

**Summary of TCGA Tumor Data
Ingested into Broad GDAC Pipeline
January 14, 2010 Run**

Tumor Type	Biospecimen	Any_Level_1	Clinical	CNA	Methylation	mRNA	miR	MAF
BRCA	346	186	244	265	186	280	0	0
COAD	203	151	130	137	167	155	0	64
GBM	508	448	490	466	288	444	415	169
HNSC	39	0	0	0	0	0	0	0
KIRC	355	39	19	254	219	41	0	0
KIRP	48	39	0	16	36	41	0	0
LAML	202	0	0	0	188	0	0	0
LGG	30	0	0	0	0	0	0	0
LUAD	128	21	11	56	128	33	0	0
LUSC	160	116	42	117	133	116	0	0
OV	584	570	532	519	425	519	566	384
READ	79	52	72	51	69	69	0	12
STAD	82	35	0	81	82	0	0	0
UCEC	145	24	0	114	70	0	0	0
Totals	2909	1681	1540	2076	1991	1698	981	629