

Table 2. *O. fasciatus* transcriptome assembly statistics

	Full Assembly	Normalized Assembly	Non-Normalized Assembly
Assembled reads (base pairs)	1,773,450 (508,738,047)	389,605 (84,353,140)	336,568 (108,372,883)
Isogroups ("genes")	16,629	10,581	7,591
Isotigs ("transcripts")	21,097	11,353	8,346
Isotig N50	1,735	846	1,162
Mean # isotigs per isogroup	1.3	1.1	1.1
Contigs ("exons")	22,235	11,839	8,731
Mean # contigs per isotig	1.9	1.2	1.3
Singletons (singletons after secondary cap3 assembly)	178,770 (112,531)	110,265 (N/A)	52,585 (N/A)