

Table 1. Summary of the sequencing and mapping the data to the *C. elegans* transcriptome

	N2 (wildtype)	Stressed N2	Unstressed <i>aak-2</i>	Stressed <i>aak-2</i>
Total sequence reads	12,446,501	8,419,728	11,860,313	9,086,838
Unambiguously mapped to <i>C. elegans</i> transcriptome (%)	7,486,131	5,408,283	8,418,505	5,857,401
	60.15%	64.23%	70.98%	64.46%
*Genes identified from mapping (%)	20,267	20,258	19,892	20,441
	85.55%	85.51%	83.96%	86.28%

*WormBase version WS180 was used for this analysis. Total number of genes in WS180 is 23,691.