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DNA vaccination via RALA nanoparticles in a microneedle delivery system induces a potent immune response against the endogenous prostate cancer stem cell antigen

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1 **DNA Vaccination via RALA Nanoparticles in a Microneedle**
2 **Delivery System Induces a Potent Immune Response against the**
3 **Endogenous Prostate Cancer Stem Cell Antigen**

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17

18 **Abstract**

19 Castrate resistant prostate cancer (CRPC) remains a major challenge for healthcare
20 professionals. Immunotherapeutic approaches, including DNA vaccination, hold the potential
21 to harness the host's own immune system to mount a cell-mediated, anti-tumour response,
22 capable of clearing disseminated tumour deposits. These anti-cancer vaccines represent a
23 promising strategy for patients with advanced disease, however, to date DNA vaccines have
24 demonstrated limited efficacy in clinical trials, owing to the lack of a suitable DNA delivery
25 system. This study was designed to evaluate the efficacy of a two-tier delivery system
26 incorporating cationic RALA/pDNA nanoparticles (NPs) into a dissolvable microneedle (MN)
27 patch for the purposes of DNA vaccination against prostate cancer. Application of NP-loaded
28 MN patches successfully resulted in endogenous production of the encoded prostate stem cell
29 antigen (PSCA). Furthermore, immunisation with RALA/pPSCA loaded MNs elicited a
30 tumour-specific immune response against TRAMP C-1 tumours *ex vivo*. Finally, vaccination
31 with RALA/pPSCA loaded MNs demonstrated anti-tumour activity in both prophylactic and
32 therapeutic prostate cancer models *in vivo*. This is further evidence that this two-tier MN
33 delivery system is a robust platform for prostate cancer DNA vaccination.

34

35 **Key Words:** DNA Vaccine; Microneedle; PSCA; RALA; CTL; Nanoparticle; Prostate Cancer

36 **1. Introduction**

37 As the second leading cause of cancer-related deaths of males within the UK, prostate cancer
38 represents a major challenge to healthcare professionals.[1] There is no available cure for
39 metastatic or castrate resistant disease, and therefore efficacious, new treatments are urgently
40 required.[2] Within the past couple of decades there has been growing interest in anti-cancer
41 vaccines which harness the body's immune system to target cells expressing immunogenic
42 tumour-associated antigens (TAAs).[3] Of these, DNA vaccines provide a relatively simple
43 and effective means of inducing a cellular immune response against encoded TAAs.[3]
44 However, despite producing promising results in preclinical studies, prostate cancer DNA
45 vaccines have demonstrated mixed efficacy in clinical trials.[3] Further to this, the recent
46 failure of PROSTVAC™ (Bavarian Nordic, US) to achieve the primary outcome measure of
47 increasing overall survival in a Phase III clinical trial for men with asymptomatic or minimally
48 symptomatic mCRPC has raised doubts about the efficacy of DNA vaccines as a
49 monotherapy.[4] Thus there is a need to develop novel DNA delivery systems to improve the
50 efficacy of such vaccines.

51 To develop a clinically relevant prostate cancer DNA vaccine several key issues need to be
52 considered including the choice of target antigen. Numerous immunogenic proteins are
53 expressed nearly exclusively by the prostate or malignant prostate tissue, providing multiple
54 potential targets for immunotherapy. One TAA is the Prostate Stem Cell Antigen (PSCA),[5]
55 a GPI-anchored cell surface antigen expressed in normal and malignant prostate tissue.[5]
56 PSCA expression correlates with prostate cancer grade and stage, and 94% of primary prostate
57 tumours (105/112), and 100% of secondary bone metastases (9/9) express PSCA, making it a
58 particularly strong target for advanced disease.[6] As well as being expressed by humans, a
59 murine homologue of PSCA has been identified,[7] and the establishment of mPSCA-
60 expressing murine prostate cancer cell lines from transgenic mice has provided a
61 physiologically relevant model to study the anti-tumour efficacy of prostate cancer DNA
62 vaccines.[7,8]

63 Apart from the TAA of choice, the effectiveness of a DNA vaccine is also dictated by the
64 success of the delivery system. As such, microneedles (MNs) represent an ideal delivery
65 platform for DNA vaccination purposes, as these micro-projections localise the genetic cargo
66 across the stratum corneum (SC) to a rich network of antigen-presenting cells (APCs) in the
67 dermis and epidermis.[9] The transfection of APCs and subsequent display of antigenic
68 peptides on Major Histocompatibility Complex (MHC) I and II molecules is central to the

69 stimulation and expansion of TAA-specific CD8⁺ and CD4⁺ T cells, the effectors of the
70 adaptive immune system.[3] Additionally, this physical delivery approach can easily be
71 combined with a vector-based strategy to enhance cellular uptake within this highly
72 immunogenic area. Previously, a two-tier delivery system immunised mice against the HPV-
73 16 E6 and E7 cervical cancer TAAs using nanoparticles (NP) loaded into a polymeric MN.
74 Immunisation via the NP-MN delivery platform resulted in tumour retardation in both
75 prophylactic and therapeutic tumour models,¹¹[10] but was limited by a low delivery dose and
76 poor compatibility of the NP cargo within the PVP matrix. Thorough screening of a range of
77 polymers to form the MN matrix and lyophilising NPs prior to MN formation has increased
78 both the functionality and the loading of NP cargo within the polymer matrix.[11,12] In this
79 investigation, the NP-MN platform was employed to immunise mice with encoded mPSCA,
80 and it was reported for the first time for prostate cancer DNA vaccination. Application of the
81 NP-MN system successfully resulted in local mPSCA expression within the treated ears of
82 mice. Furthermore, immunisation of mice with the NP-MN system induced a tumour-specific
83 cellular immune response, and inhibited the growth of TRAMP C-1 tumours in both
84 prophylactic and therapeutic challenge models.

85 **Materials and Methods**

86 **2.1 Preparation of RALA/pPSCA Nanoparticles (NP) and Lipofectamine/pPSCA NPs**

87 The RALA peptide (WEARLARALARALARHLARALARALRACEA) was supplied as a
88 lyophilised powder from Biomatik Corporation (USA). The powder was stored at -20°C and
89 reconstituted in DNase/RNase free water (Life Technologies, UK) for experimentation. A
90 plasmid encoding the murine prostate TAA, PSCA (pPSCA), was purchased from Origene
91 (USA) within the pCMV-AC-GFP expression vector, resulting in expression of PSCA with a
92 C-terminal turbo-GFP tag to allow visualisation and quantification of successful transfection.
93 pPSCA was transformed and propagated in *Escherichia coli* DH5 α cells (Life Technologies,
94 UK) as per manufacturer's instructions. The plasmid was isolated and purified using PureLink
95 HiPure Plasmid Filter Maxiprep Kits (Life Technologies, UK) prior to use.

96 RALA/pPSCA nanoparticles (NP) were formed at various N:P ratios as described
97 previously.[13] Briefly, the desired quantity of plasmid was added to DNase/RNase free water,
98 prior to addition of the desired quantity of RALA solution. NPs were left for 30 min at room
99 temperature (RT) prior to experimentation to ensure complete complexation.

100 Lipofectamine 2000 was purchased from Thermofischer Scientific (UK) as a positive control.
101 Lipofectamine/pPSCA lipoplexes were formed as per the manufacturer's instructions. Briefly,
102 3 µg of pPSCA and 6 µL of Lipofectamine 2000 were conditioned separately in 150 µL of
103 Opti-MEM media for 5 min at RT, and subsequently mixed together and allowed to incubate
104 for a further 5 min prior to use.

105 **2.2 Lyophilisation protocol for NPs**

106 To produce lyophilised NPs (L-NPs), NP solutions containing 5% w/v Trehalose were placed
107 into a VirTis Wizard 2.0 (SP Scientific, USA) and subjected to freezing at -40°C for 1 h, before
108 primary drying for 3 h at -35°C (120 mTorr), -30°C for 4 h (190 mTorr) and -25°C for 4 h (190
109 mTorr). Finally, samples underwent secondary drying for 18 h at 20°C (190 mTorr for 8 h and
110 50 mTorr for 10 h).

111 **2.3 Gel Retardation Assay**

112 RALA/pPSCA complexes were prepared at N:P ratios 0.25 - 12 using 1 µg of pDNA in a total
113 volume of 20 µL. 10 µL of sample was supplemented with 2 µL 5X Nucleic Acid Loading
114 Buffer (Bio-Rad, USA), and loaded on to a 1% Agarose gel incorporating 0.2 µg/mL EtBr as
115 a DNA intercalating agent. Samples were electrophoresed at 100 V for 1 h in 1X TAE buffer
116 and then visualised under UV light using a Multispectrum Bioimaging System (UVP, UK).

117 **2.4 Picogreen Encapsulation Assay**

118 RALA/pPSCA complexes were prepared at N:P ratios 0.5 - 12 using 1 µg of pDNA in a total
119 volume of 1000 µL DNase/RNase free water. 50 µL samples were added to black 96 well plates
120 in triplicate and supplemented with 50 µL Quanti-iT Picogreen reagent. Samples were left to
121 incubate for 30 min at RT prior to quantification of fluorescent emission at 520 nm using a
122 Synergy 2 Multi-Mode Microplate Reader (BioTek Instruments Inc, UK) following excitation
123 at 480 nm. The percentage of encapsulated pPSCA was calculated relative to the fluorescence
124 of a pDNA only sample.

125 **2.5 Size and Zeta Potential Analysis of RALA/pPSCA complexes**

126 RALA/pPSCA complexes were prepared for N:P ratios 1-15, in a volume of 50 µL
127 DNase/RNase free water. The hydrodynamic size and zeta potential of complexes was
128 determined using a Malvern Zetasizer NanoZS (Malvern Instrument Ltd, UK) via Dynamic
129 Light Scattering and Laser Doppler Velocimetry respectively. For stability studies, NPs were

130 prepared at N:P 10, and incubated at RT for up to 28 days prior to analysis. For temperature
131 stability studies the hydrodynamic size of NPs was determined over an increasing temperature
132 range from 4-37°C.

133 **2.6 Transmission Electron Microscopy (TEM)**

134 NPs (N:P 10) were prepared using 1 µg of pPSCA in a total volume of 30 µL. Samples were
135 coated on to Carbon coated Cu grids (Taab Laboratories Equipment Ltd, UK) by placing the
136 grid face down on to a 10 µL NP solution for 10 min. The grid was then dried overnight and
137 stained the following day with UranylLess electromicroscopy stain (Electron Microscopy
138 Services, USA) for 2 min prior to rinsing with distilled water and drying at RT. The grid was
139 then imaged using a JEM-1440Plus TEM (Joel, USA) at an accelerating voltage of 120 kV.

140 **2.7 Serum Stability Study**

141 NPs (N:P 10) were prepared using 1 µg of pPSCA in a total volume of 50 µL. Samples were
142 incubated alone or with 10% Foetal Calf Serum (FCS) at 37°C for 1-6 h. Samples were
143 subsequently either untreated or incubated with 5 µL of 10% Sodium Dodecyl Sulphate (SDS)
144 for 10 min at RT to dissociate NPs and free pPSCA. 20 µL samples were subsequently
145 electrophoresed and visualised on an agarose as described in Section 2.3.

146 **2.8 Cell Culture**

147 The immortalised murine dendritic cell line, DC 2.4, was maintained in RPMI media
148 supplemented with 10% FCS, 1% L-glutamine, 1% HEPES and 1% Non Essential Amino
149 Acids (Life Technologies, UK). The human epithelial Human Embryonic Kidney 293 (HEK-
150 293) cell line was maintained in DMEM supplemented with 10% FCS (Life Technologies,
151 UK). The murine prostate cancer cell line, Transgenic adenocarcinoma Mouse Prostate Cell
152 Line 1 (TRAMP C-1), was maintained in DMEM, adjusted to contain 4mM L-glutamine, 1.5
153 g/L sodium bicarbonate and 4.5 g/L glucose (ATCC, USA), supplemented with 0.005 mg/mL
154 Bovine Insulin, 10 nM Dehydroisoandrosterone (Sigma, UK), 5% FCS (Life Technologies,
155 UK) and 5% Nu-Serum IV (VWR, UK). For *ex vivo* experimentation TRAMP C-1 media was
156 further supplemented with 1% Penicillin/Streptomycin (Life Technologies, UK).

157 **2.9 Transfection studies**

158 For cytotoxicity studies HEK-293 and DC 2.4 cells were seeded into 96 well plates at densities
159 of 15,000 and 20,000 cells/well respectively and left to adhere overnight. Prior to transfection

160 cells were conditioned for 2 h in Opti-MEM (Life Technologies, UK). Cells were then
161 supplemented with 0.5 µg/well pPSCA complexed to RALA (N:P 0-12) for 6 h before being
162 replaced in complete media. 24 h following transfection, cells were supplemented with 10%
163 CellTiter 96 Aqueous One Solution Reagent (MTS) (Promega, UK) for 2 h, and the absorbance
164 at 490 nm was subsequently measured using an EL808 96-well plate reader (BioTek
165 Instruments Inc, UK). The percentage viability was determined relative to untreated controls.

166 For transfection studies HEK-293 and DC 2.4 cells were seeded into 6 well plates at densities
167 of 300,000 and 400,000 cells/well respectively and left to adhere overnight. Cells were
168 conditioned and treated as above with 3 µg/well pPSCA complexed to RALA (N:P 0-12). For
169 polymerase chain reaction (PCR) experiments, cells were transfected as above with freshly
170 prepared RALA NPs (N:P 10), Lipofectamine/pPSCA, or with L-NPs (N:P 10) which had been
171 reconstituted in 20% w/w 9-10 kDa PVA solution (Sigma, UK), dried overnight in non-
172 indented silicon moulds and dissolved in Opti-MEM prior to transfection.

173 For quantification of GFP expression, cells were harvested in 4% Formaldehyde (Sigma, UK)
174 48 h post-transfection and GFP expression was quantified via flow cytometry using a BD
175 Accuri C6 Plus Flow Cytometer (BD Biosciences, UK). Data was analysed using the machine's
176 in-built software and fluorescent intensity is reported at 4% gating to untreated cells.

177 For determination of PSCA expression via amplification PCR, cells were harvested in TRIzol
178 reagent (Thermofischer Scientific, UK) and mRNA was extracted as per the manufacturer's
179 instructions. Prior to RT-PCR reactions, mRNA was treated to remove any contaminating
180 pDNA from samples using the DNase I, RNase-free kit (Thermofischer Scientific, UK) as per
181 the manufacturer's instructions. 1000 ng of mRNA was then reverse transcribed to cDNA by
182 supplementing samples with 10 µL of Reverse Transcription mix (4 µL 5X First Strand Buffer,
183 1 µL 10 mM dNTP mix, 1 µL Dilute Random Primer, 2 µL 0.1 M DTT, 1 µL RNase OUT
184 40U/mL, 1 µL M-MLV Reverse Transcriptase, [Invitrogen, UK]) and heating to 25°C for 5
185 min, 37°C for 90 min, and 70°C for 10 min, before cooling to 4°C in a TC-312 Thermocycler
186 (Bibby Scientific Ltd, UK). Following reverse transcription, 3 µL of cDNA sample was
187 transferred to a fresh RNase-free PCR tubes with 11 µL of PCR Mastermix, containing forward
188 (5'-CTG GCC ACC TAC TTA GCC CT-3') and reverse (5'-GCG ATG TAA AGC AAC TGT
189 GC-3') primers for the murine PSCA (mPSCA) gene. PSCA was amplified via PCR using the
190 following conditions: Initial denaturation at 95°C for 2 min, followed by 30 cycles of
191 amplification at 95°C for 1 min, 57°C for 1 min and 72°C for 1 min, and a final extension period

192 of 5 min at 72°C. Following amplification, samples were electrophoresed and visualised on a
193 2% Agarose gel as per Section 2.4. mRNA from TRAMP C-1 cells was harvested and amplified
194 as above to serve as a positive control for mPSCA expression, while untreated cells were
195 harvested and amplified to serve as a negative control for gene expression.

196

197 **2.10 Microneedle (MN) manufacture**

198 Microneedles were manufactured as previously described.[12] Briefly, L-NPs (N:P 10),
199 containing 100 µg of pPSCA, were reconstituted in 30 mg of 20% w/w 9-10 kDa PVA solution
200 and mixed manually until a homogenous solution was formed. The solution was then
201 transferred to a silicon microneedle (MN) master-mould containing 361 inverted conical
202 projections with dimensions of 300 µm diameter and 600 µm length, spaced 50 µm apart.[14]
203 The solution was centrifuged at 4,000 rpm for 5 min to fill the conical indents with NP-PVA
204 solution, following which 500 mg of inert 20% w/w 9-10 kDa PVA solution was added to the
205 mould and re-centrifuged to form a baseplate. MNs were left to dry for 48 h at RT before
206 carefully peeling from the mould and removal of excess baseplate and sidewalls with scalpel.

207

208 **2.11 Determination of mPSCA expression *in vivo***

209 All *in vivo* experiments were compliant with the UK Scientific Act 1986, and carried out under
210 project license 2794. Experimental procedures utilised male C57 BL/6 mice (Charles River,
211 UK) 6-8 weeks old which were housed at 21°C and 50% humidity, and given unlimited access
212 to food and water. NP loaded MNs were applied to the left ears (LE) of anaesthetised male C57
213 BL/6 mice for 5 min using manual pressure, and adhered in place for a further 24 h using 3M
214 surgical tape (Micropore Ltd, UK). Right ears (RE) were left untreated to serve as a negative
215 control. 24 h following MN removal mice were sacrificed, ears were harvested and placed into
216 RNALater solution (Sigma, UK) for 24 h at RT. Samples were then stored at -80°C until thawed
217 for mRNA extraction. For mRNA extraction, samples were placed into Round-Bottomed 2.0
218 mL Safe-Lock Eppendorf containing 500 µL of TRIzol Reagent and two 7 mm Stainless Steel
219 beads (Qiagen, UK). Samples were homogenised in a TissueLyser LT (Qiagen, UK) using and
220 oscillation rate of 50 osc/sec for 10 min. Following homogenisation, the mRNA extraction
221 procedure and subsequent determination of mPSCA expression via amplification PCR
222 followed the procedure outlined in section 2.9.

223

224

225 **2.12 Immunisation Schedule**

226 C57 BL/6 were immunised with approximately 100 µg of pPSCA and received a total of 3
227 immunisations, two weeks apart. Immunisation was either with naked pPSCA via i.m. injection
228 (DNA-IM), naked pPSCA via MN (DNA-MN), NP via i.m. injection (NP-IM) or NP-loaded
229 MN (NP-MN). For i.m. injections mice received a single injection into the medial thigh muscle
230 with cargo which had been lyophilised and reconstituted in 50 µL sterile water. For MN
231 treatment mice were immunised with two MNs (one on each ear), which were applied as
232 previously described,[12] and have previously been demonstrated to deliver approximately 50
233 µg of pDNA per patch.[12]

234 **2.13 Quantification of IFN-γ secretion**

235 13 days following the third immunisation, prior to splenocyte harvest, TRAMP C-1 cells were
236 plated into 24 well plates at densities of 50,000 cells/well and allowed to adhere overnight. The
237 following morning cells were irradiated for 5 min using a 160 kVp x-ray source (Faxitron X-
238 ray, USA) with a 0.8 mm filter, to serve as a source of re-stimulation for PSCA-specific T cells.
239 14 days following the third immunisation, mice were sacrificed and the spleens were
240 aseptically harvested, mechanically homogenised and resuspended in Red Blood Cell Lysing
241 Buffer Hybri-Max (Sigma Aldrich, Uk) for 5 min at RT to allow lysis of red blood cells within
242 the splenocyte mixture. Cells were subsequently co-cultured at a ratio of 50:1 with irradiated
243 TRAMP C-1 cells for up to 6 days in the presence of 20 units of murine interleukin-2 (mIL-2)
244 (Peprotech, UK) at 37°C. To allow discernment of baseline levels of cell lysis, T cells from
245 control mice were also incubated without the presence of irradiated TRAMP C-1 cells. 4 days
246 following re-stimulation in the presence of irradiated TRAMP C-1 cells, the supernatant from
247 cells was harvested and IFN-γ within the supernatant was quantified via sandwich ELISA,
248 using the Murine IFN-γ Standard ABTS ELISA Development Kit (Peprotech, UK) as per the
249 manufacturer's instructions.

250 **2.14 LDH Cytotoxicity Assay**

251 14 days following the third immunisation, the spleen from each mouse was isolated, processed
252 and stimulated as above. 6 days following re-stimulation viable T cells were harvested and
253 resuspended in 1% BSA assay medium. Cells were incubated 10:1 with viable TRAMP C-1
254 cells which had been seeded at a density of 10,000 cells/well in a 96 well plate the previous
255 day. Following 5 h incubation at 37°C, the supernatant from each well was collected and the

256 cell specific TRAMP C-1 lysis was determined by measuring the Lactate dehydrogenase
257 (LDH) release from damaged cells, using a Cytotoxicity Detection Kit [LDH] (Roche, UK).

258 **2.15 Prophylactic Tumour Challenge Study**

259 C57 BL/6 mice (N=6 or 7) underwent immunisation as in section 2.12. 14 days following the
260 third immunisation, mice were challenged with a s.c. injection of 5×10^6 TRAMP C-1 cells
261 suspended in a PBS/Matrigel (BD Bioscience, UK) mixture, into the left flank. Mice were
262 monitored for tumour growth by palpation, and measuring three times weekly. Tumour volume
263 was calculated using the following formula: $V = \frac{4}{3}\pi r^3$. Experimental endpoint was reached
264 when tumours attained a geometric mean diameter (GMD) of 10, or 100 days following tumour
265 challenge. Humane endpoint was reached when mice lost <20% of their body weight, tumours
266 became ulcerated, or mice developed persistent or severe ill health (poor conditioning,
267 appearance, gait).

268 **2.16 Therapeutic Tumour Challenge Study**

269 C57 BL/6 mice (N=5/6) were challenged with a s.c. injection of 5×10^6 TRAMP C-1 cells
270 suspended in a PBS/Matrigel (BD Bioscience, UK) mixture, into the left flank. 7 days
271 following implantation, when tumours had reached a palpable size (~2 mm), mice were
272 immunised as in section 2.12. Tumour development was measured three times weekly using
273 callipers and tumour volume was calculated as in section 2.15. Experimental endpoint was
274 reached when tumours attained a GMD of 12, or 100 days following tumour challenge.
275 Humane endpoint was reached when mice lost <20% of their body weight, tumours became
276 ulcerated, or mice developed persistent or severe ill health (poor conditioning, appearance,
277 gait).

278 **2.17 Statistical Analysis**

279 Unless otherwise stated experiments have three independent replicates and are expressed as
280 mean \pm standard error mean (SEM). Unless otherwise stated statistical differences were
281 analysed using a One-Way ANOVA with a P value ≤ 0.05 considered significant.

282 **3. Results**

283 **3.1 Physicochemical Characterisation and stability of RALA/pPSCA NPs**

284 Gel retardation analysis indicated that RALA encapsulated free pPSCA from an N:P ratio of 2
285 onwards, as shown by the inability of pPSCA to migrate through an agarose gel under current

286 from an N:P ratio of 2 onwards (Figure 1-A). This was further supported by the PicoGreen
287 encapsulation assay which indicated that at an N:P ratio of 0.5, $70.93\% \pm 10.27$ of pPSCA was
288 free in solution and able to bind PicoGreen, and that at N:P 1 the amount of free pDNA had
289 decreased to $37.97\% \pm 13.31$, this was then further reduced to $<10\%$ of pPSCA being free to
290 bind PicoGreen from an N:P ≥ 2 (Figure 1-B). However, size and charge analysis revealed that
291 RALA did not complex plasmid into cationic NPs until an N:P ratio ≥ 3 (Figure 1-C). This is
292 due to the nearly neutral charge of complexes at N:P 2, which results in large aggregates due
293 to a lack of charge repulsion. At N:P ≥ 4 NP mean hydrodynamic sizes were consistently <100
294 nm and ranged from 47.27 – 93.09 nm, with mean complex zeta potentials ranged from 4.534
295 – 13.57 mV.

296 NPs were stable over a 28 day incubation period at RT, with no significant differences in
297 hydrodynamic size or zeta potential being observed ($p > 0.05$, Figure 2-A). We did note an
298 increase in particle size at Day 14, with a correspondingly lower zeta potential, although this
299 was not significant. This is not typical for RALA-NPs, as previous studies have demonstrated
300 that the size of RALA NPs has remained < 100 nm consistently for up to 30 days.[10,15] NP
301 hydrodynamic size also remained stable over a range of temperatures from 4-37°C (Figure 2-
302 B). Finally, the stability of pPSCA within NPs was tested via serum stability (Figure 3-B). NPs
303 protected pPSCA from degradation by serum nucleases for up to 6 h at 37°C in the presence of
304 FCS.

305 **3.2 *In vitro* assessment of RALA/pPSCA NPs**

306 The effect of RALA/pPSCA NPs on the viability of HEK-293 and DC 2.4 cells was assessed
307 using the MTS assay. With both cell lines there was a trend that transfecting cells with
308 increasing N:P ratios resulted in decreasing cell viabilities, however, over the range of NP
309 ratios examined (N:P 0-12) no significant decrease in cell viability was observed in either cell
310 line (Figure 3-A).

311 Flow cytometry was used to evaluate the percentage of HEK-293 and DC 2.4 cells producing
312 GFP-tagged mPSCA following transfection at varying N:P ratios (Figure 3-B). Analysis
313 revealed that complexing pPSCA to RALA resulted in significantly higher transfection
314 efficacies at N:P ratios ≥ 4 and ≥ 10 , in HEK-293 cells and DC 2.4 cells respectively, compared
315 to using naked pPSCA (N:P 0) alone.

316 As well as quantifying protein production via flow cytometry, the production of mPSCA within
317 transfected HEK-293 (Figure 3-Ci) and DC 2.4 cells (Figure 3-Cii) was verified using end point

318 RT-PCR. Amplification of cDNA from untreated TRAMP C-1 cells, with mouse specific
319 PSCA primers generated a cDNA band of the expected 130 bp size. Expression of mPSCA was
320 not detected in untreated HEK-293 and DC 2.4 cells, as determined by a lack of visible cDNA
321 band following amplification of cDNA. Conversely, amplification of cDNA from HEK-293
322 and DC 2.4 cells transfected with freshly prepared RALA/pPSCA NPs and
323 Lipofectamine/pPSCA NPs resulted in generation of cDNA bands of the expected size,
324 indicating successful mPSCA mRNA production following transfection.

325 To demonstrate that RALA/pPSCA NPs remained functional following the MN manufacturing
326 process, cells were also transfected with L-NPs which had been dried in 9-10 kDa PVA solution
327 (Figure 3-D). As above, amplification of cDNA harvested from untreated HEK-293 and DC
328 2.4 cells did not generate a gene expression product, however, cDNA harvested from HEK-
329 293 and DC 2.4 cells following transfection yielded a 130 bp product indicating the presence
330 of mPSCA. Therefore, it can be concluded that NPs remain functional following lyophilisation
331 and incorporation into 9-10 kDa PVA gels.

332 **3.3 RALA/pPSCA-MN (NP-MN) application generates gene expression *in vivo* and elicits** 333 **an antigen-specific immune response**

334 Prior to using NP-MNs for immunisation of mice, it was established that NP-MN application
335 to the back of mouse ears generated local expression of the antigenic mPSCA. mPSCA was
336 detected via end point RT-PCR within the left ears of mice following NP-MN application, but
337 not in the untreated right ears of mice, indicating that gene expression was confined to the local
338 area of MN application (Figure 4-B).

339 IFN- γ and cytotoxicity assays were carried out to examine the antigen-specific cell-mediated
340 immune response following immunisation. The presence of IFN- γ in the supernatant of T cell-
341 enriched splenocytes from immunised and naive mice, following stimulation with irradiated
342 TRAMP C-1 cells was determined using quantitative sandwich ELISA (Figure 4-C).
343 Immunisation with NP-MN resulted in increased secretion of IFN- γ from stimulated T cell-
344 enriched splenocytes compared to naive mice. However, increased levels of IFN- γ secretion
345 into the supernatant were not detected from splenocytes harvested from mice immunised with
346 DNA-MN. The results indicate that immunisation of mice with RALA/pPSCA NPs results in
347 a greater release of IFN- γ from splenocytes in response to stimulation with tumour cells,
348 consistent with a PSCA-specific T_{H1}-predominant response.

349 The cytolytic activity of splenic T cells, from immunised and untreated control mice, against
350 TRAMP C-1 tumour cells was then assessed using the LDH release assay (Figure 4-D). Splenic
351 T cells from mice immunised with NP-MN (21.91%, $p < 0.05$) lysed significantly greater
352 numbers of TRAMP C-1 cells when co-cultured at a ratio of 10:1 than those from naive mice
353 (7.36%), indicating a robust cytotoxic immune response following NP immunisation.
354 Conversely, immunisation of mice with pPSCA alone did not result in a significant increase in
355 the TRAMP C-1 specific CD8⁺ T cell population compared to control (12.02%, $p > 0.05$).

356 **3.4 NP-MN Immunisation delays tumour uptake and slows tumour progression in a** 357 **prophylactic prostate cancer model**

358 In the prophylactic challenge model, following tumour implantation mice were monitored three
359 times weekly for the formation of palpable tumours (Figure 5-B). Mice receiving no
360 immunisation (control) had all developed palpable tumours 7 days following implantation.
361 However, immunisation with DNA-IM, DNA-MN and NP-IM delayed time until tumour
362 development, with an average time to tumour development of 11.17, 7.57 and 11.17 days
363 respectively. The greatest delays in tumour development were observed in mice immunised
364 with NP-MN, where the average time to tumour formation was 16.2 days. Furthermore, 100
365 days following immunisation with NP-MN 1 mouse remained tumour-free, indicating that
366 immunisation with NP-MN was able to delay, and in some cases prevent, tumour development.

367 Immunisation of mice with NP-MN was also able to significantly lower the tumour burden
368 compared to control (unimmunised mice) (Figure 5-C -, Two-Way ANOVA, $p < 0.05$). For
369 example, at 70 days post-challenge control mice had an average tumour volume of 393.8 mm³,
370 whilst mice immunised with NP-MN, had an average tumour volume of 124.8 mm³.

371 Furthermore, immunisation with NP-MN was able to significantly extend survival of mice
372 compared to control (Figure 5-D, $p < 0.05$, Mantel-Cox Test). Indeed, 2/6 mice immunised via
373 NP-MN survived greater than 100 days post-challenge, (1 other mouse was censored at day 75
374 due to the development of a non-healing ulcer prior to experimental end-point).

375 **3.5 NP-MN Immunisation delays tumour progression and improves survival in a** 376 **therapeutic prostate cancer model**

377 In the therapeutic challenge model immunisation with NP-MN was able to significantly lower
378 the tumour burden compared to control (Figure 6-B, Two-Way ANOVA, $p < 0.05$). For
379 example, at day 63 following tumour challenge, the average tumour volume of mice immunised

380 with NP-MN was 347.7 mm³, whilst control mice, and mice immunised with DNA-MN, DNA-
381 IM and NP-IM had average tumour volumes of 589.9, 544.5, 478.5 and 493.7 mm³ respectively
382 (Figure 6-B). Additionally, immunisation with NP-MN was able to significantly extend the
383 survival of mice by ~10 days compared to control (81.33 vs 71 days respectively) (Figure 6-C,
384 $p < 0.05$, Mantel-Cox Test).

385 **4. Discussion**

386 It has been demonstrated for the first time that MNs may be used to enhance the therapeutic
387 efficacy of DNA vaccines targeting PSCA in a murine model of prostate cancer. In order to be
388 successful in the clinic, DNA vaccines must overcome the barriers to gene delivery to generate
389 sufficient TAA expression *in vivo* to elicit the expansion of an antigen-specific T cell response.
390 This delivery of pTAA is thought to be the main stumbling block for many DNA vaccines due
391 to the vulnerability of pDNA to degradation along with a reduced capacity to transfect cells
392 independently. To this end non-viral vectors which are capable of simultaneously masking the
393 anionic charge of pDNA, complexing pDNA into the NP size range and mediating cellular
394 uptake are being developed.[16] To be efficacious *in vivo*, vectors must retain these
395 characteristics at body temperature, and protect pDNA from degradation via serum
396 nucleases.[17–20] Therefore the first aim of this study was to determine that complexing
397 pPSCA to the RALA delivery peptide resulted in NPs with ideal properties for mediating
398 transfection. The RALA peptide has previously been demonstrated to form nanoscale
399 complexes with pDNA, which are stable at RT, resistant to degradation within the serum, and
400 enhance transfection efficacy *in vitro* and *in vivo*. [13] Here complexation of pPSCA with
401 RALA also resulted in NPs with ideal physical properties for cellular uptake. Furthermore, NPs
402 were stable at body temperature, and were able to protect pPSCA from degradation for up to 6
403 h within serum. Therefore the functionality of RALA/pPSCA NPs was confirmed *in vitro*
404 utilising HEK-293 cells (as a model human cell line for transfection studies), [21] and DC 2.4
405 cells (as DCs are an abundant cell line within the skin and are the key activators of potent
406 cellular immune responses). [22–24] As expected, flow cytometric and PCR analysis confirmed
407 that complexing pDNA to RALA resulted in significantly increased gene expression within
408 both cell lines.

409 In addition to ensuring sufficient cellular uptake following delivery, the cargo delivery site is
410 of key importance to DNA vaccine efficacy. Central to the induction of an adaptive immune
411 response is the priming, stimulation and subsequent expansion of an effector population of

412 antigen-specific T cells. This process is reliant on the presentation of antigenic peptides by
413 APCs on MHC class I and II molecules, to CD8⁺ and CD4⁺ T cells respectively, either
414 following direct transfection or capture of antigenic peptide produced by bystander cells.[3]
415 Given the fundamental role of APCs in DNA vaccination, it is illogical that the majority of
416 DNA vaccines targeting prostate cancer are delivered i.m. or s.c. clinically,[3] where there is a
417 relative paucity of APCs, and this may help to explain the limited efficacy of these therapies to
418 date. A more rational approach is targeting cargo to the skin, which harbours a high density of
419 “professional” APCs including specialised DCs which are adept at T cell priming. [23,25,26]
420 Previously, we reported the development of a novel two-tier NP-MN delivery platform, which
421 was able to induce a greater immune response to the encoded cargo than delivering the same
422 cargo via i.m. injection.[12] By utilising lyophilisation to concentrate NPs prior to
423 incorporation into the dissolvable PVA MN matrix we were able to load 57 µg of pDNA into
424 one device, of which 50 µg was able to be delivered over a 24 h period *in vivo*. Despite
425 undergoing the repeated concentration and drying steps of lyophilisation and MN manufacture,
426 NPs released from the device retained a high (75%) bioactivity compared to fresh NPs, making
427 the device ideal for delivery of high quantities of pDNA intradermally, which has been difficult
428 to achieve previously.[27] The NP-MN device was able to penetrate neonatal porcine skin and
429 withstand the mechanical forces needed for manual application, and perhaps more importantly
430 was able to retain mechanical robustness and cargo functionality over a 28 day period,
431 indicating that the device was ideal for DNA vaccination purposes.[12] Therefore, having
432 established the functionality of RALA/pPSCA NPs, the L-NPs were incorporated into MNs to
433 target gene expression to the epidermal and dermal layers of the skin which are abundant in
434 specialised APCs. Lyophilisation and incorporation into the dissolvable PVA matrix did not
435 compromise the ability of NPs to transfect the human epithelial or murine DC lines, and hence
436 NPs released from dissolvable PVA MNs were expected to be capable of eliciting PSCA
437 expression *in vivo*. Prior to utilising NP-MNs for immunisation studies, it was confirmed that
438 the platform was capable of generating gene expression within the ear pinna of mice, which
439 was selected as the site of immunisation due to the relative low density of hair follicles (which
440 compromise MN penetration) and high abundance of APCs.[28] Application of NP-loaded
441 MNs for 24 h resulted in antigen expression within the treated area, and so this is expected to
442 result in antigen presentation via DCs to naive CD8⁺ and CD4⁺ T cells, generating an anti-
443 mPSCA response.

444 Having established the ability of the delivery system to induce antigen expression *in vivo*, the
445 activation of an antigen-specific T cell response was evaluated. Central to the induction of a
446 potent, anti-tumour response is the generation of tumour-specific cytotoxic T lymphocytes
447 (CTLs) which are capable of directly lysing cancer cells. Cytotoxic T cell responses are also
448 heavily dependent on CD4⁺ T helper cells, and so anti-tumour immunity requires simultaneous
449 expansion of an antigen-specific T_{H1} CD4⁺ T cell population, to stimulate CD8⁺ T cells.[29–
450 31] IFN- γ secretion from the splenocytes of vaccinated animals was assessed to determine the
451 induction of a T_{H1} response. IFN- γ is a proinflammatory cytokine released from activated DCs,
452 CD4⁺ T_{H1} cells and CD8⁺ T cells in response to antigenic stimulation, and is critical for tumour
453 control through the activation of macrophages and neutrophils, which contribute to tumour
454 clearance, and the activation of CTLs.[30,32,33] As such, the greater release of IFN- γ from
455 splenocytes of mice vaccinated with both NP-NP and NP-IM indicated the induction of an
456 antigen-specific T_{H1}-predominant response. Additionally, the cytolytic activity of tumour-
457 specific CTLs from vaccinated animals was demonstrated via measurement of TRAMP C-1
458 tumour cell lysis *in vitro*.

459 Given that the induction of PSCA-specific CD4⁺ T_{H1} and CD8⁺ T cells has been correlated
460 previously with increased survival of tumour bearing mice,[34–36] and both NP-IM and NP-
461 MN elicited antigen-specific T cells responses, it was expected that immunisation of mice with
462 NP-MN or NP-IM injection would confer protection against C57 BL/6 mice against tumour
463 challenge. This would also be in agreement with our previous studies which demonstrated that
464 immunisation with both NP-MN and NP-IM could increase survival of tumour-bearing mice
465 in a cervical cancer model.[12] Indeed in the prophylactic arm of the tumour challenge studies,
466 immunisation with NP-MN resulted in increased time until tumour uptake (with 1/6 mice not
467 developing a tumour) and prolonged survival of tumour-bearing mice. However, surprisingly,
468 immunisation of mice with IM injection was only able to delay time until tumour uptake, and
469 was not sufficient to increase survival. Previously, Johnson *et al* (2007) had reported that as
470 many as 6 i.d. immunisations with a plasmid expressing rat PAP (pTVG-RP) were necessary
471 to generate an anti-PAP response in Lewis rats,[37] therefore a higher number of
472 immunisations may be needed to prevent tumour uptake in all vaccinated animals.
473 Alternatively, the superior efficacy of the NP MN system may be due to the differing release
474 kinetics of the NPs from the MN system which may produce a more persistent immune
475 response. Chen *et al* (2013) demonstrated that sustained release of peptide antigen from a depot
476 of embeddable chitosan needles elicited a stronger antibody response against the model antigen

477 than the same dose delivered by i.m. injection.[38] This response persisted for at least 6 weeks,
478 and was in agreement with other studies establishing that the slower release of antigen could
479 increase the longevity and potency of the immune response. [39,40] Correspondingly, in the
480 therapeutic arm of the tumour challenge studies, immunisation with NP-MN significantly
481 retarded the growth of established tumours, and prolonged survival. These results were
482 unexpected given that immunisation via both MN and IM injection were capable of generating
483 increased numbers of antigen-specific CTLs, which were capable of lysing TRAMP C-1 target
484 cells *ex vivo*. A key reason for this disparity may be the time taken for each of the immunisation
485 routes to prime CD4+ and CD8+ T cells. Indeed the high APC population in the mouse ear
486 pinna, and the drainage of these cells to a common lymph node may facilitate faster T cell
487 priming.[28]

488 An alternative explanation for the lack of efficacy of the NP-IM treatment arm may be that
489 although CTLs are generated following immunisation, they are unable to overcome the
490 mechanisms of immune evasion employed by the established tumour. These mechanisms
491 include the downregulation of MHC complex presentation, immune-editing, immune
492 suppression via regulatory T cells, secretion of immunosuppressive cytokines, and
493 upregulation of co-inhibitory signalling pathways.[41,42] These mechanisms also likely
494 contributed to NP-MN immunisation only modestly retarding the growth of PSCA-expressing
495 tumours *in vivo*, indicating that this vaccination platform is insufficient to induce tumour
496 regression in isolation. This is perhaps because most prostate cancer DNA vaccines are
497 delivered simultaneously with adjuvants to skew the immune response,[3] and hence the
498 platform's efficacy is likely to be improved significantly from combination with immune-
499 modulating therapies in future studies. Several strategies have been successfully employed to
500 counteract the escape mechanisms of the tumour microenvironment, including the use of
501 multivalent strategies,[43] monoclonal antibodies targeting regulatory T cells,[44] adjuvants to
502 skew the immune response,[34] and checkpoint inhibitors.[44] Given the complexity and
503 heterogeneity of the tumour microenvironment a combination of these approaches is likely to
504 be needed simultaneously to DNA vaccine therapy to achieve a synergistic effect yielding
505 clinically meaningful results. One of the key advantages of DNA vaccines is that they carry the
506 potential to encode multiple biological cargos within the same plasmid backbone. Therefore,
507 multiple immunogenic peptides, biological adjuvants, and monoclonal antibodies could be
508 simultaneously generated from one cost-effective and stable platform, giving this device great
509 adaptability for future studies.

510 In conclusion, it has been successfully demonstrated for the first time that cutaneous
511 administration of DNA vaccine via microneedles is capable of eliciting an immune response
512 towards an endogenous prostate cancer TAA. Furthermore, NP-MN immunisation was able to
513 protect against challenge with a syngeneic prostate cancer allograft, in both prophylactic and
514 therapeutic studies. The ability of the NP-MN immunisation to generate a tumour-specific CTL
515 response in the absence of additional adjuvants is encouraging, and the amenability of this
516 platform to combination with other anti-cancer strategies makes it a promising candidate for
517 future studies.

518

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521 **Statement of Significance**

522 This research explains the development and utilisation of our unique microneedle (MN) DNA
523 delivery system, which enables penetration through the *stratum corneum* and deposition within
524 the highly immunogenic skin layers via a dissolvable MN matrix, and facilitates cellular uptake
525 via complexation of pDNA cargo into nanoparticles (NPs) with the RALA delivery peptide.
526 We report for the first time on using the NP-MN platform to immunise mice with encoded
527 Prostate Stem Cell Antigen (mPSCA) for prostate cancer DNA vaccination. Application of the
528 NP-MN system resulted in local mPSCA expression in vivo. Furthermore, immunisation with
529 the NP-MN system induced a tumour-specific cellular immune response, and inhibited the
530 growth of TRAMP C-1 prostate tumours in both prophylactic and therapeutic challenge models
531 in vivo.

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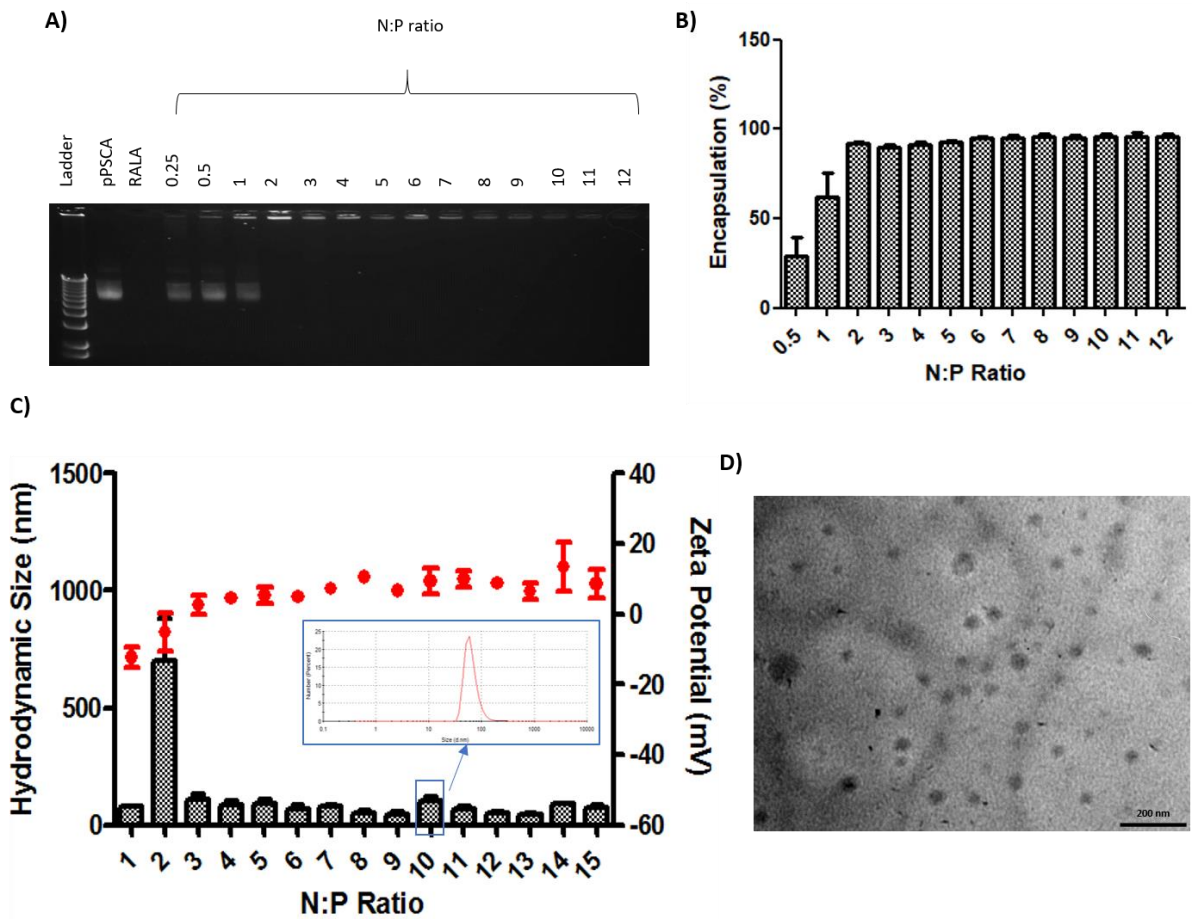
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721 **FIGURES**

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726 **Figure 1: Physicochemical characteristics of RALA/pPSCA Nanoparticles (NPs).** (A) Gel Retardation
727 Assay of NPs over a range of N:P ratios (N:P 0-12), Lane 1: DNA Ladder; Lane 2: pTAA only; Lane 3:
728 RALA only Lanes 4-17: RALA/pPSCA complexes prepared at N:P ratios 0.25-12; (B) Picogreen
729 Encapsulation Assay of NPs over a range of N:P ratios (0-12); (C) Size and Zeta Potential of NPs over a
730 range of N:P ratios (0-15), with an inlaid Size Distribution by Number graph of NPs prepared at N:P 10;
731 (D) TEM image of RALA/pPSCA NP (N:P 10), scale bar is 200 nm.

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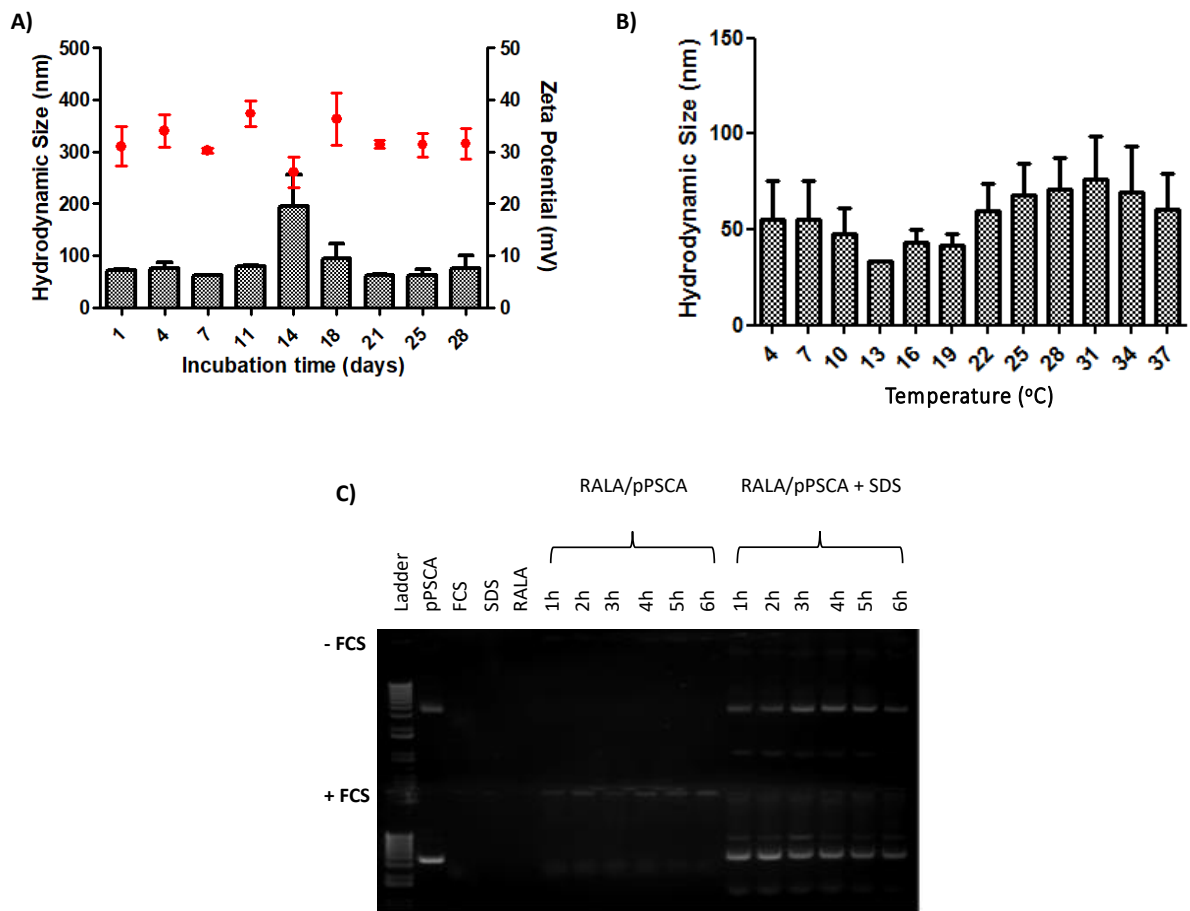
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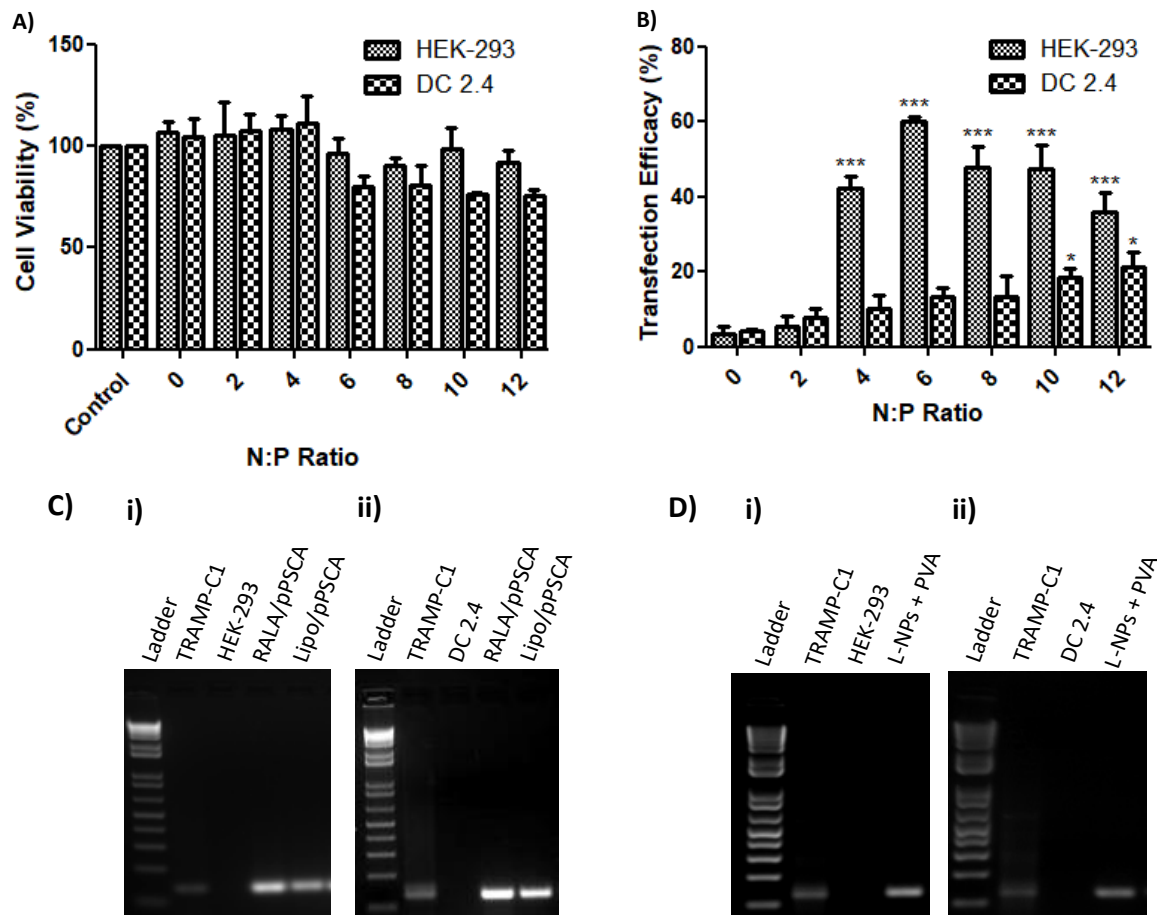
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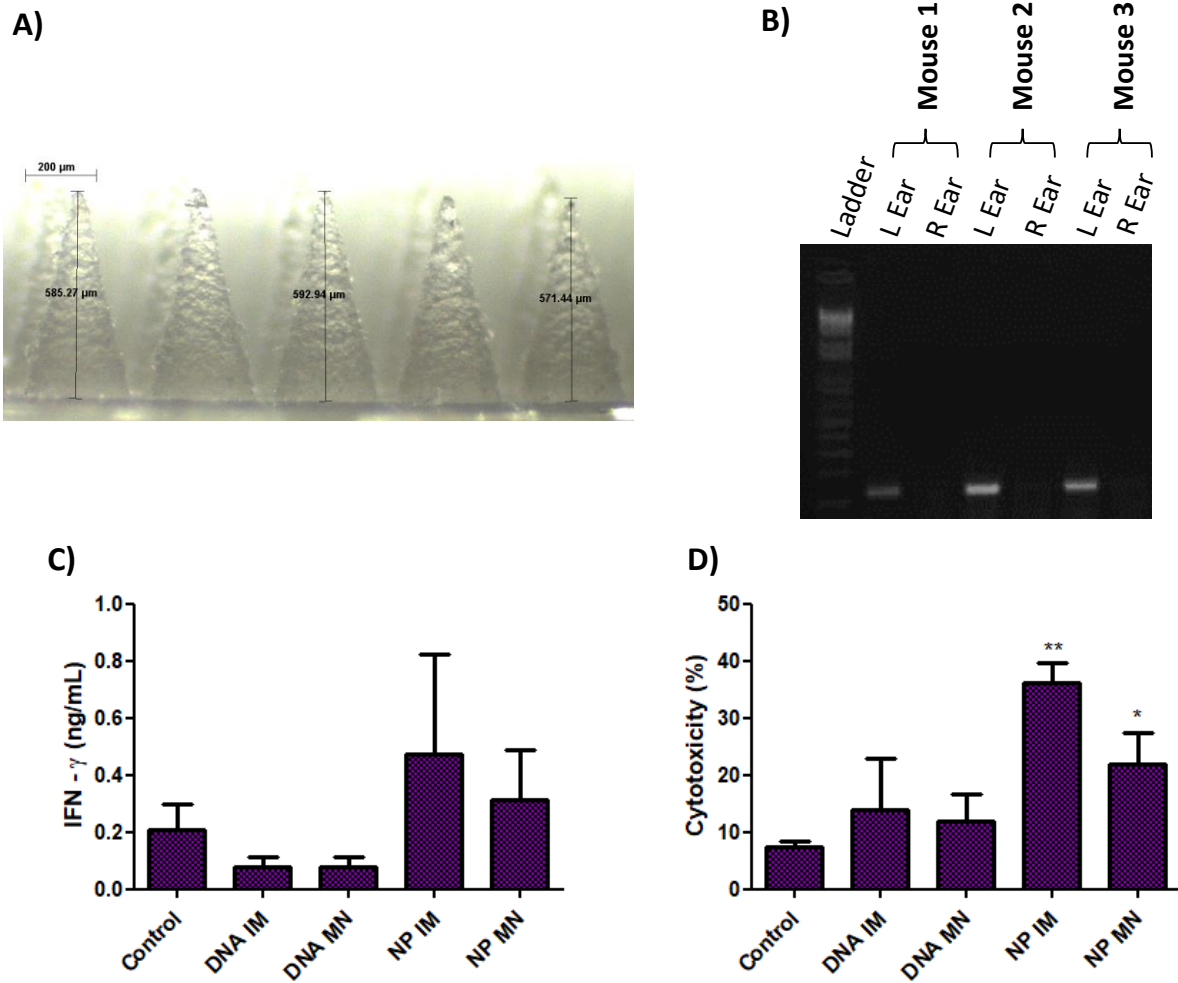
Figure 2: Stability of RALA/pPSCA Nanoparticles (NPs). (A) Hydrodynamic size and Size of RALA/pPSCA NPs (N:P 10) over time. (B) Hydrodynamic Size and PDI of RALA/pPSCA NPs (N:P 10) over range of temperatures (4°C - 37°C); (C) Serum Stability study of RALA/pPSCA NPs. Lane 1: DNA Ladder; Lane 2: pPSCA only; Lane 3: Foetal Calf Serum (FCS); Lane 4: SDS; Lane 5: RALA peptide; Lanes 6-11: RALA/pPSCA (N:P 10) and incubated in water (Row 1) or 10% FCS (Row 2) for 1-6 h at 37°C; Lanes 12-17: RALA/pPSCA (N:P 10) were incubated in water (Row 1) or 10% FCS (Row 2) for 1-6 h at 37°C, and subsequently decomplexed with 10% SDS for 10 min at room temperature prior to electrophoresis.



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760 **Figure 3: In vitro characterisation of RALA/pPSCA Nanoparticles (NPs).** (A) Cytotoxicity analysis and
 761 (B) Transfection Efficacy of RALA/pPSCA NPs over a range of N:P ratios (N:P 0-12) in the HEK-293 and
 762 DC 2.4 cell lines. (C) Confirmation of PSCA mRNA expression within the i) HEK-293 and ii) DC 2.4 cell
 763 lines via reverse transcription (RT)-PCR following transfection of cells with RALA/pPSCA or
 764 Lipofectamine/pPSCA complexes. Lane 1: DNA Ladder; Lane 2: TRAMP-C1; Lane 3: untreated HEK-293
 765 or DC 2.4 cells; Lane 4 and 5: HEK-293 or DC 2.4 cells transfected with RALA/pPSCA or
 766 Lipofectamine/pPSCA respectively. (D) Confirmation of PSCA mRNA expression within the i) HEK-293
 767 and ii) DC 2.4 cell lines via RT-PCR following transfection of cells with lyophilised RALA/pPSCA (N:P 10)
 768 NPs (L-NP) reconstituted and dried in PVA. Lane 1: DNA Ladder; Lane 2: TRAMP-C1; Lane 3: untreated
 769 HEK-293 or DC 2.4 cells; Lane 4: : HEK-293 or DC 2.4 cells transfected with L-NPs reconstituted and
 770 dried within PVA (L-NPs + PVA).

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778 **Figure 4: *In vivo* evaluation of PVA microneedles (MNs) loaded with L-NPs (NP-MN).** (A) Image of 9-
 779 10 kDa PVA MN loaded with L-NPs (NP-MN); (B) Confirmation of PSCA mRNA expression within the
 780 right ears of mice treated with NP-MNs encapsulating via RT-PCR. Lane 1: DNA Ladder; Lanes 2, 4 and
 781 6: cDNA from left (L) ears of mice 48 h following application of NP-MN; Lane 3, 5 and 7: cDNA from
 782 untreated right (R) ears of mice. (C) Tumour-specific IFN- γ release following stimulation of splenocytes
 783 from mice immunised three times at two-weekly intervals with pPSCA (DNA) or RALA/pPSCA (NP) via
 784 microneedle (MN) or intramuscular injection (IM). (D) TRAMP-C1 specific toxicity following exposure
 785 to stimulated splenocytes from mice immunised as above.

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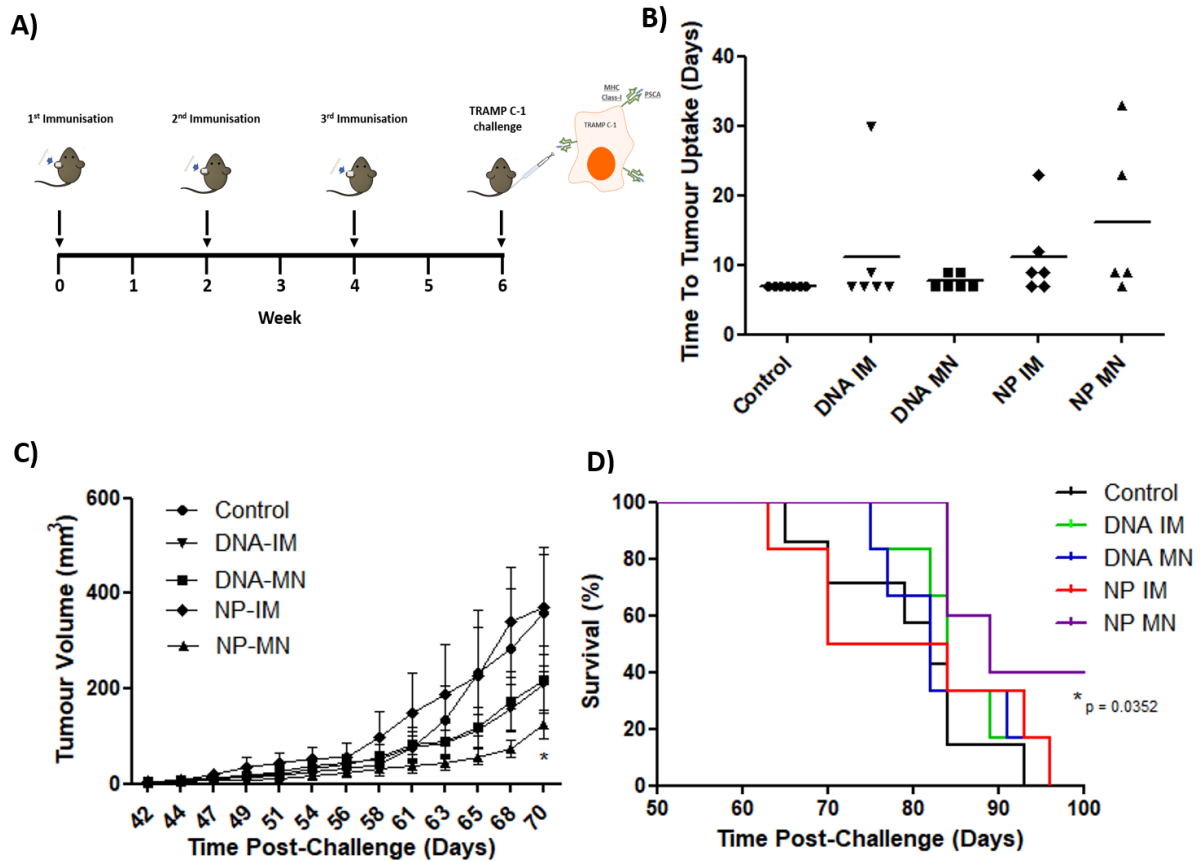
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795 **Figure 5: Prophylactic Efficacy of Immunisation with PVA MNs loaded with L-NPs (NP-MN).** (A)
 796 Schematic of Prophylactic Immunisation Schedule. C57 BL/6 mice were immunised three times at two-
 797 weekly intervals with pPSCA (DNA) or RALA/pPSCA (NP) via microneedle (MN) or intramuscular
 798 injection (IM). Two weeks following the final immunisation mice were challenged s.c. with 5×10^6
 799 TRAMP C-1 cells in PBS/Matrigel; (B) Scatter plot of time to tumour formation in immunised mice; (C)
 800 Average tumour volume of control and immunised mice; (D) Kaplan-Meier Survival Curve. N = 6/7.

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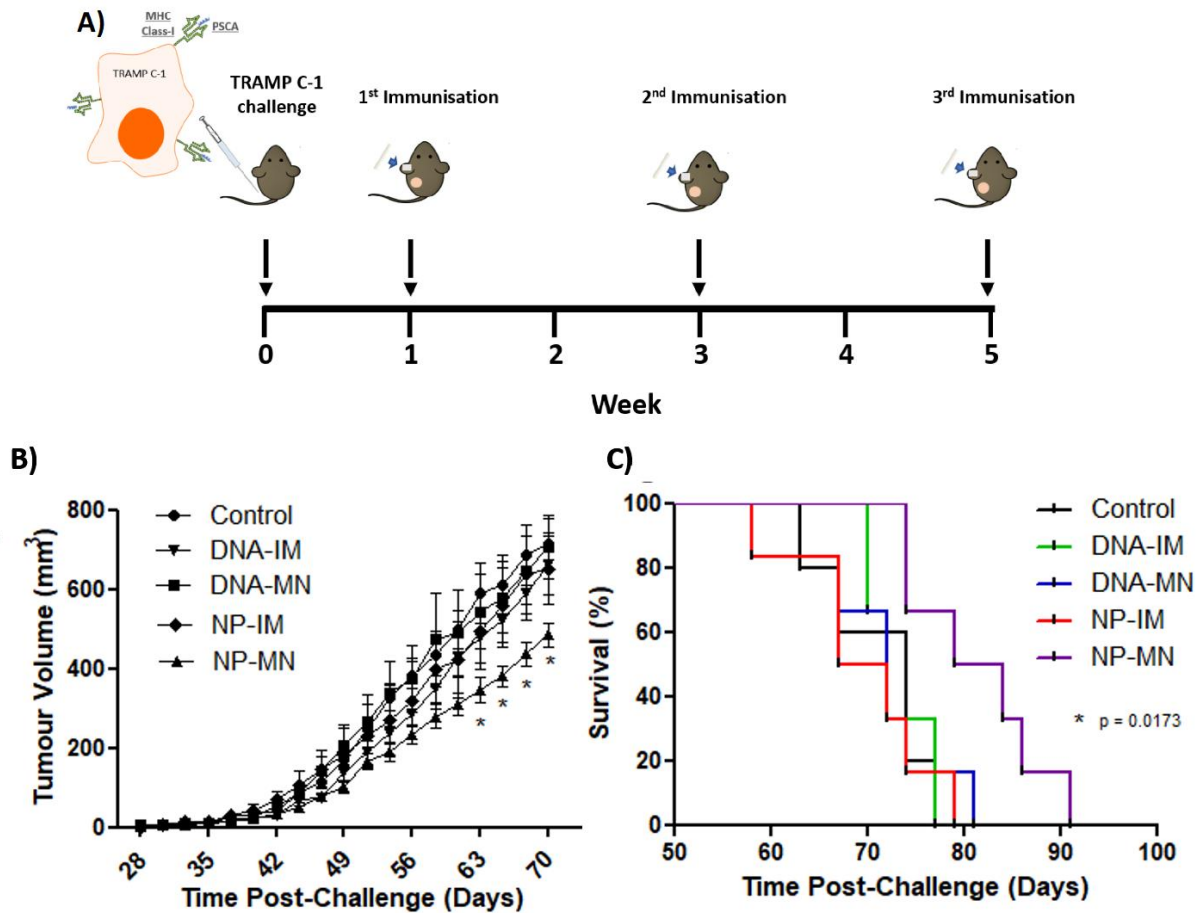
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813 **Figure 6: Therapeutic Efficacy of Immunisation with PVA MNs loaded with L-NPs (NP-MN).** (A)
 814 Schematic of Therapeutic Immunisation Schedule. C57 BL/6 mice were challenged s.c. on the flank
 815 with 5×10^6 TRAMP C-1 cells in PBS/matrigel. One week following tumour challenge mice were
 816 immunised three times at two-weekly intervals with DNA-IM, DNA-MN, NP-IM or NP-MN. (B) Average
 817 tumour volume of control and immunised mice; (C) Kaplan-Meier Survival Curve. N = 5/6.

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