

Supplemental Materials

Supplementary 1 Text: The details list of participation Hospitals and the region they represented in China.

Region of China	Hospital
Southwest	Second Affiliated Hospital of Chongqing Medical University
	First Affiliated Hospital of Army Medical University
	First Affiliated Hospital of Chongqing Medical University
	Guizhou Provincial People's Hospital
	West China Hospital of Sichuan University
Northwest	First Affiliated Hospital of Zunyi Medical University
	First Hospital of Xinjiang Medical University
	Lanzhou General Hospital of People's Liberation Army
Northeast	Tangdu Hospital, Fourth Military Medical University
	First Affiliated Hospital of Harbin Medical University
North	First Affiliated Hospital of Dalian Medical University
	Peking Univesity Jishuitan Hospital
	Tianjin Medical University General Hospital
Eastern	Third Hospital of Hebei Medical University
	Shanghai Jiao Tong University Affiliated Sixth People's Hospital
	Nanjing Drum Tower Hospital of Nanjing University Medical School
	First Affiliated Hospital of Nanjing Medical University
Central	Zhejiang Provincial People's Hospital of Hangzhou Medical College
	Second Xiangya Hospital, Central South University
South	Wuhan General Hospital of People's Liberation Army
	Nanfang Hospital of Southern Medical University

Supplementary 2 Text: Inclusion and exclusion criteria

Inclusion criteria

- A.Post-traumatic osteomyelitis was definitely diagnosed by surgeons
- B.All ages
- C.Traumatic fractures of limbs (humerus, ulnar and radius, clavicle, femur, tibia and fibula, patella, calcaneus, talus)
- D.Complete clinical data records;
- E.Complete sampling and surgical procedure records
- F.Complete imagological, surgical or histopathological data
- G.Bacterial culture and drug sensitivity test to confirm infection;
- H.Approved by Ethics Review Committee

Exclusion criteria

- A.Post-traumatic osteomyelitis could not be diagnosed definitely
- B.There is clear clinical evidence of infection, but without positive result of laboratory bacterial culture
- C.Without the results of drug sensitivity
- D.Don't approved by Ethics Review Committee

Supplemental Table 1. Testing Bacteria Library List.

Gram Positive bacteria	Gram Negative bacteria	
MSSA	Pseudomonas aeruginosa	Haemophilus haemolyticus
MRSA	Enterobacter cloacae	Enterobacter asburiae
Enterococcus faecalis	Escherichia coli	Aeromonas salmonicida
Staphylococcus epidermidis	Acinetobacter baumannii	Burkholderia cepacia
S. haemolyticus	Klebsiella pneumoniae	Leclercia adecarboxylata
Enterococcus faecium	Serratia marcescens	Enterobacterium intermediates
Gram-positive bacillus	Proteus mirabilis	Serratia odorifera
Staphylococcus simulans	proteusbacillus vulgaris	Enterobacter amnigenus type 2
Enterococcus avium	Stenotrophomonas maltophilia	Achromobacter denitrificans
coagulase negative staphylococcus	Aeromonas hydrophila	Achromobacter sp
Streptococcus haemolyticus	Klebsiella oxytoca	Cedecea lapagei
S. capitis	Enterobacter aerogenes	Myroides odoratimimus
S. lugdunensis	Aeromonas sobria	Cupriavidus
Streptococcus pyogenes	Acinetobacter baumannii-Acinetobacter calcoaceticus complex	Providencia stuartii
Staphylococcus warneri	pseudomonas putida	Vibrio fluvialis
Enterococcus Gallinarum	Proteus vulgaris	Hafnia alvei
Enterococcus casseliflavus	Citrobacter freundii	carbapenem-resistant Acinetobacter baumannii
Streptococcus agalactiae	Morganella morganiiF	Doblin salmonella
Staphylococcus cohnii	Enterobacter carcinoma	Moraxella osloensis
Micrococcus luteus	Acinetobacter lwoffii	Enterobacter hormaechei
Streptococcus mitis	Fluorescence/foul-smelling pseudomonas	Cedecea davisae
Bacillus cereus	Flavobacterium indologenes	Aeromonas caviae
Streptococcus pneumoniae	Acinetobacter junii	Flavobacterium odoratum
S. saprophyticus	Serratia marcescens subspecies	Raoultella planticola
Streptococcus anginosus	Klebsiella pneumoniae subspecies	Serratia fonticola
staphylococcus xylosus	Xylose oxidizes noncolorific bacillus	Acinetobacter calcoaceticus
S. hominis	Citrobacter braakii	Ochrobactrum anthropi
Staphylococcus lugdunensis	Acinetobacter haemolyticus	Serratia plymuthica
Enterococcus casseliflavus/gsllinarum	Baumann/calcium acetate acinetobacter complex	Alcaligenes xylosoxidans subsp xylosoxidans
Kocuria kristinae	Pseudomonas stutzeri	<u>Alcaligenes xylosoxidans</u>
Streptococcus dysgalactiae	Morganella morganii	Total=96
Group d streptococcus	Prpteus penneri	
Streptococcus constellatus	Enterobacter agglomerans	
S. auricularis	Lycidine klebsiella	
Staphylococcus chromogenes	Proteus penneri	
Enterococcus hirae	Alcaligenes faecalis	
Streptococcus epiphylsis	Pasteurella multocida	
Staphylococcus schleiferi	Citrobacter farmeri	
Enterococcus durans	Klebsiella plantica	
Staphylococcus squirrel	Baumann/acetate acinetobacter haemolyticus	
S. cystobacteria	Sphingomonas paucimobilis	
Staphylococcus kloosii	Serratia rubidaea	
Staphylococcus intermedius/Staphylococ	Serratia liquefaciens	
Granulicatella para-adiacens	Flavobacterium breve	
Dermacoccus nishinomiyaensis	Providencia rettgeri	
Aerococcus viridans	Providencia stuartii	
Enterococcus raffinosus	Brac citrobacter	
Erysipelothrix rhusiopathiae	Achromobacter xylosoxidans	
Staphylococcus hominis	Non-decarboxylleucella	
Staphylococcus arlettae	Pseudomonas	
Streptococcus tolutte	Cozer's bacillus citrate	
Corynebacterium striatum	Malonate negative citrate bacillus	
Streptococcus	sagenopsis	
Staphylococcus hyicus	Pseudomonas alcaligenes	
Streptococcus acidominimus	Ralstonia pickettii	
Staphylococcus lentus	Aeromonas	
Enterococcus casseliflavus	Alcaligenes faecalis ssp faecalis	
Bacillus subtilis	Paratyphoid C	
Streptococcus milleri	Myroides	
Micrococcus antarcticus/Micrococcus lute	Citrobacter koseri	
Rhodococcus equi	Haemophilus parainfluenzae	
Staphylococcus caprae	Myroldes odoratus	
Bacillus thuringiensis	Gram-negative bacilli	
Corynebacterium jack	Pseudomonas shi (stutzer)	
Enterococcus	Cronobacter sakazakii	

Total=65

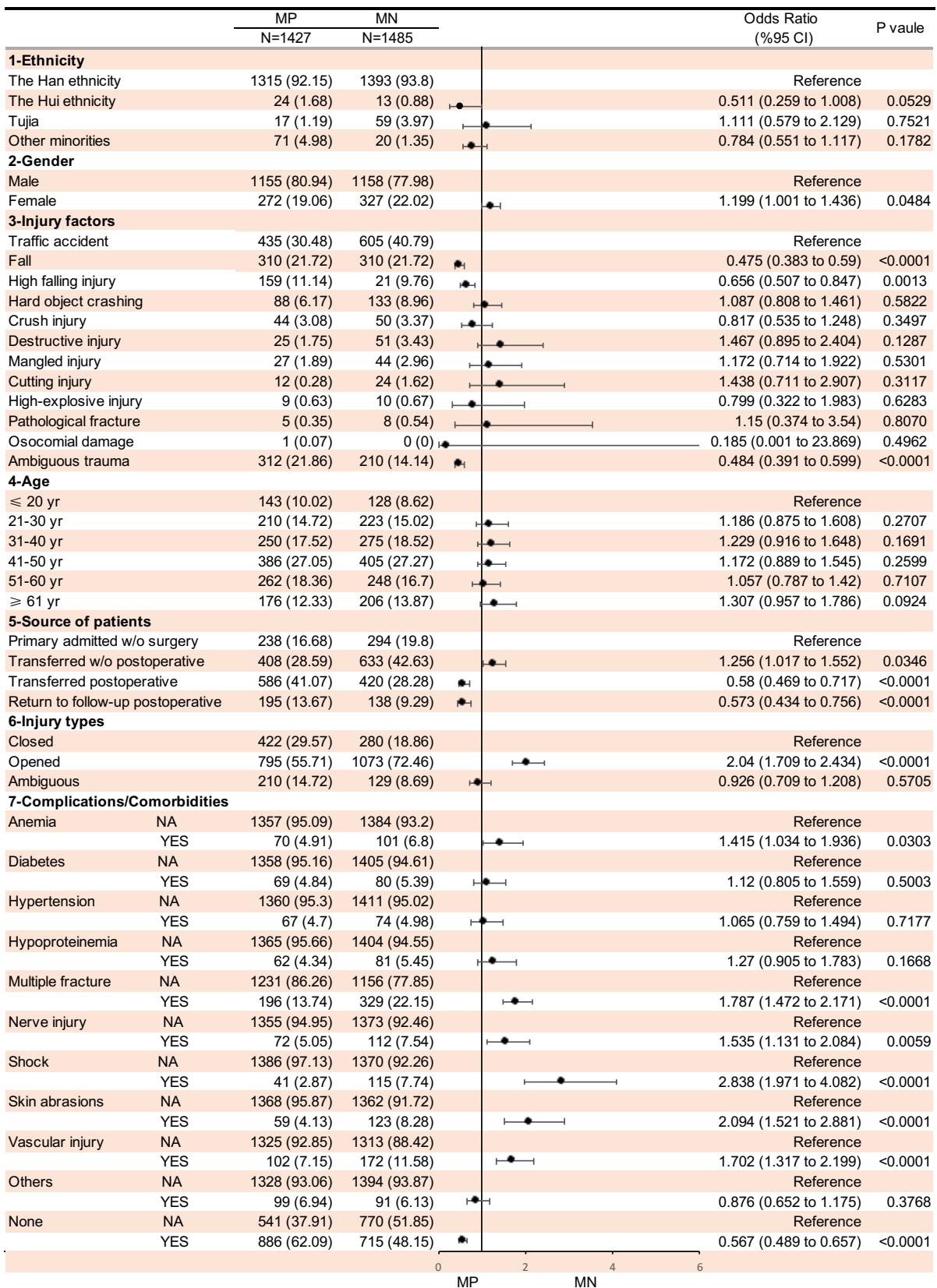
Supplemental Table 2.

The total number of age groups that participated from each geographic region of China

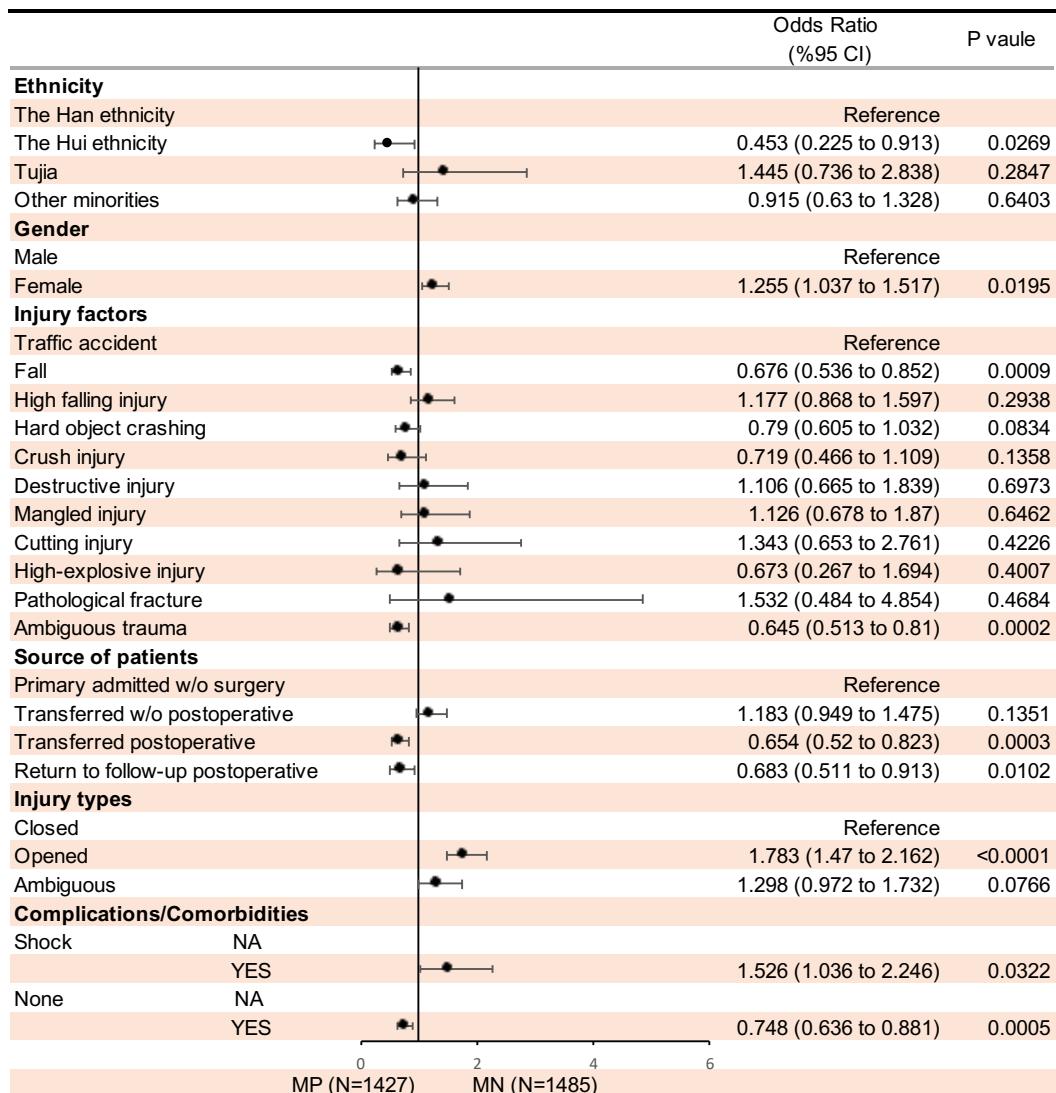
	Southwest	Northwest	Central China	North China	South China	East China	Northeast	Total
≤14	29	33	14	8	9	19	28	140
14-18	56	22	11	3	8	17	6	123
≥18	1513	383	350	295	302	170	250	3263
Total	1598	438	375	306	319	206	284	3526
Percentage	45.32%	12.42%	10.64%	8.68%	9.05%	5.84%	8.05%	100.00%

The percentage of age groups from each participated geographic region of China

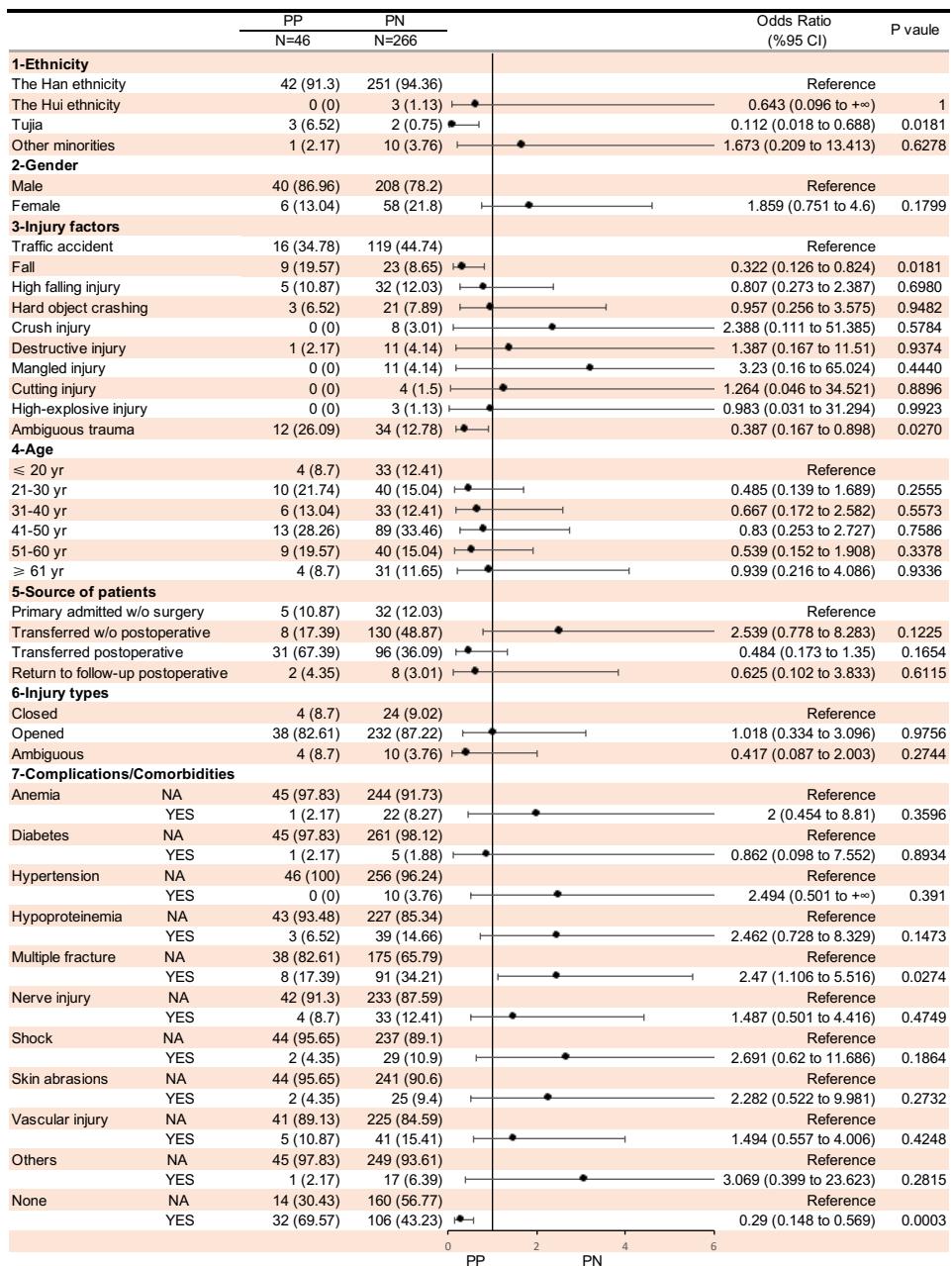
Supplemental Table 3. Univariate analysis of logistic regression was conducted to identify relative risk factors for patients who had monomicrobial positive (MP) or monomicrobial negative (MN) infection.



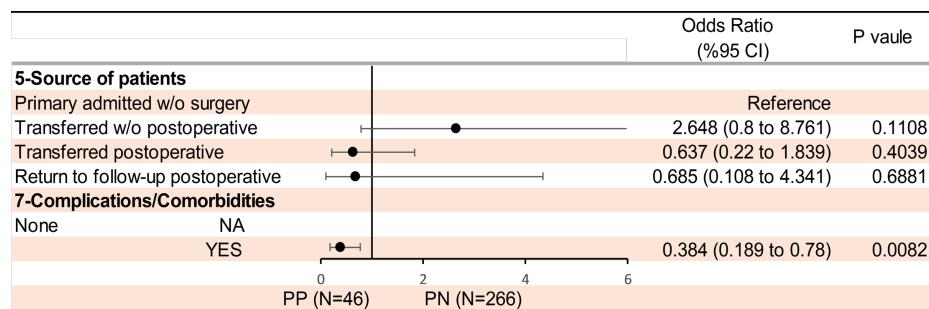
Supplemental Table 4. Multivariate analysis of logistic regression was performed to identify the relative risk factors for patients with monomicrobial positive (MP) or monomicrobial negative (MN) infections.



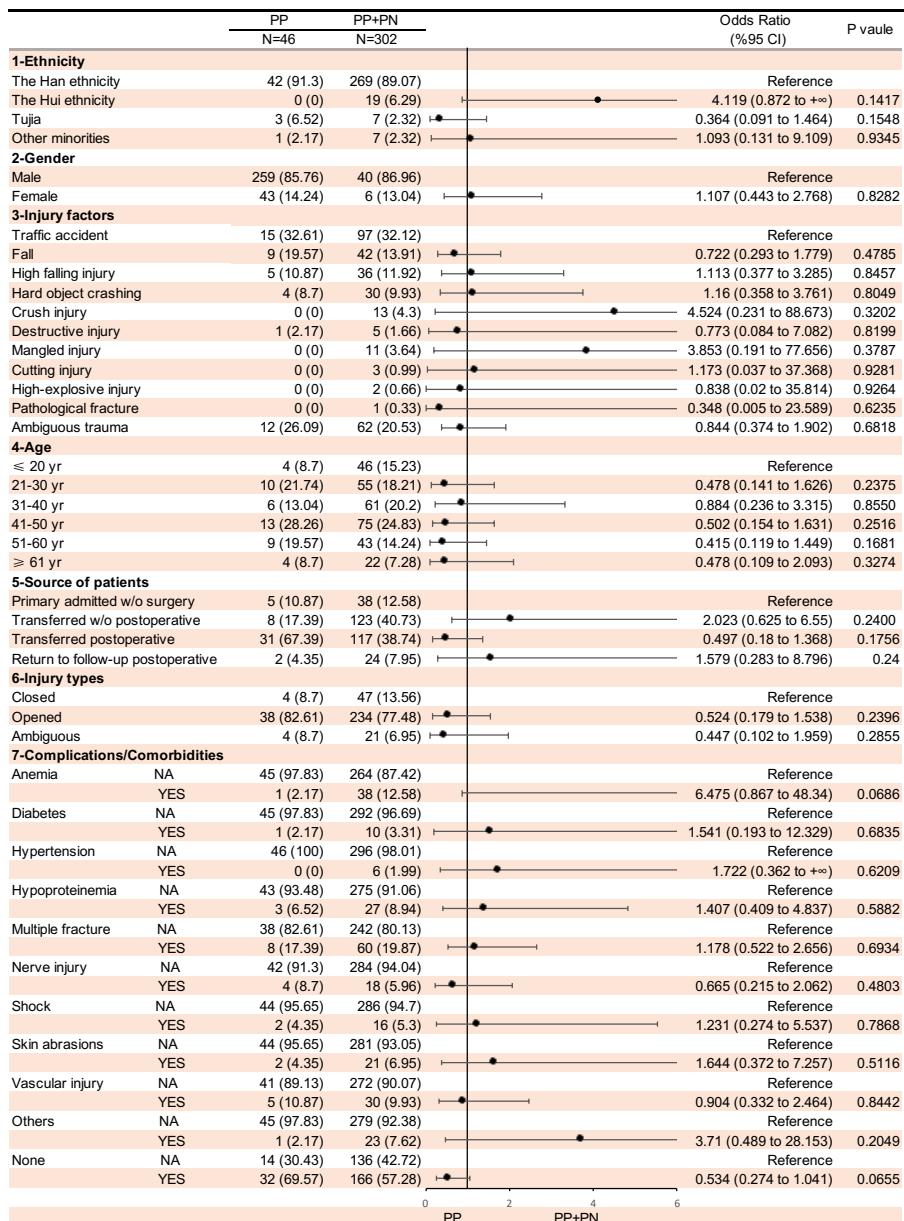
Supplemental Table 5. Univariate analysis of logistic regression was conducted to identify relative risk factors for patients with polymicrobial positive (PP) or polymicrobial negative (PN) infections.



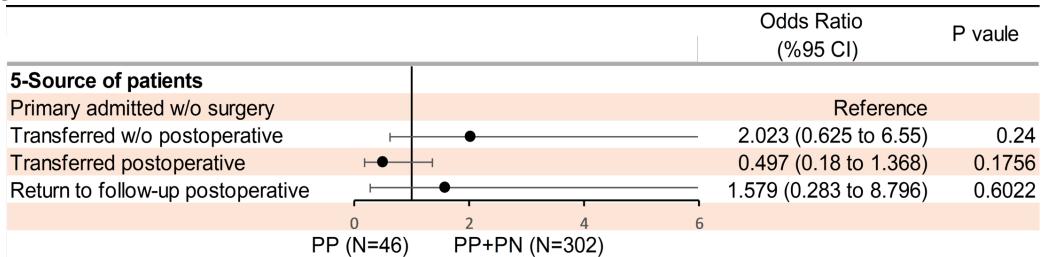
Supplemental Table 6. Multivariate analysis of logistic regression was conducted to identify the relative risk factors for patients with polymicrobial positive (PP) or polymicrobial negative (PN) infections.



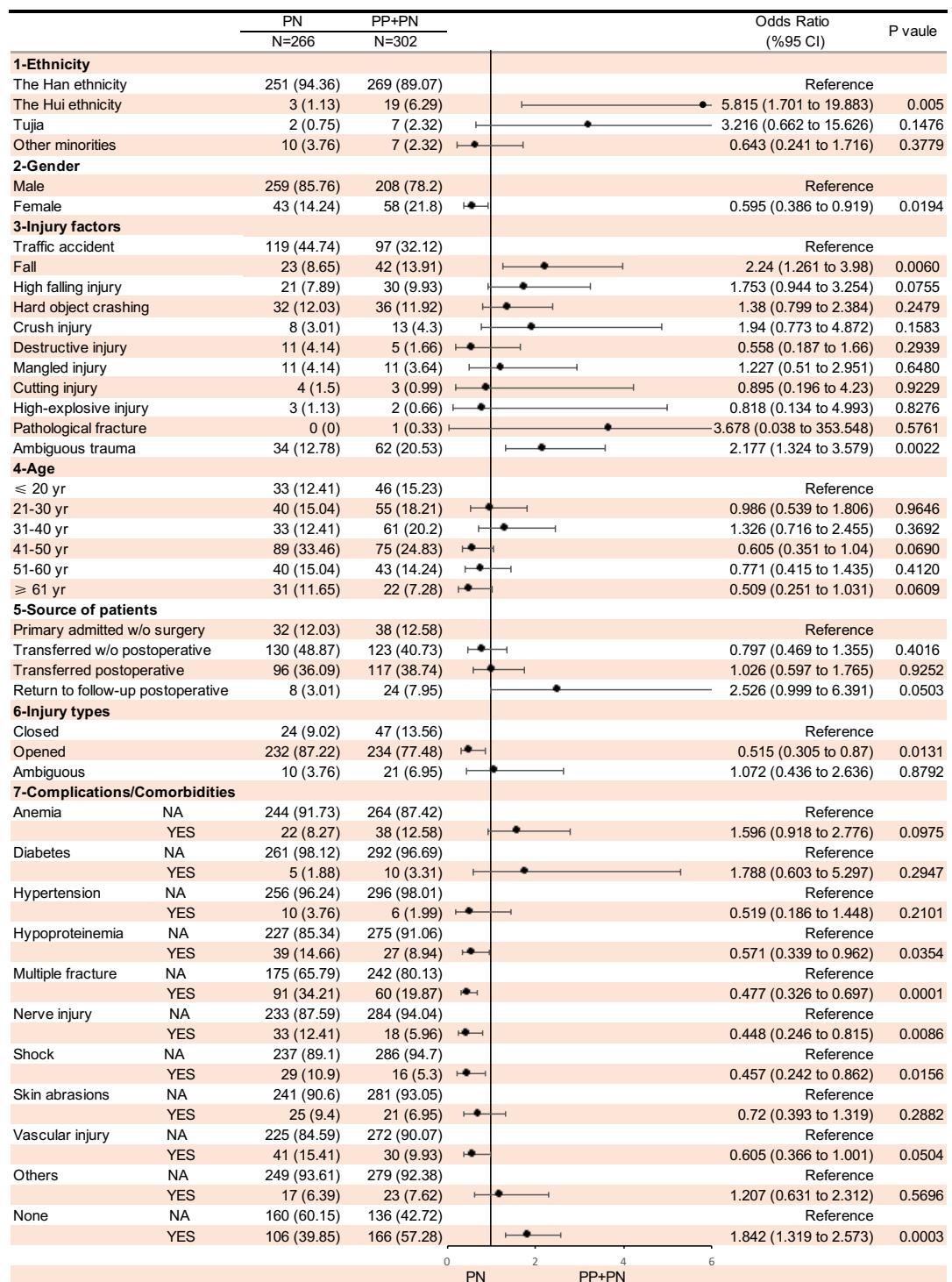
Supplemental Table 7. Univariate analysis of logistic regression was conducted to identify the relative risk factors for patients with polymicrobial positive (PP) or polymicrobial positive/negative (PP+PN) infections.



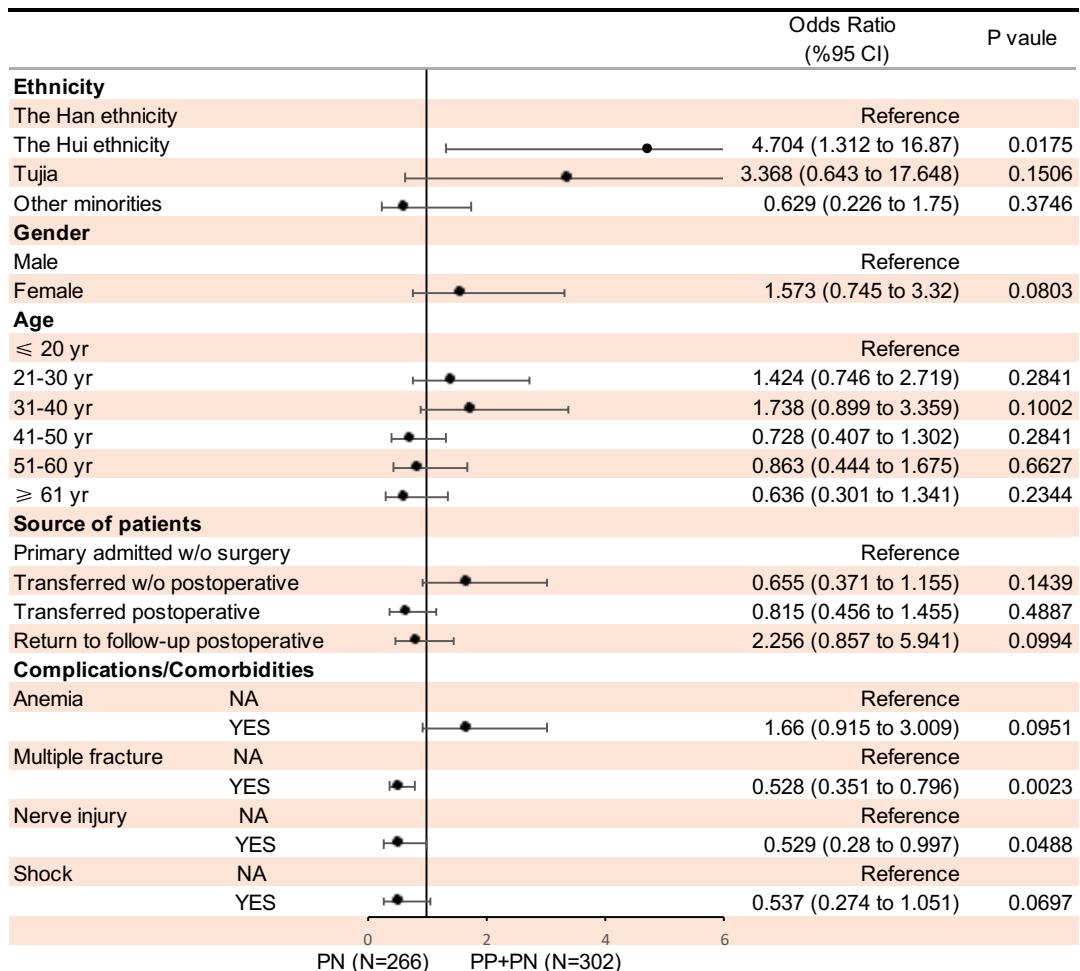
Supplemental Table 8. Multivariate logistic regression analysis was conducted to identify the relative risk factors for patients with polymicrobial positive (PP) or polymicrobial positive/negative (PP+PN) infections.



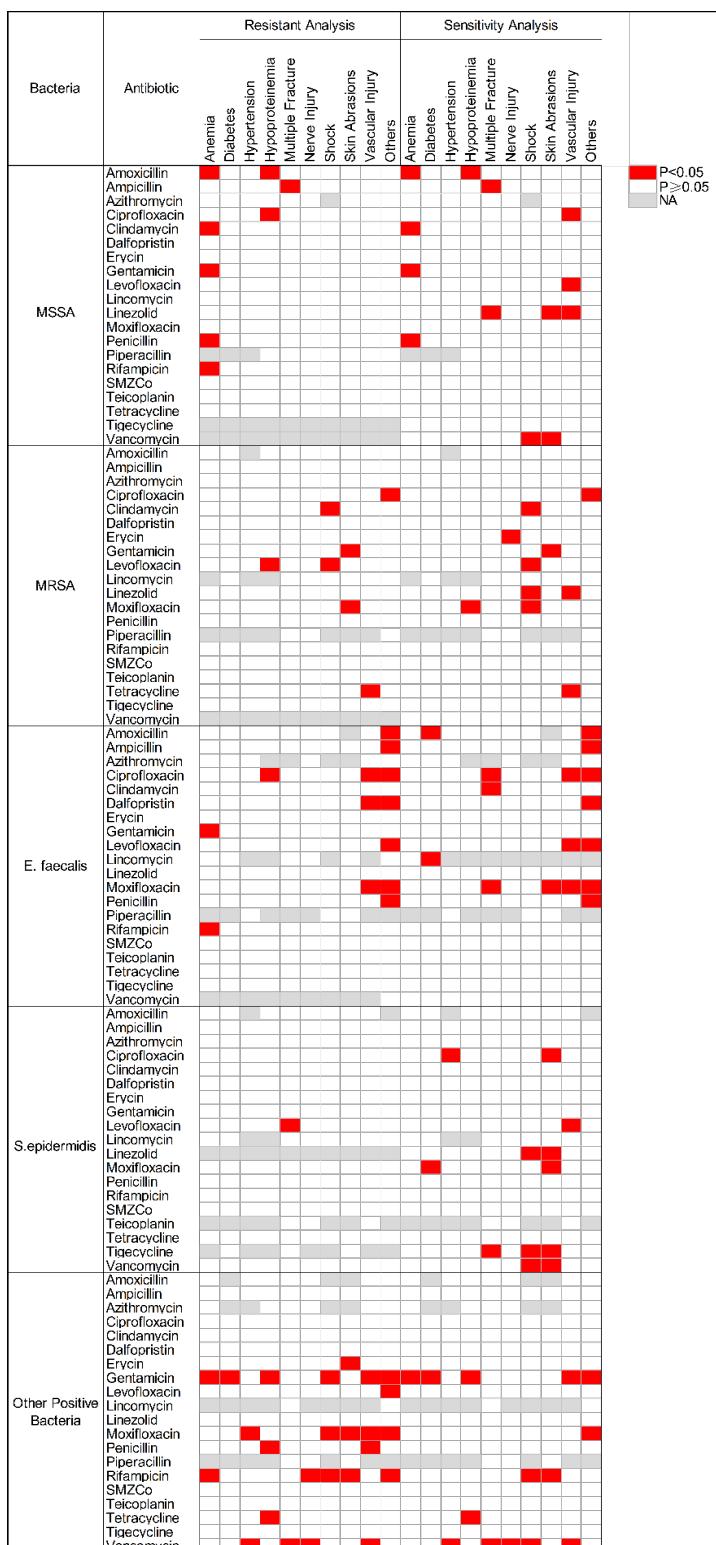
Supplemental Table 9. Univariate analysis of logistic regression was conducted to identify the relative risk factors for patients with polymicrobial negative (PN) or polymicrobial positive/negative infection (PP+PN).



Supplemental Table 10. Multivariate analysis of logistic regression was conducted to identify the relative risk factors for patients with polymicrobial negative (PN) or polymicrobial positive/negative (PP+PN) infections.



Supplemental Figure 1. The statistical analysis results show the antibiotic resistance and sensitivity of Gram-positive bacteria in twenty commonly available antibiotics, under the influence of complications or comorbidities. A p-value less than 0.05 is considered statistically significant, while NA indicates that no data was available.



P<0.05

P≥0.05

NA

Supplemental Figure 2. The statistical analysis yielded results comparing antibiotic resistance and sensitivity in Gram-negative bacteria across twenty commonly available antibiotics, while accounting for the influence of complications or comorbidities. A *p*-value of less than 0.05 was considered statistically significant, and 'NA' indicates that no data was available.

