

Genomics Track results, categorization task — IBM India Research Laboratory

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aIBMIRLmet	280	595	52	0.3200	0.8434	0.4640	17.00	4165	5644	0.7380
aIBMIRLrul	294	629	38	0.3185	0.8855	0.4685	17.00	4369	5644	0.7741
aIBMIRLsvm	297	699	35	0.2982	0.8946	0.4473	17.00	4350	5644	0.7707

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eIBMIRLmet	77	1151	28	0.0627	0.7333	0.1155	64.00	3777	6720	0.5621
eIBMIRLrul	76	1108	29	0.0642	0.7238	0.1179	64.00	3756	6720	0.5589
eIBMIRLsvm	97	1602	8	0.0571	0.9238	0.1075	64.00	4606	6720	0.6854

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gIBMIRLmet	467	1836	51	0.2028	0.9015	0.3311	11.00	3301	5698	0.5793
gIBMIRLrul	481	2073	37	0.1883	0.9286	0.3132	11.00	3218	5698	0.5648
gIBMIRLsvm	449	1721	69	0.2069	0.8668	0.3341	11.00	3218	5698	0.5648

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tIBMIRLmet	19	182	1	0.0945	0.9500	0.1719	231.00	4207	4620	0.9106
tIBMIRLrul	19	439	1	0.0415	0.9500	0.0795	231.00	3950	4620	0.8550
tIBMIRLsvm	16	504	4	0.0308	0.8000	0.0593	231.00	3192	4620	0.6909