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Broad activity of diphenyleneiodonium analogues against *Mycobacterium tuberculosis*, malaria parasites and bacterial pathogens

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Graphical abstract



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- 2 malaria parasites and bacterial pathogens
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- 23
- 24 Key words: diphenyleneiodonium, type II NADH-quinone oxidoreductase, Gram-negative,
- 25 *Plasmodium*, malaria.

26 Abbreviations: DPI, diphenyleneiodonium; NADH, nicotinamide adenine dinucleotide

27 reduced; NDH-2, type II NADH-quinone oxidoreductases

28

29 Abstract

In this study, a structure-activity relationship (SAR) compound series based on the NDH-2 30 inhibitor diphenyleneiodonium (DPI) was synthesised. Compounds were evaluated primarily 31 for in vitro efficacy against Gram-positive and Gram-negative bacteria, commonly 32 responsible for nosocomial and community acquired infections. In addition, we also assessed 33 the activity of these compounds against Mycobacterium tuberculosis (Tuberculosis) and 34 Plasmodium spp. (Malaria). This led to the discovery of highly potent compounds active 35 against bacterial pathogens and malaria parasites in the low nanomolar range, several of 36 which had favourable toxicity profiles against mammalian cells. 37

38

39 1. Introduction

40 Antibiotic resistance has evolved into a serious global health concern [1, 2]. In the United States over 23,000 people die each year due to infections with antibiotic-resistant 41 bacteria. Notwithstanding the human cost, antibiotic resistance is also a massive economic 42 burden which has been estimated to cost as much as \$20 billion USD in excess healthcare 43 expenses, with associated lost productivity estimated to be as high as \$35 billion USD/year 44 [2]. Sadly, the 'magic bullet' antimicrobial therapies we have gratuitously used over the past 45 decades are rapidly losing their calibre. Modern healthcare over the last century has been 46 founded on the basis that bacterial infections can be effectively treated using antimicrobial 47 drugs. In a world without effective antibiotics, modern medical procedures that we take for 48 granted, such as chemotherapy or simple surgery, will have increasing risk due to the threat 49 of untreatable bacterial infections. Once again common bacterial infections will more than 50 often result in death. Medicine is clearly entering a critical period, if bacteria continue 51 developing resistance to multiple antibiotics at the present rate, and at the same time the 52 53 pipeline continues to dry up, there could be catastrophic costs to healthcare and society [3].

In the developing world, infectious diseases caused by microbial pathogens remain a major disease burden with malaria (*Plasmodium* spp.; 429,000 deaths in 2015)[4] and TB (*Mycobacterium tuberculosis*; 1,800,000 deaths in 2015)[5] contributing to >2 million deaths every year globally [6]. Multi-drug resistant malaria parasites, particularly *P. falciparum* which causes the greatest burden of mortality, and *M. tuberculosis* are spreading in the Asia-Pacific region [7, 8], reducing the effectiveness of current front-line drugs and increasing potentially deadly treatment failures.

There is an urgent unmet medical need to discover new scaffolds with superior 61 activity against these problematic human pathogens. Enzymes involved in energy metabolism 62 are emerging as very important novel drug targets for anti-infective drug development [9-15]. 63 64 Encouragingly, respiratory chain inhibitors appear to be the Achilles' heels of dormant nonreplicating cells ('persisters'), that are often refractory to antibiotics and difficult to treat [16]. 65 Instead of the multi-subunit complex I respiratory enzyme found in mammalian cells, 66 protozoa, bacteria and plants possess a single sub-unit non-proton pumping, rotenone 67 insensitive alternative complex I [10, 14, 17, 18]. This type II NADH-menaquinone 68 oxidoreductase (NDH-2) contains a single non-covalently bound flavin adenine dinucleotide 69 (FAD) cofactor and catalyzes the oxidation of NADH with menaquinone [19]. The absence 70 of NDH-2 in the respiratory chain of mammalian mitochondria makes it a very attractive 71 target for antibiotic drug development. 72

Diphenyleneiodonium (DPI) is known to be a potent, yet undeveloped, inhibitor of NDH-2 [15, 20]. A few very early reports investigated the potential of iodonium compounds, including DPI, as skin antiseptics against Gram-negative bacteria [21-23]. In the present study we have synthesized a series of novel DPI analogues and evaluated their *in vitro* effectiveness against infectious human pathogens that rely on NDH-2 for their energy needs, namely Gram-positive and Gram-negative bacteria commonly responsible for nosocomial

and community acquired infections and *Mycobacterium tuberculosis*. In addition, we explored the sensitivity of *Plasmodium* spp. malaria parasites, which have a type II NADH-menaquinone oxidoreductase of unclear function, to the DPI analogues. The potential of these compounds as antimicrobial leads with a novel mode of action are discussed.

83

84 2. Results

85 *2.1 Chemistry*

The synthetic route to access the diphenyleneiodonium core typically involves 3 steps 86 as described in the literature (Scheme 1)[24]. The biphenyl amine **3** is classically formed in 87 good to excellent yield using a Suzuki-coupling reaction between an appropriately substituted 88 2-iodoaniline 1 and an appropriately substituted phenylboronic acid 2. The biphenyl amine 3 89 was subsequently transformed into the biphenyl iodide 4 through diazotisation in Sandmeyer 90 reaction, followed by oxidation of the biphenyl iodide 4 utilising *m*-CPBA under acidic 91 conditions to afford the corresponding cyclic diphenyleneiodonium compound 5 (Scheme 1). 92 The desired final products were obtained as precipitates from the reaction mixture and were 93 easily isolated in high purity and acceptable yields. This procedure was successfully adopted 94 for the synthesis of all novel DPI analogues reported herein as shown in Table 1. 95



- and conditions: a) Dioxane/H₂O (9:1), K₂CO₃, TBAB, Pd(dppf)Cl₂, 130 °C, 1 h; b) 1. THF, 4 M HCl, 0 °C, aq.
- 100 NaNO₂, 20 min. 2. aq. KI, 0 °C 10 min, r.t, 1 h; c) CH₂Cl₂, *m*-CPBA, TfOH, r.t, 1 h.
- 101

97

98

- 102 **Table 1**
- 103 Structures of DPI analogues synthesised and numbering system adopted



Compound	R	Compound	R	Compound	R	Compound	R
50	Н)				5aa	5-F, 5'-F
5n	3-F	5i	4-F	5d	5-F	5z	5-F, 5'-Cl
5m	3-Cl	5h	4-Cl	5c	5-C1	5y	5-Cl, 2'-Cl
51	3-CN	5g	4-CN			5x	5-F, 2'Cl
5 1	2.014	5 0	4.014	51	5.014	-	
5K	3-OMe	51	4-OMe	50	5-OMe	SW	5-Me, 2 ⁻ -Cl
5j	3-Me	5e	4-Me	5a	5-Me	5v	5-Cl, 2-Cl
		5p	4,5-benzo			5u	5-Cl, 2-F
				5q	5-CF3	5t	5-Cl, 2-OMe

5r

5-OCF3

5s

5-Cl, 2-Me

105	<i>Note</i> : The ring numbering system used herein is only for readers' convenience (not IUPAC nomenclature).
106	The compounds were made iteratively in two series. The first series comprises 50, 5a, 5b, 5c,
107	5d, 5e, 5f, 5g, 5h, 5i, 5j, 5k, 5l, 5m, 5n, 5p and will be referred to hereon as DPI series 1,
108	while DPI series 2 comprises 5q , 5r , 5s , 5t , 5u , 5v , 5w , 5y , 5x , 5z , 5aa .

109

110 2.2 Gram-positive and -negative bacteria MIC determination

MIC determinations for DPI series 1 against several problematic Gram-positive and negative bacterial species revealed very potent activity for several analogues. In support of
our recent report [25], DPI itself (50) displayed good activity against *Pseudomonas aeuruginosa* (MICs generally 2.3-75 μM) and *Acinetobacter baumannii* (MICs 0.58-19 μM).
Activity was slightly weaker against *Klebsiella pneumoniae* (MICs 37 and 75 μM),
vancomycin-resistant *Enterococcus faecium* (MICs 37 μM) but better against drug resistant
strains of *Staphylococcus aureus* (MICs 4.7 and 19 μM) (Table 2).

Several analogues had very similar profiles to DPI, these being 5a, 5h, 5i, 5l, 5n and 118 **5p**, indicating that activity was not greatly influence by installation of a 5-Me, 4-Cl, 4-F, 3-119 CN, 3-F or a fused phenyl ring. One analogue lost significant activity in a broad spectrum 120 sense, this being **5b**, suggesting that the electron donating methoxy group in the 5-position is 121 particularly disfavoured. There were six other analogues (5e, 5f, 5g, 5j and 5k) that were 122 broadly similar in profile to DPI with the exception of selectively weaker activity against 123 certain strains, an example being **5f** (4-OMe) and **5k** (3-OMe) with strikingly weaker activity 124 against Pa QLD PSA (70 µM), Pa M146201 and Ab07AC-366 (Table 2). In contrast, 5c 125 126 displayed exquisitely potent activity across a broad spectrum of strains, with potency greater than the detectable limit for 11 bacterial strains. 127

The stability of the DPI analogues were investigated by incubating of analogue (**5j**) in bacterial culture media under standard assay conditions (37 °C, 200 μ L Cation-Adjusted Mueller-Hinton Broth, 96-well polypropylene microtitre plates) and tested the resulting chemical species of the compound by LCMS in the next day. The LCMS analysis showed that the compound was stable under this condition.

Further, the impact of the counter anion towards the biological activity was investigated. A small selected set of chloride counter ion analogues were synthesised, where the chloride counter ion replaces the triflate counter ion. These chloride containing compounds were tested against a selected Gram-negative bacteria (Pa ATCC 27853). The results indicate that there is no notable anion linked trend in these results, hence there appears to be no strong link between the biological activity and the presence/absence of triflate anion.

139

140 **Table 2**

MIC Activity µM (SI)									
	50	5a	5b	5c	5d	5e	5f	5g	PMB
	75	18	>70	< 0.27	2.2	>73	>70	71	
Pa ATCC 27853	(0.39)	(0.14)	(>0.55)	(<2.4)	(2.1)	(>0.52)	(>0.29)	(0.32)	0.77
	9.3	4.5	>70	0.54	4.5	36	70	18	
Pa QLD PSA 1	<mark>(3.1)</mark>	<mark>(0.58)</mark>	<mark>(>0.55)</mark>	(1.2)	<mark>(1.0)</mark>	<mark>(1.0)</mark>	<mark>(0.29)</mark>	(0.13)	<mark>0.77</mark>
	4.7	1.1	70	< 0.27	4.5	9.0	8.7	8.8	
Pa #912	<mark>(6.3)</mark>	<mark>(2.3)</mark>	<mark>(0.55)</mark>	<mark>(<2.4)</mark>	<mark>(1.0)</mark>	<mark>(4.2)</mark>	<mark>2.3)</mark>	<mark>(2.5)</mark>	<mark>1.5</mark>
	2.3	0.57	70	< 0.27	< 0.28	9.0	18	4.4	
Pa 19147nm	<mark>(13)</mark>	<mark>(4.6)</mark>	<mark>(0.55)</mark>	<mark>(<2.4)</mark>	<mark>(<16)</mark>	<mark>(4.2)</mark>	<mark>(1.2)</mark>	<mark>(5.1)</mark>	<mark>25</mark>
	4.7	1.1	70	< 0.27	< 0.28	9.0	35	8.8	
Pa 18878B klon 1	<mark>(6.3)</mark>	<mark>(2.3)</mark>	<mark>(0.55)</mark>	<mark>(<2.4)</mark>	<mark>(<16)</mark>	<mark>(4.2)</mark>	<mark>(0.58)</mark>	(2.5)	<mark>>25</mark>
	9.3	2.3	>70	0.54	1.1	36	70	8.8	
Pa M146201	<mark>(3.1)</mark>	<mark>(1.2)</mark>	<mark>(>0.55)</mark>	(1.2)	<mark>(4.1)</mark>	<mark>(1.0)</mark>	<mark>(0.29)</mark>	(2.5)	<mark>3.1</mark>
	9.3	2.3	35	< 0.27	0.56	18	35	18	
Ab ATCC 19606	(<u>3.1</u>)	(1.2)	(1.1)	<mark>(<2.4)</mark>	<mark>(8.2)</mark>	(2.2)	(0.58)	(0.13)	<mark>0.77</mark>
	4.7	2.3	70	<0.27	< 0.28	9.0	35	18	
Ab 246-01-C	(6.3)	(1.2)	<u>(0.55)</u>	(<2.4)	<u>(<16)</u>	<mark>(4.2)</mark>	(0.58)	(0.13)	<mark>0.19</mark>
	2.3	1.1	35	<0.27	0.56	9.0	18	4.4	0.00
Ab ATCC 17978	(<u>13)</u>	(2.3)	(1.1)	(<2.4)	(8.2)	(4.2)	(1.2)	(5.1)	<mark>0.38</mark>
A1 ATCC 10(0) 110	4.7	2.3	$\frac{1}{2}$	< 0.27	0.56	9.0	8.7	2.2	00
Ab ATCC 19606 coll0	(6.3)	(1.2)	(<u>2.2)</u>	(<2.4)	(8.2)	(4.2)	(2.3)	(10) 71	<mark>98</mark>
11 07 4 6 226	19	2.2	70	< 0.27	1.1	18	70	/1	<u>c 0</u>
Ab 0/AC-336	(<mark>1.6)</mark> 0.59	(1.2) 0.57	(0.55)	(<2.4)	(4.1)	(<u>2.2)</u>	(0.29)	(0.32)	<mark>6.2</mark>
Ab ATCC 17078 col10	0.58	(1.5)	$\frac{33}{(1,1)}$	< 0.27	< 0.28	1.1	$\frac{4.4}{(4.6)}$	$\frac{2.2}{(10)}$	12
Ab ATCC 17978 coll0	(<mark>50)</mark> 27	(4.0) 18	(1.1)	(<2.4) 4 3	(<10) 4 5	(<u>33)</u> 72	(4.6) 70	(10) 25	12
Kn ATCC 12992	$\frac{31}{(0.78)}$	$\frac{10}{(0,14)}$	>/U	$\frac{4.3}{(0.15)}$	$\frac{4.3}{(1.0)}$	$\frac{12}{(0.52)}$	$\frac{70}{(0.20)}$	$\frac{33}{(0.62)}$	12
KP ATCC 13003	37	36	(<u>>0.55)</u> 70	86	<u>(1.0)</u> 9.0	$\frac{(0.52)}{72}$	(0.29) 70	(0.05) 71	12
Kp M320445	(0.78)	(0.07)	(0.55)	(0.08)	(0.51)	(0.52)	(0.29)	(0.32)	<mark>0.38</mark>

141 Gram-positive and -negative bacteria MIC determination

	75	72	>70	8.6	9.0	>72	>70	>71	
Kp #1	<mark>(0.78)</mark>	<mark>(0.04)</mark>	<mark>(>0.55)</mark>	<mark>(0.08)</mark>	(0.51)	<mark>(>0.52)</mark>	<mark>(>0.29)</mark>	(>0.32)	<mark>98</mark>
-	75	>72	>70	8.6	9.0	>72	>70	>71	
Кр 224-11-С	<mark>(0.78)</mark>	<mark>(>0.04)</mark>	<mark>(>0.55)</mark>	<mark>(0.08)</mark>	<mark>(0.51)</mark>	<mark>(>0.52)</mark>	<mark>(>0.29)</mark>	<mark>(>0.32)</mark>	<mark>>25</mark>
	37	>72	>70	8.6	9.0	72	>70	71	
Кр 248-33-D	<mark>(0.78)</mark>	<mark>(>0.04)</mark>	<mark>(>0.55)</mark>	<mark>(0.08)</mark>	<mark>(0.51)</mark>	<mark>(0.52)</mark>	<mark>(>0.29)</mark>	<mark>(0.32)</mark>	<mark>12</mark>
	37	18	70	0.54	1.1	72	>70	71	
Ec N2381	<mark>(0.78)</mark>	<mark>(0.14)</mark>	<mark>(0.55)</mark>	<mark>(1.2)</mark>	<mark>(4.2)</mark>	<mark>(0.52)</mark>	<mark>(>0.29)</mark>	<mark>(0.32)</mark>	<mark>1.5</mark>
	37	>72	>70	0.54	1.1	>72	>70	>71	
Ec N4149	<mark>(0.78)</mark>	<mark>(>0.04)</mark>	<mark>(>0.55)</mark>	<mark>(1.2)</mark>	<mark>(4.2)</mark>	<mark>(>0.52)</mark>	<mark>(>0.29)</mark>	<mark>(>0.32)</mark>	<mark>1.5</mark>
	37	18	>70	2.2	4.5	72	>70	71	
Ec N11281	<mark>(0.78)</mark>	<mark>(0.14)</mark>	<mark>(>0.55)</mark>	<mark>(0.31)</mark>	<mark>(1.0)</mark>	<mark>(0.52)</mark>	<mark>(>0.29)</mark>	<mark>(0.32)</mark>	<mark>1.5</mark>
	4.7	2.3	8.7	< 0.27	0.56	4.5	4.4	1.1	
VRE ATCC 700221	<mark>(6.3)</mark>	<mark>(1.2)</mark>	<mark>(4.4)</mark>	<mark>(<2.4)</mark>	<mark>(8.2)</mark>	<mark>(8.3)</mark>	<mark>(4.6)</mark>	(<mark>20)</mark>	<mark>>25</mark>
	4.7	36	35	1.1	2.2	9.0	18	1.1	
MRSA ATCC 43300	<mark>(6.3)</mark>	<mark>(0.07)</mark>	<mark>(1.1)</mark>	<mark>(0.61)</mark>	<mark>(2.1)</mark>	<mark>(4.2)</mark>	(1.5)	(20)	<mark>>25</mark>
	19	36	35	2.3	2.2	18	35	8.8	
VISA ATCC 700698	<mark>(1.6)</mark>	<mark>(0.07)</mark>	<mark>(1.1)</mark>	<mark>(0.29)</mark>	<mark>(2.1)</mark>	<mark>(2.2)</mark>	<mark>(0.58)</mark>	(2.5)	<mark>>25</mark>
	19	36	35	1.1	2.2	18	35	8.8	
VRSA ATCC 700699	<mark>(1.6)</mark>	<mark>(0.07)</mark>	<mark>(1.1)</mark>	<mark>(0.61)</mark>	<mark>(2.1)</mark>	(2.2)	(0.58)	<mark>(2.5)</mark>	<mark>>25</mark>

VRSA ATCC 700699 (1. 142 Table 2. (continued)

				MIC Acti	vity µM <mark>(SI</mark>)				
	5h	5i	5j	5k	51	5m	5n	5p	PMB
	8.7	18	72	>70	35	8.7	8.9	8.4	
Pa ATCC 27853	(1.9)	(0.67)	(0.05)	(>0.11)	(0.09)	(0.13)	(0.66)	(0.09)	0.77
	8.7	18	>72	70	8.8	2.2	8.9	4.2	
Pa QLD PSA 1	(1.9)	(0.67)	(>0.05)	(0.11)	(0.37)	(0.51)	(0.66)	(0.17)	0.77
-	4.3	2.2	4.5	>70	4.4	1.1	1.1	2.1	
Pa #912	(3.7)	(5.3)	(0.81)	(>0.11)	(0.75)	(1.0)	(5.3)	(0.34)	1.5
	4.3	1.1	4.5	8.7	2.2	0.54	0.56	2.1	
Pa 19147nm	(3.7)	(11)	(0.81)	(0.81)	(1.5)	(2.0)	(11)	(0.34)	25
	2.2	2.2	4.5	18	4.4	0.54	1.1	2.1	
Pa 18878B klon 1	(7.5)	(5.3)	(0.81)	(0.39)	(0.75)	(2.0)	(5.3)	(0.34)	>25
	43	4 5	36	70	18	2.2	2.2	8.4	
Pa M146201	(3.7)	(2,7)	$\frac{00}{(0,11)}$	(0,11)	(0.19)	$\frac{2.2}{(0.51)}$	$\frac{2.2}{(2.6)}$	(0, 09)	31
1 4 1011 10201	43	4 5	91	18	18	2.2	4 5	4.2	<u></u>
Ab ATCC 19606	$\frac{(37)}{(37)}$	$\frac{(27)}{(27)}$	(0.41)	$\frac{10}{(0.39)}$	$\frac{10}{(0.19)}$	$\frac{2.2}{(0.51)}$	$\frac{(2.6)}{(2.6)}$	$\frac{12}{(0.17)}$	0.77
no mee 19000	43	4.5	18	35	18	2.2	<u>(2.0)</u> 4.5	4.2	0.77
Ab 246-01-C	$\frac{(37)}{(37)}$	$\frac{(2,7)}{(2,7)}$	(0.21)	$\frac{33}{(0.21)}$	$\frac{10}{(0.19)}$	$\frac{2.2}{(0.51)}$	$\frac{(2.6)}{(2.6)}$	$\frac{1.2}{(0.17)}$	0.19
710 240 01 C	$\frac{(3.7)}{2.2}$	$\frac{(2.7)}{2.2}$	4.5	18	4.4	0.54	1.1	1 1	0.17
Ab ATCC 17978	$\frac{2.2}{(7.5)}$	$\frac{2.2}{(5.3)}$	(0.81)	$\frac{10}{(0.30)}$	$\frac{4.4}{(0.75)}$	$\frac{0.54}{(2.0)}$	$\frac{1.1}{(5.3)}$	$\frac{1.1}{(0.60)}$	0.38
A0 ATCC 17978	(7.5)	2.2	(0.01)	(0.39)	(0.75)	(2.0) 1.1	(3.3)	(0.09)	0.50
Ab ATCC 10606 col10	$\frac{2.2}{(7.5)}$	$\frac{2.2}{(5.2)}$	(0.81)	$\frac{2.2}{(2.1)}$	$\frac{2.2}{(1.5)}$	(1.0)	$\frac{2.2}{(2.6)}$	1.1	08
A0 A1CC 19000 collo	(7.5) 9.7	(J.J) 0.1	(0.01) 10	(3.1) > 70	(1.5) 5 71	(1.0)	(2.0)	(0.09)	<mark>90</mark>
AL 07 AC 226	8./	9.1	18	>/0	>/1	2.2	8.9	8.4	C D
AD 0/AC-330	(1.9)	(1.5)	(0.21)	(>0.11)	(<u>>0.05)</u>	(0.51)	(0.00)	(0.09)	0.2
AL ATCC 17078110	0.5	0.56	1.1	2.2	2.2	< 0.27	0.56	(1.4)	10
Ab ATCC 1/9/8 coll0	(<u>30)</u>	(2.0)	(<u>3.2)</u>	(<u>3.1</u>)	(1.5)	(<4.1)	(11) 26	(1.4)	12
K ATCC 12992	8.7	18	72 (0.05)	>/0	18	8./	36	8.4	10
Kp ATCC 13883	(1.9)	(0.67)	(0.05)	(>0.11)	(0.19)	(0.13)	(0.16)	(0.09)	12
VF	35	18	36	70	35	8.7	36	17	0.00
Кр М320445	(0.47)	(0.67)	(0.11)	(0.11)	<mark>(0.09)</mark>	(0.13)	(0.16)	<mark>(0.04)</mark>	<mark>0.38</mark>
	35	36	72	>70	>71	17	72	34	
Kp #1	(0.47)	(0.33)	(0.05)	(>0.11)	(>0.05)	<u>(0.06)</u>	<mark>(0.08)</mark>	(0.02)	<mark>98</mark>
	35	36	72	>70	>71	17	72	34	
Кр 224-11-С	(0.47)	(0.33)	(0.05)	(>0.11)	<u>(>0.05)</u>	<mark>(0.06)</mark>	<mark>(0.08)</mark>	(0.02)	<mark>>25</mark>
	17	18	72	>70	>71	17	36	34	
Кр 248-33-D	(0.93)	<mark>(0.67)</mark>	<u>(0.05)</u>	(>0.11)	<u>(>0.05)</u>	(0.06)	(0.16)	(0.02)	<u>12</u>
	17	18	72	35	71	8.7	18	8.4	
Ec N2381	<mark>(0.93)</mark>	<mark>(0.67)</mark>	(0.05)	(0.21)	(0.05)	<u>(0.13)</u>	<mark>(0.08)</mark>	<mark>(0.09)</mark>	1.5
	17	36	72	>70	71	8.7	18	34	
Ec N4149	<mark>(0.93)</mark>	<mark>(0.33)</mark>	<mark>(0.05)</mark>	<mark>(>0.11)</mark>	<mark>(0.05)</mark>	<mark>(0.13)</mark>	<mark>(0.08)</mark>	<mark>(0.02)</mark>	1.5
	17	18	36	>70	71	4.3	18	17	
Ec N11281	<mark>(0.93)</mark>	<mark>(0.67)</mark>	<mark>(0.11)</mark>	<mark>(>0.11)</mark>	(0.05)	<mark>(0.26)</mark>	<mark>(0.08)</mark>	<mark>(0.04)</mark>	1.5
	8.7	4.5	4.5	4.4	2.2	1.1	2.2	1.1	
VRE ATCC 700221	<mark>(1.9)</mark>	<mark>(2.7)</mark>	<mark>(0.81)</mark>	<mark>(1.6)</mark>	<mark>(1.5)</mark>	<mark>(1.0)</mark>	<mark>(2.6)</mark>	<mark>(0.69)</mark>	<mark>>25</mark>
	2.2	4.5	4.5	4.4	4.4	2.2	2.2	2.1	
MRSA ATCC 43300	<mark>(7.5)</mark>	(2.7)	<mark>(0.81)</mark>	<mark>(1.6)</mark>	<mark>(0.75)</mark>	(0.51)	<mark>(2.6)</mark>	<mark>(0.34)</mark>	<mark>>25</mark>
	4.3	18	9.1	8.7	8.8	2.2	8.9	4.2	
VISA ATCC 700698	<mark>(3.7)</mark>	<mark>(0.67)</mark>	(0.41)	<mark>(0.78)</mark>	(0.37)	(0.51)	<mark>(0.66)</mark>	(0.17)	<mark>>25</mark>

	4.3	9.1	9.05	8.7	8.8	2.2	8.9	4.2	
VRSA ATCC 700699	<mark>(3.7)</mark>	(1.3)	<mark>(0.398)</mark>	<mark>(0.78)</mark>	<mark>(0.37)</mark>	(0.51)	<mark>(0.66)</mark>	(0.17)	<mark>>25</mark>
(SI) = colocitivity index relative to UEDC2 colls and was coloulated by CC = (10/EDS)/MIC									

- 143 (SI) = selectivity index relative to HEPG2 cells and was calculated by CC_{50} (1%FBS)/MIC.
- 144

145 2.3 Inhibition of Mycobacterium tuberculosis growth

Shown in Table 3 are the results of testing DPI series 1 compounds and DPI series 2 146 compounds against *M. tuberculosis*. It can be seen in Table 3 that all compounds displayed 147 anti-mycobacterial activity and that the weakest compound, 5b, nicely matched the 148 observation that this compound was also the weakest in the bacterial panel. Further, 5e, 5f, 5j 149 and **5k** are also common to the groupings of compounds that are slightly less potent than DPI. 150 The observation that 5c was very potent in the bacterial panel but also amongst the 151 most potent compounds against M. tuberculosis led us to synthesise a set of second 152 generation analogues based around the structure of 5c. This compound contains a 5-chloro 153 substituent and so our DPI series 2 focuses on analogues with a halogen in a 5-position. 154 Gratifyingly, as show in Table 3, these compounds were uniformly extremely potent. In 155 particular, 5s, with 5-Cl, 2-Me substitution, was extremely potent, and 3-fold more potent 156 than **50** (DPI itself), with an MIC of $0.13 \,\mu$ M. 157

158

159 **Table 3**

160 Inhibition of *Mycobacterium tuberculosis* growth

DPI Series 1	MIC μM (SI)	DPI Series 2	MIC μM <mark>(SI)</mark> (SI) ^a
50	0.30 (96)	50	0.26 (0.09) (0.36) ^ª
(50	0.14 (10)	5	0.25 (0.21) (0.00)3
(58	0.14 (19)	51	0.23 (0.21) (0.99)
5b	4.4 <mark>(8.8)</mark>	55	0.13 <mark>(0.74)</mark> (2.8) ^a
5c	0.28 <mark>(2.4)</mark>	5t	0.51 <mark>(0.31)</mark> (0.44) ^a
5d	0.29 <mark>(16)</mark>	5u	0.27 <mark>(3.4)</mark> (10) ^a
5e	0.57 <mark>(66)</mark>	5v	0.50 <mark>(1.8)</mark> (3.7) ^a
5f	1.1 <mark>(19)</mark>	5w	0.27 <mark>(0.22)</mark> (1.8) ^a

	A	CCEPTED N	MANUSCRIPT	
5g	0.29 <mark>(78)</mark>	5x	0.27 <mark>(0.19)</mark> (1.4) ^a	
5h	0.54 <mark>(30)</mark>	5y	0.26 <mark>(0.32)</mark> (2.1) ^a	
5i	0.29 <mark>(41)</mark>	5z	1.0 <mark>(0.38)</mark> (0.80) ^a	
5j	0.57 <mark>(6.4)</mark>	5aa	0.28 <mark>(0.32)</mark> (0.58) ^ª	
5k	0.28 <mark>(24)</mark>			
51	0.29 <mark>(12)</mark>			
5m	0.28 <mark>(3.9)</mark>			
5n	0.29 <mark>(20)</mark>			
5р	0.52 <mark>(2.4)</mark>			
Rifampicin	0.30		0.61	\bigcirc

161 (SI) = selectivity index relative to HEPG2 cells and was calculated by CC_{50} (1%FBS)/MIC.

162 $(SI)^{a}$ = selectivity index relative to HEPG2 cells and was calculated by CC₅₀ (10% FBS)/MIC.

163

164 2.4 In vitro activity of DPI against Plasmodium spp.

DPI has been shown to inhibit P. falciparum growth in vitro, however, the reported 165 166 IC_{50} values and the proposed mechanism of action vary [26, 27]. In this study, the growth inhibitory IC₅₀ for D10 (chloroquine sensitive; 0.13 μ M) parasites after 90 hours of treatment 167 (early ring stage to late trophozoite stage next cycle) was 3.3-fold higher than that achieved 168 169 for the CS2 chloroquine resistant line (0.04 µM; Table 4 and Suppl. Fig. 1A), indicating that DPI is inhibitory to both chloroquine resistant and sensitive strains in vitro. The zoonotic 170 human pathogen P. knowlesi YH1[28], an emerging pathogen in Southeast Asia and a 171 laboratory adapted model for the major human pathogen P. vivax, had an IC_{50} 5.6 fold higher 172 than that of the chloroquine sensitive D10 line at 0.74 µM (Table 4 and Suppl. Fig. 1A). 173 Whether this represents a difference in parasite sensitivities, or is due to the shorter lifecycle 174 of P. knowlesi (around 32 hours in these assays) and therefore reduced exposure time is not 175 clear. 176

177 The *in vitro* growth inhibitory activity after 90 hours of treatment of the 26 DPI 178 analogues was tested against the D10 chloroquine sensitive line (Table 5, Suppl. Fig. 1B and

179 1C). The activity of the analogues ranged from >3-fold decrease in growth inhibitory IC₅₀ compared to DPI (to below 0.045 µM), to a loss of growth inhibition up to a concentration of 180 1 µM. Six of the 26 analogues had a greater than 2-fold increase in growth inhibitory activity. 181 P. knowlesi growth inhibition was also tested against three analogues which displayed 182 increased potency against P. falciparum. Analogues 5c (P.k. YH1 0.08 µM versus D10 0.05 183 μ M, p>0.054, Table 5), 5d (*P.k.* YH1 0.08 μ M versus D10 0.05 μ M, p>0.032) and 5g (*P.k.* 184 YH1 0.11 μ M versus D10 0.04 μ M, p>0.0016), showed significantly improved potency over 185 DPI and inhibited *P. knowlesi* growth to a similar extent to *P. falciparum*. 186

Next we attempted to make the D10 chloroquine sensitive line resistant to DPI. After 187 6 days of treatment with 0.2 µM DPI, the DPI concentration was raised to 0.4 µM for a 188 further 6 days. The DPI concentration was then reduced back to 0.2 µM and parasites grown 189 continuously under drug pressure. After a further 18 days of culture, viable parasites were 190 visible and maintained under drug pressure until a stable population could be obtained (called 191 D10-DPI^R). A sub culture of DPI selected parasites was made and the drug pressure was 192 removed to test whether a tolerant, rather than resistant, parasite population had been selected 193 for (line with DPI drug pressure removed for between 2-5 weeks is called D10-DPI^{off}). There 194 was no statistical difference between the growth inhibitory IC_{50} for the D10-DPI^r line (0.36) 195 μ M) and the D10-DPI^{off} line (0.46 μ M; p=0.264). In contrast, there was a ~2-fold difference 196 between both the D10-DPI^r (p=0.011) and D10-DPI^{off} (p=0.005) lines compared to D10 (0.19) 197 µM, Fig. 2) in parallel experiments. Sequence comparison between the D10-PfPHG parental, 198 D10-DPI^r and D10-DPI^{off} lines showed no mutations in PfNDH2, suggesting it is unlikely 199 that mutations in PfNDH2 are causing the reduced sensitivity to DPI. The sequence of Type 200 II *Pf* dihydroorotate dehydrogenase, reported as an alternative target of PfNDH2 inhibitors 201 [27], was also found to have no mutations associated with reduced sensitivity to DPI. It 202 remains to be determined the mechanism by which resistance to DPI occurs in these 203

204 parasites. Attempts to select drug resistant parasites at a DPI concentration of 0.8 μ M 205 between days 6 and 12 were unsuccessful.

206 Changes in sensitivity of the D10-DPI^R line to more potent DPI analogues **5q**, **5c**, **5d** 207 and **5g** were assessed by comparing the growth inhibitory IC₅₀ between the resistant and 208 parental lines (Table 5). The D10-DPI^R line was significantly more tolerant of analogue **5g** 209 (D10-DPI^R 0.09 μ M versus D10 0.04 μ M, p=0.003), but not of **5c** (D10-DPI^R 0.07 μ M versus 210 D10 0.05 μ M, p=0.165), **5d** (D10-DPI^R 0.08 μ M versus D10 0.05 μ M, p=0.0567) and **5q** 211 (D10-DPI^R 0.14 μ M versus D10 0.13 μ M, p=0.091). These data indicate that selection for a 212 DPI resistant line did not necessarily confer resistance to analogues of DPI.

In general, the potencies of DPI analogues in series 1 for bacterial species and M. 213 tuberculosis (e.g increased potency for 5c; reduced potency for 5b) were reflected in the 214 relative potency for *P. falciparum*. However, some discrepancies were observed with 5g and 215 51, both potent inhibitors of *P. falciparum* that were very poor inhibitors of bacteria. This 216 result suggests that the hydrophilic electron withdrawing groups at the 3- and 4-position are 217 particularly disfavoured against bacterial isolates. In terms of series 2 compounds, there were 218 broad similarities between the IC₅₀ for *P. falciparum* and *M. tuberculosis* (R^2 0.27, p=0.101), 219 with the exception of 5u and 5v which were inhibitory to *M. tuberculosis* but not *P*. 220 221 falciparum growth, suggesting that 2 electron withdrawing substituents at the 2- and 5position on the same phenyl ring are not tolerated for *P. falciparum* growth inhibition. When 222 5u and 5v were removed from the comparison, the correlation between growth inhibitory 223 IC₅₀s for *P. falciparum* and *M. tuberculosis* improved significantly (R^2 0.455, p=0.046), 224 confirming the broad similarity of drug potency for series 2 compounds against these 225 important human pathogens. 226

227

229 **Table 4**

Parasite line	IC ₅₀ -90 hour
	(µM)
<i>Pf</i> D10 (chloroquine sensitive)	0.13
Pf CS2 (chloroquine resistant)	0.04
P. knowlesi YH1	0.74

230 In vitro activity of DPI against P. falciparum and P. knowlesi malaria

231 232

233 **Table 5**

234 In vitro activity of DPI analogues against P. falciparum and P. knowlesi malaria

DPI	IC ₅₀ µM <mark>(SI)</mark>	IC ₅₀ µM <mark>(SI)</mark>	IC ₅₀ µM <mark>(SI)</mark>	DPI	IC ₅₀ µM (SI)	IC ₅₀ µM (SI)
Series 1	<i>Pf</i> D10	<i>Pf</i> D10 ^r	Pk YH1	Series 2	(SI) ^a	(SI) ^a
					<i>Pf</i> D10	<i>Pf</i> D10 ^r
50	0.13 <mark>(221)</mark>			5q	0.13 <mark>(0.18)</mark> (0.72) ^a	0.14 <mark>(0.18)</mark> (0.70) ^a
5a	0.07 <mark>(37)</mark>			5r	0.13 <mark>(0.39)</mark> (1.9) ^a	
5b	0.17 <mark>(225)</mark>			55	0.11 <mark>(0.82)</mark> (3.1) ^a	
5c	0.05 <mark>(13)</mark>	0.07 <mark>(8.9)</mark>	0.08 (8.5)	5t	0.13 (1.2) (1.75) ^a	
5d	0.05 <mark>(96)</mark>	0.08 <mark>(61)</mark>	0.08 <mark>(59)</mark>	5u	>1 <mark>(>0.93)</mark> (>2.7) ^a	
5e	0.08 <mark>(500)</mark>			5v	>1 <mark>(>0.89)</mark> (>1.9) ^a	
5f	0.10 <mark>(213)</mark>			5w	0.11 <mark>(0.63)</mark> <mark>(5.1)</mark> ª	
5g	0.04 <mark>(531)</mark>	0.09 <mark>(256)</mark>	0.11 <mark>(232)</mark>	5x	0.19 <mark>(0.28)</mark> (2.1) ^a	
5h	0.08 <mark>(194)</mark>			5y	0.36 <mark>(0.23)</mark> (1.5) ^a	
5i	0.10 <mark>(125)</mark>			5z	>1 <mark>(>0.39)</mark> (0.83) ^a	
5j	0.23 <mark>(15.8)</mark>			5aa	0.84 <mark>(0.11)</mark> (0.19) ^a	
51	0.05 <mark>(62)</mark>					
5m	0.07 <mark>(15)</mark>					





Supplementary Fig 1: *P. falciparum* growth inhibitory activity of DPI and analogues *in vitro*. (A) Dose response curves demonstrate that DPI is inhibitory to *P. falciparum* growth *in vitro*, with an IC₅₀ <0.2 μ M after 90 hours of parasite treatment for both chloroquine sensitive (D10) and chloroquine resistant (CS2) lines. Analogues of DPI in many cases demonstrated improved growth inhibitory activity against D10 parasites with (B) 11 out of 15 analogues for series 1, and (C) 1 out of 11 analogues for series 2 demonstrating >2 fold reduction in growth inhibitory IC₅₀.



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Fig 2. Parasite line selected for resistance to DPI shows reduced drug sensitivity. D10 parasites selected for
resistance to DPI during continuous culture (D10-DPI^R) exhibited ~2 fold reduction in sensitivity to DPI, even
after extended removal of drug pressure (D10-DPI^{off}, 2-5 weeks without drug pressure) when compared to D10
parental parasites.

260

262 In parallel with the above assays, we assessed both series of DPI analogues for cytotoxicity against HEPG2 cells. As shown in Table 6, DPI series 1 displayed a wide range 263 of cytotoxicity in 1% FBS, from as low as 0.66 µM for 5c to as high as 38 µM for 5b. Ten of 264 the 15 series 1 compounds had improved antimalarial activity in the low nM range (Table. 5; 265 5a, 5d, 5e, 5f, 5g, 5h, 5i, 5k, 5l, 5m) whilst maintaining a CC_{50} above 1 μ M. For M. 266 tuberculosis (Table. 4), 8 out of 15 compounds met this criteria (5a, 5d, 5g, 5i, 5k, 5l, 5m, 267 5n), confirming that improved DPI analogue activity against broad pathogens is not 268 absolutely at the expense of mammalian cell toxicity. As seen for compound 5c from which 269 series 2 was derived, analogues with a halogen in a 5-position were significantly more 270 cytotoxic (Table 6). However, this is at conditions artificially low in serum concentration 271 (1%). When retested in the presence of 10% FBS, cytotoxicity decreased up to 8-fold and 272 toxicity would decrease further in vivo where serum concentration is 100%. Nevertheless, 273 cytotoxicity, particularly for the series 2 compounds, is a clear potential liability and will 274 need to be addressed in future DPI analogue optimisation studies. 275

- 276
- 277 **Table 6**
- 278 Cytotoxicity Assay

	CC50[µM]			CC ₅₀ [µM]	
DPI Series 1	1%FBS	10%FBS	DPI Series 2	1%FBS	10%FBS
50	29.2		5q	0.02361	0.0947
5a	2.6		5r	0.05026	0.2508
5b	38.3		5s	0.09295	0.3488
5c	0.66		5t	0.1504	0.2242
5d	4.6		5u	0.9303	2.724
5e	37.5		5v	0.8938	1.848
5f	20.2		5w	0.0601	0.4863
5g	22.3		5x	0.05238	0.3864

	A	CCEPTED MANUSCRIPT		
5h	16.1	5y	0.08457	0.5436
5i	11.9	5z	0.3915	0.8331
5j	3.6	5aa	0.08862	0.161
5k	6.8			
51	3.3			
5m	1.1			
5n	5.9			
5p	0.72			

281 **3. Discussion**

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There has been a steady decline in the number of FDA approved antibiotics, with only 282 283 10 antibiotics considered "New Molecular Entities" being approved by the FDA from 2004 to 284 2012 [29, 30]. This is dwarfed in comparison to the 20 new classes of antibiotics developed between 1930 and 1962, and the 30 new antibiotics approved between 1983 and 1992 [29]. 285 286 There were no new classes of antimicrobials discovered between 1968 and 2000, and the two novel classes discovered in 2000 and 2003 (daptomycin and linezolid) are only effective 287 against Gram-positive bacteria [29]. The lack of new antibiotics renders physicians impotent 288 to treat emerging resistance to existing antibiotics. Globally, spreading resistance to 289 antimalarial and anti-tubercular drugs is of urgent concern [7, 8], with increasing treatment 290 failures and the potential for increased mortality in the years ahead unless alternative 291 treatments become available. The present report helps address this urgent need for new anti-292 infectives by evaluating the *in vitro* antibacterial and antimalarial activity of a novel series of 293 294 DPI compounds.

In view of the increasing incidence of MDR pathogens there is an urgent need for new antibiotics with novel modes of action. Agents that selectively target respiratory enzymes that are unique to the pathogen such as NDH-2 in the electron transport chain of bacteria and

298 certain parasites will offer a superior approach for treating persistent infections, greatly reduced treatment periods and provide a high selectivity for the pathogen versus the host. DPI 299 has been show to inactivate flavin enzymes via a radical reaction mechanism which leads to 300 the covalent modification of the reduced flavin cofactor [31-33]. The reduced flavin transfers 301 an electron to DPI, generating semiquinone and a diphenyliodyl radical. The free radical 302 fragments to give iodobenzene and a phenyl radical, the latter undergoes recombination with 303 the flavin semiguinone to form various phenyl adducts [32]. We have previously shown that 304 DPI inhibits NDH-2 activity in isolated *Escherichia coli* membranes [34]. Coincidently, we 305 also demonstrated that a secondary mode of action of the polymyxin lipopeptide antibiotics 306 against Gram-negative bacteria involves the inhibition of NDH-2 activity [34]. 307

The unique structure of the Gram-negative cell wall provides an often impermeable barrier to antibiotics, particularly hydrophobic compounds [35]. A major advantage of diaryliodonium compounds is their amphipathic character which allows them to easily cross even the most formidable membrane barrier such as the Gram-negative cell wall. Despite their apparent attractiveness as anti-infective agents, iodonium compounds have undergone limited drug development.

A few very early reports investigated the potential of iodonium compounds including 314 DPI as skin antiseptics against Gram-negative bacteria [21-23]. The compounds per se or as 315 emulsions with pine oils showed bactericidal activity. The MIC values reported against both 316 Gram-positive and Gram-negative bacteria obtained in the present study were much lower 317 (~10-fold) compared to those of early reports, suggesting DPI is much more active than 318 previously thought [36-38]. This discrepancy may have arisen due to the fact these authors 319 employed DPI dissolved in aqueous solutions without any co-solvent and then attempted to 320 estimate the concentration using the dipicrylamine chemical reactivity technique [36-38]. 321

Acute toxicity studies with dogs (received an intraperitonal dose of 70 mg/kg) and 322 monkeys (received intraperitonal doses of doses of 30 and 50 mg/kg) indicated DPI has some 323 toxic effects on the central nervous system and skeletal muscle at high concentrations [39, 324 40]. DPI has also been shown to act as a primary eye irritant and a moderate irritant via 325 dermal exposure in humans [41, 42]. In oral exposure studies with dogs at levels of 100 326 mg/kg, principle toxicity included frequent vomiting, reduced cardiac efficiency and 327 electrolyte imbalance [43, 44]. DPI has also been reported to induce hypoglycaemia in rats at 328 high concentrations, which can be minimized by fortifying the drinking water with glucose 329 [45]. The LD50 of DPI administered orally to rats is 60 mg/kg. Based on these animal 330 toxicity studies it appears that DPI related toxicity occurs at very high concentrations well 331 beyond the very low µM levels required to kill the tested bacterial species *in vitro*. This offers 332 some assurance of safety and a high degree of pathogen versus host selectivity, however, this 333 will also be dependent upon the pharmacokinetics of DPI. Encouragingly, tissue distribution 334 studies in rats that were administered [¹²⁵I]-DPI revealed a predominant localization of 335 radioactivity in the liver, kidneys, heart and adipose tissue [46]. 336

From a medicinal chemistry perspective, an iodonium chemotype would usually be 337 viewed with great caution. However, data achieved in this study which shows analogues can 338 be synthesised with improved pathogen killing activity that maintain desirably low 339 mammalian cell toxicity suggests such a compound could be considered worthy of further 340 elaboration to investigate structure-activity relationships. For the first time, we have reported 341 here a detailed assessment of the SAR across a range of microorganisms for a novel set of 342 DPI analogues. We show that some of our novel DPI analogues are much more potent than 343 DPI itself with IC₅₀ values in the low nanomolar range. However, some compounds, 344 particularly those with a halogen in a 5-position in series 2, are relatively cytotoxic to 345

mammalian cells and need to be further modified to determine if cytotoxicity can be reducedwhile potency against bacterial and parasite pathogens is maintained.

While DPI is one of the less cytotoxic compounds against HEPG2 cells, it is one of 348 the most potently cytotoxic compounds against THP1 cells. Therefore there seems to be 349 something of a disconnect between our observation of *in vitro* cytotoxicity and the historical 350 use of DPI in vivo with an apparently acceptable, albeit moderate, toxicity profile. For this 351 reason, while cytotoxicity-associated liability would necessarily require monitoring, it is 352 possible that the extreme potency of some of our DPI analogues could find an application in 353 certain antibacterial settings after further improvements in the cytotoxicity profile. Whether 354 sufficiently divergent SAR can be established to build on the antibacterial potency we have 355 achieved and minimise mammalian cytotoxicity for further development of this class for in 356 vivo use will be the focus of future research. 357

358

359 4. Conclusion

A series of DPI analogues were synthesised and subsequently assessed for their biological activity. Several of these compounds exhibited high potency, with low nanomolar activity against problematic Gram-negative and Gram-positive bacteria, *Mycobacterium tuberculosis* and *Plasmodium* spp., protozoan parasites.

364

365 **5. Experimental**

366 5.1 Chemistry

All non-aqueous reactions were performed under an atmosphere of nitrogen, unless otherwise specified. Commercially available reagents were used without further purification. Thin-layer chromatography (TLC) was performed on silica gel $60F_{254}$ pre-coated aluminum sheets (0.25 mm, Merck). Automated flash column chromatography was performed with a Biotage Isolera

371 One instrument using Biotage SNAP cartridges packed with silica (KP-SilTM). Nuclear Magnetic Resonance (NMR) spectra were recorded at 400.13 Hz on an Avance III Nanobay 372 400 MHz Bruker spectrometer coupled to the BACS 60 automatic sample changer. Proton 373 resonances are annotated as: chemical shift (δ), multiplicity (s, singlet; d, doublet; m, 374 multiplet), coupling constant (J, Hz), and the number of protons. Mass spectrometry was 375 performed with an Agilent 6224 TOF LC/MS coupled to an Agilent 1290 Infinity (Agilent, 376 Palo Alto, CA). All data were acquired and reference mass corrected via a dual-spray 377 electrospray ionisation (ESI) source. Analytical HPLC was acquired on an Agilent 1260 378 Infinity analytical HPLC coupled with a G1322A degasser, G1312B binary pump, G1367E 379 high-performance autosampler, and G4212B diode array detector. Conditions were as 380 follows: Zorbax Eclipse Plus C18 rapid resolution column (4.6×100 mm) with UV detection 381 at 254 and 214 nm, 30 °C; the sample was eluted using a gradient of 5–100% solvent B in 382 solvent A, where solvent A was 0.1% aq. TFA and solvent B was 0.1% TFA in CH₃CN 383 (5-100% B [9 min], 100% B [1min]; 0.5 mL/min). 384

385

386 *5.1.1 General Method A: Preparation of Biphenylamine Derivatives* **3a-3aa**.

Substituted 2-iodoaniline 1 (1.0 eq.), substituted phenylboronic acid 2 (1.2 eq.), K_2CO_3 (3.0 387 eq.), tetrabutylammonium bromide (0.1 eq.), PdCl₂(dppf) (0.1 eq.) and dioxane/H₂O (9:1) 388 (0.5 M) were added to a 10 mL microwave-vial. The vial was sealed with a cap and placed in 389 a Cem Discover-microwave cavity. After irradiation at 130 °C for 1 h and subsequent 390 cooling, the solvent was removed *in vacuo*. The residue was taken up into EtOAc (30 mL) 391 and washed once with water and brine. The organic layer was dried over MgSO₄, filtered, and 392 concentrated. The crude product was purified by flash column chromatography using 0-10% 393 EtOAc/petroleum benzine to give the biphenylamine product **3a-3aa**. 394

- 5.1.2 The following compounds have all been reported previously and their ¹H spectra data
 showed good agreement with the literature data:
- 398 2'-Methyl-[1,1'-biphenyl]-2-amine (**3a**)[47], 2'-Methoxy-[1,1'-biphenyl]-2-amine (**3b**)[48], 2'
- 399 Fluoro-[1,1'-biphenyl]-2-amine (3c)[49], 2'-Chloro-[1,1'-biphenyl]-2-amine (3d)[50], 3'-
- 400 Methyl-[1,1'-biphenyl]-2-amine (3e)[51], 3'-Methoxy-[1,1'-biphenyl]-2-amine (3f)[51], 3'-
- 401 Fluoro-[1,1'-biphenyl]-2-amine (3h)[51], 3'-Chloro-[1,1'-biphenyl]-2-amine (3i)[51], 4'-
- 402 Methyl-[1,1'-biphenyl]-2-amine (**3j**)[51], 4'-Methoxy-[1,1'-biphenyl]-2-amine (**3k**)[51], 4'-
- 403 Fluoro-[1,1'-biphenyl]-2-amine (**3m**)[51], 4'-Chloro-[1,1'-biphenyl]-2-amine (**3n**)[51], [1,1'-
- 404 Biphenyl]-2-amine (**30**)[52], 2-(Naphthalen-2-yl)aniline (**3p**)[53], (2'-Amino-[1,1'-
- 405 biphenyl]-2-yl)-2,2,2-trifluoroethan-1-one (3r)[54].
- 406

5.1.3 2'-Amino-[1,1'-biphenyl]-3-carbonitrile (3g). Light yellow solid (0.29 g, 65% yield).
¹H NMR (400 MHz, CDCl₃) δ 7.81 (ddd, J = 2.2, 1.7, 0.8 Hz, 1H), 7.78 – 7.72 (m, 1H), 7.68
- 7.63 (m, 1H), 7.57 (td, J = 7.7, 0.5 Hz, 1H), 7.24 (ddd, J = 8.0, 7.4, 1.6 Hz, 1H), 7.12 (dd, J
= 7.6, 1.4 Hz, 1H), 6.90 (td, J = 7.5, 1.1 Hz, 1H), 6.85 (dd, J = 8.0, 0.8 Hz, 1H), 4.19 (br s,
2H). ¹³C NMR (101 MHz, CDCl₃) δ 143.42, 140.94, 133.64, 132.64, 130.72, 130.31, 129.68,
129.48, 125.02, 118.95, 118.72, 116.04, 112.95. LCMS Rt 3.21 min, *m*/z 195.1 [M + H]⁺.

5.1.4 2'-Amino-[1,1'-biphenyl]-4-carbonitrile (3l). Brown oil (0.27 g, 61% yield). ¹H NMR
(400 MHz, CDCl₃) δ 7.75 (dq, J = 5.4, 1.7 Hz, 2H), 7.65 – 7.61 (m, 2H), 7.26 (d, J = 7.7 Hz,
1H), 7.15 (d, J = 7.6 Hz, 1H), 6.93 (t, J = 7.5 Hz, 1H), 6.87 (d, J = 8.0 Hz, 1H), 4.41 (s, 2H).
¹³C NMR (101 MHz, CDCl₃) δ 144.68, 143.40, 132.71, 130.33, 129.93, 129.72, 125.55,
119.10, 118.94, 116.19, 110.96. LCMS Rt 3.22 min, *m/z* 195.1 [M + H]⁺.

5.1.5 2'-(Trifluoromethyl)-[1,1'-biphenyl]-2-amine (3q). Dark brown oil (0.39 g, 74% yield).
¹H NMR (400 MHz, CDCl₃) δ 7.47 – 7.34 (m, 4H), 7.25 (td, J = 7.8, 1.6 Hz, 1H), 7.13 (dd, J
= 7.5, 1.4 Hz, 1H), 6.92 (t, J = 8.8 Hz, 2H), 4.42 (s, 2H). ¹³C NMR (101 MHz, CDCl₃) δ
147.05, 144.04, 132.84, 132.52, 130.97, 129.25, 127.39, 122.54, 121.83, 121.53, 119.26,
118.50, 115.80. LCMS Rt 3.39 min, m/z 239.3 [M + H]⁺.

425

5.1.6 2'-Chloro-5'-methyl-[1,1'-biphenyl]-2-amine (3s). Brown solid (0.39 g, 79% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.40 (d, J = 8.1 Hz, 1H), 7.24 (ddd, J = 8.0, 7.4, 1.6 Hz, 1H), 7.20
- 7.11 (m, 2H), 7.08 (dd, J = 7.5, 1.5 Hz, 1H), 6.91 – 6.82 (m, 2H), 3.72 (s, 2H), 2.37 (s, 3H).
¹³C NMR (101 MHz, CDCl₃) δ 143.87, 137.70, 137.24, 132.61, 130.81, 130.49, 129.93,
129.69, 129.15, 125.66, 118.46, 115.63, 20.94. LCMS Rt 3.41 min, *m/z* 218.1 [M + H]⁺.

431

5.1.7 2'-Chloro-5'-methoxy-[1,1'-biphenyl]-2-amine (3t). Brown oil (0.46 g, 87% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.46 – 7.35 (m, 1H), 7.26 – 7.19 (m, 1H), 7.09 (dd, J = 7.5, 1.5
Hz, 1H), 6.95 – 6.78 (m, 4H), 3.83 (s, 3H), 3.62 (s, 2H). ¹³C NMR (101 MHz, CDCl₃) δ
158.70, 143.81, 138.86, 130.69, 130.41, 129.30, 125.55, 125.25, 118.45, 116.77, 115.69,
115.36, 55.73. LCMS Rt 3.35 min, m/z 234.1 [M + H]⁺.

437

^{5.1.8 2&#}x27;,5'-Dichloro-[1,1'-biphenyl]-2-amine (3u). Colourless oil (0.43 g, 80% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.46 (d, J = 8.6 Hz, 1H), 7.38 (s, 1H), 7.32 (dd, J = 8.5, 2.6 Hz,
1H), 7.29 – 7.24 (m, 1H), 7.07 (dd, J = 7.6, 1.5 Hz, 1H), 6.89 (ddd, J = 10.7, 8.5, 4.6 Hz, 2H),
4.11 (s, 2H). ¹³C NMR (101 MHz, CDCl₃) δ 143.72, 139.70, 133.05, 132.46, 131.93, 131.12,
130.36, 129.70, 129.22, 124.17, 118.58, 115.83. LCMS Rt 3.49 min, *m/z* 238.0 [M + H]⁺.

5.1.9 2'-Chloro-5'-fluoro-[1,1'-biphenyl]-2-amine (3v). Yellow oil (0.38 g, 76% yield). ¹H
NMR (400 MHz, CDCl₃) δ ¹H NMR (400 MHz, CDCl₃) δ 7.46 – 7.35 (m, 3H), 7.34 – 7.28
(m, 1H), 7.16 (td, J = 8.2, 6.3 Hz, 1H), 6.63 (dd, J = 11.1, 8.5 Hz, 2H), 5.07 – 3.98 (m, 2H).
¹³C NMR (101 MHz, CDCl₃) δ 162.72, 160.26, 143.69, 139.91, 139.83, 131.31, 131.23,
130.32, 129.64, 129.02, 124.38, 118.98, 118.76, 118.53, 116.36, 116.13, 115.82. LCMS Rt
3.88 min, m/z 222.1 [M + H]⁺.

450

5.1.10 2'-Chloro-3-methyl-[1,1'-biphenyl]-2-amine (3w). Brown oil (0.32 g, 79% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.59 – 7.48 (m, 1H), 7.36 (q, J = 2.9 Hz, 3H), 7.16 (dd, J = 7.4,
0.6 Hz, 1H), 6.98 (dd, J = 7.5, 1.3 Hz, 1H), 6.85 (t, J = 7.5 Hz, 1H), 4.03 (s, 2H), 2.30 (s,
3H). ¹³C NMR (101 MHz, CDCl₃) δ 142.05, 138.37, 134.14, 132.11, 130.36, 130.06, 129.13,
128.28, 127.33, 125.13, 122.59, 117.96, 17.95. LCMS Rt 3.44 min, *m/z* 218.1 [M + H]⁺.

5.1.11 2'-Chloro-3-fluoro-[1,1'-biphenyl]-2-amine (3x). Brown oil (0.31 g, 69% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.57 – 7.50 (m, 1H), 7.38 – 7.32 (m, 3H), 7.15 (dd, J = 7.4, 0.8
Hz, 1H), 6.97 (dd, J = 7.6, 1.1 Hz, 1H), 6.84 (t, J = 7.5 Hz, 1H), 3.87 (s, 2H). ¹³C NMR (101
MHz, CDCl₃) δ 161.82, 159.40, 145.77, 145.72, 134.89, 134.44, 130.50, 130.32, 129.74,
129.63, 128.64, 128.24, 114.18, 110.99, 110.97, 105.30, 105.07. LCMS Rt 3.40 min, *m/z*222.1 [M + H]⁺.

463

5.1.12 2',6-Difluoro-[1,1'-biphenyl]-2-amine (3y). Brown solid (0.36 g, 85% yield). ¹H NMR
(400 MHz, CDCl₃) δ 7.46 – 7.37 (m, 2H), 7.27 – 7.14 (m, 3H), 6.66 – 6.59 (m, 2H), 4.12 (s,
2H). ¹³C NMR (101 MHz, CDCl₃) δ 162.23, 161.67, 159.80, 159.20, 146.27, 146.22, 132.47,
132.44, 130.38, 130.30, 130.06, 129.95, 124.65, 124.61, 120.03, 119.86, 116.47, 116.25,
110.99, 110.97, 109.29, 109.09, 105.16, 104.93. LCMS Rt 3.28 min, *m/z* 206.1 [M + H]⁺.

5.1.13 2'-Chloro-6-fluoro-[1,1'-biphenyl]-2-amine (3z).Brown oil (0.27 g, 77% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.58 – 7.50 (m, 1H), 7.37 (t, J = 4.0 Hz, 3H), 7.07 (ddd, J = 10.8,
8.1, 1.3 Hz, 1H), 6.89 (d, J = 7.3 Hz, 1H), 6.80 (td, J = 7.9, 5.2 Hz, 1H), 3.56 (s, 2H). ¹³C
NMR (101 MHz, CDCl₃) δ 153.01, 150.63, 136.86, 136.83, 133.89, 132.84, 132.72, 131.89,
130.16, 129.53, 127.41, 127.31, 127.27, 125.81, 125.78, 117.67, 117.59, 114.83, 114.64.
LCMS Rt 3.41 min, m/z 222.1 [M + H]⁺.

476

5.1.14 2',3-Dichloro-[1,1'-biphenyl]-2-amine (3aa). Brown solid (0.28 g, 75% yield). ¹H
NMR (400 MHz, CDCl₃) δ 7.45 (d, J = 8.6 Hz, 1H), 7.39 (d, J = 2.5 Hz, 1H), 7.34 – 7.23 (m,
2H), 7.08 (dd, J = 7.6, 1.5 Hz, 1H), 6.91 (dd, J = 13.8, 7.5 Hz, 2H), 4.48 (s, 2H). ¹³C NMR
(101 MHz, CDCl₃) δ 143.73, 139.69, 133.02, 132.43, 131.91, 131.10, 130.34, 129.68,
129.19, 124.12, 118.53, 115.80. LCMS Rt 3.47 min, *m/z* 238.0 [M + H]⁺.

482

483 5.2.1 General Method B: Synthesis of Cyclic Diphenyleniodonium Trifluoromethanesulfonate
484 Derivatives 5a-5aa.

The preparation was performed according to the literature procedure.[24] To a stirred 485 solution of biphenylamine 3a-3aa (1.0 eq.) in THF (0.1 M) was added 4 M HCl (3 mL), and 486 the solution was cooled in an ice water bath. A solution of NaNO₂ (1.2 eq.) in H₂O (3 mL) 487 was added dropwise. After 20 min, a solution of KI (2.5 eq.) in H₂O (3 mL) was added, and 488 stirred for 10 min in an ice water bath. Then the solution was slowly warmed to room 489 temperature and stirred for 1 h before an aqueous solution of 20% Na₂S₂O₃ was added until 490 491 the colour of the mixture didn't change. The phases were separated, and the aqueous phase extracted with EtOAc (15 mL x 3). Then the combined organic layers were washed with H₂O 492 and brine, dried over MgSO₄, and concentrated. The residue was purified by flash column 493

494 chromatography using 0-5% EtOAc/petroleum benzine to give the biphenyliodide 4. These495 compounds were used directly in the next step.

To a stirred solution of biphenyliodide **4** (1.0 eq.) in anhydrous CH_2Cl_2 (0.2 M) was added *m*-497 CPBA (75%, 1.5 eq.), TfOH (0.003 eq.). The solution was stirred for 1 h at room 498 temperature. The solvent was removed by rotary evaporation and Et_2O (2 mL) was added to 499 the remained solid. The mixture was stirred for 20 min, and then filtered. The obtained solid 500 was washed with Et_2O (3x), dried in a vacuum oven to afford the desired cyclic 501 diphenyleniodonium trifluoromethanesulfonate **5a-5aa**.

502

503 5.2.2 The following compounds have all been reported previously and their ¹H spectra data
504 showed good agreement with the literature data:

505	1-Methyldibenzo[b,d]iodol-5-ium	trifluorometh	anesulfonate	(5a)[55],	2-
506	Methoxydibenzo[b,d]iodol-5-ium	trifluorometh	anesulfonate	(5k)[24]	3-
507	Methoxydibenzo[b,d]iodol-5-ium	trifluorometha	anesulfonate	(5k)[24],	3-
508	Cyanodibenzo[<i>b</i> , <i>d</i>]iodol-5-ium	trifluoromethar	nesulfonate	(5l)[24],	3-
509	Chlorodibenzo[<i>b</i> , <i>d</i>]iodol-5-ium	trifluoromethan	esulfonate	(5m)[55],	3-
510	Fluorodibenzo[b,d]iodol-5-ium trifle	uoromethanesulfon	ate (5n)[55], Dib	benzo[b,d]iodol-:	5-ium
511	trifluoromethanesulfonate	(50) [24],	Benzo[b]naphtl	ho[1,2-d]iodol-1	l-ium
512	trifluoromethanesulfonate (5p)[24]				

513

5.2.3 *1-Methoxydibenzo[b,d]iodol-5-ium trifluoromethanesulfonate* (5b). Off-white solid
(0.05 g, 22% yield). ¹H NMR (400 MHz, DMSO) δ 8.83 (dd, J = 8.1, 1.4 Hz, 1H), 8.28 (dd, J
= 8.2, 1.1 Hz, 1H), 7.91 – 7.88 (m, 1H), 7.86 – 7.82 (m, 1H), 7.74 – 7.60 (m, 2H), 7.58 – 7.50
(m, 1H), 4.11 (s, 3H). ¹³C NMR (101 MHz, DMSO) δ 160.26, 141.91, 131.48, 131.30,
130.96, 130.76, 130.34, 130.07, 122.81, 122.49, 120.56, 114.16, 57.10. LCMS Rt 2.89 min,

519 *m/z* 309.0 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₁₀IO [M- OTf]⁺, 308.9771, found
520 308.9771. HPLC purity >95%, Rt 4.59 min.

521

522 5.2.4 1-Chlorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5c). Colourless solid (0.3
g, 87% yield). ¹H NMR (400 MHz, DMSO) δ 9.14 (d, J = 7.6 Hz, 1H), 8.30 (t, J = 7.6 Hz,
24 2H), 7.98 (d, J = 7.8 Hz, 1H), 7.91 (t, J = 7.4 Hz, 1H), 7.80 (t, J = 7.3 Hz, 1H), 7.66 (t, J =
8.0 Hz, 1H). ¹³C NMR (101 MHz, DMSO) δ 140.74, 137.85, 134.34, 133.35, 131.92, 131.21,
131.09, 131.01, 130.92, 130.38, 123.50, 121.66. LCMS Rt 2.99 min, *m/z* 312.9 [M - OTf]⁺.
HRMS (ESI) calcd for C₁₂H₇CII [M- OTf]⁺, 312.9275, found 312.9276. HPLC purity >95%,
Rt 4.86 min.

529

5.2.5 1-Fluorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5d). Colourless solid (0.2
g, 66% yield). ¹H NMR (400 MHz, DMSO) δ 8.42 (d, J = 7.9 Hz, 1H), 8.25 (dd, J = 8.2, 0.7
Hz, 1H), 8.10 (dd, J = 8.0, 0.7 Hz, 1H), 7.89 (t, J = 7.6 Hz, 1H), 7.83 – 7.68 (m, 3H). ¹³C
NMR (101 MHz, DMSO) δ 160.05, 139.50, 139.45, 132.09, 132.00, 131.59, 131.54, 131.08,
130.51, 130.29, 130.12, 127.41, 122.03, 121.28, 118.89, 118.68. LCMS rt 2.81 min, *m/z*297.0 [M - OTf]⁺. HRMS (ESI) calcd for C₁₂H₇FI [M- OTf]⁺, 296.9571, found 296.9570.
HPLC purity >95%, Rt 4.53 min.

537

5.2.6 2-Methyldibenzo[b,d]iodol-5-ium trifluromethanesulfonate (5e). Off-white solid (0.2 g,
71% yield). ¹H NMR (400 MHz, DMSO) δ 8.46 (d, J = 7.5 Hz, 1H), 8.34 (s, 1H), 8.21 (d, J =
7.9 Hz, 1H), 8.07 (d, J = 8.2 Hz, 1H), 7.86 (t, J = 7.0 Hz, 1H), 7.71 (t, J = 7.4 Hz, 1H), 7.55
(d, J = 8.2 Hz, 1H), 2.52 (s, 3H). ¹³C NMR (101 MHz, DMSO) δ 142.19, 142.15, 141.43,
132.53, 131.48, 131.17, 131.08, 130.62, 127.82, 127.37, 122.16, 118.42, 21.22. LCMS rt 2.88

543 min, m/z 293.0 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₁₀I [M- OTf]⁺, 292.9822, found 544 292.9821. HPLC purity >95%, Rt 4.61 min.

545

5.2.7 2-Cyanodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5g). Off-white solid (0.04
g, 39% yield). ¹H NMR (400 MHz, DMSO) δ 9.10 (s, 1H), 8.63 (s, 1H), 8.41 (s, 1H), 8.19 (d,
J = 46.2 Hz, 2H), 7.86 (d, J = 51.6 Hz, 2H). ¹³C NMR (101 MHz, DMSO) δ 143.53, 140.76,
133.69, 132.51, 132.25, 131.37, 131.01, 128.10, 127.31, 123.01, 118.26, 114.26. LCMS rt
2.70 min, *m*/z 303.9 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₇IN [M- OTf]⁺, 303.9618, found
303.9618. HPLC purity >95%, Rt 4.14 min.

552

5.2.8 2-Chlorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5h). Off-white solid (0.13
g, 69% yield). ¹H NMR (400 MHz, DMSO) δ 8.68 (d, J = 2.2 Hz, 1H), 8.61 – 8.57 (m, 1H),
8.21 (dd, J = 15.0, 8.3 Hz, 2H), 7.88 (t, J = 7.2 Hz, 1H), 7.81 – 7.72 (m, 2H). ¹³C NMR (101
MHz, DMSO) δ 144.32, 141.02, 136.87, 132.53, 132.18, 131.26, 131.12, 131.03, 128.03,
127.17, 122.70, 119.99. LCMS rt 2.91 min, *m/z* 312.9 [M - OTf]⁺. HRMS (ESI) calcd for
C₁₂H₇CII [M- OTf]⁺, 312.9275, found 312.9276. HPLC purity >95%, Rt 4.82 min.

559

560 5.2.9 2-Fluorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5i). Off-white solid (0.07 561 g, 56% yield). ¹H NMR (400 MHz, DMSO) δ 8.52 (ddd, J = 12.7, 8.9, 2.0 Hz, 2H), 8.27 – 562 8.19 (m, 2H), 7.92 – 7.84 (m, 1H), 7.78 – 7.71 (m, 1H), 7.62 (td, J = 8.8, 2.8 Hz, 1H). ¹³C 563 NMR (101 MHz, DMSO) δ 165.72, 163.79, 163.26, 144.88, 144.79, 141.28, 141.25 ($J_{CF} =$ 564 3.1 Hz), 133.02, 132.93, 132.13, 131.26, 131.08, 127.98, 122.70, 119.10, 118.86, 116.05, 565 116.03, 114.59, 114.34. LCMS rt 2.81 min, m/z 297.0 [M - OTf]⁺. HRMS (ESI) calcd for 566 $C_{12}H_7FI$ [M-OTf]⁺, 296.9571, found 296.9570. HPLC purity >95%, Rt 4.39 min.

568	5.2.10 3-Methyldibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5j). Off-white solid (0.09
569	g, 67% yield). ¹ H NMR (400 MHz, DMSO) δ 8.43 (dd, J = 7.9, 1.2 Hz, 1H), 8.36 (d, J = 8.1
570	Hz, 1H), 8.19 (dd, <i>J</i> = 8.2, 0.8 Hz, 1H), 7.99 (s, 1H), 7.87 – 7.81 (m, 1H), 7.69 (dd, <i>J</i> = 11.1,
571	4.3 Hz, 2H), 2.50 (s, 3H). ¹³ C NMR (101 MHz, DMSO) δ 142.21, 142.03, 139.63, 132.17,
572	131.16, 131.13, 131.02, 130.74, 127.16, 127.10, 122.14, 121.79, 21.68. LCMS Rt 2.89 min,
573	m/z 293.0 [M - OTf] ⁺ . HRMS (ESI) calcd for C ₁₃ H ₁₀ I [M- OTf] ⁺ , 292.9822, found 292.9821.
574	HPLC purity >95%, Rt 4.70 min.

575

576 <u>5.2.1.1 DPI Series 2 analogues</u>

577

5.2.1.2 1-(Trifluoromethyl)dibenzo[b,d]iodol-5-ium trifluoromethanesulfonate 578 (**5q**). Colourless solid (0.11 g, 77% yield). ¹H NMR (400 MHz, DMSO) δ 8.63 (d, J = 8.2 Hz, 1H), 579 8.50 (d, J = 8.3 Hz, 1H), 8.42 - 8.34 (m, 2H), 7.99 (ddd, J = 8.5, 7.3, 1.3 Hz, 1H), 7.92 (t, J = 580 8.0 Hz, 1H), 7.86 – 7.79 (m, 1H). ¹³C NMR (101 MHz, DMSO) δ 139.44, 139.19, 135.83, 581 132.10, 131.50, 130.83, 130.26, 130.20, 130.13, 124.90, 121.99. LCMS Rt 2.89 min, m/z 582 347.0 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₇F₃I [M- OTf]⁺, 346.9539, found 346.9541. 583 HPLC purity >95%, Rt 5.11 min. 584

585

5.2.1.3 1-(2,2,2-Trifluoroacetyl)dibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5r).
Colourless solid (0.12 g, 69% yield). ¹H NMR (400 MHz, DMSO) δ 8.52 (d, J = 8.1 Hz, 1H),
8.33 (d, J = 8.2 Hz, 2H), 7.97 (dd, J = 12.0, 4.6 Hz, 2H), 7.82 (ddd, J = 8.5, 3.6, 2.0 Hz, 2H).
¹³C NMR (101 MHz, DMSO) δ 147.67, 139.48, 134.51, 131.96, 131.76, 131.58, 131.31,
130.44, 130.16, 125.94, 124.48, 124.08, 123.04, 122.73, 121.89, 121.84, 119.53, 119.31,
116.73, 116.33, 79.75, 79.42, 79.09, 31.13. LCMS Rt 2.92 min, *m/z* 362.9 [M - OTf]⁺.

592 HRMS (ESI) calcd for $C_{13}H_7F_3IO [M -OTf]^+$, 362.9488, found 362.9490. HPLC purity 593 >95%, Rt 5.31 min.

594

5.2.1.4 1-Chloro-4-methyldibenzo[b,d]iodol-5-ium trifluoromethanesulfonate 595 (5s).Colourless solid (0.14 g, 78% yield). ¹H NMR (400 MHz, DMSO) δ 9.15 (dd, J = 8.2, 1.4596 Hz, 1H), 8.45 (dd, J = 8.3, 1.0 Hz, 1H), 7.97 – 7.89 (m, 2H), 7.87 – 7.76 (m, 1H), 7.53 (d, J = 597 8.2 Hz, 1H), 2.72 (s, 3H). ¹³C NMR (101 MHz, DMSO) δ 141.78, 139.12, 137.52, 134.42, 598 131.98, 131.81, 131.31, 131.28, 131.21, 130.47, 128.25, 121.35, 25.70. LCMS Rt 2.97 min, 599 m/z 326.9 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₉ClI [M- OTf]⁺, 326.9432, found 600 326.9433. HPLC purity >95%, Rt 5.04 min. 601

602

1-Chloro-4-methoxydibenzo[b,d]iodol-5-ium trifluoromethanesulfonate 603 5.2.1.5 (5t).Colourless solid (0.18 g, 67% yield). ¹H NMR (400 MHz, DMSO) δ 9.08 (d, J = 8.1 Hz, 1H), 604 8.42 (d, J = 8.3 Hz, 1H), 7.93 (dd, J = 8.1, 4.6 Hz, 2H), 7.80 (t, J = 7.8 Hz, 1H), 7.36 (d, J = 605 8.8 Hz, 1H), 4.09 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ 161.18, 146.20, 143.05, 140.51, 606 136.81, 136.69, 135.91, 135.60, 128.92, 127.50, 126.54, 124.30, 117.41, 63.08. LCMS Rt 607 2.87 min, m/z 342.9 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₉ClIO [M- OTf]⁺, 342.9381, 608 found 342.9382. HPLC purity >95%, Rt 5.07 min. 609

610

5.2.1.6 1-Chloro-4-fluorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5u). Colourless
solid (0.2 g, 72% yield). ¹H NMR (400 MHz, DMSO) δ 9.16 (dd, J = 8.2, 1.4 Hz, 1H), 8.41
(dd, J = 8.3, 1.0 Hz, 1H), 8.04 (dd, J = 8.9, 5.1 Hz, 1H), 7.99 – 7.94 (m, 1H), 7.90 – 7.81 (m,
1H), 7.66 (dd, J = 8.9, 7.2 Hz, 1H). ¹³C NMR (101 MHz, DMSO) δ 160.83, 158.37, 140.48,
139.97, 139.93, 136.41, 136.34, 132.52, 131.80, 131.32, 131.11, 128.36, 128.33, 125.95,
122.74, 122.55, 119.54, 117.32, 117.10, 111.26, 110.97. LCMS Rt 2.87 min, *m/z* 330.9 [M -

- 617 $OTf]^+$. HRMS (ESI) calcd for $C_{12}H_6ClFI$ [M- $OTf]^+$, 330.9181, found 330.9181. HPLC purity 618 >95%, Rt 4.60 min.
- 619

5.2.1.7 1,4-Dichlorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5ν). Colourless solid (0.11 g, 71% yield). ¹H NMR (400 MHz, DMSO) δ 9.14 (dd, J = 8.2, 1.4 Hz, 1H), 8.46 (dd, J = 8.2, 1.1 Hz, 1H), 7.98 (ddd, J = 11.2, 9.2, 4.9 Hz, 2H), 7.91 – 7.76 (m, 2H). ¹³C NMR (101 MHz, DMSO) δ 141.60, 139.14, 135.90, 132.52, 132.48, 131.89, 131.71, 131.60, 131.46, 129.73, 128.01, 122.64. LCMS Rt 2.76 min, m/z 346.9 [M - OTf]⁺. HRMS (ESI) calcd for C₁₂H₆Cl₂I [M- OTf]⁺, 346.8886, found 346.8889. HPLC purity >95%, Rt 4.87 min.

626

5.2.1.8 6-Chloro-1-methyldibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5w).627 Colourless solid (0.25 g, 69% yield). ¹H NMR (400 MHz, DMSO) δ 8.90 (d, J = 7.6 Hz, 1H), 628 8.43 (d, J = 8.0 Hz, 1H), 7.99 (d, J = 7.6 Hz, 1H), 7.80 (t, J = 7.5 Hz, 1H), 7.66 (d, J = 6.2 629 Hz, 2H), 2.72 (s, 3H). ¹³C NMR (101 MHz, DMSO) δ 140.98, 139.78, 138.64, 134.66, 630 133.53, 132.18, 131.36, 131.05, 130.79, 128.55, 126.22, 122.92, 26.14. LCMS Rt 2.78 min, 631 m/z 327.0 [M - OTf]⁺. HRMS (ESI) calcd for C₁₃H₉ClI [M- OTf]⁺, 326.9432, found 632 326.9433. HPLC purity >95%, Rt 5.02 min. 633

634

5.2.1.9 6-Chloro-1-fluorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (**5***x*). Colourless solid (0.12 g, 69% yield). ¹H NMR (400 MHz, DMSO) δ 8.38 (t, J = 2.1 Hz, 1H), 8.26 (d, J =8.8 Hz, 1H), 8.13 (dd, J = 8.0, 1.0 Hz, 1H), 7.89 – 7.72 (m, 3H). ¹³C NMR (101 MHz, DMSO) δ 160.23, 141.35, 136.71, 132.87, 132.78, 132.60, 131.16, 129.37, 129.19, 127.44, 122.76, 119.48, 118.99, 118.78. LCMS Rt 2.93 min, m/z 330.9 [M - OTf]⁺. HRMS (ESI) calcd for C₁₂H₆CIFI⁺ [M- OTf]⁺, 330.9181, found 330.9182. HPLC purity >95%, Rt 5.15 min.

5.2.1.10 1,6-Dichlorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5y). Colourless
solid (0.09 g, 65% yield). ¹H NMR (400 MHz, DMSO) δ 9.17 – 9.05 (m, 1H), 8.48 (dd, J =
8.4, 1.0 Hz, 1H), 8.07 (dt, J = 24.6, 12.3 Hz, 1H), 8.02 – 7.91 (m, 2H), 7.73 (t, J = 8.2 Hz,
1H). ¹³C NMR (101 MHz, DMSO) δ 142.60, 138.43, 134.68, 134.16, 133.36, 133.11, 131.33,
131.11, 130.57, 129.31, 125.55, 124.31, 122.75, 119.54. LCMS Rt 2.95 min, *m/z* 346.9 [M OTf]⁺. HRMS (ESI) calcd for C₁₂H₆Cl₂I [M- OTf]⁺, 346.8886, found 346.8887. HPLC purity

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650

>95%, Rt 4.87 min.

5.2.1.11 1-Chloro-9-fluorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5z).651 Colourless solid (0.1 g, 70% yield). ¹H NMR (400 MHz, DMSO) δ 8.25 (dd, J = 8.1, 0.9 Hz, 652 1H), 8.16 (dd, J = 7.7, 1.1 Hz, 1H), 8.01 (dd, J = 8.0, 0.9 Hz, 1H), 7.91 – 7.68 (m, 3H). ¹³C 653 NMR (101 MHz, DMSO) δ 161.52, 158.91, 137.37, 137.31, 134.20, 134.01, 133.28, 133.20, 654 132.18, 129.78, 128.71, 128.55, 127.31, 122.77, 121.78, 121.74, 119.79, 119.54. LCMS Rt 655 2.93 min, m/z 330.9 [M - OTf]⁺. HRMS (ESI) calcd for C₁₂H₆ClFI [M- OTf]⁺, 330.9181, 656 found 330.9182. HPLC purity >95%, Rt 4.96 min. 657

658

5.2.1.12 1,9-Difluorodibenzo[b,d]iodol-5-ium trifluoromethanesulfonate (5aa). Off-white
solid (0.12 g, 66% yield). ¹H NMR (400 MHz, DMSO) δ 8.15 (dd, J = 7.6, 1.4 Hz, 2H), 7.89
- 7.71 (m, 4H). ¹³C NMR (101 MHz, DMSO) δ 161.26, 158.67, 133.07, 127.36, 122.74,
121.66, 119.80, 119.67, 119.54. LCMS Rt 2.89 min, *m/z* 314.9 [M - OTf]⁺. HRMS (ESI)
calcd for C₁₂H₆F₂I [M- OTf]⁺, 314.9477, found 314.9477. HPLC purity >95%, Rt 4.61 min.

665 5.3 Biological Asssay

666 *5.3.1 Organisms*

Pseudomonas aeruginosa, Klebsiella pneumoniae, Escherichia coli, Enterococcus faecalis, Enterococcus faecium, Staphylococcus aureus, Acinetobacter baumannii and 668 Mycobacterium tuberculosis H37Rv strains employed in this study were clinical isolates or 669 obtained from the American Type Culture Collection (Rockville, MD, USA). Bacterial 670 isolates (excluding *M. tuberculosis*) were stored in tryptone soy broth (Oxoid) with 20% 671 glycerol (Ajax Finechem, Seven Hills, NSW, Australia) at -80 °C. M. tuberculosis H37Rv 672 (ATCC 25618) was propagated in 37 °C in 7H9 media (BD Diagnostic Systems, Sparks, MD, 673 USA) supplemented with 10% albumin-dextrose-catalase (ADC), 0.5% glycerol and 0.02% 674 tyloxapol. Isolates were stored at -80 °C in the same media with 30% glycerol. 675

P. falciparum (D10-PfPHG[56], CS2-PHG[57]) and P. knowlesi (YH1[28]) parasites were 676 cultured in human O⁺ erythrocytes according to the method of Trager and Jensen.[58] 677 Briefly, parasites were grown in RPMI-HEPES culture medium (pH 7.4) supplemented with 678 50 µM hypoxanthine, 25 mM NaHCO₃, 20 µM gentamicin and 0.5% Albumax II (Gibco, 679 Melbourne, VIC, Australia). Cultures were maintained in airtight boxes in a 37 °C incubator 680 in an atmosphere of 1% O₂, 4% CO₂ and 95% N₂. 681

682

5.3.2 Gram-positive and Gram-negative bacterial panel assay 683

The MICs of test compounds against P. aeuruginosa, K. pneumoniae, E. coli, E. 684 faecalis, E. faecium, S. aureus and A. baumannii strains were determined by the broth 685 microdilution method according to the guidelines of the Clinical and Laboratory Standards 686 Institute [25]. Experiments were performed with Cation-Adjusted Mueller-Hinton Broth 687 (CaMHB) in 96-well polypropylene microtitre plates. Wells were inoculated with 100 µL of 688 bacterial suspension prepared in CaMHB (containing $\sim 10^6$ colony forming units (cfu) per 689 mL) and 100 µL of CaMHB containing increasing concentrations of DPI (0 to 128 µM). The 690 MICs were defined as the lowest concentration at which visible growth was inhibited 691

692 following 18 h incubation at 37 °C. The IC_{50} of the antibiotic polymixin B (Sigma) was 693 assayed as a growth inhibitory positive control.

694

695 5.3.3 Mycobacterium tuberculosis growth inhibition

For MIC determination compounds were serially diluted in 96 well tissue culture 696 plates in 10 µL of purified H₂O in triplicate (0 to 10 µM). M. tuberculosis grown in 697 supplemented 7H9 media (90 µL) was adjusted to an OD_{600nm} of 0.001, added to wells and 698 incubated for 7 days at 37 °C. Resazurin (10 µL; 0.05% w/v; Sigma-Aldrich, Australia) was 699 then added, incubated for 4-24 h at 37 °C and fluorescence measured at 590 nm using a 700 FLUOstar Omega microplate reader (BMG Labtech, Germany). The MICs were defined as 701 702 the lowest concentration at which bacterial growth was completely inhibited compared to non-treated bacteria. The IC₅₀ of the antibiotic rifampicin (Sigma) was assayed as a growth 703 inhibitory positive control. 704

705

706 5.3.4 Plasmodium spp. growth inhibition assays

Parasites were synchronized to early ring stages using a combination of heparin 707 synchronization [59] and sorbitol lysis. Malaria growth inhibition assays using ring stage 708 parasites were setup in a 96-well round bottom plate as described previously [56] at 1% 709 parasitaemia and 1% haematocrit in a final volume of 45 µL. A 10x final concentration of 710 DPI and controls were added to make the final volume to 50 µL. P. falciparum assays were 711 cultured for 90 h, through the next cycle of replication, until the parasites were mainly mature 712 trophozoites (36-42 h post-invasion). P. knowlesi assays were cultured for 50 h, again until 713 the parasites were late trophozoites of the next growth cycle. Assays were stained with 10 714 µg/mL ethidium bromide (EtBr, Bio-Rad, Melbourne, VIC, Australia) for 1 h and then 715 washed prior to flow cytometry (Becton Dickinson LSR) assessment of parasitaemia. GFP 716

and non-GFP fluorescent parasites were gated and counted according to established protocols [60]. Flow cytometry data was analysed using FlowJo software (Tree Star, St, Ashland, OR, USA). The IC₅₀ of parasite growth inhibition was determined using Graphpad PRISM (Graphpad Software, La Jolla, CA, USA) following the recommended protocol for non-linear regression of a log(inhibitor) *vs*. response curve [60]. The IC₅₀ of the antimalarial chloroquine (Sigma) was assayed as a growth inhibitory positive control.

723

724 5.3.5 P. falciparum resistance selection and sequencing

The central 1567bp (18bp-1585bp) of the *Pf*NDH2 sequence was PCR amplified from 725 the D10-PfPHG parental, D10-DPI^r and D10-DPI^{off} lines using the primers PfNDH2 F 726 GGTTAATATATAATGTTAGTAAAGTTCAGG and *Pf*NDH2 R 727 CATTTTTTTTTTTTTGATGAAAGGAC. The sequence of Type II Pf dihydroorotate 728 dehydrogenase was amplified from the target lines using the primers PfDHOD F 729 GTGTGATAGATAGCTCCAGTCG and PfDHOD R GCACTTATGTGTCGCCCG. Sanger 730 sequencing of the resulting PCR products was conducted at the Australian Genome Research 731 Facility, with alignments compared using Geneious (Biomatters, New Zealand). 732

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734 5.3.6 Cytotoxicity Assay

HepG2 cells (ATCC HB-8065) were seeded as 4000 cells per well in a 384-well plate in DMEM medium (GIBCO-Invitrogen #11995-073), with 1% FBS or 10% FBS as specified. Cells were incubated for 24 h at 37 °C, 5% CO₂ to allow cells to attach to the plates. Compounds were added into each well with a series of concentrations from 300 μ M to 0.14 μ M in 3-fold dilution, the cells were then incubated for 24 h at 37 °C, 5% CO₂. After the incubation, 10 μ M resazurin (dissolved in PBS) was added to each well. The plates were then incubated for 2 h at 37 °C, 5% CO₂. The fluorescence intensity was read using a Polarstar

742	Omega plate reader with excitation/emission 560/590. The data was analysed using Prism			
743	software. Results are presented as the average percentage of control \pm SD for each set of			
744	duplicate wells using the following equation: Percentage Viability = ($FI_{Compound}$ –			
745	FI _{Negative} /FI _{Untreated} –FI _{Negative}) x 100.			
746				
747	Conflict of interest			
748	The authors declare they have no conflict of interest.			
749				
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755				
756	Appendix A. Supplementary data			
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Highlights

- A series of compounds based on the NDH-2 inhibitor diphenyleneiodonium (DPI) were synthesised.
- Compound **5s** and **5g** exhibited low nanomolar activity against *M. tuberculosis and P.*

falciparum respectively.