

Genomics Track results, categorization task — California State University San Marcos

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
Tcsusm1	1	40	19	0.0244	0.0500	0.0328	231.00	191	4620	0.0413
Tcsusm2	1	38	19	0.0256	0.0500	0.0339	231.00	193	4620	0.0418