

Genomics Track results, categorization task — Tsinghua University

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
THUIRgA0p9x	288	244	44	0.5414	0.8675	0.6667	17.00	4652	5644	0.8242
THUIRgenA1p1	299	311	33	0.4902	0.9006	0.6348	17.00	4772	5644	0.8455

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
THUIRgenE1p8	97	637	8	0.1322	0.9238	0.2312	64.00	5571	6720	0.8290

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
THUIRgenG1p1	337	1508	181	0.1827	0.6506	0.2852	11.00	2199	5698	0.3859
THUIRgenGMNG	351	1315	167	0.2107	0.6776	0.3214	11.00	2546	5698	0.4468

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
THUIRgenT1p5	19	873	1	0.0213	0.9500	0.0417	231.00	3516	4620	0.7610