

DDS-03: A Highly Potent and Selective Novel Small Molecule Series to treat Multi-Drug Resistant *Neisseria gonorrhoeae*

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Background: The emergence and spread of multi-drug resistance to antibiotics used against the Gram-negative bacterial pathogen, *Neisseria gonorrhoeae*, the causative agent of the sexually transmitted infection gonorrhoea has resulted in a dramatic loss of effective regimens for the condition. Currently, ceftriaxone (injectable) is the only viable monotherapy option available. A robust pipeline of novel antibiotics is required to provide a sustainable defence against *N. gonorrhoeae* infections worldwide.

To this end, Summit is developing new mechanism oral pathogen specific antibiotics for the treatment of *N. gonorrhoeae*. Here we report on a novel chemotype with excellent levels of activity and remarkable selectivity associated with a novel mechanism of action as established using our Discuva Platform. The DDS-03 chemotype and mechanism is distinct to our other compound series targeting *N. gonorrhoeae*, SMT-571 (see poster #647).

Methods: Agar-based MICs were established according to CLSI guidelines with ceftriaxone as a comparator. Agar MICs and mutational frequencies were performed using WHO *N. gonorrhoeae* reference strains. *In vitro* kill kinetic assays were conducted in broth using the multi-drug resistant clinical isolate [WHO-M]. *In vitro* ADME and toxicological assays were run using standardised protocols internally and at established service providers. Mechanism of action studies were performed using Summit's proprietary transposon and bioinformatics based technologies.

Results:

- DDS-03 series exhibits excellent activity across *N. gonorrhoeae* clinical isolates (WHO Panel), including the highly resistant WHO-X Japanese strain.
- Examples from the DDS-03 series show no shift or a 2-fold shift in the presence of 4% serum.
- A panel of 96 clinical isolates, accessed through the NIAID preclinical services, returned MIC values of 0.01 µg/ml.
- DDS-03 series exhibits remarkable pathogen specificity.
- This series is bactericidal and exhibits low levels of mutational frequency (~2.8x10⁻⁸–3x10⁻⁹) → defined mutants do not exhibit cross-resistance to azithromycin and doxycycline.
- Examples from the DDS-03 series do not show any evidence of synergy or antagonism in combination with azithromycin and doxycycline.

ACTIVITY	FA1090	WHO-M	WHO-L	WHO-N	WHO-O	WHO-G	WHO-F	WHO-K	WHO-X
MIC (µg/ml)	0.001	0.002	0.001	0.002	0.001	0.002	0.001	0.002	0.001

Table 1: DDS-03a is active against multi-drug resistant *N. gonorrhoeae* clinical isolates (WHO Panel).

Compound	MIC ₉₀	Number of Isolates
DDS-03b	0.01 µg/ml	96

Table 2: DDS-03 series exhibits potent activity across 96 clinical isolates (supported by NIAID).

Genus	Species	Strain	MIC (µg/ml)		Genus	Species	Strain	MIC (µg/ml)	
			DDS-03c	Ciprofloxacin				DDS-03c	Ciprofloxacin
<i>Staphylococcus</i>	<i>aureus</i>	BAA 1720	>64	>32	<i>Escherichia</i>	<i>coli</i>	ATCC 25922	>64	≤0.63
<i>Staphylococcus</i>	<i>epidermidis</i>	NRS7	>64	8	<i>Haemophilus</i>	<i>influenzae</i>	A1 1246001	>64	-
<i>Bacillus</i>	<i>thuringiensis</i>	ATCC 35646	>64	0.25	<i>Legionella</i>	<i>pneumophila</i>	ATCC 33152	>64	0.5
<i>Enterococcus</i>	<i>faecalis</i>	ATCC 51299	>64	0.5	<i>Neisseria</i>	<i>gonorrhoeae</i>	ATCC 49226 (DSM 9189)	≤0.125	0.004
<i>Enterococcus</i>	<i>faecium</i>	NCTC 8619	>64	4	<i>Streptococcus</i>	<i>pneumoniae</i>	4478-07	>64	-
<i>Enterobacter</i>	<i>cloacae</i>	NCTC 13464	>64	0.25	<i>Streptococcus</i>	<i>pyogenes</i>	EUP_112	>64	-
<i>Enterobacter</i>	<i>aerogenes</i>	DSM 12058	>64	>32	<i>Lactobacillus</i>	<i>rhamnosus</i>	ATCC 53103	>64	-
<i>Moraxella</i>	<i>catarrhalis</i>	DSM 11994	>64	≤0.06	<i>Neisseria</i>	<i>meningitidis</i>	ATCC 13077 (DSM 10036)	≤0.125	-
<i>Acinetobacter</i>	<i>calcoaceticus</i>	DSM 16962	>64	0.125	<i>Clostridium</i>	<i>difficile</i>	ATCC 43255	>64	-
<i>Acinetobacter</i>	<i>junii</i>	DSM 14968	>64	0.125	<i>Clostridium</i>	<i>perfringens</i>	ATCC 8237	>64	-
<i>Acinetobacter</i>	<i>ursingii</i>	DSM 16037	>64	1	<i>Peptostreptococcus</i>	<i>anaerobius</i>	DSM 20357	>64	-
<i>Salmonella</i>	<i>enterica</i>	NCTC 6017	>64	≤0.63	<i>Bacteroides</i>	<i>fragilis</i>	ATCC 25285	>64	-
<i>Pseudomonas</i>	<i>aeruginosa</i>	EUP_10.1	>64	0.25	<i>Clostridium</i>	<i>innocuum</i>	DSM 1286	>64	-
<i>Proteus</i>	<i>mirabilis</i>	DSM 30116	>64	≤0.63	<i>Roseburia</i>	<i>faecis</i>	DSM 16840	>64	-
<i>Providencia</i>	<i>rettgeri</i>	DSM 1131	>64	≤0.63	<i>Bifidobacterium</i>	<i>bifidum</i>	ATCC 11863	>64	-

Table 3: A representative from the DDS-03 series has been profiled against a panel of 30 bacteria.

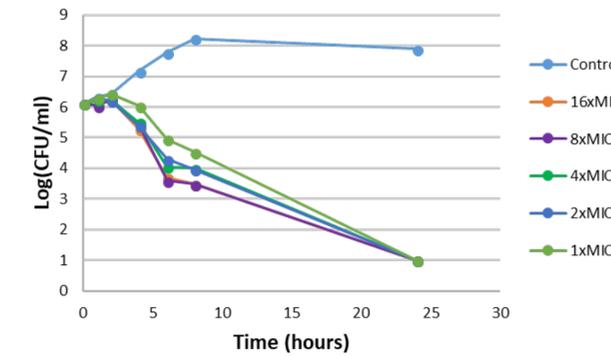


Figure 1: Time-kill studies confirmed a bactericidal killing profile against the multi-drug resistant clinical isolate WHO-M.

Compound	FoR (4xMIC)	
	24h	48h
DDS-03e	3.77x10 ⁻⁸	>2.9x10 ⁻⁷
DDS-03f	2.88x10 ⁻⁹	2.88x10 ⁻⁹
DDS-03g	2.65x10 ⁻⁹	3.57x10 ⁻⁸

Table 4: DDS-03 series exhibit low levels of mutational frequency.

RBC	Cytotox	Mitotox	GSH (50 µM)	Ames (25 µM)
>200 µM	>100 µM	Negative	Negative	Negative

Table 5: The DDS-03 chemotype has a clean ADME and toxicology profile.

Discuva Technology: Target Identification

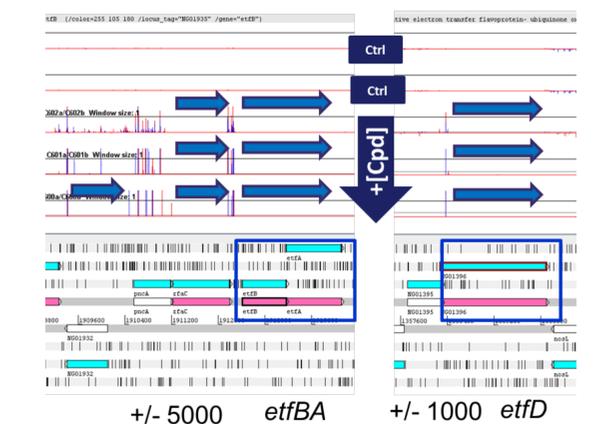


Figure 2: Discuva Technology identified the electron transfer proteins (etf) A/B/D as the mechanism of action for the DDS-03 series.

- Single dose pharmacokinetic (SDPK) studies where compounds were administered 1mg/kg IV and 10mg/kg PO have returned highly favourable oral C_{Max}/MIC ratios for representative examples from the series.
- DDS-03 series has a clean toxicology profile in RBC, cytotoxicity and mitotoxicity.
- DDS-03 series is negative in Ames assay and no adducts were observed in a glutathione incubation assay in the presence and absence of S9 fraction.
- Our Discuva Platform identified the electron transfer proteins (etf) A/B/D as the mechanism of action for the DDS-03 series which was further confirmed by whole genome sequencing of the resistant mutants.

Conclusions: Summit Therapeutics is developing a small molecule oral antibiotic with exceptional levels of potency, selectivity and a novel mechanism of action for the treatment of *N. gonorrhoeae* infections. The series is undergoing further optimization.

Acknowledgment:

