

**Genomics Track results, categorization task — Fudan University (Web Info. Mining)**

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**allele subtask**

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
aFduMarsI	299	330	33	0.4754	0.9006	0.6223	17.00	4753	5644	0.8421
aFduMarsII	305	422	27	0.4195	0.9187	0.5760	17.00	4763	5644	0.8439
aFduMarsIII	302	626	30	0.3254	0.9096	0.4794	17.00	4508	5644	0.7987

**expression subtask**

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
eFduMarsI	98	418	7	0.1899	0.9333	0.3156	64.00	5854	6720	0.8711
eFduMarsII	98	418	7	0.1899	0.9333	0.3156	64.00	5854	6720	0.8711
eFduMarsIII	100	1159	5	0.0794	0.9524	0.1466	64.00	5241	6720	0.7799

**go subtask**

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
gFduMarsI	403	1121	115	0.2644	0.7780	0.3947	11.00	3312	5698	0.5813
gFduMarsII	459	1704	59	0.2122	0.8861	0.3424	11.00	3345	5698	0.5870
gFduMarsIII	471	1995	47	0.1910	0.9093	0.3157	11.00	3186	5698	0.5591

**tumor subtask**

Run	TP	FP	FN	Precision	Recall	F beta	Utility Factor	Raw	Max	Normalized
tFduMarsI	19	160	1	0.1061	0.9500	0.1910	231.00	4229	4620	0.9154
tFduMarsII	19	173	1	0.0990	0.9500	0.1792	231.00	4216	4620	0.9126
tFduMarsIII	20	680	0	0.0286	1.0000	0.0556	231.00	3940	4620	0.8528