

Genomics Track results, categorization task — UCHSC at Fitzsimons (University of Colorado School of**allele subtask**

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aUCHSCnb1En3	254	246	78	0.5080	0.7651	0.6106	17.00	4072	5644	0.7215
aUCHSCnb1En4	215	138	117	0.6091	0.6476	0.6277	17.00	3517	5644	0.6231
aUCHSCsvm	148	38	184	0.7957	0.4458	0.5714	17.00	2478	5644	0.4391

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eUCHSCnb1En3	36	27	69	0.5714	0.3429	0.4286	64.00	2277	6720	0.3388
eUCHSCnb1En4	39	36	66	0.5200	0.3714	0.4333	64.00	2460	6720	0.3661
eUCHSCsvm	4	0	101	1.0000	0.0381	0.0734	64.00	256	6720	0.0381

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gUCHSCnb1En3	177	241	341	0.4234	0.3417	0.3782	11.00	1706	5698	0.2994
gUCHSCnb1En4	92	74	426	0.5542	0.1776	0.2690	11.00	938	5698	0.1646
gUCHSCsvm	95	139	423	0.4060	0.1834	0.2527	11.00	906	5698	0.1590

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tUCHSCnb1En3	6	25	14	0.1935	0.3000	0.2353	231.00	1361	4620	0.2946
tUCHSCnb1En4	3	5	17	0.3750	0.1500	0.2143	231.00	688	4620	0.1489
tUCHSCsvm	1	0	19	1.0000	0.0500	0.0952	231.00	231	4620	0.0500