

RESEARCH ARTICLE

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"Affect of anaerobiosis on the antibiotic susceptibility of *H. influenzae*"

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Abstract

Background: *Haemophilus influenzae* is a human-restricted facultative anaerobe which resides mostly in the oropharynx. The majority of isolates recovered from the throat are unencapsulated commensals (NTHi), but depending on host susceptibility they cause bronchitis, otitis media and on occasion bacteremia and meningitis. Because of the variable oxygen availability in the various niche permitting bacterium replication, the organism must thrive in well oxygenated surfaces, such as pharyngeal epithelium to anoxic environments like the bottom of a Biofilm and in airway mucus. Other reports indicate that *H. influenzae* use aerobic respiration, anaerobic respiration and fermentation to generate ATP. To gain insight in to the activity of several classes of antibiotics against five well-characterized unencapsulated *H. influenzae* in room air, in 5% CO₂ and under strict anaerobiosis. We also tested for the role of oxidative killing by all cidal antibiotics.

Results: In comparison to room air, testing in 5% CO₂ had minimal effects on the susceptibility to aminoglycosides, cephalosporins, tetracycline and chloramphenicol: the MIC of rifampin and ciprofloxacin increased eight fold with certain strains in 5% CO₂. All antibiotics, except trimethoprim were cidal under both growth conditions. Aminoglycosides remained bactericidal in a strict anaerobic environment, while a reliable MBC was obtained with trimethoprim only under anaerobic conditions. Kinetic analysis of the cidal action of spectinomycin and tetracycline indicated slower killing anaerobically. An oxidative mechanism for aerobic killing could not be demonstrated.

Conclusions: We conclude that β-lactams, cephalosporins, macrolides, tetracycline's, aminoglycosides, chloramphenicol, rifampin and ciprofloxacin are bactericidal against five well-characterizes *H. influenzae* in an aerobic and anaerobic environment. The activity of trimethoprim was increased in anaerobic conditions.

Keywords: *Haemophilus influenzae*, Antibiotic activity, Anaerobiosis

Background

Haemophilus influenzae is classified by Bergey's Manual of Determinative Microbiology (9th Edition, pg 195) as a "nonmotile Gram negative coccobacillus which is a facultative anaerobe having both a respiratory and fermentative type of metabolism" [1]. Multiple studies examining *H. influenzae* gene content or expression under various environmental conditions have suggested that the organism prefers an anaerobic environment, and may use anaerobic respiration and fermentation depending on the local oxygen availability. In well aerated, shaking sBHI broth cultures with *H. influenzae* Rd KW20 incubated at 37°C in room air there is oxygen consumption during

logarithmic growth [2]. The partial pressure of oxygen in the media during logarithmic growth averaged 64 Torr during the 120 min of logarithmic growth but varying from 145 to 53 Torr: 64 Torr is approximately an O₂ concentration of 0.8 mM. With the increased consumption of oxygen, there is a reciprocal increase in the pCO₂; from, < 4 Torr to a peak value of 13.8 Torr at a time when the O₂ concentration was at its nadir [2]. During the lag and stationary phases the media pO₂ ranged from 128 to 145 Torr [2]. During overnight incubation of strain Rd KW20 in room air the pH of sBHI decreases from 7.39 to 5.55 indicating fermentation. In an aerated and shaken culture, as well as incubation in a BD GasPak, genes of the nitrate operon (*napFDAGHBC*) and the nitrite complex (*nrfABCD*) were detected [3] indicating ongoing anaerobic respiration. As the O₂ concentration in the media decrease the high affinity *cydAB*

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is responsible for aerobic respiration. The simultaneous detection of the presence of respiratory and fermentative enzymes in anaerobic and microaerophilic environments indicates that all of the pathways are active in *H. influenzae*.

Others studying *H. influenzae* colonization mechanisms found that strain Eagan and a mutant deficient in MnSOD both grew best at the lowest oxygen supply rate tested, an environment supplying 3 mMol O₂/liter /hr [4]. Although the MnSOD mutant had decreased colonization capacity of the infant rat nasopharynx, its invasive capacity was unchanged, suggesting a preference for anaerobic or micro aerobic lifestyle after invasion. Studies [5] of the anoxic redox control two-component system, ArcAB indicated that the loss of a functional ArcA response regulator decreased the serum resistance of the type b strain. They found that fumarate reductase was more highly expressed in the *arcA* + parent, while L-Lactate dehydrogenase was derepressed in the *arcA* mutant. Unfortunately the details of the techniques to create an anaerobic environment was filling flasks and sealing them making the environment micro aerobic.

Nontypeable *H. influenzae* form biofilms in the middle ear of children with otitis media [6], a finding that has been reproduced in animal models of otitis media [7]. Bacterial metabolism is not uniform in all portions of the Biofilm with the basilar portion being hypoxic. One explanation for the antibiotic resistance of bacterial biofilms is that the basilar portions are anoxic [8]. Thus understanding the response of *H. influenzae* to antibiotics in an anoxic environment will clarify the role of ambient oxygen concentration on antibiotic activity.

It has been recently proposed that all classes of bactericidal antibiotics kill *E. coli* via an oxidative mechanism, but bacteriostatic antibiotics do not elicit an oxidative respiratory burst [9]. It is not certain how a cidal antibiotic increases aerobic respiration, particularly in a bacterium such as *Haemophilus influenzae* which has uses multiple mechanisms to generate ATP depending on the O₂ availability. Since all antibiotics which have been tested, except trimethoprim, appear cidal against *H. influenzae* we determined the activity of multiple antibiotics against a well-defined panel of *H. influenzae* in room air, in 5% CO₂ and in a Coy anaerobic chamber. The antibiotics were chosen to represent different classes and modes of action and were not chosen to inform clinicians. This data should not be used for clinical decisions.

We examined five *H. influenzae* strains whose genome has been sequenced and annotated so that additional insight into antibiotic cidal mechanisms can be explored. We found that all classes of antibiotics were bactericidal and could not detect an oxidative process in bacterial killing.

Results

Using the aerobic and anaerobic techniques described above we first determined the susceptibility of the reference *H. influenzae* strain for testing antibiotic susceptibility, ATCC 49766 to ampicillin, chloramphenicol and tetracycline. With ampicillin the aerobic MIC/MBC was 0.15/5.0 and 1.5/4.0 µg/ml anaerobically, the aerobic chloramphenicol MIC/MBC was 0.2/ 2.6 and the anaerobic MIC/MBC was 0.4/3.0 µg/ml. With tetracycline the MIC/MBC was 0.25/1.0 µg/ml aerobically and 0.30/2.20 µg/ml anaerobically. These results indicated that anaerobic susceptibility testing with NTHi whose genome sequence is available was feasible. Recognizing that the intrinsic error of MIC testing is 100% [10], we emphasized only those results which were more than double, or 50% of the value obtained in room air.

Aerobic susceptibility

The aerobic susceptibility of the sequenced isolates is shown in Table 1 and Table 2 while Table 3 and Table 4 depict the susceptibility in 5% CO₂. The MIC and MBC for the aminoglycosides was slightly increased in the presence of 5% CO₂ (Table 3 and Table 4). With strain Rd KW20 the erythromycin MBC increased six-fold when tested in the presence of 5% CO₂ (compare Table 2, versus Table 4). All tested strains had an increase in the MIC and MBC with ciprofloxacin (Table 2 versus Table 4).

Anaerobic susceptibility

Under anaerobic conditions using sBHI with 0.1% ferric nitrate the MBC for tetracycline increased with all strains except with strain R2866, with the values not significantly different with incubation in either environment. Anaerobic incubation had the most dramatic effect on the susceptibility to trimethoprim. In room air or 5% CO₂ the MBC was difficult to measure accurately: the values reported met the criteria of 99.9% kill, but there was poor reproducibility among the replicates. The MBC value of 1800 µg/ml was the highest trimethoprim concentration tested with the MIC not significantly different except with strain R3932 (anaerobically adapted R3642). Under anaerobic conditions the trimethoprim end-points could easily be identified with all strains except R3392 (anaerobically adapted R3642). In room air the MIC was 450 µg/ml and increased on average one well to 788 µg/ml with the MBC being 1800 µg/ml in room air and 5% CO₂. Current reviews indicate that bacteria grown in an anaerobic environment are resistant to aminoglycosides as they do not have the required respiration to facilitate drug uptake [11]. However, kanamycin and tobramycin maintained their bactericidal activity under anaerobic conditions, with the MIC and MBC values in the clinical susceptibility range for all strains.

Table 1 Aerobic susceptibility in air

H. influenzae	μg/ml													
	Amp		Tet		Spec		Tmp		Kan		Tobr		Cam	
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC
Rd KW20	0.33	0.54	0.29	2.31	22.5	22.5	5.25	225	1.40	1.40	0.70	2.8	0.18	0.26
R2846	0.44	0.50	0.20	2.31	4.22	8.44	1.98	113	0.70	1.40	0.35	1.05	0.26	0.80
R2866	900	>1800	0.29	1.16	16.88	16.88	0.88	113	1.40	1.40	0.70	1.41	0.18	0.70
R3642	675	>1800	0.29	2.31	5.63	14.07	450	1800	1.06	2.11	0.53	0.70	0.14	1.05
R3661	0.33	0.59	0.29	3.72	16.85	16.85	17.58	1800	1.41	2.80	0.70	1.75	0.18	1.05

*produces TEM β-lactamase.

The results of the anaerobic susceptibility testing are shown in Table 5 and Table 6.

Strains R2866 (R3930 for the anaerobic adapted derivative) and R3642 (R3931 for the anaerobically adapted derivative) have a TEM-like β-lactamase and are resistant to ampicillin, and as expected they are susceptible to Cefaclor and cefuroxime (Table 6). Of the two cephalosporin's tested, Cefaclor was the least potent in all of the environments.

Antibiotic cidal activity

Determining the rate of bacterial killing by antibiotics is a sensitive way of testing antibiotic activity. We chose tetracycline and spectinomycin concentrations which were approximately twice the MBC determined with the micro plate assay. With incubation in room air there was killing of 99.9% of the initial inoculum in four hours with each strain with each antibiotic (Figure 1A). Anaerobic killing by tetracycline was faster with strains R3931 (adapted R2846) and R3930 (adapted R2866) at concentrations which were approximately twice the anaerobic MBC, with the 99.9% kill being achieved at 3 to 4 hours (Figure 1B). With strain R3930 (anaerobic R2866) it took 6 hours before 99.9% of the inoculum was killed by 20 μg/ml spectinomycin (Figure 1B).

Oxidative cidal mechanism

We could not detect hydroxyl ion during the killing of strains R2846 or R2866 by tetracycline, kanamycin or

spectinomycin in room air at 37°C with 5 μM 3' (p-hydroxyphenyl) fluorescein (HPF) (Figure 2). As a positive control we conducted a killing curve with 1 mM H₂O₂: there was 99.9% killing of both strains within 1 hour of incubation. We also used 3'(p-aminophenyl) fluorescein and could not detect hydroxyl ion during the cidal activity. The killing experiments were repeated with tetracycline, Kanamycin or spectinomycin in the presence of 0.5 mM 2,2'-dipyridyl, an iron chelator which was added simultaneously with antibiotics: Dipyridyl alone had no affect on the rate of loss of viability eliminating a hydroxyl mediated Fenton reaction, a key step in the Collins hypothesis. Additionally we repeated the killing experiments with the same antibiotics in the presence of 150 mM thiourea which quenches hydroxyl ion as it is formed, and found no effect.

Membrane permeability

The current model for aminoglycosides action is its binding to a site in the ribosome which causes misreading of the mRNA leading to misfolded proteins. These proteins accumulate in the periplasmic space and outer membrane creating channels that permit a larger influx of the aminoglycoside [12]. We found that outer membrane permeability of strain R2866 increased during the cidal action of kanamycin at 1 μg/ml; there was no detectable change in permeability during cell killing with tetracycline at 2 μg/ml (Figure 3). Tetracycline prevents the association of aminoacyl tRNA with the ribosome,

Table 2 Aerobic susceptibility in air

H. influenzae	μg/ml										ng/ml			
	Cfrx		Cclr		Clar		Erm		Rif		Cip			
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC
Rd KW20	0.51	3.00	11.72	5.63	5.63	16.88	0.48	1.93	70.0	70.0	0.71	3.94		
R2846	0.68	1.56	11.72	15.94	0.26	1.41	0.26	0.44	26.5	70.0	2.25	4.50		
R2866	0.72	1.56	15.94	24.37	2.81	5.63	0.26	0.44	70.0	105.5	1.69	3.10		
R3642	1.10	2.81	15.94	30.94	2.81	11.25	0.35	0.97	70.0	105.5	3.38	3.38		
R3661	0.85	4.07	12.19	14.06	4.22	22.50	0.53	1.41	35.0	70.0	1.41	4.50		

Amp ampicillin, Tet tetracycline, Cip ciprofloxacin, Clar clarithromycin, Cfrx cefuroxime, Spec spectinomycin, Cclr Cefaclor, Tmp trimethoprim, Rif rifampin, Kan Kanamycin, Tobr tobramycin, Cam chloramphenicol, Erm erythromycin.

Table 3 Aerobic susceptibility in 5% CO₂

H. influenzae	μg/ml													
	Amp		Tet		Spec		Tmp		Kan		Tobr		Cam	
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC
Rd KW20	1.17	1.41	0.35	5.63	22.5	33.85	9.10	338	4.22	5.63	2.82	4.22	0.27	0.8
R2846	0.44	0.50	0.32	0.94	10.3	18.8	1.10	84.4	1.41	4.22	1.41	2.81	0.27	0.8
R2866	1350	>1800	0.32	2.81	26.3	33.8	0.88	113	1.4	1.4	2.11	2.81	0.27	0.8
R3642	900	>1800	0.25	0.32	5.63	14.1	788	1800	1.41	4.22	1.41	2.11	0.27	1.4
R3661	1.41	2.81	0.29	2.83	18.8	22.5	4.39	1575	2.81	5.63	2.11	4.22	0.35	1.4

which stops protein synthesis [13] and has no immediate effect on the cell membrane. We had previously shown that therapeutic tobramycin concentrations caused mistranslation in aminoglycoside susceptible *H. influenzae* [14].

Discussion

Anaerobic growth

In our prior studies on the anaerobic growth of *H. influenzae* strain Rd KW20 we used a disposable GasPak (Becton Dickson) anaerobic chamber after bubbling N₂ through the media and monitoring anaerobiosis with the GasPak colorimetric indicator. In that study [3] after 24 hours of growth the bacterial density ranged from 1.9 × 10⁹ to 6.5 × 10⁹ CFU/ml in both microaerophilic and "anaerobic" environments. The starting pH of sBHI broth was 7.34 ± 0.12 (n = 9) while after microaerophilic incubation it was 5.97 ± 0.15 (n = 4), after incubation in the anaerobic chamber it was 4.69 ± 0.59 (n = 6). The majority of the proteins detected were present under both growth conditions [3]. There was slightly more acetate, 0.60 mM in the media after anaerobic growth, compared to micro aerobic growth 0.46 mM. This data suggests that *H. influenzae* uses fermentation pathways in both growth environments however the acetate concentration was equivalent with incubation in both environments: 0.46 mM aerobic and 0.60 mM after anaerobic growth [3].

In *H. influenzae* genes, *fnr* and *arcA*, which are transcriptional regulators homologous to *E. coli* genes which

modify expression of other genes involved in growth in environments with varying oxygen availability. Deletion of *arcA* increased the expression of enzymes in the respiratory chain and TCA cycle when grown under micro aerobic conditions [15]. In *H. influenzae* having mechanisms which combat oxidative and nitrosative stress are important for its life cycle [16,17]. In addition, *fnr* is required for defense against nitric oxide donors [18]. *H. influenzae* with inactivated *fnr* were unable to metabolize nitrite added to the media indicating that *nrfABCD* was not functional. In a study of the defense of nontypeable *H. influenzae* against reactive nitrogen species Harrington et al. [18] found that placing 7.5 ml of media in a 60 ml DeLong flask resulted in an oxygen saturation (ie dissolved oxygen concentration) of 80%. This corresponds to an O₂ concentration of 5.7 mg/L or 0.178 mM. This concentration is micro aerobic [19] as the oxygen concentration in well aerated water at 25°C is at 8 mg/L (0.25 mM). However, the micro aerobic condition completely suppressed the expression of a *nrf* fusion [18]. When the Delong flask was filled with 60 ml the oxygen saturation was 38%, 2.6 mg/L or 0.081 mM and the *nrf* fusion was expressed over nine fold greater even though strict anaerobiosis was not achieved.

In silico reviews of *H. influenzae* metabolism concluded that the gene content favored an anaerobic environment [20]. In the study integrating enzyme activity and proteomics, periplasmic nitrate reductase (*napB*) and the cytoplasmic nitrite Reductase (*nrfA*) we found

Table 4 Aerobic susceptibility in 5% CO₂

H. influenzae	ng/ml											
	Cfrx		Cclr		Clar		Erm		Rif		Cip	
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC
Rd KW20	2.11	2.34	15.47	23	18.58	30	3.26	11.69	122.5	140	8.29	10.17
R2846	1.14	4.22	11.7	18.7	0.35	14.10	0.49	0.57	70.0	87.5	8.29	12.04
R2866	1.41	2.81	15.94	24.4	6.57	11.25	3.53	0.57	157	157	16.58	16.58
R3642	1.88	5.63	23.4	30.9	7.5	30.0	0.35	1.41	87.5	105	10.6	16.58
R3661	2.34	5.63	15.9	18.75	11.25	30.0	0.61	1.10	87.5	105	16.58	20.3

Amp ampicillin, Tet tetracycline, Cip ciprofloxacin, Clar clarithromycin, Cfrx cefuroxime, Spec spectinomycin, Cclr Cefaclor, TMP trimethoprim, Rif rifampin, Kan Kanamycin, Tobr tobramycin, Cam chloramphenicol, Erm erythromycin.

Table 5 Anaerobic susceptibility

H. influenzae	μg/ml													
	Amp		Tet		Spec		Tmp		Kan		Tobr		Cam	
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC
R3929	0.53	4.22	0.18	15.9	25.31	26.72	3.52	42.2	1.06	2.11	1.06	2.81	0.53	2.81
R3931	1.41	4.22	0.35	2.11	18.28	45.0	0.88	5.25	3.17	18.7	4.13	6.56	0.53	2.81
R3930	1575	1800	0.27	1.05	2.11	8.44	0.88	84.4	1.23	2.46	1.23	2.81	0.53	2.81
R3932	1350	1800	0.35	2.81	1.46	2.34	562	675	1.29	1.64	1.28	1.87	0.70	2.81
R3933	1.76	4.22	0.28	8.44	9.92	33.75	3.5	28.10	1.99	3.28	1.64	3.75	0.53	2.81

that both enzymes were expressed in anaerobic and micro aerobic growth [2]. However, in the current studies we found that the addition of 10 mM KN₃ or 10 mM NaNO₂ did not increase the growth rate, or the final bacterial density after overnight incubation at 37°C in the anaerobe chamber (3–5 ×10⁹ CFU/ml) of strains R3930 or R3931. In silico reviews of *H. influenzae* metabolism concluded that the gene content favored an anaerobic environment [20]. In the study integrating enzyme activity and proteomics, periplasmic nitrate reductase (*napB*) and the cytoplasmic nitrite Reductase (*nrfA*) we found that both enzymes were expressed in anaerobic and micro aerobic growth [2]. However, in the current studies we found that the addition of 10 mM KN₃ or 10 mM NaNO₂ did not increase the growth rate, or the final bacterial density after overnight incubation at 37°C in the anaerobe chamber (3–5 ×10⁹ CFU/ml) of strains R3930 or R3931.

We previously examined the proteome of *H. influenzae* Rd KW20 in late log phase by liquid chromatography coupled with ion trap tandem mass spectrometry [3] and found that the proteome was nearly identical after growth in a Becton Dickson GasPak or in room air. On obtaining these results we serially measured the partial pressure of O₂ during incubation of 10³ CFU/ml in 50 ml of sBHI at 37°C in a 250 ml baffled Erlenmeyer flask agitating the flask at 200 rpm in room air. During the logarithmic growth in sBHI, at an A_{650 nm} between 0.28 and 0.64, the pO₂ in the media decreased from 125–

145 Torr to 53 Torr and remained at 68 Torr during the log phase; but reaching 135 Torr at on entrance to stationary phase, an A_{650nm} of 1.10. Thus the incubation conditions were microaerophilic for the majority of growth during incubation in room air. Additional data measuring enzyme activity in cell lysates as well as organic acids indicated that there was fumarate reductase activity in cells grown in both anaerobic and micro aerobic conditions [2]. Insertional mutagenesis of *frdA* resulted in restricted growth in micro aerobic and anaerobic environment. All of the data indicates that *H. influenzae* Rd KW20 uses classic anaerobic respiration, micro aerobic respiration and fermentative pathways for growth in environments with differing oxygen availability. By using multiple pathways the bacterium can adapt to the different oxygen supply in the human respiratory tract and blood stream. This may represent “hybrid” metabolism in which membrane ArcB located on the membrane mediates changes in gene expression through ArcA when the oxygen supply is too low to penetrate to the cytoplasm and inactivate the direct oxygen sensor, FNR [21].

The marked increase in the potency of trimethoprim under anaerobic conditions (Table 5) suggests that anaerobic growth is more dependent on tetrahydrofolate than aerobic growth. The proteins which were most highly induced in *H. influenzae* strain Rd KW20 with TMP treatment were 5-methylenetetrahydropteroyltri-gutamate-homocysteine methyl transferase (*metE*) and

Table 6 Anaerobic susceptibility

H. influenzae	μg/ml										ng/ml			
	Cfrx		Cclr		Clar		Erm		Rif		Cip			
	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC	MIC	MBC
R3929	0.35	1.78	22.50	45	8.44	45.0	0.35	1.06	33.8	67.5	3.7	5.98		
R3931	0.68	1.56	11.25	45	11.25	33.75	0.53	1.41	78.8	123.8	7.0	7.14		
R3930	0.48	1.05	16.88	45	2.81	16.88	0.35	0.88	107	315	4.4	7.39		
R3932	1.10	19.69	8.44	45	11.25	33.75	0.35	1.41	14.1	28.1	15.5	28.48		
R3933	1.23	7.51	16.88	45	11.25	33.75	0.35	1.41	56.3	135	6.70	8.79		

Amp ampicillin, Tet tetracycline, Cip ciprofloxacin, Clar clarithromycin, Cfrx cefuroxime, Spec spectinomycin, Cclr Cefaclor, Tmp trimethoprim, Rif rifampin, Kan Kanamycin, Tobr tobramycin, Cam chloramphenicol, Erm erythromycin.

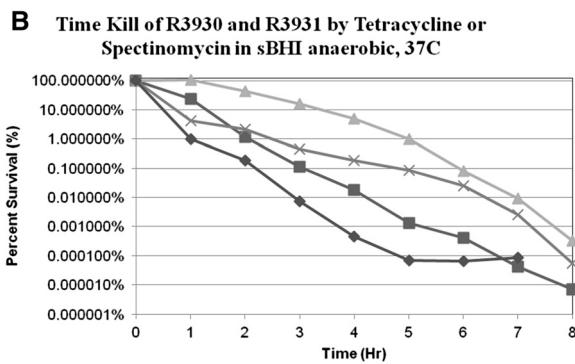
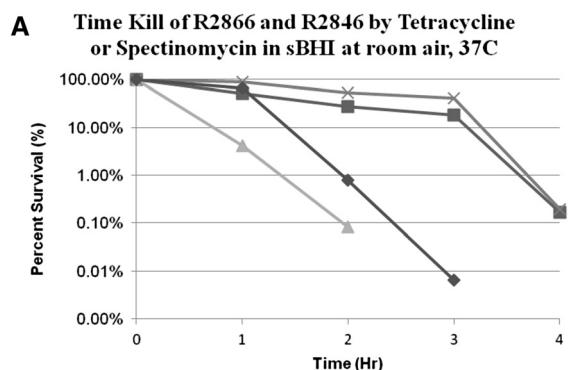


Figure 1 A. Strain R2866 was incubated in room air with tetracycline at 2 $\mu\text{g}/\text{ml}$ (■), or spectinomycin at 20 $\mu\text{g}/\text{ml}$ (▲) with the number of CFU determined on aliquots obtained hourly. Strain R2846 was incubated with 3 $\mu\text{g}/\text{ml}$ tetracycline (♦) or spectinomycin at 8 $\mu\text{g}/\text{ml}$ (×) with the surviving CFU determined on aliquots obtained hourly. B. Strain R2866 was incubated anaerobically with tetracycline at 2 $\mu\text{g}/\text{ml}$ (■), or spectinomycin at 20 $\mu\text{g}/\text{ml}$ (▲) with the number of CFU determined on aliquots obtained hourly. Strain R2846 was incubated with 3 $\mu\text{g}/\text{ml}$ tetracycline (♦) or spectinomycin at 8 $\mu\text{g}/\text{ml}$ (×) with the surviving CFU determined on aliquots obtained hourly.

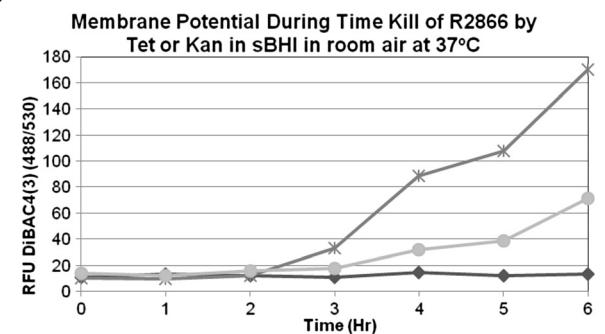


Figure 3 Strain R2866 was incubated in room air with Kanamycin at 1 $\mu\text{g}/\text{ml}$ (×) or tetracycline at 2 $\mu\text{g}/\text{ml}$ (♦) in sBHI containing 10 μM bis(1,2-dibutylbarbituric acid) trimethine oxonol [DiBAC_{4nm}(3)] and the fluorescence measured at 530. Strain R2866 gradually lost membrane permeability during the incubation (●).

serine hydroxymethyl transferase (*glyA*): both enzymes are involved in the pathway for the synthesis of L-Met, L-Ser, L-Thr and Gly [22].

Anaerobic killing

One mechanism cited for the decreased activity of antibiotics in an anaerobic environment is the decreased uptake of an aminoglycoside (streptomycin) into the bacterial cell [23].

Testing of a facultative anaerobe such as *E. coli* under strict anaerobic conditions has recently been reported [24,25]. Both of these publications demonstrated that killing by clinically useful concentrations of ampicillin, kanamycin and norfloxacin occurred in Coy chambers. They also found that the hydroxyl ion scavenger, thiourea had no effect on the rate of killing, as we found with *H. influenzae*. Keren et al. also found that testing *E. coli* BW25113 under anaerobic conditions the norfloxacin MICV was 0.25 $\mu\text{g}/\text{ml}$ while under aerobic incubation the MIC was 0.125 $\mu\text{g}/\text{ml}$ [25].

We found *H. influenzae* strain R3642 (anaerobically adapted R3932) was resistant to ciprofloxacin MIC = 15.5 ng/ml with a MBC of 28.5 ng/ml, whereas aerobically grown strains were susceptible. Others have found that the activity of levofloxacin, a fluoroquinolone with a mechanism of action identical to ciprofloxacin, against *P. aeruginosa* was not affected by anaerobic incubation. We have no explanation for the differing results.

Hydroxyl ion formation

In contrast to the report of the Collins group which showed that 5 $\mu\text{g}/\text{ml}$ of kanamycin killed 99.9% of *E. coli* MG1655 in three hours [9], Keren et al. found that kanamycin at 10 $\mu\text{g}/\text{ml}$ killed 90% of *E. coli* BW25113 in the same duration of incubation [25]. We could detect hydroxyl ion during the killing of *H. influenzae* by 1 mM

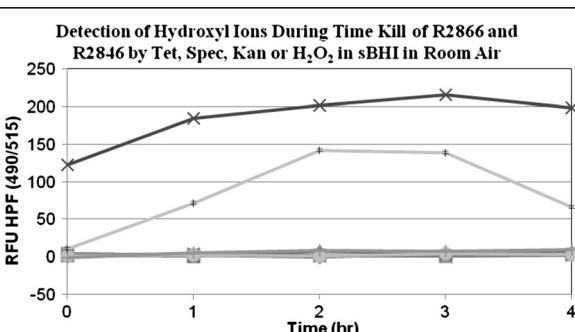


Figure 2 Hydroxyl ions were detected during an aerobic time-kill assay in room air by measuring the fluorescence at 516_{nm} of 3' (p-hydroxyphenyl), fluorescein (HPF, 5 μM). Positive controls consisted of 1 mM H₂O₂ added to the media with strain R2866 (X) and R2846 (#). Hydroxyl ions were not detected during incubation of R2866 with 2 $\mu\text{g}/\text{ml}$ tetracycline (■) or Kanamycin at 1.5 $\mu\text{g}/\text{ml}$ (▲) or with R2846 in media containing tetracycline at 3 $\mu\text{g}/\text{ml}$ (●) or Kanamycin at 1.5 $\mu\text{g}/\text{ml}$ (▼).

H_2O_2 (Figure 3), but not during killing by tetracycline and spectinomycin.

These studies do not find evidence for oxidative killing of five well-characterized strains of *H. influenzae* and are in agreement with two recently published papers demonstrating the absence of oxidative killing of *E. coli* [24,25].

Conclusions

We conclude that β -lactams, cephalosporins, macrolides, tetracycline's, aminoglycosides, chloramphenicol, rifampin and ciprofloxacin are bactericidal against five well-characterizes *H. influenzae* in an aerobic and anaerobic environment. The activity of trimethoprim was increased in anaerobic conditions.

Methods

Antibiotics

Table 7 lists the antibiotics used, their supplier and catalog number. Bacteria were stored at $-80^{\circ}C$ in sterile skim milk, and were cultured at $37^{\circ}C$ in 5% CO_2 on chocolate agar supplemented with 1% (v/v) IsoviteX (Becton Dickinson and Co., Franklin Lakes, NJ) or in Difco brain heart infusion broth (Becton Dickinson and Co.) supplemented (sBHI) with hemin HCl ($10 \mu\text{g ml}^{-1}$; Sigma Chemical Co., St. Louis, MO) and β -NAD $^{+}$ ($10 \mu\text{g ml}^{-1}$; Sigma) and with agar (Bacto-Agar, Difco) for solid sBHI plates. To detect hydroxyl ion we used 3'-(p-amino phenyl fluorescein) (APF, Invitrogen) at $5 \mu\text{M}$, while outer membrane integrity was assessed with bis (1,2-dibutylbarbituric acid) trimethine oxonol [DiBAC₄ (3)] also at $10 \mu\text{M}$.

Table 7 Antibiotics used

Antibiotic	Catalog number	Lot number	Vendor
Ampicillin	BP1760-25	965826	Fisher Scientific
Cefaclor	SI-114 8A	XY5511 AMX	Eli Lilly
Cefuroxime	C4417-1G	110M0794V	Sigma
Chloramphenicol	C-0378	44H0560	Sigma
Ciprofloxacin HCl	86393-320	9867	Miles laboratories
Clarithromycin	C4502	n/a	LKT Laboratories
Erythromycin	329815	3415B88	EMD Millipore
Kanamycin Sulfate	K4000-36	n/a	Sigma
Rifampicin	557303	644293	Calbiochem
Spectinomycin Sulfate	S-9137	81 K12755	Sigma
Tetracycline HCl	T-3383-25	20 K1279	Sigma
Tobramycin	T-1783	124 K0948	Sigma
Trimethoprim	T-7883	27H1015	Sigma

Bacteria

We first tested the aerobic susceptibility of the ATCC reference strain, *H. influenzae* ATCC 49766 with ampicillin, tetracycline and chloramphenicol with the microtiter plate system to verify the reliability of the technique before examining the test strains. The five test strains were *H. influenzae* Rd KW20 [26]; R2866, a nontypeable strain isolated from the blood of a normal infant with meningitis [27]; R2846 identified as strain 12 on primary isolation from middle ear exudate [28]. Strain R3642 originally identified as 86-028NP was isolated from the nasopharynx of a child with otitis media [29]. R3661 was isolated from middle ear fluid and sequenced by D. Dyer at the University of Oklahoma identified there as 3224A (<http://www.microgen.ouhsc.edu>). We annotated the R3661 genome with Glimmer v3.02 (<http://www.ccb.umd.edu/software/glimmer/>) as we had done with strains R2846 and R2866. Our annotation required that on BLASTn searches homology to other orfs, usually *E. coli* to be 80% or more and the predicted protein mass be at least 75% identical to the homolog. The genomic sequence of these of the five test strains is available at NCBI <http://www.ncbi.nlm.nih.gov/genome/genomes/165?subset=>.

We chose these five strains as they were sequenced to high coverage, and annotated by hand with reference to biochemical functions when available. With reliable genomic data future experiments concerning the antibiotic activity under anaerobic conditions can be investigated. We did not seek to determine the clinical utility of the antibiotics tested.

All strains were stored at $-70^{\circ}C$ in skim milk and cultured at $37^{\circ}C$ over night in 5% CO_2 on chocolate agar (aerobic) or sBHI agar (anaerobic) prior to use. Antibiotic susceptibility was determined in BHI broth supplemented with β -NAD $^{+}$ and hemin chloride both at $10 \mu\text{g/ml}$ (sBHI broth).

Aerobic susceptibility

We used the technique specified in the CLSI monograph for testing aerobic bacteria (M07-A9) adapted to a microtiter format as previously described [30] with an inoculum of 3 to 5×10^5 CFU/ml: $150 \mu\text{L}$ of BHI broth supplemented with hemin Cl and β -NAD + (sBHI) was inoculated into each well of a 96 well round bottom Nunc Polystyrene plate (Sigma Cat# P5366). The plate was covered with a lid and incubated at $37^{\circ}C$ in room air for 18 to 24 hours. Although the CLSI monograph M07-A8 indicates that testing of *H. influenzae* antibiotic susceptibility be performed in ambient air, it also notes that certain fastidious organisms be tested in 5% CO_2 . We therefore also tested the antibiotic susceptibility in 5% CO_2 . The plates were read in a Spectra Max 190 spectrophotometer with the $A_{600\text{nm}}$ of the initial inoculum was 0.089 to 0.124 with overnight growth in

antibiotic-free wells giving an average $A_{600\text{nm}}$ of 0.313 ± 0.741 . Converting the $A_{600\text{nm}}$ in the microtiter plate to a 1.0 cm light path an $A_{600\text{nm}}$ of 0.313 in the microtiter plate is equal to $A_{600\text{nm}}$ of 0.865 in a spectrophotometer with a 1 cm light path or $\sim 3 \times 10^9$ CFU/ml. The MIC was defined as the antibiotic concentration with an $A_{600\text{nm}} < 0.10$, an approximate 90% kill, while subculture of the wells with antibiotic concentrations greater than the MIC permitted calculation of the MBC, a 99.9% kill. Each strain was tested in duplicate and each run was repeated three times, with the mean value of MIC and MBC reported in the tables.

Anaerobic susceptibility

Because of the variation in the time required for the BBL GasPak system to reach an O_2 concentration of 0.5% [31] we performed the anaerobic experiments in a Coy anaerobic chamber with an O_2 concentration of < 0.002 ppm but containing 5% CO_2 . Growth of the five anaerobically adapted *H. influenzae* in sBHI broth containing 0.1% ferric nitrate at 37°C in the Coy chamber increased from a starting $A_{600\text{nm}}$ of $.0045 \pm 0.008$ to an average of 0.449 ± 0.060 . We adapted CLSI protocol M11-A7 (2007) to the 96 well microtiter plate used for aerobic testing. The test bacteria were passed twice on sBHI agar incubating each subculture overnight at 37°C in the anaerobic chamber. After obtaining good anaerobic growth, the anaerobically adapted strains were renumbered; Rd KW20, R3929; R2866, R3930; R2846, R3931; R3642, R3932 and R3661, R3933 and frozen at $-70^\circ C$ in skim milk. The anaerobically adapted strains were thawed in the Coy chamber, plated on sBHI agar and incubated overnight in the Coy chamber. The bacteria were harvested from these plates and added to sBHI broth which had been stored overnight in the Coy anaerobic chamber at 37°C containing 5% CO_2 . The O_2 concentration was < 0.002 ppm throughout the incubation. MIC and MBC were determined as described for aerobic incubation. The microtiter plates were removed from the Coy at the end of anaerobic incubation and the $A_{600\text{nm}}$ read in the spectrophotometer.

Aerobic killing kinetics

H. influenzae strains R2846 and R2866 were grown overnight on sBHI agar incubated at 37°C in 5% CO_2 and harvested into sBHI broth. The bacterial density was adjusted to 0.2 at 600nm , approximately 10^8 CFU/ml and 1 ml of the bacterial suspension was placed in a 125 ml baffled Erlenmeyer flask containing 10 ml of sBHI; antibiotics were added and incubated at 37°C in room air with shaking at 200 rpm. Aliquots were removed at zero time, 1, 2, 3 and 4 hours of incubation, diluted and plated on sBHI agar and incubated overnight at 37°C in 5% CO_2 to determine bacterial density. The

concentration of tetracycline and spectinomycin selected was approximately twice the MBC value determined in the plate assay. To allow for comparison of the activities we expressed the CFU at each time point as a percentage of the starting inoculum. The strains were tested in duplicate and the assay was repeated twice and the averages plotted.

Hydroxyl radical measurement

To determine if the aerobic killing of *H. influenzae* by bactericidal antibiotics in room air was via the formation of a hydroxyl radical we measured the oxidation of 5 μM hydroxyphenyl fluorescein (HPF) during the cidal action of tetracycline, kanamycin, spectinomycin and H_2O_2 with a Becton Dickinson FACS caliber flow cytometer with excitation at 490nm and emission at 515nm . We also repeated the bactericidal assay in the presence of 3'-(p-amino phenyl fluorescein) (APF) recording fluorescence as described for HPF. We used both strains at a starting density of $2 - 4 \times 10^8$ CFU/ml to obtain a reliable signal and subcultured each strain at 1, 2, 3 and 4 hours of incubation in room air at 37°C in sBHI broth agitating the culture at 200 rpm. To determine if iron was involved in the oxidative bactericidal reaction we repeated the bactericidal assay in the presence of 0.5 mM 2, 2'Dipyridyl and 150 mM thiourea.

Outer membrane integrity

To confirm that outer membrane integrity was affected during a kanamycin kill curve in room air we added 10 mM bis(1,2-dibutylbarbituric acid) trimethine oxonol (DiBAC₄(3)) to the cidal assay. DiBAC₄(3) can enter depolarized cells where it binds to hydrophobic molecules. When cells are lose membrane potential they will continuously take up the dye and increase the cells fluorescence with excitation at 530nm and emission maximum at 560nm .

Anaerobic killing kinetics

Strains R3930 and R3931 were harvested from chocolate agar plates after overnight growth in the Coy chamber at 37°C and 5% CO_2 with an O_2 concentration < 0.002 ppm. The bacteria were suspended in sBHI broth containing 0.1% ferric nitrate which had been stored anaerobically for a minimum of two hours and adjusting the density to an $A_{600\text{nm}}$ of ~ 0.2 . Tetracycline and spectinomycin were added at approximately twice the anaerobic MBC determined in the micro plate reader and incubated at 37°C in the Coy. The flasks were put in the 37°C incubator that is housed in the Coy chamber with the 125 ml baffled Erlenmeyer flask on a magnetic stirrer. After addition of the antibiotics, producing a final volume of 10 ml, an aliquot was taken and additional aliquots were taken from flasks in the Coy chamber after 1, 2, 3, 4,

and 6 hours of incubation, diluted and plated on chocolate agar plates which were incubated in 5% CO₂ at 37°C to determine bacterial density. We assumed that determining bacterial density under these conditions after bacterial killing occurred accurately indicated bacterial density.

Availability of supporting data

Raw data on individual strains with specific antibiotics and incubation conditions will be made available on request.

Abbreviations

sBHI: brain-heart infusion broth supplemented with hemin Cl and β-NAD⁺; MIC: Minimum inhibitory concentration; MBC: Minimum bactericidal concentration; Torr: a unit of pressure equal to 1 mmHg or 0.0193 psi; pO₂: the partial pressure of oxygen; pCO₂: the partial pressure of carbon dioxide gas; mM: millimolar; NTHi: nontypeable *H. influenzae*; MnSOD: manganese-dependent superoxide dismutase.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

HKS, EC and VN carried out the experiments. KLN designed the experimental protocols and performed certain experiments. ALS conceived the hypothesis, designed certain experiments and authored the manuscript. All authors read and approved the final manuscript.

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ALS is professor of Pediatrics and Microbiology at the University of Washington School of Medicine. KLN is a senior technician in ALS laboratory. HKS, VN and EC are all technicians in various laboratories.

Acknowledgements

This work was supported in part by National Institute of Allergy and Infectious Disease under award R01AI46512, and the Seed Fund of Seattle Children's Research Institute. The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of health.

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Received: 29 April 2013 Accepted: 21 June 2013

Published: 26 June 2013

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doi:10.1186/1756-0500-6-241

Cite this article as: Smith et al.: "Affect of anaerobiosis on the antibiotic susceptibility of *H. influenzae*". *BMC Research Notes* 2013 6:241.

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