

nGASP Results*

Sn - Sensitivity

Sp - Specificity

MAX (MEDIAN)	Nucleotide		Exon		Transcript		Gene	
	Sn	Sp	Sn	Sp	Sn	Sp	Sn	Sp
Cat 1 ab initio	98.3 (96.8)	91.6 (88.6)	86.4 (83.7)	78.7 (72.8)	53.9 (45.7)	40.7 (29.0)	64.6 (55.3)	40.8 (30.5)
Cat 2 multi genome	97.7 (97.4)	90.9 (90.0)	85.8 (83.5)	78.4 (72.8)	52.6 (52.1)	40.6 (29.7)	63.6 (61.7)	40.6 (29.7)
Cat 3 expression	99.0 (98.2)	95.4 (90.5)	92.5 (88.5)	80.9 (72.2)	68.2 (55.6)	53.1 (33.6)	81.0 (68.8)	53.1 (34.8)
Cat 4 combiners	99.4 (98.9)	93.2 (88.5)	94.3 (90.5)	87.4 (77.4)	75.1 (63.2)	59.0 (39.8)	84.3 (77.9)	59.1 (41.2)

*cross-validation in progress

16th International *C. elegans* Meeting
UCLA

