allele subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aQUNB8	281	602	51	0.3182	0.8464	0.4626	17.00	4175	5644	0.7397
aQUT11	257	422	75	0.3785	0.7741	0.5084	17.00	3947	5644	0.6993
aQUT14	288	516	44	0.3582	0.8675	0.5070	17.00	4380	5644	0.7760

expression subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eQUNB11	67	550	38	0.1086	0.6381	0.1856	64.00	3738	6720	0.5563
eQUNB19	48	376	57	0.1132	0.4571	0.1815	64.00	2696	6720	0.4012
eQUT18	55	514	50	0.0967	0.5238	0.1632	64.00	3006	6720	0.4473

go subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gQUNB12	342	1791	176	0.1603	0.6602	0.2580	11.00	1971	5698	0.3459
gQUNB15	294	1105	224	0.2102	0.5676	0.3067	11.00	2129	5698	0.3736
gQUT22	319	1442	199	0.1811	0.6158	0.2799	11.00	2067	5698	0.3628

tumor subtask

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tQUNB3	18	721	2	0.0244	0.9000	0.0474	231.00	3437	4620	0.7439
tQUT10	20	1498	0	0.0132	1.0000	0.0260	231.00	3122	4620	0.6758
tQUT14	13	29	7	0.3095	0.6500	0.4194	231.00	2974	4620	0.6437