

## Article

# Role of Bioaerosols on the Short-Distance Transmission of Multidrug-Resistant Methicillin-Resistant *Staphylococcus aureus* (MRSA) in a Chicken Farm Environment

Bing-Mu Hsu <sup>1,†</sup>, Jung-Sheng Chen <sup>2</sup>, Gwo-Jong Hsu <sup>3,†</sup>, Suprokash Koner <sup>1,4,†</sup>, Viji Nagarajan <sup>1</sup> and Hsin-Chi Tsai <sup>5,6,\*</sup>

<sup>1</sup> Department of Earth and Environmental Sciences, National Chung Cheng University, Chiayi County 621, Taiwan; bmhsu@eq.ccu.edu.tw (B.-M.H.); suprokashkoner22@gmail.com (S.K.); mathumitha08@gmail.com (V.N.)

<sup>2</sup> Department of Medical Research, E-Da Hospital, Kaohsiung City 824, Taiwan; nicky071214@gmail.com

<sup>3</sup> Division of Infectious Diseases, Ditmanson Medical Foundation, Chia-Yi Christian Hospital, Chiayi City 600, Taiwan; cych01347@gmail.com

<sup>4</sup> Department of Biomedical Sciences, National Chung Cheng University, Chiayi County 621, Taiwan

<sup>5</sup> Department of Psychiatry, School of Medicine, Tzu Chi University, Hualien County 970, Taiwan

<sup>6</sup> Department of Psychiatry, Tzu-Chi General Hospital, Hualien County 970, Taiwan

\* Correspondence: cssbmw45@gmail.com; Tel.: +886-38561825

† These authors contributed equally to this work.

## Supplementary Materials

**Table S1.** The details of bioaerosol sampling of chicken farm environment.

Sampling point	Global positioning system (GPS)	Samples type	Air sampling duration	Air flow speed	Total Volume of air sampling
1 <sup>st</sup> chicken shed	23°58'73.80" N, 120°49'09.34" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
2 <sup>nd</sup> chicken shed	23°58'78.32" N, 120°49'07.84" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
Exposure square	23°58'75.76" N, 120°49'07.95" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
3m Downwind	23°58'76.18" N, 120°49'08.88" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
5m Downwind	23°58'76.31" N, 120°49'10.00" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
20m Downwind	23°58'63.17" N, 120°49'11.04" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
50m Downwind	23°58'58.66" N, 120°49'16.07" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter
50m Upwind	23°58'78.25" N, 120°48'93.12" E	Bioaerosol	10 min	28.3 (L/min)	283 Litter

**Table S2.** MRSA strain identification, *Spa* typing, *SCCmec* typing, and virulence factors detecting genes primers list and PCR conditions.

Target gene	Size	Sequence (5' to 3')	Reaction Materials Final Volume: 25 µl	PCR Condition	Reference
nuc	270	nuc-F 5'-GCGATTGATGGTGATACGGTT-3'	DNA: 100-300 ng	Pre-denaturation: 95°C 5 min	[1,2]
mecA	448	nuc-R 5'-AGCCAAGCCTTGACGAATAAAGC-3'	Primer: 400 nM	Denaturation: 94°C 60s	
		mecA-F 5'-CTCAGGTACTGCTATCCACC-3'	nuc FR & mecA FR	Annealing: 55°C 60s	
		mecA-R 5'-CACTTGGTATATCTTCACC-3'	Master mix: 5 µl	Extension: 72°C 60s	
				D.A.E. Cycles: 30 cycles	
				Final extension: 72°C 10 min	

Spa	270	Spa-1113F: 5'- TAAAGACGATCCTTCGGTGAGC-3' Spa-1514R: 5'- CAGCAGTAGTGCCGTTTGCTT-3'	DNA: 100-300 ng	Pre-denaturation: 80°C 5 min	https://spa-server.ridom.de/
			Primer: 200 nM Spa FR	Denaturation: 94°C 45s Annealing: 60°C 45s Extension: 72°C 90s	
			Master mix: 5 µl	D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	
SCCmec I	495	CIF2 F2: 5'-TTCGAGTTGCTGATGAAGAAGG-3' CIF2 R2: 5'-ATTTACCACAAGGACTACCAGC-3' KDP F1: 5'-AATCATCTGCCATTGGTGATGC-3' KDP R1: 5'-CGAATGAAGTGAAAGAAAGTGG-3'	DNA: 100-300 ng Primer: 400 nM CIF-FR, 200 nM KDP-FR, 200 nM RIFF3R9	Pre-denaturation: 94°C 4 min Denaturation: 94°C 30s Annealing: 53°C 30s Extension: 72°C 1 min	[3]
SCCmec II, III	284 209	MECI P2: 5'-ATCAAGACTTGCATTGAGGC-3' MECI P3: 5'-GCGGTTTCAATTCATTGTC-3'	400 nM MECI-FR, 400 nM RIFF10R13, 800 nM DCS-FR	D.A.E. Cycles: 30 cycles Final extension: 72°C 4 min	
SCCmec III	414	RIF F3: 5'-GTGATTGTTGAGATATGTGG-3' RIF R9: 5'-CGCTTTATCTGTATCTATCGC-3' RIF F10: 5'-TTCTTAAGTACACGCTGAATCG-3' RIF R13: 5'-GTCACAGTAATCCATCAATGC-3' DCS F2: 5'-CATCCTATGATAGCTTGGTC-3' DCS R1: 5'-CTAAATCATAGCCATGACCG-3'	Master mix: 5 µl		
SCCmec V	325	Type V-F: 5'-GAACATTGTTACTTAAATGAGCG-3' Type V-R: 5'-TGAAAGTTGTACCCTTGACACC-3'	DNA: 100-300 ng Primer: 100 nM V-FR Master mix: 5 µl	Pre-denaturation: 94°C 5 min Denaturation-1: 94°C 45s Annealing-1: 65°C 45s Extension-1: 72°C 1.5 min D.A.E.-1 Cycles: 10 cycles Denaturation-2: 94°C 45s Annealing-2: 55°C 45s Extension-2: 72°C 1.5 min D.A.E.-2 Cycles: 25 cycles Final extension: 72°C 10 min	[4]
SCCmec VI	134	ccrB4-F: 5'-CGAAGTATAGACACTGGAGCGATA-3' ccrB4-R: 5'-GCGACTCTCTTGCGCTTTA-3'	DNA: 100-300 ng Primer: 100 nM FR, Master mix: 5 µl	Pre-denaturation: 95°C 10 min Denaturation: 95°C 30s Annealing: 50°C 30s Extension: 72°C 30s D.A.E. Cycles: 40 cycles Final extension: 72°C 5 min	[5]

SCCmec VII SCCmec VIII	473 138	Type VII F: 5'- GTGACGTTGATATTGCAGTGGT-3' Type VII R: 5'-TGAAGAAGTTTGTTCGCGT-3' Type VIII F: 5'- AGCGACGATGAACAACACCGCTACTTACTC AA-3' Type VIII R: 5'- TTGGTTGAGAATGAGAACAGTGGTAAGATC- 3'	DNA: 100-300 ng Primer: 400 nM FR Master mix: 5 μl	Pre-denaturation: 95°C 2 min Denaturation: 95°C 30s Annealing: 54°C 1 min Extension: 72°C 1 min 20s D.A.E. Cycles: 35 cycles Final extension: 72°C 7 min	[6]
PVL	433	PVL-1: 5'- ATCATTAGGTAAAATGTCTGGACATGATCC A-3' PVL-2: 5'- GCATCAAGTGTATTGGATAGCAAAAGC-3'	DNA: 100-300 ng Primer: 400 nM FR Master mix: 5 μl	Pre-denaturation: 94°C 5 min Denaturation: 94°C 40s Annealing: 53°C 40s Extension: 72°C 1 min D.A.E. Cycles: 35 cycles Final extension: 72°C 10 min	[7]
<i>entA</i> <i>entB</i> <i>entC</i> <i>entD</i> <i>entE</i> <i>tsst-1</i> <i>eta</i> <i>etb</i>	121 478 459 384 495 271 464 200	<i>entA</i> -F: 5'-TTGGAACGGTTAAAACGAA-3' <i>entA</i> -R: 5'-GAACCTTCCCATCAAAAACA-3' <i>entB</i> -F: 5'-TCGCATCAAACCTGACAAACG-3' <i>entB</i> -R: 5'-GCAGGTACTCTATAAGTGCC-3' <i>entC</i> -F: 5'-GGAGGAATAACAAAACATGAAGG- 3' <i>entC</i> -R: 5'-AAAGGCAAGCACCGAAGTAC-3' <i>entD</i> -F: 5'-TGGTGGTGAAATAGATAGGAC-3' <i>entD</i> -R: 5'-TGAAGGTGCTCTGTGGATAAT-3' <i>entE</i> -F: 5'-TGGTAGCGAGAAAAGCGAAG-3' <i>entE</i> -R: 5'-TGTAATAATGCCTTGCCTGAA-3' <i>tsst-1</i> -F: 5'-CTGGTATAGTAGTGGGTCTG-3' <i>tsst-1</i> -R: 5'-AGGTAGTTCTATTGGAGTAGG-3' <i>eta</i> -F: 5'-TTTGCTTTCTTGATTGGATTG-3' <i>eta</i> -R: 5'-GATGTGTTCCGTTTGATTGAC-3' <i>etb</i> -F: 5'-ACGGCTATATACATTCAATT-3' <i>etb</i> -R: 5'-TCCATCGATAATATACCTAA-3'	DNA: 100-300 ng Primer: 400 nM Master mix: 5 μl	Pre-denaturation: 94°C 5 min Denaturation: 94°C 1 min Annealing: 2 min Extension: 72°C 1 min D.A.E. Cycles: 35 cycles Final extension: 72°C 5 min AnnealingTemp. <i>entA</i> : 50°C <i>entB</i> : 55°C <i>entC</i> : 59°C <i>entD</i> : 51°C <i>entE</i> : 55.5°C <i>tsst-1</i> : 54°C <i>eta</i> : 54°C <i>etb</i> : 50.9°C	[8]

**Table S3.** Pearson correlation coefficient (r) value and level of significant for environmental parameters and total bacteria count.

Correlations test					
	Ammonia	Methylamine	Sampling Distance	Wind speed (m/s)	Total bacteria count
Ammonia	1.000	0.961**	-0.092	-0.447	0.865**
Methylamine	0.961**	1.000	-0.089	-0.443	0.940**
Sampling Distance	-0.092	-0.089	1.000	-0.490	-0.048
Wind speed (m/s)	-0.447	-0.443	-0.490	1.000	-0.476
Total bacteria count	0.865**	0.940**	-0.048	-0.476	1.000

\*\* . Correlation is significant at the 0.01 level (2-tailed).

**Table S4.** Standard score of Zone of inhibition diameter measurement in disk diffusion method to determine the microbial resistance property.

Antibiotic name	Zone of inhibition diameter measurement value (mm)		
	Resistant	Intermediate	Susceptible
Chloramphenicol	≤12	13-17	≥18
Ciprofloxacin	≤15	16-20	≥21
Clindamycin	≤14	15-20	≥21
Erythromycin	≤13	14-22	≥23
Gentamicin	≤12	13-14	≥15
Rifampicin	≤16	17-19	≥20
Tetracycline	≤14	15-18	≥19
Sulfamethoxazole-Trimethoprim	≤10	11-15	≥16

A reference documents CLSI M100-S27 which is modified in 2017 has been used to determine the score of Zone of inhibition diameter measurement [9]

**Table S5.** The results of Chi-square test between sampling point with MRSA strain characterization

Chi-Square test			
Variables	Degree of freedom (df)	Sampling point	
		Pearson Chi-Square	Asymptotic Significance (2-sided)
SCCmec typing	7	23.000 <sup>a</sup>	0.002
MRSA grouping	7	23.000 <sup>a</sup>	0.002
Spa type	7	23.000 <sup>a</sup>	0.002
Virulence factors	21	53.959 <sup>b</sup>	0.000
Multiple drug resistance	14	26.680 <sup>c</sup>	0.021

a. 16 cells (100.0%) have expected count less than 5. The minimum expected count is .09.

b. 32 cells (100.0%) have expected count less than 5. The minimum expected count is .09.

c. 24 cells (100.0%) have expected count less than 5. The minimum expected count is .04.

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