

Samples		Number of reads (raw sequence)	Number of reads that matched to the <i>C. elegans</i> genome	Known miRNAs	Known 21U-RNAs	tRNAs	rRNAs	snoRNAs	Other reads	
wild-type N2 hermaphrodites	Embryo	14251981	5742750	1905997	70888	10308	1220025	5532	2530000	
			%	33.19	1.23	0.18	21.24	0.10	44.00	
	mid-L1	10013600	5617234	3432825	48300	4274	293218	2462	1836155	
			%	61.11	0.86	0.08	5.22	0.04	32.69	
	mid-L2	10031813	6047597	4016452	25978	2702	300577	1652	1700236	
			%	66.41	0.43	0.04	4.97	0.03	28.11	
	mid-L3	9191014	4948026	3125869	32624	2135	213152	1723	1572523	
			%	63.17	0.66	0.04	4.31	0.03	31.78	
	mid-L4	8917357	6072252	4162442	249374	1890	83359	611	1574576	
			%	68.55	4.11	0.03	1.37	0.01	25.93	
	young adult	9287316	5975243	3854485	407042	4925	102615	702	1605474	
			%	64.51	6.81	0.08	1.72	0.01	26.87	
	males <i>dpy-28;him-8</i>	young adult	11985021	7602104	3863173	28273	47937	1134185	5532	2523004
				%	50.82	0.37	0.63	14.92	0.07	33.19
	Total number of sequence reads		73678102	42005206						