

Genomics Track results, categorization task — LangPower Computing, Inc.

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
LPC6	143	191	189	0.4281	0.4307	0.4294	17.00	2240	5644	0.3969

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
LPC7	11	20	9	0.3548	0.5500	0.4314	231.00	2521	4620	0.5457