

Genomics Track results, categorization task — Erasmus MC

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
ABPLUS	296	932	36	0.2410	0.8916	0.3795	17.00	4100	5644	0.7264
FTA	130	235	202	0.3562	0.3916	0.3730	17.00	1975	5644	0.3499

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
ABPLUSE	86	937	19	0.0841	0.8190	0.1525	64.00	4567	6720	0.6796
FTE	43	472	62	0.0835	0.4095	0.1387	64.00	2280	6720	0.3393

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
ABPLUSG	376	1350	142	0.2178	0.7259	0.3351	11.00	2786	5698	0.4889
FTG	149	525	369	0.2211	0.2876	0.2500	11.00	1114	5698	0.1955

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
ABPLUST	19	587	1	0.0314	0.9500	0.0607	231.00	3802	4620	0.8229
FTT	10	102	10	0.0893	0.5000	0.1515	231.00	2208	4620	0.4779