

Table S1 Observed valid sequence and OUT numbers in different samples

sample	Total reads	Identified reads	OUTs
C-0-D <sub>0</sub>	14892	14466	193
C-0-D <sub>12</sub>	19965	19802	41

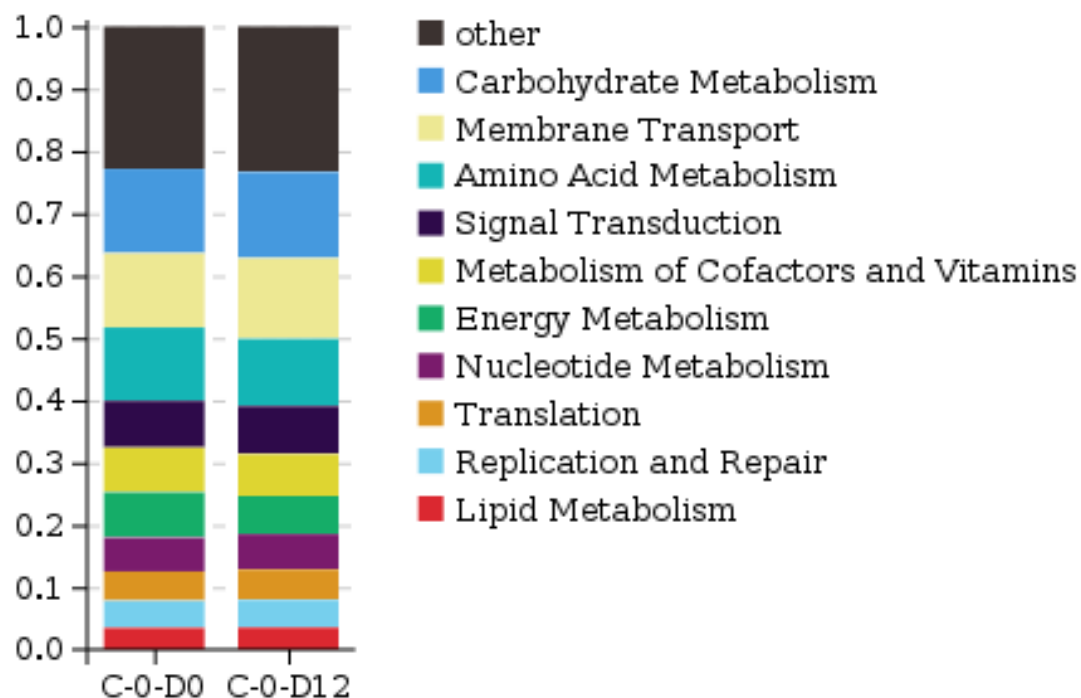


Figure.S1 Function stacking map

Table S2 Techniques in monitoring spoilage microbiota

Techniques	Crustaceans	Bacteria	Level	Alpha diversity	References
16s rRNA high-throughput Next-generation sequencing	<i>Procambarus clarkia</i>	<i>Aeromonas</i> , <i>Macrococcus</i> , <i>Vibrio</i> sp., <i>Acinetobacter</i> , <i>Citrobacter</i> , <i>Enterobacter</i>	Genus	Shannon index 2.5 Simpson index 0.148	[52]
	<i>Procambarus clarkii</i>	<i>Shewanella</i> sp., <i>Carnobacterium</i> sp., <i>Brochothrix</i> sp., <i>Psychrobacter</i> sp., <i>Vagococcus</i> sp., <i>Acinetobacter</i> sp.	Genus	Shannon index 6.58 Simpson index 0.97	[53]
	<i>Litopenaeus vannamei</i>	<i>Vibrio</i> , <i>Candidatus</i> , <i>Bacilloplama</i> , <i>Photobacterium</i> , <i>Candidatus</i> , <i>Bacilloplama</i>	Genus		[54]
	<i>Penaeus vannamei</i>	<i>Proteobacteria</i> <i>Sphingomonas</i> , <i>Carnobacterium</i> , <i>Psychrobacter</i> , <i>Psychro-</i> <i>bacter</i> , <i>Pseudomonas</i> , <i>Shewanella</i> , <i>Acinetobacter</i> , <i>Vibrio</i>	Genus		[55]
	<i>Litopenaeus vannamei</i>	<i>Pseudoalteromonas</i> , <i>Aliivibrio</i> , <i>Photobacterium</i>	Genus		[56]
	<i>Parapenaeus Longirostris</i>	<i>Psychrobacter</i> , <i>Brevundimonas</i> , <i>Stenotrophomonas</i> , <i>Staphylococcus</i> , <i>Legionella</i> , <i>Acinetobacter</i> , <i>Bacillus</i> , <i>Escherichia-Shigella</i> , <i>Enterococcus</i> , <i>Enterobacter</i> , <i>Klebsiella</i>	Genus		[57]
	<i>Solenocera melantho</i>	<i>Vagococcus</i> , <i>Shewanella</i> , <i>Pseudomonas</i> , <i>Psychrobacter</i> , <i>Trichococcus</i>	Genus		[58]
	<i>Pacific White Shrimp</i>	<i>Acinetobacter</i> , <i>Psychrobacter</i> , <i>Shewanella</i> , <i>Carnobacterium</i> , <i>Pseudomonas</i> , <i>Vibrio</i>	Genus		[59]
	<i>Ctenopharyngodon idellus</i>	<i>Pseudomonas</i> , <i>Aeromonas</i> , <i>Shewanella</i>	Genus		[60]
	<i>Procambarus clarkii</i>	<i>Aeromonas jandaei</i>	Species		[61]
16S rRNA sequencing	<i>Litopenaeus Vannamei</i>	<i>lactic acid bacteria</i> , <i>Pseudomonas</i> spp., <i>Shewanella</i> spp., <i>Aeromonas</i> spp.	Genus		[62]

	<i>Litopenaeus Vannamei</i>	<i>Enterobacter, Acinetobacter, Pseudomonas, Aeromonas, Aeromonas, Enterococcus</i>	Genus		[63]
	<i>Penaeus notialis</i>	<i>lactic acid bacteria, Enterobacteriaceae, Pseudomonas spp.</i>	Genus		[64]
<b>Culture dependent API kits</b>	<i>Fenneropenaeus indicus</i>	<i>Actinomyces, Sphingobacterium multivorum, Vibrio proteolyticus, Staphylococcus cohnii, S. auricularis</i>	Species	Simpson index 0.9	[65]
<b>DGGE</b>	<i>Exopalaemon Modemus</i>	<i>Pseudomonas sp., Shewanella, Flavobacterium, Staphylococcus</i>	Genus		[66]
	<i>Crangon crangon</i>	<i>lactic acid bacteria, Pseudomonas sp., Shewanella, Flavobacterium</i>			
		<i>Psychrobacter immobilis, Psychrobacter cibarius. Pseudoalteromonas nigrifaciens, Pseudoalteromonas elyakovii, Pseudoalteromonas paragorgicola</i>	Species		[67]
		<i>Planococcus, Exiguobacterium, Carnobacterium, Pseudomonas, Chryseobacterium Staphylococcus</i>			
<b>PMA combined with DGGE</b>	<i>Litopenaeus vannamei</i>	<i>Acinetobacter sp., Vibrio sp., Aeromonas sp., Lactococcus sp., Exiguobacterium sp., Kurthia sp..</i>	Genus	Shannon index nearly 2.8	[60]
<b>TTGE</b>	<i>Litopenaeus vannamei</i>	<i>C. maltaromaticum and S. baltica.</i>	Species		[61]

