Genomics Track results, categorization task — University of Illinois at Urbana-Champaign

allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aMUSCUIUC1	201	100	131	0.6678	0.6054	0.6351	17.00	3317	5644	0.5877
aMUSCUIUC2	258	211	74	0.5501	0.7771	0.6442	17.00	4175	5644	0.7397
aMUSCUIUC3	268	358	64	0.4281	0.8072	0.5595	17.00	4198	5644	0.7438

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eMUSCUIUC1	49	167	56	0.2269	0.4667	0.3053	64.00	2969	6720	0.4418
eMUSCUIUC2	0	50	105	0.0000	0.0000	0.0000	64.00	-50	6720	-0.0074
eMUSCUIUC3	50	268	55	0.1572	0.4762	0.2364	64.00	2932	6720	0.4363

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gMUSCUIUC1	145	224	373	0.3930	0.2799	0.3269	11.00	1371	5698	0.2406
gMUSCUIUC2	90	1185	428	0.0706	0.1737	0.1004	11.00	-195	5698	-0.0342
gMUSCUIUC3	179	1831	339	0.0891	0.3456	0.1416	11.00	138	5698	0.0242

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tMUSCUIUC1	12	23	8	0.3429	0.6000	0.4364	231.00	2749	4620	0.5950
tMUSCUIUC2	14	155	6	0.0828	0.7000	0.1481	231.00	3079	4620	0.6665
tMUSCUIUC3	14	30	6	0.3182	0.7000	0.4375	231.00	3204	4620	0.6935