

**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

**Summary of TCGA Tumor Data
Ingested into Broad GDAC Pipeline
December 23, 2010 Run**

Tumor Type	Biospecimen	Any_Level_1	Clinical	CNA	Methylation	mRNA	miR	MAF
BRCA	319	186	194	176	186	280	0	0
COAD	203	151	130	137	167	155	0	0
GBM	0	448	0	466	288	444	415	0
HNSC	39	0	0	0	0	0	0	0
KIRC	355	39	19	73	219	41	0	0
KIRP	48	39	0	39	36	41	0	0
LAML	202	0	0	0	188	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	50	69	69	0	0
STAD	82	35	0	81	82	0	0	0
UCEC	125	24	0	68	70	0	0	0
Totals	2324	1681	1000	1782	1991	1698	981	384



Workspaces | Samples | Individuals | Pipelines | Annotation Setup | Connectors | Help Workspace: prod__2011_01_14__ov__01

PIPELINE WORKFLOW STATUS

Show only jobs in: For workflow:

Start Jobs With Priority:

	Pipeline	Reports	Job Count	Not Ready	Analysis Ready	In Process	Failed	Complete
1	convertCNLevelIIIData		1	0%	0%	0%	0%	100%
2	Find miR Direct Targets	view	1	0%	0%	0%	0%	100%
3	GDAC CNMF miRNA clustering	view	1	0%	0%	0%	0%	100%
4	GDAC Correlate Expression with Methylation	view	1	0%	0%	0%	0%	100%
5	GDAC median mRNA Expression		1	0%	0%	0%	0%	100%
6	GDAC miRNAConsensusClustering	view	1	0%	0%	0%	0%	100%
7	GDAC clinicalDataMergerPipeline clinical		1	0%	0%	0%	0%	100%
8	Gistic2	view	1	0%	0%	0%	0%	100%
9	MutSig	view	1	0%	0%	0%	0%	100%
10	Paradigm		1	0%	0%	0%	0%	100%
11	GDAC CNMF mRNA clustering	view	1	0%	0%	0%	0%	100%
12	GDAC mRNAConsensusClustering	view	1	0%	0%	0%	0%	100%
13	GDAC clinicalDataPickerPipeline clinical		1	0%	0%	0%	0%	100%
14	MakeReducedSegment		1	0%	0%	0%	0%	100%
15	MutationAssessor	view	1	0%	0%	0%	0%	100%
16	Correlate gene mutation status with Clinical data	view	1	0%	0%	0%	0%	100%
17	Correlate microRNA Expression with Clinical Data	view	1	0%	0%	0%	0%	100%
18	Correlate miRNA CNMF clustering with Clinical data	view	1	0%	0%	0%	0%	100%
19	Correlate miRNA consensus clustering with Clinical data	view	1	0%	0%	0%	0%	100%
20	Correlate mRNA CNMF clustering with Clinical data	view	1	0%	0%	0%	0%	100%
21	Correlate mRNA consensus clustering with Clinical data	view	1	0%	0%	0%	0%	100%
22	Correlate mRNA Expression with Clinical data	view	1	0%	0%	0%	0%	100%
23	GetCopyNumberExpCor	view	1	0%	0%	0%	0%	100%
24	GetCopyNumberExpCorMiRNA	view	1	0%	0%	0%	0%	100%
25	GDAC_geneListPathwayEnrichmentPipeline	view	1	0%	0%	0%	0%	100%
Summary			25	0	0	0	0	25



Workspaces | Samples | Individuals | Pipelines | Annotation Setup | Connectors | Help Workspace: prod__2011_01_14_gbm__02

PIPELINE WORKFLOW STATUS

Show only jobs in: For workflow:

Start Jobs With Priority:

	Pipeline	Reports	Job Count	Not Ready	Analysis Ready	In Process	Failed	Complete
1	convertCNLevelIIIData		1	0%	0%	0%	0%	100%
2	Find miR Direct Targets	view	1	0%	0%	0%	0%	100%
3	GDAC CNMF miRNA clustering	view	1	0%	0%	0%	0%	100%
4	GDAC Correlate Expression with Methylation	view	1	0%	0%	0%	0%	100%
5	GDAC median mRNA Expression		1	0%	0%	0%	0%	100%
6	GDAC miRNAConsensusClustering	view	1	0%	0%	0%	0%	100%
7	GDAC clinicalDataMergerPipeline clinical		1	0%	0%	0%	0%	100%
8	Gistic2	view	1	0%	0%	0%	0%	100%
9	MutSig	view	1	0%	0%	0%	0%	100%
10	Paradigm		1	0%	0%	0%	0%	100%
11	GDAC CNMF mRNA clustering	view	1	0%	0%	0%	0%	100%
12	GDAC mRNAConsensusClustering	view	1	0%	0%	0%	0%	100%
13	GDAC clinicalDataPickerPipeline clinical		1	0%	0%	0%	0%	100%
14	MakeReducedSegment		1	0%	0%	0%	0%	100%
15	MutationAssessor	view	1	0%	0%	0%	0%	100%
16	Correlate gene mutation status with Clinical data	view	1	0%	0%	0%	0%	100%
17	Correlate microRNA Expression with Clinical Data	view	1	0%	0%	0%	0%	100%
18	Correlate miRNA CNMF clustering with Clinical data	view	1	0%	0%	0%	0%	100%
19	Correlate miRNA consensus clustering with Clinical data	view	1	0%	0%	0%	0%	100%
20	Correlate mRNA CNMF clustering with Clinical data	view	1	0%	0%	0%	0%	100%
21	Correlate mRNA consensus clustering with Clinical data	view	1	0%	0%	0%	0%	100%
22	Correlate mRNA Expression with Clinical data	view	1	0%	0%	0%	0%	100%
23	GetCopyNumberExpCor	view	1	0%	0%	0%	0%	100%
24	GetCopyNumberExpCorMiRNA	view	1	0%	0%	0%	0%	100%
25	GDAC_geneListPathwayEnrichmentPipeline	view	1	0%	0%	0%	0%	100%
Summary			25	0	0	0	0	25