

Table 2 Genetic distance data (Tamura 3-parameter model) between mtCOI sequences

Sl. no.	Species	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
1	<i>O. venusta (Lak)</i>																				
2	<i>O. venusta typica (Med)</i>	43%																			
3	<i>O. media (Lak)</i>	46%	55%																		
4	<i>O. curta</i>	38%	62%	33%																	
5	<i>O. mediterranea</i>	47%	52%	46%	41%																
6	<i>O. cf. mediterranea</i>	46%	54%	38%	35%	18%															
7	<i>O. parabathyalis</i>	41%	48%	35%	36%	31%	28%														
8	<i>O. prendeli</i>	40%	55%	43%	39%	33%	32%	26%													
9	<i>O. scottodicarloi</i>	41%	58%	32%	23%	34%	34%	30%	34%												
10	<i>O. shmelevi</i>	46%	49%	44%	42%	45%	45%	34%	39%	39%											
11	<i>O. waldemari</i>	39%	61%	33%	2%	40%	35%	34%	37%	25%	43%										
12	<i>Oncaea. sp. 7</i>	39%	54%	39%	35%	34%	35%	22%	29%	32%	35%	33%									
13	<i>D. rigida</i>	64%	43%	59%	54%	63%	62%	47%	54%	50%	49%	56%	60%								
14	<i>O. dissimiis</i>	48%	50%	49%	53%	55%	49%	45%	53%	49%	48%	51%	46%	48%							
15	<i>O. nana</i>	39%	54%	50%	43%	46%	37%	33%	36%	41%	50%	45%	35%	57%	52%						
16	<i>O. simiis</i>	41%	31%	52%	52%	59%	53%	47%	49%	57%	52%	55%	58%	45%	55%	59%					
17	<i>O. simplex</i>	44%	50%	44%	47%	48%	36%	34%	34%	49%	43%	46%	41%	52%	54%	36%	46%				
18	<i>O. oculata</i>	53%	48%	51%	56%	61%	59%	57%	59%	62%	50%	59%	61%	41%	51%	55%	47%	49%			
19	<i>O. attenuata</i>	45%	52%	50%	47%	48%	46%	44%	47%	52%	49%	49%	44%	42%	51%	43%	55%	42%	45%		
20	<i>P. nana</i>	390%	488%	464%	465%	395%	431%	299%	328%	406%	396%	348%	515%	297%	811%	297%	795%	364%	473%	476%	