Supplemental Figure S2. Allelic expression bias and gene expression level Gene expression levels of each sample were arbitrarily grouped into 5 category (1<CPM<10; 10<CPM<30, 30<CPM<100, 100<CPM<300, CPM>300), number of genes that exhibited higher *P. latipinna* allele expression and number of genes that exhibited higher *P. mexicana* allele expression were used to calculate the ratio for each expression level group. For each sample, ratio values were listed as low to high expression groups.