



Supplement of

Influence of terrestrial and marine air mass on the constituents and intermixing of bioaerosols over a coastal atmosphere

Qun He et al.

Correspondence to: Min Wei (minwei@sdnu.edu.cn)

The copyright of individual parts of the supplement might differ from the article licence.

Table S1 Sample classification and description influenced by different air masses

Group	Date	Samples									
		19-Jan	20-Jan	1-Mar	9-Mar	11-Mar	12-Mar	22-Mar	23-Mar	24-Mar	25-Mar
TE	PM _{2.5}	51.35	124.67	30.58	16.5	65.63	38.21	53.05	77.46	84.42	60.83
	Sample	WH_1	WH_2	WH_7	WH_8	WH_10	WH_11	WH_15	WH_16	WH_17	WH_18
MA	Date	21-Jan	16-Mar	21-Mar	29-Mar	30-Mar	31-Mar				
	PM _{2.5}	7.92	12.38	15.37	25.38	25.42	28.83				
MIX	Sample	WH_3	WH_13	WH_14	WH_22	WH_23	WH_24				
	Date	26-Feb	27-Feb	28-Feb	10-Mar	14-Mar	26-Mar	27-Mar	28-Mar		
PM _{2.5} concentration ($\mu\text{g}/\text{m}^3$)	PM _{2.5}	41.12	60.61	41	51.63	36.42	52.58	64.69	48.58		
	Sample	WH_4	WH_5	WH_6	WH_9	WH_12	WH_19	WH_20	WH_21		

TE, terrestrial air mass; MA, marine air mass; MIX, mix air mass.

Table S2 Indicator bacterial and fungal species influenced by different air masses

Group	Bacteria			Fungi		
	TE	MA	MIX	TE	MA	MIX
TE/MA/MIX	<i>Enhydrobacter</i>	<i>Comamonas</i>			<i>Aspergillus</i>	<i>Malassezia</i>
	<i>Cellulosimicrobium</i>	<i>Streptococcus</i>	<i>Enhydrobacter</i>	<i>Cladosporium</i>	<i>Coprinellus</i>	<i>Alternaria</i>
	<i>Pleomorphomonas</i>	<i>Novosphingobium</i>	<i>Lactobacillus</i>			
TE/MA		<i>Aerococcus</i>				
		<i>Comamonas</i>				
	<i>Lactobacillus</i>	<i>Streptococcus</i>				
	<i>Deinococcus</i>	<i>Novosphingobium</i>		<i>Cladosporium</i>		
	<i>Enhydrobacter</i>	<i>Aerococcus</i>		<i>Alternaria</i>	<i>Coprinellus</i>	
	<i>Sphingomonas</i>	<i>Delftia</i>		<i>Malassezia</i>		
MA/MIX		<i>Erysipelothrix</i>				
		<i>Chryseobacterium</i>				
			<i>Enhydrobacter</i>			
			<i>Lactobacillus</i>			
		<i>Comamonas</i>	<i>Cloacibacterium</i>		<i>Aspergillus</i>	<i>Malassezia</i>
		<i>Streptococcus</i>	<i>Cellulosimicrobium</i>		<i>Coprinellus</i>	<i>Alternaria</i>
		<i>Novosphingobium</i>	<i>Pleomorphomonas</i>			
		<i>Aerococcus</i>	<i>Deinococcus</i>			
			<i>Sphingomonas</i>			

TE, terrestrial air mass; MA, marine air mass; MIX, mix air mass.

Table S3 Ecological functions of bacterial and fungal communities influenced by different air masses

Group	Bacteria			Fungi		
	TE	MA	MIX	TE	MA	MIX
TE/MA/MIX	animal_parasites_or_symbionts		aerobic_chemoheterotrophy	Animal Pathogen-Undefined		Animal Pathogen-
	human_pathogens_all	hydrocarbon_degradation	animal_parasites_or_symbionts	Saprotroph	Dung	Undefined Saprotroph
	I	on	human_pathogens_animal	Animal Pathogen-Endophyte-Plant	Plant	Animal Pathogen-Endophyte-Plant
		human_gut	II	Pathogen-Wood	Wood	Pathogen-Wood
		mammal_gut		Saprotroph		Saprotroph
				Animal Pathogen-Endophyte-Lichen		
TE/MA		aromatic_compound_degradation		Parasite-Plant		
	animal_parasites_or_symbionts	human_gut		Pathogen-Wood		
	human_pathogens_all	mammal_gut		Saprotroph	Dung	Saprotroph-
	I	hydrocarbon_degradation		Animal Pathogen-Endophyte-Plant	Plant	Saprotroph-
	plant_pathogen	on		Pathogen-Wood	Wood	Wood Saprotroph
		aromatic_hydrocarbon_degradation		Saprotroph		
		plastic_degradation		Animal Pathogen-Undefined		
				Saprotroph		

MA/MIX

aerobic_chemoheterotr			Animal Pathogen-
ophy	hydrocarbon_degra	Dung Saprotoph-	Undefined Saprotoph
animal_parasites_or_sy	dation	Plant Saprotoph-	Animal Pathogen-
mbionts	human_gut	Wood Saprotoph	Endophyte-Plant
human_pathogens_all	mammal_gut		Pathogen-Wood
nitrate_reduction			Saprotoph
			Animal Pathogen-
			Endophyte-Lichen
			Parasite-Plant
			Pathogen-Wood
			Saprotoph

TE, terrestrial air mass; MA, marine air mass; MIX, mix air mass.

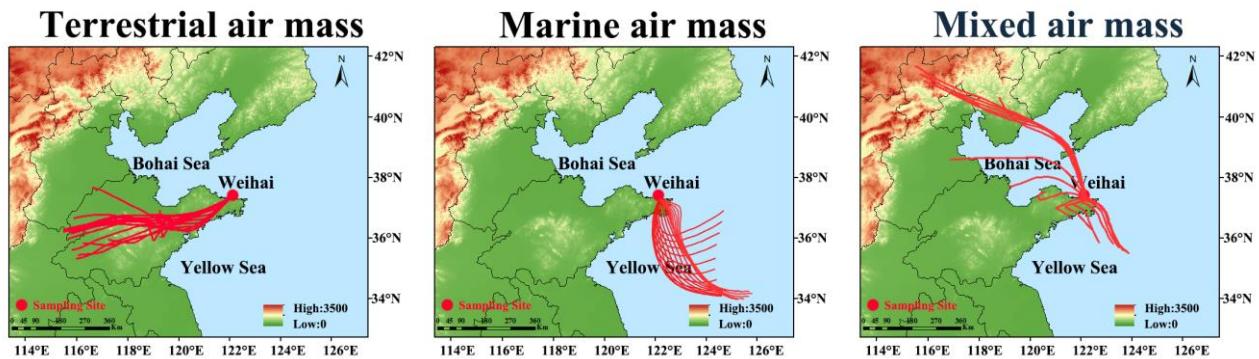


Fig. S1 The Sampling site in the coastal city of Weihai, Northern China, and the typical terrestrial, marine, and mixed air-mass are indicated.

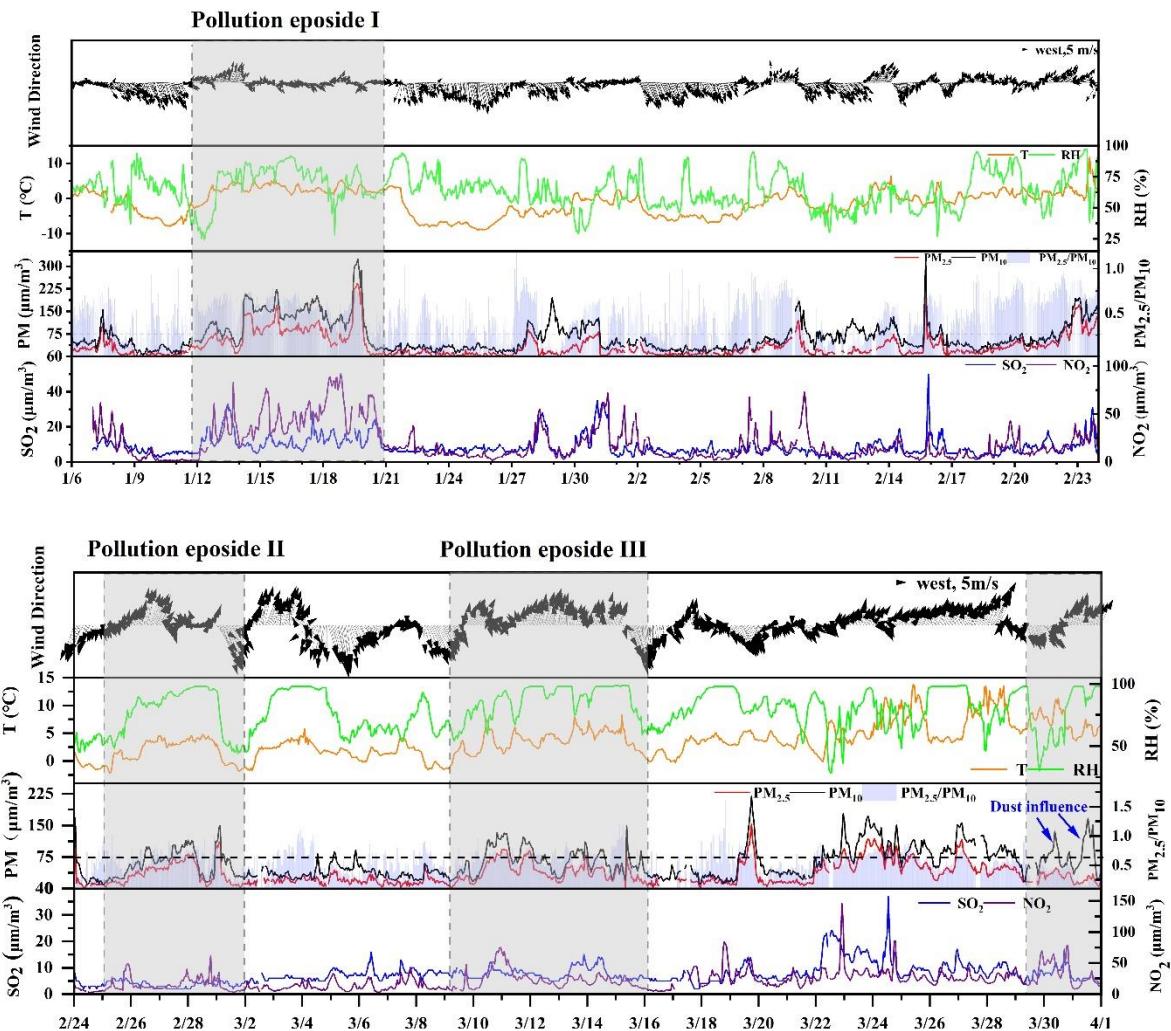


Fig. S2 Air pollutants and meteorological condition of three typical pollution episodes during the sampling period

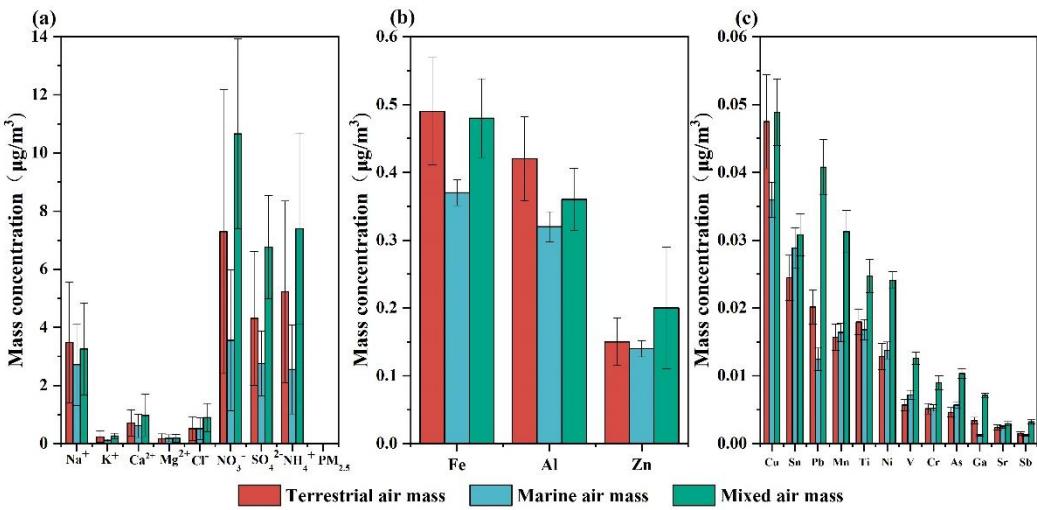


Fig.S3. Water-soluble ions (a) and metal element concentration (b, c) in $\text{PM}_{2.5}$ influenced by terrestrial, marine and mix air mass.

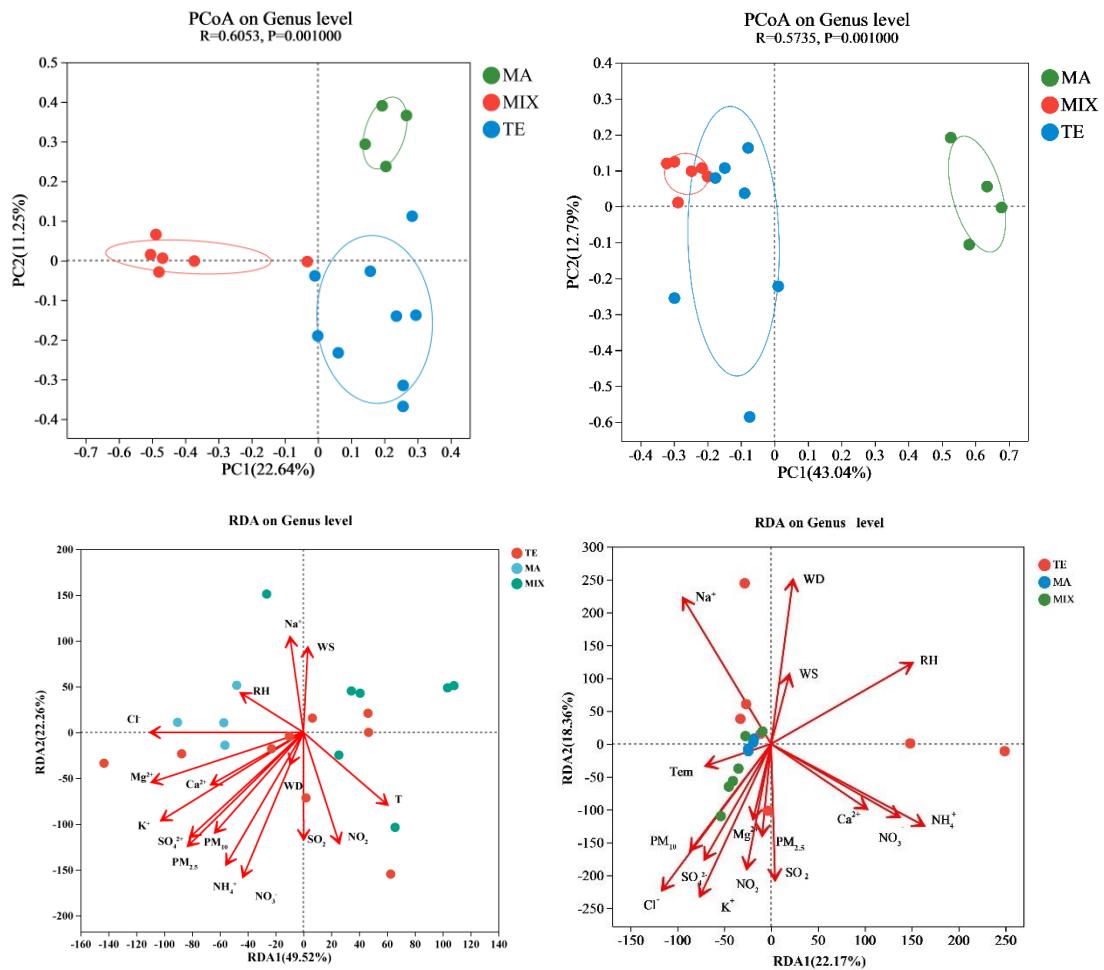
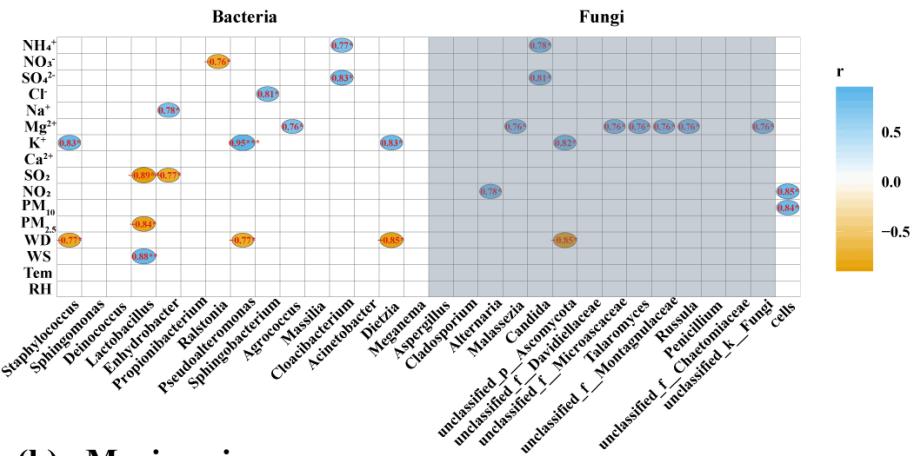
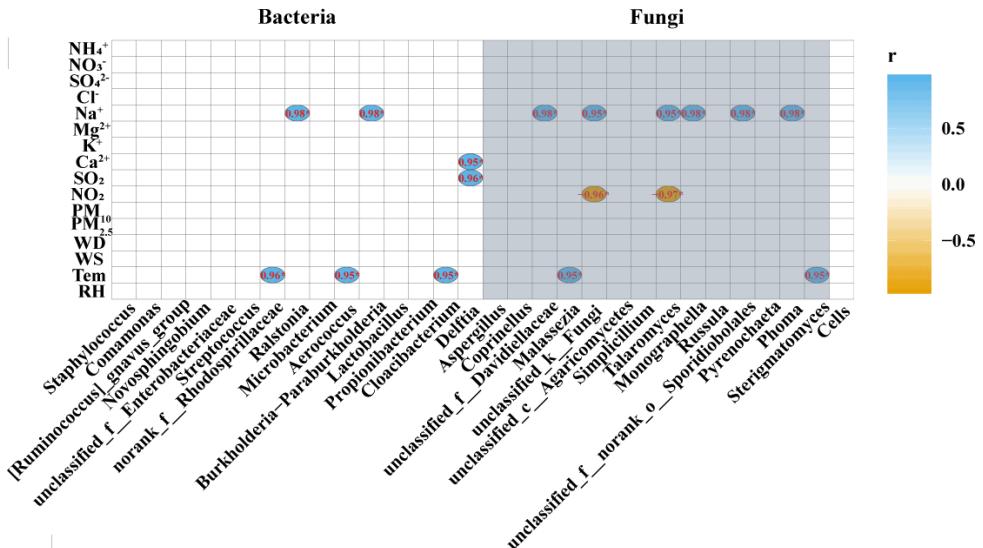


Fig. S4 PCoA and RDA analysis of bacterial and fungal community with environmental parameters.

(a) Terrestrial air mass



(b) Marine air mass



(c) Mixed air mass

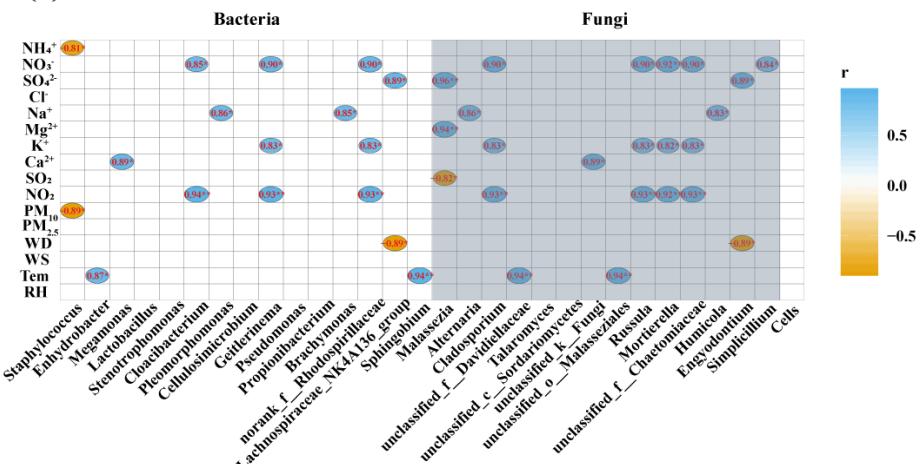


Fig. S5 Relationships between microorganisms and environmental factors under the influence of different air masses, continental air masses (a) marine air masses (b) and mixed air masses (c).