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Title: Development of an Opsonophagocytic Killing Assay for Group A Streptococcus

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Corresponding Author: Dr. Scott Jones, Ph.D.

Corresponding Author's Institution: University College London

First Author: Scott Jones, Ph.D.

Order of Authors: Scott Jones, Ph.D.; Nicole J Moreland; Marta Zancolli; Jeremy Raynes; Jacelyn M S Loh; Pierre R Smeesters; Shiranee Sriskandan; Jonathan R Carapetis; John D Fraser; David Goldblatt

Abstract: Group A Streptococcus (GAS) or Streptococcus pyogenes is responsible for an estimated 500,000 deaths worldwide each year. Protection against GAS infection is thought to be mediated by phagocytosis, enhanced by bacteria-specific antibody. There are no licenced GAS vaccines, despite many promising candidates in preclinical and early stage clinical development, the most advanced of which are based on the GAS M-protein. Vaccine progress has been hindered, in part, by the lack of a standardised functional assay suitable for vaccine evaluation. Current assays, developed over 50 years ago, rely on non-immune human whole blood as a source of neutrophils and complement. Variations in complement and neutrophil activity between donors result in variable data that is difficult to interpret. We have developed an opsonophagocytic killing assay (OPKA) for GAS that utilises dimethylformamide (DMF)-differentiated human promyelocytic leukemia cells (HL-60) as a source of neutrophils and baby rabbit complement, thus removing the major sources of variation in current assays. We have standardised the OPKA for several clinically relevant GAS strain types (emm1, emm6 and emm12) and have shown antibody-specific killing for each emm-type using M-protein specific rabbit antisera. Specificity was demonstrated by pre-incubation of the antisera with homologous M-protein antigens that blocked antibody-specific killing. Additional qualifications of the GAS OPKA, including the assessment of the accuracy, precision, linearity and the lower limit of quantification, were also performed. This GAS OPKA assay has the potential to provide a robust and reproducible platform to accelerate GAS vaccine development.

Highlights (for review)

1. Opsonophagocytic killing assay (OPKA) developed for a number of clinically relevant Group A *Streptococcus* (GAS).
2. Homologous rabbit full-length M-protein antisera and human IVIg shown to effectively kill GAS strains expressing *emm*-types 1, 6 and 12.
3. Killing of bacteria is effectively inhibited by homologous antigen but not heterologous antigens proving assay specificity.
4. GAS OPKA shown to be highly precise with intra-assay and inter-assay coefficients of variation less than 30%.

1 Group A *Streptococcus* (GAS) or *Streptococcus pyogenes* is responsible for an estimated
2 500,000 deaths worldwide each year. Protection against GAS infection is thought to be mediated by
3 phagocytosis, enhanced by bacteria-specific antibody. There are no licenced GAS vaccines, despite
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12 current assays. We have standardised the OPKA for several clinically relevant GAS strain types
13 (*emm1*, *emm6* and *emm12*) and have shown antibody-specific killing for each *emm*-type using M-
14 protein specific rabbit antisera. Specificity was demonstrated by pre-incubation of the antisera with
15 homologous M-protein antigens that blocked antibody-specific killing. Additional qualifications of
16 the GAS OPKA, including the assessment of the accuracy, precision, linearity and the lower limit of
17 quantification, were also performed. This GAS OPKA assay has the potential to provide a robust and
18 reproducible platform to accelerate GAS vaccine development.

DEVELOPMENT OF AN OPSONOPHAGOCYtic KILLING ASSAY FOR GROUP A STREPTOCOCCUS

Scott Jones^{1a}, Nicole J Moreland², Marta Zancoli^{1*}, Jeremy Raynes², Jacelyn MS Loh², Pierre R Smeesters^{3,4}, Shiranee Sriskandan⁵, Jonathan R Carapetis⁶, John D Fraser², David Goldblatt¹

^a Corresponding author: scott.jones@ucl.ac.uk

¹Immunobiology, UCL Great Ormond Street Institute of Child Health, 30 Guilford Street, London WC1N 1EH, United Kingdom.

²Department of Molecular Medicine & Pathology, School of Medical Sciences, The University of Auckland, Private Bag 92019, Auckland, New Zealand.

³Molecular bacteriology laboratory, Université Libre de Bruxelles and Academic Children Hospital, Brussels, Belgium

⁴Murdoch Children's Research Institute and University of Melbourne, Melbourne, Australia

⁵Faculty of Medicine, Imperial College London, Commonwealth Building, Hammersmith Hospital, Du Cane Road, London W12 0NN, United Kingdom.

⁶Telethon Kids Institute, University of Western Australia and Perth Children's Hospital, Perth, Australia.

* Current address: Theradex (Europe) Ltd, 2nd Floor The Pinnacle, Station Way, Crawley, West Sussex, RH10 1JH, United Kingdom.

1 1. INTRODUCTION

2 Group A *Streptococcus* (GAS) or *Streptococcus pyogenes* is a species of Gram-positive
3 bacteria responsible for a significant worldwide burden of disease [1-4]. Clinical syndromes resulting
4 from GAS infection include superficial infections, such as pyoderma and pharyngitis, severe invasive
5 GAS disease (iGAS), acute rheumatic fever (ARF) and associated rheumatic heart disease (RHD), and
6 post-streptococcal glomerulonephritis. Conservative estimates suggest that more than 33 million
7 people are afflicted with severe GAS disease that results in roughly 500,000 deaths each year, with
8 RHD responsible for two thirds of this mortality [5].

9 Despite the substantial morbidity and mortality GAS infection poses, there are currently no
10 licenced vaccines against GAS [6, 7]. Epidemiological data reveals that the majority of infections
11 occur within the ages of 1-15, suggesting that immunity to GAS develops with age [8, 9]. First
12 identified by Rebecca Lancefield, the M-protein is a major GAS virulence factor that comprises a
13 hypervariable N-terminus, which forms the basis for GAS strain typing (known as *emm*-typing), and a
14 conserved C-terminus [10, 11]. The M-protein has multiple functions, including an ability to subvert
15 phagocytosis by binding multiple host proteins [11, 12]. Antibodies specific to the M protein have
16 long been known produce a protective immune response and as such the M protein is being
17 extensively investigated as a vaccine antigen [10, 13]. M-protein based vaccines are generally
18 composed of either the hypervariable N-terminal region of multiple *emm*-types or the conserved C-
19 terminal region and have been shown in early stage clinical trials to be safe and well tolerated [10,
20 14-18]. Non-M-protein based vaccines are also in development, though none have yet been tested in
21 human trials. These tend to comprise a combination of conserved GAS antigens that have been
22 identified through reverse vaccinology or studies on GAS virulence [19-21]. Such antigens include
23 C5a peptidase, serine esterase, GAS carbohydrate, fibronectin-binding protein, serum opacity factor,
24 SpyCep and T-antigens [6, 7, 14, 22-25].

25 The progress of GAS vaccine development has been hindered, in part, by the lack of a
26 reliable and fully standardised functional assay that assesses immunity to GAS. The bactericidal assay
27 used most widely to assess the immunogenicity of GAS vaccine antigens is known as the Lancefield
28 assay [13, 26-29]. The Lancefield assay measures the growth and survival of GAS in fresh human
29 blood from an immune individual (direct) or non-immune whole blood supplemented with serum
30 from an immune individual or animal immunised with an antigen of interest (indirect) [28, 29].
31 Natural variations in complement and neutrophil activity that exist between individuals often result
32 in variable data that is difficult to interpret, and does not allow for comparison of results between
33 laboratories [21, 30]. In addition, widespread natural exposure to GAS renders identifying a non-
34 immune donor (i.e. an individual with minimal reactivity to the bacterium) laborious [24, 29].

35 The need for improved assays to measure GAS vaccine efficiency has been highlighted as a
36 priority by the international vaccine community and was recently recognized by the Coalition to
37 Accelerate New Vaccines Against Streptococcus (CANVAS) [6, 14, 18, 24]. CANVAS is an Australian
38 and New Zealand government-sponsored GAS vaccine development programme that aims to
39 overcome key developmental hurdles for GAS vaccines. The coalition was formed in response to the
40 unacceptably high rates of ARF and RHD in Indigenous populations in Australia and New Zealand [31,
41 32]. To overcome the issues of variability associated with the traditional Lancefield methods, we
42 have developed a GAS opsonophagocytic killing assay (OPKA) that uses an exogenous source of both
43 complement and phagocytic cells. This assay is adapted from a validated pneumococcal OPKA which
44 makes use of baby rabbit complement (BRC) as the exogenous source of complement and
45 dimethylformamide (DMF) -differentiated human promyelocytic leukemia cells (HL-60) as the
46 exogenous source of phagocytic cells [33-36]. Using high titre anti-M protein rabbit sera and human
47 intravenous immunoglobulin, we have developed and optimised an OPKA for a number of clinically
48 relevant GAS strains.

49 2. MATERIALS AND METHODS

50 2.1. Bacterial Strains

51 The following GAS strains were used in this study: *emm1* (strains 43, 02-12 and GAS05134),
52 *emm12* (strains 611020, 611025 and GAS09437) and *emm6* (strain GASOPA6_02). These *emm*-types
53 were selected because they belong to distinct *emm*-protein clusters in the recently described *emm*
54 cluster typing system [37]. Additional strain details are listed in **Table 1**. Working stocks of bacterial
55 strains were generated by streaking a fleck of frozen master stocks onto a horse blood agar plate
56 (Fisher Scientific, Leicestershire, UK), which was incubated overnight at 37°C, 5% CO₂. Todd-Hewitt
57 Broth (THB; Sigma-Aldrich Company Ltd, Dorset, UK) was inoculated with a single colony and grown
58 at 37°C, 5% CO₂ to an OD₆₀₀ (optical density at 600nm) of 0.6-0.7; considered to be the late-log phase
59 of growth. Once the required OD₆₀₀ was reached, cultures were mixed 1:1 with STGG medium (3%
60 w/v tryptone soya broth (Thermo Fisher Scientific Oxoid Ltd, Basingstoke, UK), 0.5% w/v D-glucose
61 (Sigma-Aldrich Company Ltd), 10% v/v glycerol (Sigma-Aldrich Company Ltd) in sterile, pyrogen-free
62 water (Baxter Healthcare UK, Newbury, UK) and stored in 0.5ml aliquots at -80°C until required.

63 2.2. Antisera

64 Two sets of sera were used to optimise the GAS OPKA: human intravenous immunoglobulin
65 (IVIg; Euglobulin®; Baxter, Berkshire, UK) and rabbit anti-full-length M-protein sera. To generate M-
66 proteins for rabbit vaccinations *emm1* (SF370) and *emm6* (MGAS10394) were amplified from
67 genomic DNA. The primer pairs ctagGGATCCaacggtgatgtaatcctagg and
68 ctagGAATTCctgtctcttagtttccttcattgg were used for *emm1*, and cgcGGATCCagagtgtttcctagggggacg and
69 ctagGAATTCctgtctcttagtttccttcattgg for *emm6*. The genes were cloned into pET32a3c sequences and
70 the sequences were confirmed by Sanger sequencing. For *emm12* (HKU16) the gene was synthesised
71 (Life technologies) and inserted into pET151/D-TOPO. Each of the three proteins were expressed
72 with the signal sequence and transmembrane regions removed (M1, aaN42-Q449; M6, aaR43-T358;
73 M12, aaD42-Q540) in BL21(DE3) pLysS *E.coli* and purified by immobilised-metal affinity

74 chromatography (IMAC) as previously described [37, 38]. The His₆ tag was cleaved from the
75 recombinant M proteins with 5 µg/ml 3C-His₆ protease (M1 and M6) or a 1:100 ratio of rTEV-His₆
76 protease to M protein (M12) for 16 hours at 4°C. The M-proteins were separated from the cleaved
77 His₆ tag and the rTEV-His₆ protease by IMAC and further purified by gel filtration. Female New
78 Zealand White rabbits were immunised subcutaneously with 200µg of the purified, recombinant M1,
79 M6 or M12 proteins in Incomplete Freund's adjuvant (Sigma-Aldrich) at 0, 2 and 4 weeks prior to
80 exsanguination. The reactivity of immune sera to the recombinant M proteins was confirmed by
81 ELISA (endpoint titre >100,000).

82 2.3. HL-60 Cells

83 Master stocks of HL-60 cells were prepared from frozen commercial stock (ATCC Standards
84 UK, Middlesex, UK). Working stocks of HL-60 cells were differentiated into neutrophil-like cells by
85 culture in 0.8% DMF (Sigma-Aldrich Company Ltd) in M2 medium (10% v/v FetalClone1 (Hyclone, GE
86 Healthcare Life Sciences, Buckinghamshire, UK), 1% L-Glutamine (Invitrogen, Life Technologies Ltd,
87 Paisley, UK), in RPMI 1640 (Invitrogen)) at a concentration of 4x10⁵ cells per ml. Cells were
88 differentiated at 37°C, 5% CO₂ for five to six days. On the day of the assay, cells were washed with
89 HBSS without Ca/Mg (Invitrogen), followed by HBSS with Ca/Mg (Invitrogen) and re-suspended at
90 1x10⁷ cells per ml in fresh opsonisation buffer (10% v/v defined FBS (Hyclone), 0.1% w/v gelatin
91 (Sigma-Aldrich Company Ltd) in HBSS with Ca/Mg). The phenotype of differentiated HL-60 cells was
92 assessed by flow cytometry using mouse anti-human CD35 FITC conjugated antibody (Serotec, Bio-
93 Rad, Kidlington, UK) and/or mouse anti-human CD71 PE-conjugated antibody (Serotec).
94 Differentiated cells were accepted for use in the assay if the up-regulation of CD35 was ≥55% of the
95 cell population and CD71 expressing was down-regulated by ≥15% when compared to the working
96 stock preparation.

97 2.4. Complement

98 In order to reduce levels of non-specific killing of bacteria in the assay, new batches of baby
99 rabbit complement were pre-absorbed as previously described with minor changes [39]. Briefly,
100 200ml of three to four-week baby rabbit complement (Pel-Freez, AR, USA) was defrosted in ice-cold
101 water with constant agitation. Whilst still in ice-cold water, the defrosted complement was
102 inoculated with 10 colonies per 50ml of an overnight horse-blood agar culture of *Staphylococcus*
103 *aureus* (strain 6538) and incubated on a roller for 30 minutes at 4°C. The complement was then
104 vacuum filtered with a 0.22µM, aliquoted on ice and stored at -80°C until required.

105 2.5. Opsonophagocytic Killing Assay

106 Working stocks of bacteria were defrosted and washed and diluted in OPS buffer to 120,000
107 CFU per ml. Bacteria were added to samples serially diluted (in duplicate) with OPS buffer in a
108 round-bottomed 96-well plate and incubated for 30 minutes at room temperature on a mini-orbital
109 shaker at 700rpm. Diluted active or heat-inactivated BRC was added to each well followed by DMF-
110 differentiated HL-60 cells and the plate was incubated for an additional 90 minutes at 37°C, 5% CO₂
111 on a mini-orbital shaker at 700rpm. The ratio between HL-60 cells and bacteria in each assay was
112 approximately between 500-2000 HL-60 cells to one bacterial cell. Two complement only controls
113 (referred to as control A and control B hereafter) were included on each plate to calculate the level
114 of non-specific killing in an assay. Control A contains bacteria, differentiated HL-60 cells and heat-
115 inactivated complement only and control B contains bacteria, differentiated HL-60 cells and activate
116 complement only.

117 After the second incubation, plates were placed on ice for at least thirty minutes to stop the
118 reaction. After mixing well, 10µl of each well was 'spotted' onto Todd-Hewitt-yeast agar plates (THB
119 supplement with 0.5% w/v yeast extract (Sigma-Aldrich Company Ltd) and 1.5% w/v bacteriological
120 agar (Sigma-Aldrich Company Ltd), which was tilted to allow the spots to spread across one quarter
121 of the plate and left to dry at room temperature. Once dry, 20ml autoclaved overlay agar (THB

122 supplement with w/v yeast extract, 0.75% w/v bacteriological agar and 0.0025% 2, 3, 5-
123 tetraphenyltetrazolium chloride (Sigma-Aldrich Company Ltd) was poured over each plate and
124 allowed to set. Plates were then incubated overnight at 37°C, 5% CO₂.

125 The number of CFU per spot was enumerated using an automated colony counter
126 (ProtoCOL3; Synbiosis, Cambridge, UK). The percentage killing at each dilution of a sample was
127 calculated as: $(\text{CFU}[\text{control B}] - \text{CFU}[\text{antisera}]) / \text{CFU}[\text{control B}] * 100$ and the dilution of the sample
128 resulting in 50% killing was calculated as the opsonic index (OI) or opsonic titre. The percentage of
129 non-specific killing in each assay was calculated as: $1 - (\text{CFU}[\text{control B}] / \text{CFU}[\text{control A}]) * 100$. Assays
130 were accepted if levels of non-specific killing were $\leq 35\%$ and the CFU of both complement controls
131 were between 50-200. Furthermore, serum samples were only reported as positive if the maximum
132 level of killing achieved was $\geq 70\%$. These acceptance criteria are comparable to those used for the
133 validated pneumococcal OPKA and are included to ensure the reproducibility and quality of assay
134 results [36].

135 2.6 Determination of Assay Specificity

136 Rabbit anti-full-length M-protein sera were pre-incubated with a titration of full-length M-
137 protein or OPS buffer only (0% inhibition control) for thirty minutes at room temperature. Washed
138 and diluted bacteria were then added to the pre-incubated samples and the assay was continued as
139 previously described. The percentage inhibition of killing at each concentration of antigen was
140 calculated as: $(\text{CFU}[\text{antisera:antigen}] - \text{CFU}[0\% \text{ inhibition control}]) / (\text{CFU}[\text{control B}] - \text{CFU}[0\% \text{ inhibition}$
141 $\text{control}]) * 100$.

142 **3. RESULTS**

143 The GAS OPKA was adapted from a validated pneumococcal OPKA with several modifications
144 with respect to incubation times and the preparation and amount of exogenous BRC added [33, 34].

145 3.1 GAS OPKA Optimisation

146 Prior to performing the GAS OPKA, the dilution of bacterial working stocks resulting in spots
147 containing between 150 and 200 CFU was determined for each strain. The concentration of BRC is
148 critical to maintain non-specific killing below 35%, the acceptance criteria for the assay. To optimise
149 the assay with respect to non-specific killing, bacteria were incubated with a serial dilution of pre-
150 absorbed BRC in the presence of differentiated HL-60 cells. Representative complement titration
151 data of CFU at each concentration of heat-inactivated or active BRC and the level of non-specific
152 killing for GAS strain M1(43) is shown in **Figure 1**. In this example, the concentration of complement
153 resulting in <35% non-specific killing was 2.1%. Across all seven strains used in this study the optimal
154 concentration of baby rabbit complement was 2.1% for GAS strains M1(43), M1(02-12),
155 M1(GAS05134), M12(611020), M12(GAS09437) and 3.1% for GAS strains M12(611025) and
156 M6(GASOPA6_02) (Table 2).

157 3.2 GAS OPKA

158 Once assay conditions were optimised for each GAS strain, the ability of human IVIg and
159 rabbit anti-full-length M-protein sera to kill GAS was assessed. Representative data for two GAS
160 strains M1(43) and M12(611020) tested against human IVIg, rabbit antisera from the homologous
161 M-type (M1 or M12), matched pre-immunisation sera and rabbit anti-sera for the heterologous M-
162 types are shown in **Figure 2**. There was a marked reduction in CFU of M1(43) in the presence of the
163 M1 anti-sera and IVIg, but not for the pre-immune sera or the M6 and M12 antisera. Similarly, there
164 was a marked reduction in CFU for the M12(611020) strain in the presence of M12 anti-sera and
165 IVIg, but not for pre-immune sera, or the M1 and M6 antisera. This marked M type-specific killing is
166 further demonstrated by the OIs determined for each sera against the seven GAS strain as
167 summarized in **Table 2**. Killing of all strains by homologous full-length M-protein antisera and human
168 IVIg had an average OI of 4089 and 6497, respectively. In contrast no killing by heterologous antisera
169 or pre-immunisation sera was observed for any of the strains tested.

170 3.3 GAS OPKA Specificity

171 The specificity of the GAS OPKA was determined by assessing the ability of homologous and
172 heterologous purified M-protein to inhibit the killing of bacteria by rabbit anti-full-length M-protein
173 sera. Representative data of percentage inhibition of killing by rabbit anti-full-length M1 sera, pre-
174 incubated with full-length M1, M6 or M12, run against the GAS strain M1(43) is shown in **Figure 3a**.
175 In this example, maximum inhibition of killing achieved by the homologous antigen (M1) was 86.5%
176 and the maximum inhibition of killing by the heterologous antigens (M6 and M12) was 9.7% and
177 10.7%, respectively. By fitting a sigmoidal standard curve to log[x] transformed data, the
178 concentration of homologous M1 antigen resulting in 50% inhibition of killing of strain M1(43) in this
179 example was interpolated to be 72.4ng/ml. Across all the seven strains investigated the average
180 concentration of homologous antigen resulting in 50% inhibition of killing was 131ng/ml \pm 76.25. The
181 maximum inhibition of killing by homologous and heterologous antigens for each GAS strain is
182 detailed in **Table 3**.

183 3.4 GAS OPKA Precision

184 GAS OPKA precision was assessed by running rabbit anti-full-length M-protein antisera three
185 times either on the same plate with one control (intra-assay coefficient of variation) or on three
186 different plates with individual controls (inter-assay coefficient of variation) with each GAS strain.
187 Representative data from each of the three runs used to calculate the intra-assay coefficient of
188 variation and inter-assay coefficient of variation for GAS strain M1(43) are shown in **Figure 3a** and
189 **Figure 3b**, respectively. The intra-assay and inter-assay coefficients of variation for each optimised
190 strain are detailed in **Table 3**. The intra-assay coefficient of variation of the GAS OPKA was assessed
191 to be 15.4% and the inter-assay coefficient of variation of the GAS OPKA was assessed to be 17.9%.

192 3.5 Relative Accuracy

193 The relative accuracy of the GAS OPKA was assessed by running rabbit full-length M-protein
194 antisera through the GAS OPKA at four different starting dilutions. The percent agreement (whether

195 OIs are within a factor of three of each other) and coefficient of variation between the OIs for each
196 of these four samples was calculated for each strain. Representative data for the calculation of the
197 relative accuracy of GAS strain M1(43), is shown in **Figure 3d**. In this example, rabbit anti-full-length
198 M1 was pre-diluted to a factor of 4, 16, 32 and 64 and had OIs between 2060-2967 against M1(43)
199 resulting in a coefficient of variation of 15.21%. The relative accuracy of each optimised GAS strain is
200 detailed in **Table 3**. The average coefficient of variation across all seven GAS strain studied was
201 calculated as 15.59% and OI agreement was at 100% for all strains.

202 3.6 Linearity

203 The linearity of the GAS OPKA was assessed by running non-immune rabbit serum, spiked
204 with four different concentrations of full-length M-protein antisera, through the GAS OPKA. The
205 correlation (R-squared) between initial sample dilution (log transformed) and resultant OI (log
206 transformed) of the four samples was calculated for each strain. Representative data for the
207 calculation of OPKA linearity for GAS strain M1(43) is shown in **Figure 3e**. In this example, rabbit anti-
208 full-length M1 was pre-diluted to a factor of 4, 16, 32 and 64 and had OIs of 2700, 740, 604 and 128
209 against M1(43), respectively. The correlation (R-squared) was calculated as 0.993 with a slope of -
210 1.096 ± 0.065 . The linearity for each optimised GAS strain is detailed in **Table 3**. Across all seven
211 strains, the average correlation (R-squared) between initial sample dilution (log transformed) and
212 resultant OI (log transformed) was 0.987 with an average gradient (slope) of -1.07.

213 3.7 Lower Limit of Quantification

214 The lower limit of quantification (LLOQ) of the GAS OPKA was assessed by spiking low
215 concentrations of full-length M-protein-rabbit serum into heat-inactivated non-immune rabbit
216 serum for each corresponding GAS strain to produce a sample with an opsonic index of 4-12. Each
217 sample was generated individually four times and run through the assay in duplicate. Representative
218 data for the calculation of the LLOQ of GAS strain M1(43), is shown in **Figure 3f**. In this example, the

219 average OI of these spiked samples was calculated as 8.75 ± 0.5 . The LLOQ for each of the seven GAS
220 strains is detailed in **Table 3**. The average LLOQ across the seven strains was calculated as 7.9.

221 4. DISCUSSION

222 We have optimised an OPKA for a number of clinically relevant GAS strain types (*emm1*,
223 *emm6* and *emm12*). Additionally, we have taken the first steps towards qualifying and standardising
224 the GAS OPKA; assay specificity, precision, relative accuracy, linearity and lower limit of
225 quantification [40]. Both IVIg and anti-M protein specific sera generated in rabbits were able to kill
226 each of the GAS strains used in study. The specificity of the GAS OPKA was established by measuring
227 the ability of homologous or heterologous full-length M-protein to block the ability to kill GAS. For
228 each of the strains tested, homologous antigen was able to block >70% of killing whereas
229 heterologous antigen was only able to block 10% thus proving assay specificity. The strains used in
230 this study strains belong to three distinct *emm*-cluster types: this cluster system classifies M-proteins
231 based on their functional and structural features and the lack of heterologous killing observed in our
232 study is in line with the distinct cluster phylogeny of *emm1* (cluster A-C3), *emm6* (single protein
233 cluster M6) and *emm12* (cluster A-C4) [37]. The ability of certain M-proteins, including M1 used in
234 this study, to non-specifically bind human immunoglobulin via the Fc region is well described [37, 41,
235 42]. At the highest concentration tested (1.25 μ g/ml), soluble M1 protein did not significantly inhibit
236 killing by either the rabbit anti-M6 or rabbit anti-M12 serum. This suggests that the affinity of the
237 non-specific interaction between soluble M1-IgG Fc is lower than the affinity of the specific
238 interaction between antisera and surface bound M-protein such that opsonophagocytosis occurs.

239 While *emm1* is the most common GAS *emm* type globally, each of the *emm*-types used in
240 this study is typically considered to be associated with disease in the developed world [43]. As such,
241 cumulatively *emm1*, *emm6* and *emm12* contribute to approximately 34% of GAS related disease in
242 Established Market Countries but only 10% and 3% of GAS-related disease in Africa and Pacific Island
243 nations/Indigenous Australians, respectively [43]. Furthermore, there is a large degree of *emm*-type

244 diversity associated with ARF in Maori and Pacific children in New Zealand with *emm1*, *emm6* and
245 *emm12* rarely observed [44]. Optimising GAS OPKA conditions for additional strains that are clinically
246 relevant to low and middle-income countries, and Indigenous populations at high risk of GAS disease
247 in New Zealand and Australia, will form a significant part of future work.

248 The GAS strains in this study were cultured to late exponential phase (an OD₆₀₀ of 0.6-0.7)
249 following the current multiplexed opsonisation assay for *Streptococcus pneumoniae* [45]. Licenced
250 pneumococcal vaccines are composed of the polysaccharides that make up the bacterial capsule and
251 culturing bacteria to an OD₆₀₀ of 0.7 ensures sufficient capsule growth required for the assessment
252 of pneumococcal polysaccharides antibody function [46]. As with *Streptococcus pneumoniae*, GAS
253 capsule expression peaks during this exponential growth-phase [47]. For non-M protein GAS
254 antigens that may be obscured by excessive capsule expression, culturing of GAS strains to late
255 exponential phase may prove ineffective and optimisation of assays with early log-phase GAS maybe
256 be required.

257 The inter-assay and intra-assay coefficients of variation of the GAS OPKA were assessed and
258 found to be <30% for all strains which is acceptable for a dynamic biological assay such as the assay
259 described here. The average LLOQ was assessed to be 7.9, the relative accuracy coefficient of
260 variation was 15.6% and the assay linearity correlation was 0.987; all well within the acceptable
261 levels of variation for an assay of this complexity [39, 48, 49]. Full GAS OPKA standardisation will
262 require additional qualification as recommended by the ICH Harmonised Tripartite Guidelines for the
263 validation of analytical procedures [40]. These additional qualifications will need to include an
264 assessment of assay robustness and reproducibility between laboratories.

265 Correlates of protection do not currently exist for immunity to a GAS infection [9]. The
266 standardised, reproducible and specific opsonophagocytic killing assay described here should enable
267 a correlate to be developed when used in conjunction with convalescent GAS sera from individuals
268 who have recovered from a GAS infection or vaccine trials with a clinical endpoint. While our study

269 has demonstrated that the assay should enable correlates to be determined for M-protein based
270 vaccines, it also has the potential to become the gold-standard assay for vaccines comprised of
271 conserved, cell wall-associated GAS antigens. In conclusion, this standardised GAS OPKA should
272 provide a reliable method to assess GAS immunity and support the licensure of GAS vaccine
273 candidates.

AUTHORS' CONTRIBUTION

DG, NJM, JRC and JDF conceived of the presented idea and consulted throughout. SJ wrote the manuscript in consultation with NJM, SS, JRC, JDF and DG. SJ, MZ, NJM, JR and MSJL performed the experiments. GAS strains were provided by the CANVAS strain selection group (including PRS and NJM) and SS. SS also provided advice and expertise on GAS biology and human IVIg. DG supervised the findings of this work.

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CONFLICTS OF INTEREST

Authors declare no conflicts of interest.

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Table 1 GAS Strains

Summary table of the GAS strains used in this study with additional details of M-clusters, clinical syndrome association, country of origin and year of isolation.

M- type	M-cluster	Strain ID	Clinical syndrome	Country of origin	Year of isolation
1	A-C3	43	Pharyngitis	UK	2009
		02-12	Pharyngitis	Brazil	2012
		GAS05134	Acute rheumatic fever	New Zealand	2005
12	A-C4	611020	Pharyngitis	Australia	2011
		611025	Pharyngitis	Australia	2011
		GAS09437	Acute rheumatic fever	New Zealand	2009
6	Single protein M-cluster	GASOPA6_02	Pharyngitis	UK	2009

Table 2 Opsonophagocytic Killing Assay Summary

Summary table of GAS OPKA assay conditions (optimal concentration of baby rabbit complement and second incubation times), mean levels of non-specific killing and mean opsonic indexes achieved with human IVIg, pre and post rabbit anti-full-length M-protein antisera. The mean levels of non-specific killing were calculated from seven sequential assays. Mean opsonic indexes were calculated from three sequential assays.

M-Protein		1			12			6
Strain ID		43	02-12	GAS05134	611020	611025	GAS09437	GASOPA6_02
Optimal Concentration of Complement		2.1%	2.1%	2.1%	2.1%	3.1%	2.1%	3.1%
Non-specific killing (n=7)		20.3	13.4	6.87	13.5	20.2	13.0	30.6
Incubation Time (mins)		60	60	60	60	90	60	60
GMT Opsonic Index (n=3)	Pre-immunisation	<10	<10	<10	<10	<10	<10	<10
	Homologous	2307	2753	2571	7610	4685	2902	5794
	Heterologous (1)	<10	<10	<10	<10	<10	<10	<10
	Heterologous (2)	<10	<10	<10	<10	<10	<10	<10
	IVIg	4840	3443	4303	3360	12599	15663	1271

Table 3 GAS OPKA Qualifications Summary

Summary table of assay qualifications. The maximum percentage inhibition of killing by rabbit anti-full-length M-protein sera (pre-incubated with purified homologous or heterologous purified M-protein) is shown for each strain (n=1). Intra-assay and inter-assay coefficients of variation were calculated by running rabbit anti-full-length M-protein sera three times on the same assay or three individual assays, respectively. The coefficient of variation of opsonic indexes is shown for each strain (n=3). The relative accuracy was assessed by running rabbit full-length M-protein antisera, diluted to four different factors with OPS buffer. The percent agreement (whether OIs are within a factor of 3 of each other) and coefficient of variation (%) between the OIs for each of these four samples is shown (n=4). The linearity of the GAS OPKA was assessed by running non-immune rabbit serum, spiked with four different concentrations of full-length M-protein antisera. The correlation (R-squared) between initial sample dilution (log transformed) and resultant OI (log transformed) of the four samples is shown (n=4). The lower limit of quantification (LLOQ) was assessed by running full-length M-protein-spiked rabbit serum against each GAS strain. These samples were generated by spiking heat-inactivated non-immune rabbit serum with sufficient homologous rabbit anti-full-length M-protein antisera to produce a sample with an opsonic index of 4-12. The average OI is shown for each run (n=4).

M-Protein		1			12			6
Strain ID		43	02-12	GAS05134	611020	611025	GAS09437	GASOPA6_2
Specificity (n=1)	Inhibition by homologous antigen (%)	87	73	85	83	100	85	71
	Inhibition by heterologous antigen (%)	11	12	10	9	11	11	6
Precision (n=3)	Intra-assay coefficient of variation (%)	24.7	10.8	21.3	19.5	16.2	6.7	8.9
	Inter-assay coefficient of variation (%)	26.4	20.0	18.0	21.5	20.9	9.4	9.0
Relative Accuracy (n=4)	Agreement (%)	100	100	100	100	100	100	100
	Coefficient of variation (%)	15.21	14.5	8.69	17.0	20.8	16.1	22.3
Linearity (n=4)	R-squared	0.993	0.996	0.997	0.966	0.976	0.988	0.993
	Slope	-1.10	-1.14	-1.04	-1.18	-1.01	-1.11	-0.92
LLOQ (n=4)		8.75	14.75	7.50	6.50	5.00	6.75	6.25

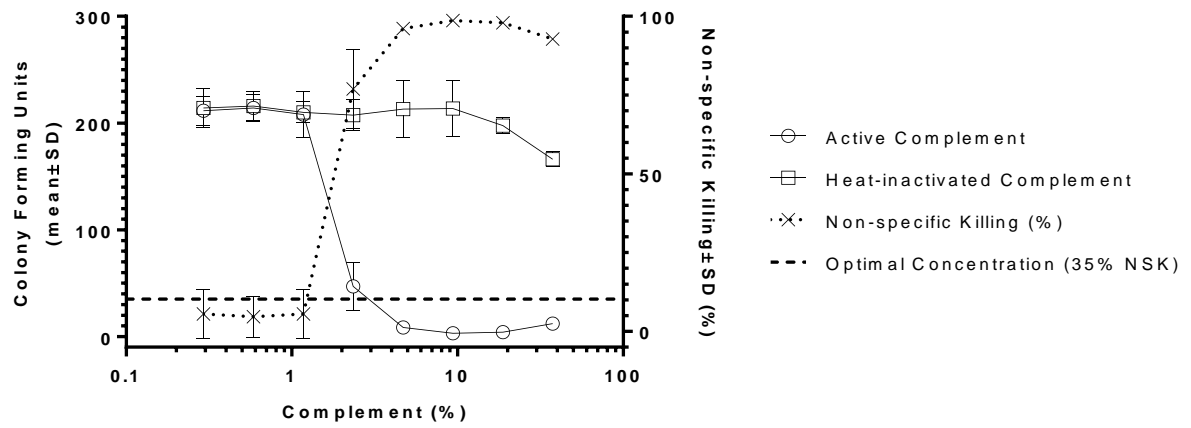


Fig 1 GAS OPKA Optimisation

The optimal concentration of BRC for GAS OPKA was calculated by incubating bacteria with a serial dilution of BRC in the presence of differentiated HL-60 cells. **b**, Representative data showing the number of CFU (left-y-axis) and percentage non-specific killing (right-y-axis, dotted line) at each dilution of heat-inactivated (open squares) or active (open circles) BRC with GAS strain M1(43) and differentiated HL-60 cells. Dilutions of BRC resulting in <35% non-specific killing were used as the optimal dilution (right-y-axis, dashed line).

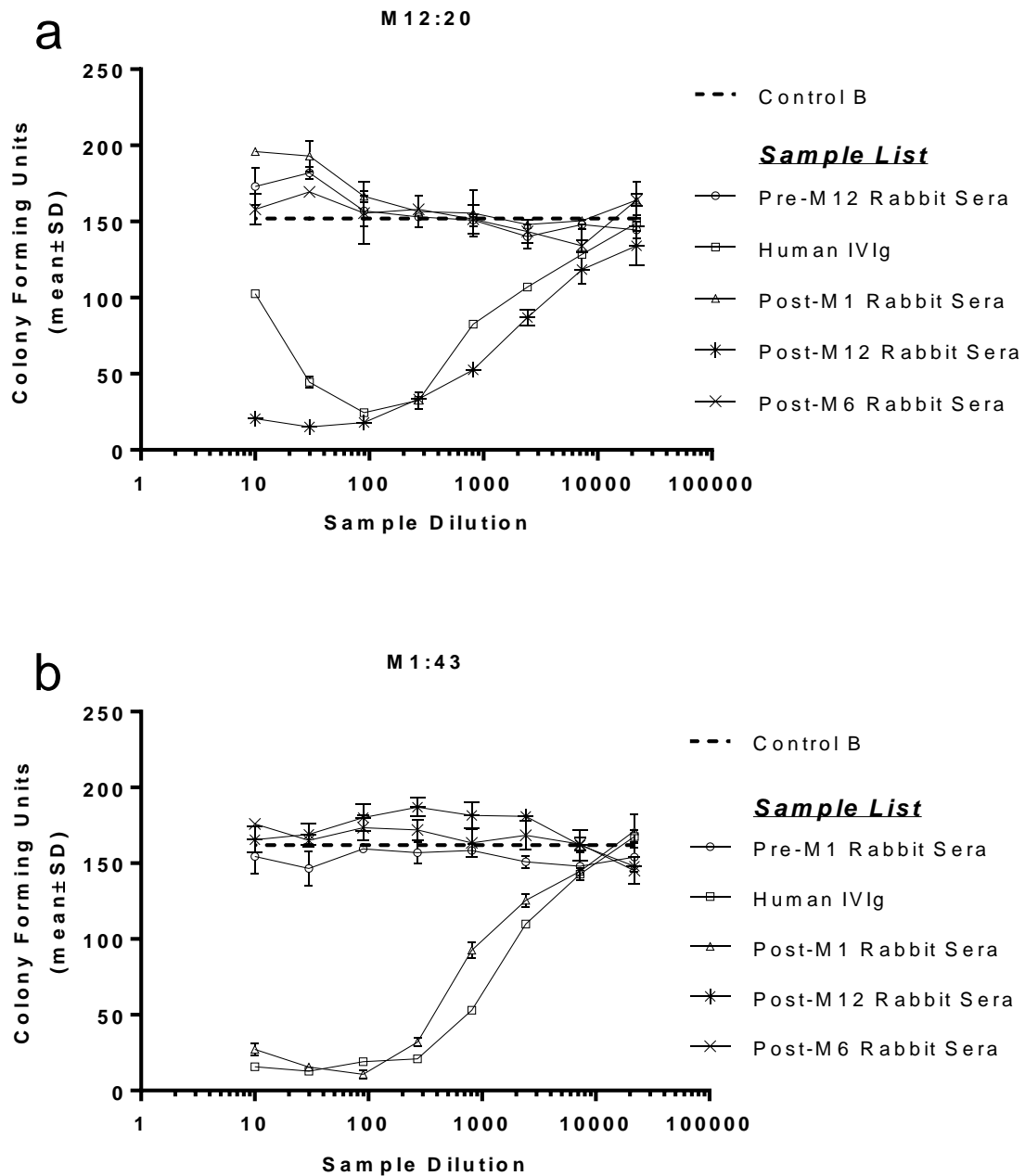


Fig 2 – Opsonophagocytic Killing Assay

Representative GAS OPKA data for GAS strains M1(43) (a) and M12(611020) (b). Both strains were run with human IVIg (open squares), rabbit anti-M1 sera (open triangles), rabbit anti-M6 sera (crosses), rabbit anti-M12 sera (stars) and either pre-M1 immunisation sera or pre-M12 immunisation sera (open circles). The average number of colony forming units (CFU) at each sample dilution is shown with the active complement control (control B – dashed line).

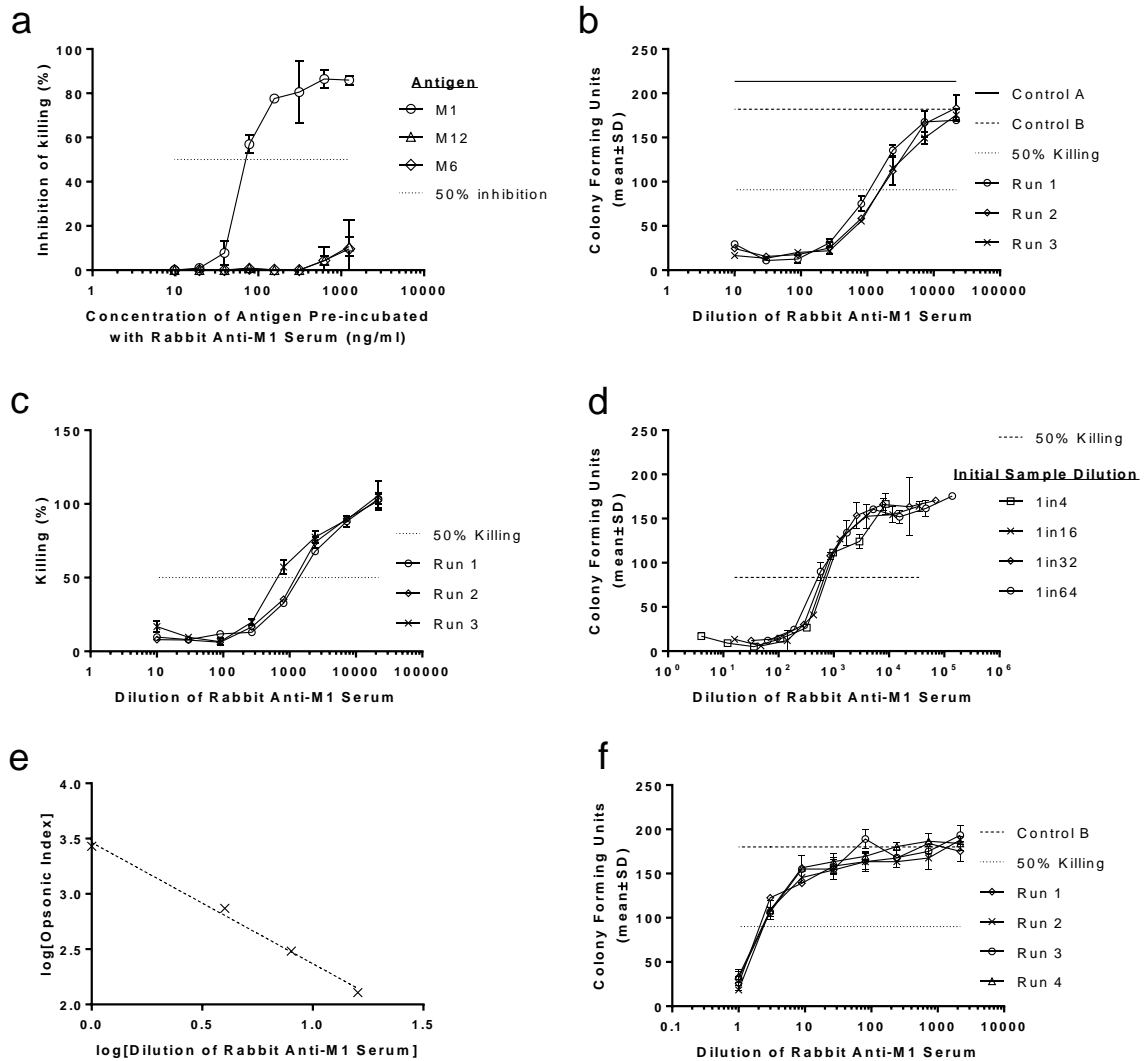


Fig 3 GAS OPKA Qualifications

a, Percentage inhibition of killing of the GAS strain M1(43) by rabbit anti-M1 serum pre-incubated with purified M1 (open circles), M6 (open diamond) or M12 (open triangle) protein. The maximum inhibition of killing achieved by homologous antigen (M1) was 86.5% whereas maximum inhibition by heterologous antigen was 9.7% (M6) and 10.7% (M12). **b**, Rabbit anti-full-length M-protein antiserum was run against GAS strain M1(43) three times on the same plate to calculate the intra-assay coefficient of variation of the opsonic index (50% killing – dotted line). The average number of colony forming units (CFU) at each sample dilution for the three runs is shown with heat-inactivated (A – solid line) and active (B – dashed line) complement controls. **c**, Rabbit anti-full-length M-protein antisera was run against GAS strain M1(43) three times on three different plates with individual controls to calculate the inter-assay coefficient of variation of the opsonic index (50% killing – dotted line). The average percentage killing of M1(43) at each sample dilution for the three runs is shown. **d**, Rabbit anti-full-length M-protein antiserum was run against GAS strain M1(43) at four different dilution on the same plate to calculate the relative accuracy. The average number of CFU at each sample dilution for the four initial dilution is shown with the 50% killing line (dashed line). **e**, Non-immune rabbit serum was spiked with four different concentrations of rabbit anti-full-length M-protein antiserum and run against GAS strain M1(43) to assess assay linearity. The opsonic index at each sample dilution (both log transformed) is shown with the line of linear regression (dashed line). **f**, Non-immune rabbit serum was spiked with sufficient rabbit anti-full-length M-protein antiserum to produce an opsonic index of 4 and run against GAS strain M1(43) to assess the lower limit of quantification. The average number of CFU at each sample dilution of four runs is shown with the 50% killing line (dotted line) and CFU with active complement (control B – dashed line).