Genomics Track results, categorization task — National Taiwan University

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aNTUMAC	315	601	17	0.3439	0.9488	0.5048	17.00	4754	5644	0.8423

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eNTUMAC	86	454	19	0.1593	0.8190	0.2667	64.00	5050	6720	0.7515

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gNTUMAC	456	1978	62	0.1873	0.8803	0.3089	11.00	3038	5698	0.5332

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tNTUMAC	18	324	2	0.0526	0.9000	0.0994	231.00	3834	4620	0.8299
tNTUMACasem	13	370	7	0.0339	0.6500	0.0645	231.00	2633	4620	0.5699
tNTUMACwj	19	348	1	0.0518	0.9500	0.0982	231.00	4041	4620	0.8747