

SASPject: A Novel Antibacterial Technology Targeting MDR *Pseudomonas aeruginosa* Demonstrating a Low Propensity for Resistance Development

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ABSTRACT

Background: Phico's SASPject technology uses a synthetic biology approach to create nano-delivery vehicles (NDV), capable of delivering a SASP gene expression system to target bacteria. For novel antibacterials to be successful it is critical that they demonstrate a low propensity for the development of resistance. The unique mode of action of SASPject, including the ability of SASP to prevent DNA replication and gene transcription in a DNA sequence-independent manner, coupled with rapid bactericidal activity suggests SASPject has a low propensity to induce resistance compared to other antibiotics. Here the potential for resistance of *Pseudomonas aeruginosa* (*Pa*) to develop in response to *Pa* SASPject therapy was measured in a 52-day passaging study

Methods: Passaging studies: *Pa* 3176 cultures at $\sim 10^5$ cfu/ml in Luria Bertani broth supplemented with magnesium, calcium, and glucose (LB+) were passaged daily after overnight growth at 37 °C in the presence of *Pa* SASPject at 1.25 x 10⁶, 6.25 x 10⁵ and 3.13 x 10⁵ U/ml. A control without added *Pa* SASPject was also passaged in parallel. Susceptibility of the cultures to SASPject was assessed daily by 3 h kill assays, with $\sim 10^5$ cfu/ml *P. aeruginosa* and $\sim 1.5 \times 10^9$ U/ml SASPject incubated statically at 37 °C.

Results: Bacterial kill to below the detection limit of 10 cfu/ml was observed in each 3 h kill assay against *Pa*. No decrease in sensitivity was observed under any of the three *Pa* SASPject concentrations measured during 52 days of monitoring, after which the experiment was stopped.

Conclusion: The unique mode of action of SASP means that mutations in bacterial DNA cannot lead to resistance as SASP inactivates bacterial DNA in a sequence-independent manner. This is supported by the data presented here, where resistance did not develop within 52 days of passaging *Pa* with *Pa* SASPject. In conclusion the SASPject technology platform displays a low propensity for resistance development in the treatment of *Pa*.

INTRODUCTION

The emergence of multidrug resistant Gram-negative bacteria such as *Pseudomonas aeruginosa*, *Escherichia coli*, *Acinetobacter baumannii*, and *Klebsiella spp.* amongst others is a global public health issue and has highlighted the urgent need for new therapeutic options, ideally with novel mechanisms of action.

SASPject™ is a new class of antibacterial currently under development (1). The SASPject platform technology comprises NDV's designed to deliver a gene encoding an antibacterial protein (SASP) into target bacteria (Figure 1). SASPs are small proteins expressed by sporulating bacteria which, when delivered into bacteria non-specifically, inactivate bacterial DNA and lead to rapid cell death. Due to the specific nature of the NDV-bacterium interaction, single-species or multi-species or genera can be selectively targeted. Consequently, non-selective inhibition of commensal bacteria, which is a significant drawback of some conventional antibiotic therapies, can be avoided.

Previously, a SASPject antibacterial (PT1.2) against *Staphylococcus aureus*, including a diverse range of antibiotic-resistant strains including MRSA and VISA/hVISA (1), has been developed. PT1.2 successfully passed through a Phase I clinical trial for nasal decolonisation of *S. aureus* in humans and demonstrated excellent *in vitro* activity (2, 3).

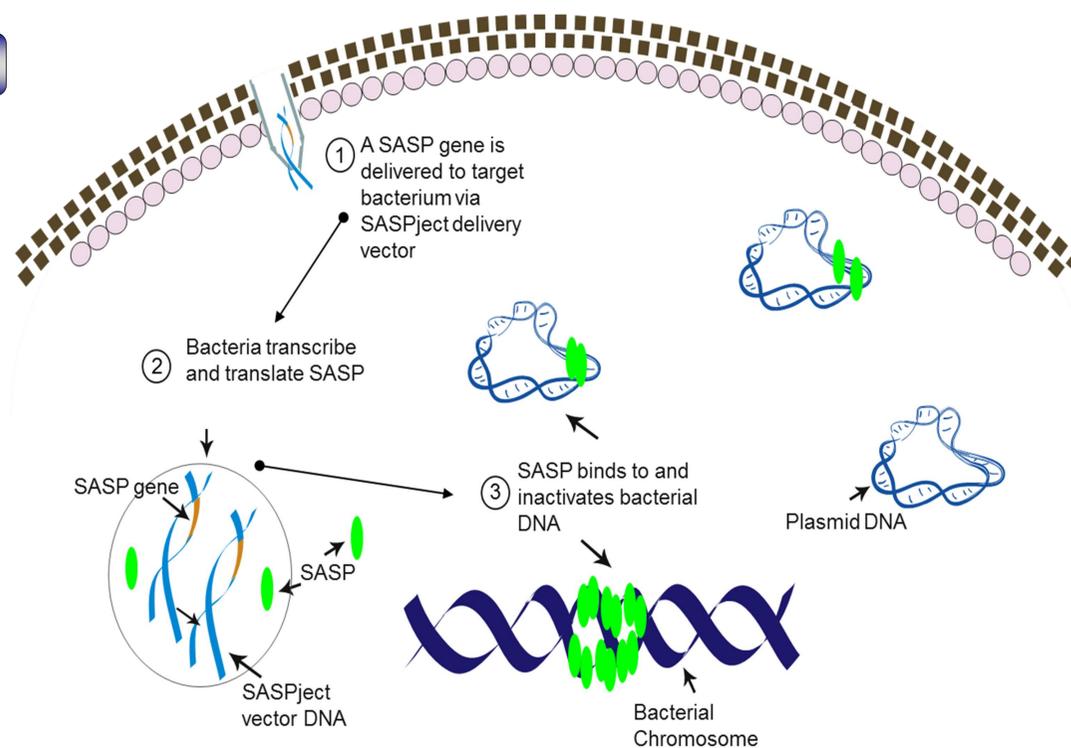


Figure 1. Mechanism of action of *Pa* SASPject

Pa SASPject has demonstrated a rapid bactericidal activity against a geographically diverse range of MDR *Pa* strains (4) supporting the previously published effective SASPject therapeutic targeting *S. aureus*, PT1.2 (2,3). Due to SASPject's unique mode of action the technology has demonstrated a low propensity for resistance development to *S. aureus* (5). In this study, *P. aeruginosa* has been serially passaged in the presence of three concentrations of *Pa* SASPject representing inhibitory to sub-inhibitory concentrations of *Pa* SASPject. The susceptibility of serially passaged *Pa* was measured at regular intervals.

METHODS

Passaging studies

P. aeruginosa Asia strain 3176 was streaked onto a fresh Luria-Bertani agar plate containing MgSO₄ (5 mM), CaCl₂ (5 mM) and Glucose (0.1% w/v) (LB+) and grown overnight at 37 °C. Four individual colonies were subsequently re-suspended into 5 ml LB+ broth and grown overnight at 37 °C with shaking (350 rpm). The following day the OD₆₀₀ of the overnight culture of *Pa* strain 3176 was measured and diluted to 10⁵ cfu/ml in fresh LB+ broth. *Pa* SASPject was added under three different exposure regimes 1.25 x 10⁶, 6.25 x 10⁵ and 3.13 x 10⁵ U/ml.

Triplicate cultures were used for each of the exposure regimes while a no exposure control culture was inoculated with *Pa* strain 3176. Cultures were incubated overnight, static at 37 °C. After overnight culture the OD₆₀₀ was measured and cultures were diluted back to 10⁵ cfu/ml and *Pa* SASPject was added at the same exposure concentrations. This procedure was repeatedly daily.

The susceptibility of the passaged cultures exposed to *Pa* SASPject were assessed on a daily basis by a 3 h kill assay. Briefly, an aliquot of the overnight culture diluted for serial passaging studies (10⁵ cfu/ml) was incubated with *Pa* SASPject (1 x 10⁹ U/ml) and incubated for 3 h statically at 37 °C. Samples were then diluted and plated onto LB+ agar plates. Following overnight incubation at 32 °C viable cells were enumerated.

RESULTS

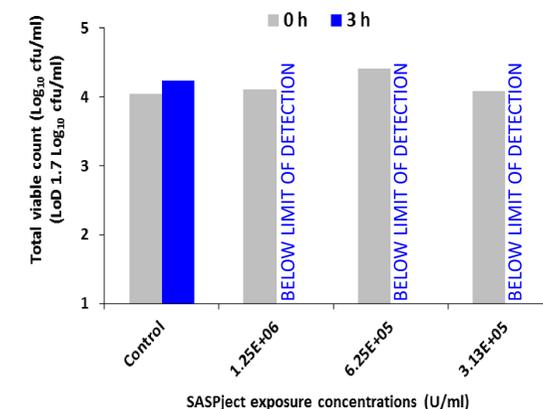


Figure 2. 3 h kill assay of *Pa* Asian strain 3176 (1 x 10⁵ cfu/ml) by *Pa* SASPject (1 x 10⁹ U/ml) after 20 days exposure to *Pa* SASPject therapeutic

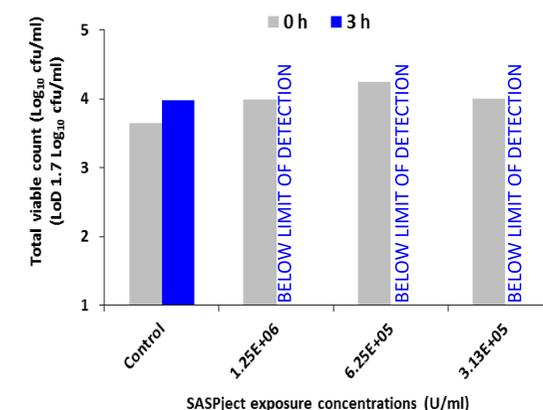


Figure 3. 3 h kill assay of *Pa* Asia strain 3176 (1 x 10⁵ cfu/ml) by *Pa* SASPject (1 x 10⁹ U/ml) after 40 days exposure to *Pa* SASPject therapeutic

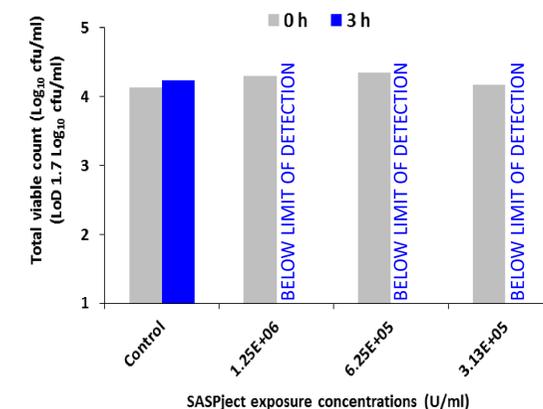


Figure 4. 3 h kill assay of *Pa* Asia strain 3176 (1 x 10⁵ cfu/ml) by *Pa* SASPject (1 x 10⁹ U/ml) after 52 days exposure to *Pa* SASPject therapeutic

- After 52 days of serial passaging the susceptibility of *P. aeruginosa* Asia strain 3176 to *Pa* SASPject was unaffected, with complete removal of the bacterial load to below the detection limit within 3 hours of exposure (Figures 2-4)

- Differing exposure concentrations of SASPject had no effect on the potential for resistance to develop in *P. aeruginosa* with removal to below the detection limit within 3 h (Figures 2-4)

CONCLUSIONS

- Pseudomonas aeruginosa* remains susceptible to *Pa* SASPject following 52 days of exposure

- Sub-inhibitory concentrations of SASPject do not increase the potential for the development of resistance of kill of *P. aeruginosa*.

- Observations reported here reflect those seen with PT1.2 against MRSA (5) and highlight that the unique mode of action of SASP means both Gram-positive and Gram-negative bacteria are extremely unlikely to develop resistance

- Further studies are required to characterise *Pa* SASPject binding and gene delivery *in vivo*

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