while the P₄-SedU₂ only group reveal no evidence of tumour growth inhibition compared to untreated control (Fig. 7 B). Throughout the course of the study, 2:1 RALA/P₄-SedU₂ NPs improved median survival from ~15 days to 29 days (Fig. 7 C) and significantly increased tumour doubling time from 9 days to 24 days (Fig. 7 D). The delay in tumour growth and improved survival of mice treated with RALA/P₄-SedU₂ NPs shows the utility of the RALA peptide used as a delivery vehicle to ensure intracellular delivery of the drug, resulting in a therapeutic effect. The free drug delivered without RALA was no more effective than control in delaying tumour growth or improving survival.

Tumour necrosis was apparent by the coring of the tumours and surrounding necrotic tissue when tumours approached 500 mm³. Coring was only observed in the 2:1 RALA/P₄-SedU₂ NPs treatment group. Necrotic coring of a tumour is indicative of down regulation of angiogenic factors which constrains the vascularisation of the TME leading to perpetual hypoxia and glucose deprivation [110,111]. These findings correlate with other studies which have shown that selenols reduce MMP-2 and VEGF [49,87–89], contributing to down-regulation of angiogenesis causing tumour necrosis and destruction of neoplastic tissue [112–114].

Increased tumour growth was observed \sim 7 days after the final treatment, indicating that prolonged administration of this drug could retard this growth and could be possible as no rate-limiting toxicity was observed. It is possible that there is an upregulation of 'rescue' factors: Notch1, Cox-2, TGF- β , IL-8, FGF which would need to be measured.

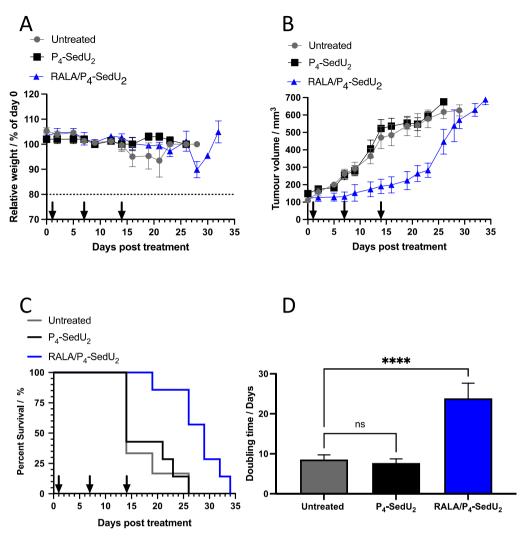


Fig. 7. Body mass, tumour growth and survival dynamics of treated BALB/c mice. A) Relative body mass of control and treated mice measured $3\times$ per week. If body mass dropped $\geq 20\%$ of starting mass, the mouse was culled. B) Tumour volume of control and treated mice over time with tumours measured $3\times$ per week. C) Kaplan-Meier curve displaying percentage survival, with terminations occurring when tumours reached 650 mm³. D) Tumour doubling time between day 0 and 14 (treatment 1 and 3). Arrows denote administration of treatments and N = 6.

Once the effects of 2:1 RALA/ P_4 -SedU₂ NPs wear off, the disproportionate presence of factors may cause elevated proliferation of the nonnecrotic neoplastic tissue [115,116], putatively causing the observed sudden increase in tumour growth.

4. Conclusions

Synthesis of the dinucleoside tetraphosphates demonstrated effective swift coupling of short lived monophosphoroselenolate species with imidazolyl activated pyrophosphates to yield stable dinucleoside tetraphosphates. Using aqueous conditions to effect quick deprotection enables the divalent metal catalysis to accomplish rapid coupling that competes with the rate of spontaneous degradation of the monophosphoroselenolate. Moreover, this results in isolation of the material by RP-HPLC, desalting, and ion exchange. The synthesised compounds were stable in standard conditions (oxic, pH 7, room temperature). Further work on the synthesis should include varying reaction times, other divalent metal additives, and ⁷⁷Se isotopic enrichment. Nucleotide modification research translates to other areas, including; anti-viral agents [16], mRNA cap analogues [117,118], systematic evolution of ligands by exponential enrichment (SELEX) [119] and mechanistic probes [120], indicating multiple biological and clinical applications.

Acknowledgements

This research was funded by the UK Department for the Economy (DfE). Graphical abstract was produced using BioRender.

CRediT authorship contribution statement

Jordan J. Wilson: Writing – original draft, Methodology, Formal analysis, Data curation, Conceptualization. Lindsey Bennie: Writing – review & editing, Methodology, Investigation. Olga Eguaogie: Methodology, Investigation. Ahmed Elkashif: Writing – review & editing, Investigation. Patrick F. Conlon: Writing – review & editing, Validation. Lynn Jena: Methodology, Investigation. Emma McErlean: Investigation, Formal analysis. Niamh Buckley: Methodology, Formal analysis. Klaudia Englert: Methodology, Formal analysis. Nicholas J. Dunne: Supervision, Formal analysis. James H.R. Tucker: Supervision, Formal analysis. Joseph S. Vyle: Writing – review & editing, Writing – original draft, Supervision, Methodology, Funding acquisition, Formal analysis, Conceptualization. Helen O. McCarthy: Writing – review & editing, Supervision, Methodology, Funding acquisition, Formal analysis, Conceptualization.

Data availability

Data will be made available on request.

Appendix A. Supplementary data

The supporting information is available free of charge at www.lin khere.com

Supporting information contains extra details on the synthesis of the dinucleoside tetraphosphates, with full reaction conditions and methods including purification. NMR, HPLC and MS details, methods, and all corresponding spectra. Includes the formulation strategy and lyophilisation protocol for the RALA NPs. Primers used for RT-qPCR. All protocols used for *in vivo* experimentation to conduct all animal studies. Supplementary data to this article can be found online at https://doi.org/10.1016/j.jconrel.2024.03.036.

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