

Table S12. Estimated accuracy of gene prediction

	High-expression transcripts		mRNA (NCBI)	
	Sensitivity	Specificity	Sensitivity	Specificity
Gene	69.58%	78.39%	21.67%	31.43%
Transcript	42.25%	75.74%	21.67%	31.43%
Exon	71.92%	92.62%	69.74%	43.09%
Nucleotide	95.27%	98.44%	94.18%	55.27%