

Table S1. List of EMBL accession numbers for the haplotypes of *Palmorchestia* populations and other species examined.

Taxa	Population code	Acc. numbers Mitochondrial	Acc. numbers Histone H3
<i>Palmorchestia hypogaea</i>	GA	AM749316-21 AM749329-30	AM748651
	AG	AM749379-83	AM748653
	AR	AM749305-08	AM748649-50
	MA	AM749302-04 AM749311-18	--
	PA	AM749372-78	--
	RA	AM749322-28 AM749331	AM748652
	BU	AM749391	--
			--
	JA	AM749384-90	--
	AV	AM749370	--
<i>P. epigaea</i>	FR	AM749364-69	AM748647
	CG	AM749332-41	AM748648
	BG	AM749340	--
		AM749342-49	AM748654-57
		AM749353-55	AM748658-59
<i>Orchestia guancha</i>		AM749358-59	AM748660-61
<i>O. gomeri</i>		AM749356	AM748665
<i>O. sp</i>		AM749350-51	AM748662-64
<i>O. gammarellus</i>		AM749357	
<i>O. stephenseni</i>		AM749360-63	
<i>Talitroides alluaudi</i>		AM749352	AM748646

Figure S1. Bayesian tree obtained in a combined analysis using mitochondrial DNA and nuclear histone H3 sequences. Values above nodes correspond to posterior probability values.

