Genomics Track results, categorization task — IBM TJ Watson Research Center (Zhang)

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aibmadz05m1	299	290	33	0.5076	0.9006	0.6493	17.00	4793	5644	0.8492
aibmadz05m2	299	296	33	0.5025	0.9006	0.6451	17.00	4787	5644	0.8482
aibmadz05s	310	354	22	0.4669	0.9337	0.6225	17.00	4916	5644	0.8710

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eibmadz05m1	92	326	13	0.2201	0.8762	0.3518	64.00	5562	6720	0.8277
eibmadz05m2	93	348	12	0.2109	0.8857	0.3407	64.00	5604	6720	0.8339
eibmadz05s	98	584	7	0.1437	0.9333	0.2490	64.00	5688	6720	0.8464

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gibmadz05m1	320	675	198	0.3216	0.6178	0.4230	11.00	2845	5698	0.4993
gibmadz05m2	322	691	196	0.3179	0.6216	0.4206	11.00	2851	5698	0.5004
gibmadz05s	302	634	216	0.3226	0.5830	0.4154	11.00	2688	5698	0.4717

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tibmadz05m1	19	263	1	0.0674	0.9500	0.1258	231.00	4126	4620	0.8931
tibmadz05m2	19	232	1	0.0757	0.9500	0.1402	231.00	4157	4620	0.8998
tibmadz05s	19	257	1	0.0688	0.9500	0.1284	231.00	4132	4620	0.8944