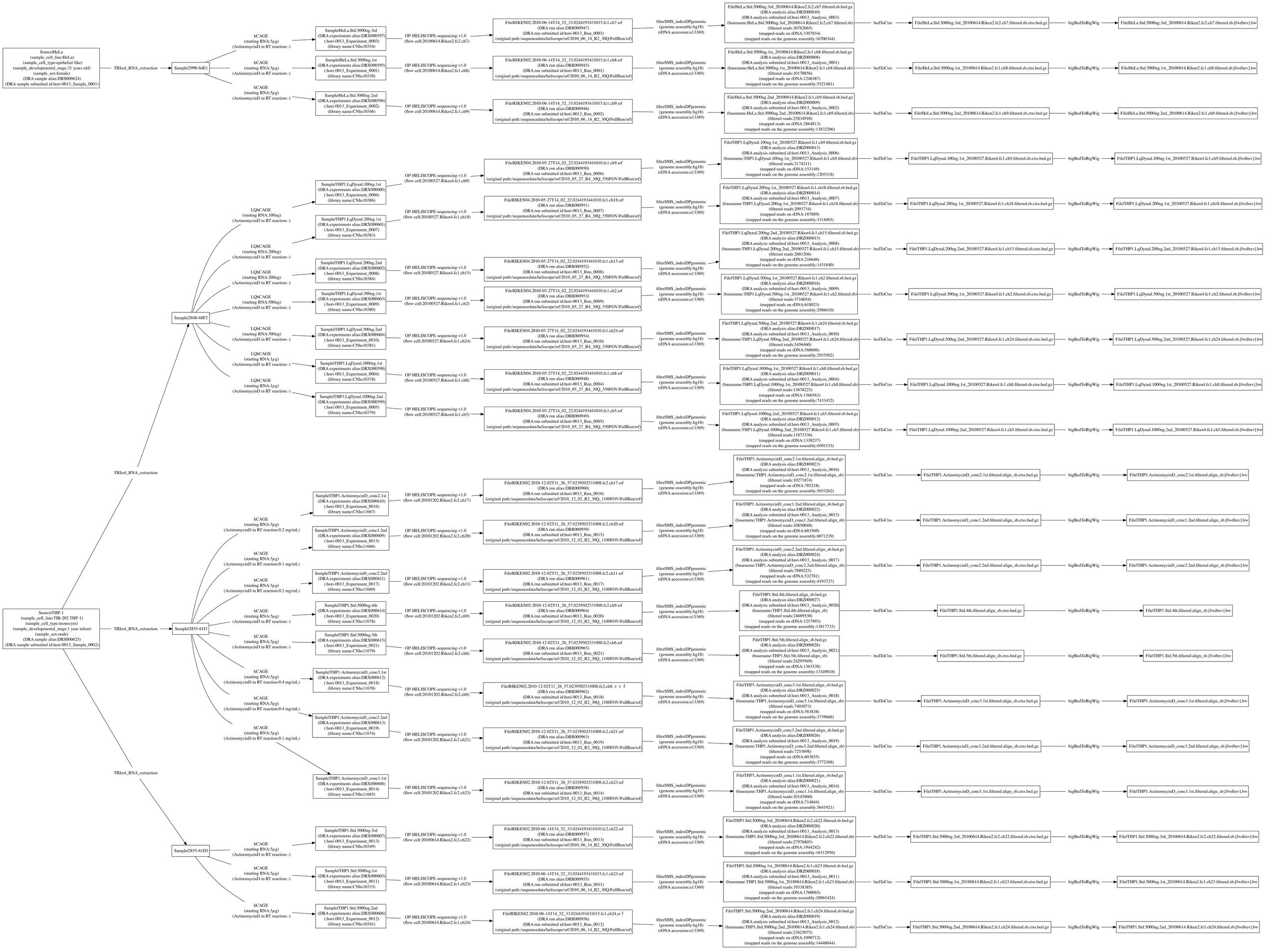
SourceHeLa
(sample_cell_line:HeLa)
(sample_cell_type:epithelial-like)
(sample_developmental_stage:31 years old)
(sample_sex:female)
(DRA sample alias:DRS000624)
(DRA sample submitted id:hori-0013_Sample_0001)
TRIzol_RNA_extraction
Sample2990-64E1
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleHeLa.Std.5000ng.3rd
(DRA experiments alias:DRX000597)
(:hori-0013_Experiment_0003)
(library name:CNhs10354)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100614.Riken2.fc2.ch7)
FileIRIKEN02.2010-06-14T14_52_33.0244193431015.fc1.ch7.srf
(DRA run alias:DRR000947)
(DRA run submitted id:hori-0013_Run_0003)
(original path:/sequencedata/heliscope/srf/2010_06_14_R2_30Q/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileIHeLa.Std.5000ng.3rd_20100614.Riken2.fc2.ch7.filtered.sb.bed.gz
(DRA analysis alias:DRZ000010)
(DRA analysis submitted id:hori-0013_Analysis_0003)
(basename:HeLa.Std.5000ng.3rd_20100614.Riken2.fc2.ch7.filtered.sb)
(filtered reads:30782065)
(mapped reads on rDNA:3307654)
(mapped reads on the genome assembly:16700344)
bedToCtss
FileIHeLa.Std.5000ng.3rd_20100614.Riken2.fc2.ch7.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileIHeLa.Std.5000ng.3rd_20100614.Riken2.fc2.ch7.filtered.sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleHeLa.Std.5000ng.1st
(DRA experiments alias:DRX000595)
(:hori-0013_Experiment_0001)
(library name:CNhs10338)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100614.Riken2.fc1.ch8)
FileIRIKEN02.2010-06-14T14_52_33.0244193431015.fc1.ch8.srf
(DRA run alias:DRR000945)
(DRA run submitted id:hori-0013_Run_0001)
(original path:/sequencedata/heliscope/srf/2010_06_14_R2_30Q/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileIHeLa.Std.5000ng.1st_20100614.Riken2.fc1.ch8.filtered.sb.bed.gz
(DRA analysis alias:DRZ000008)
(DRA analysis submitted id:hori-0013_Analysis_0001)
(basename:HeLa.Std.5000ng.1st_20100614.Riken2.fc1.ch8.filtered.sb)
(filtered reads:10170856)
(mapped reads on rDNA:1246387)
(mapped reads on the genome assembly:5521481)
bedToCtss
FileIHeLa.Std.5000ng.1st_20100614.Riken2.fc1.ch8.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileIHeLa.Std.5000ng.1st_20100614.Riken2.fc1.ch8.filtered.sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleHeLa.Std.5000ng.2nd
(DRA experiments alias:DRX000596)
(:hori-0013_Experiment_0002)
(library name:CNhs10346)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100614.Riken2.fc1.ch9)
FileIRIKEN02.2010-06-14T14_52_33.0244193431015.fc1.ch9.srf
(DRA run alias:DRR000946)
(DRA run submitted id:hori-0013_Run_0002)
(original path:/sequencedata/heliscope/srf/2010_06_14_R2_30Q/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileIHeLa.Std.5000ng.2nd_20100614.Riken2.fc1.ch9.filtered.sb.bed.gz
(DRA analysis alias:DRZ000009)
(DRA analysis submitted id:hori-0013_Analysis_0002)
(basename:HeLa.Std.5000ng.2nd_20100614.Riken2.fc1.ch9.filtered.sb)
(filtered reads:25834910)
(mapped reads on rDNA:2864813)
(mapped reads on the genome assembly:13832206)
bedToCtss
FileIHeLa.Std.5000ng.2nd_20100614.Riken2.fc1.ch9.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileIHeLa.Std.5000ng.2nd_20100614.Riken2.fc1.ch9.filtered.sb.[fwdlrev].bw
Sample2848-68F2
LQbCAGE
(starting RNA:100ng)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.100ng.1st
(DRA experiments alias:DRX000600)
(:hori-0013_Experiment_0006)
(library name:CNhs10386)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch9)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch9.srf
(DRA run alias:DRR000950)
(DRA run submitted id:hori-0013_Run_0006)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.100ng.1st_20100527.Riken4.fc1.ch9.filtered.sb.bed.gz
(DRA analysis alias:DRZ000013)
(DRA analysis submitted id:hori-0013_Analysis_0006)
(basename:THP1.LqDynal.100ng.1st_20100527.Riken4.fc1.ch9.filtered.sb)
(filtered reads:2174211)
(mapped reads on rDNA:153149)
(mapped reads on the genome assembly:1205318)
bedToCtss
FileITHP1.LqDynal.100ng.1st_20100527.Riken4.fc1.ch9.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.100ng.1st_20100527.Riken4.fc1.ch9.filtered.sb.[fwdlrev].bw
LQbCAGE
(starting RNA:200ng)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.200ng.1st
(DRA experiments alias:DRX000601)
(:hori-0013_Experiment_0007)
(library name:CNhs10383)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch18)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch18.srf
(DRA run alias:DRR000951)
(DRA run submitted id:hori-0013_Run_0007)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.200ng.1st_20100527.Riken4.fc1.ch18.filtered.sb.bed.gz
(DRA analysis alias:DRZ000014)
(DRA analysis submitted id:hori-0013_Analysis_0007)
(basename:THP1.LqDynal.200ng.1st_20100527.Riken4.fc1.ch18.filtered.sb)
(filtered reads:2093734)
(mapped reads on rDNA:187889)
(mapped reads on the genome assembly:1114465)
bedToCtss
FileITHP1.LqDynal.200ng.1st_20100527.Riken4.fc1.ch18.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.200ng.1st_20100527.Riken4.fc1.ch18.filtered.sb.[fwdlrev].bw
LQbCAGE
(starting RNA:200ng)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.200ng.2nd
(DRA experiments alias:DRX000602)
(:hori-0013_Experiment_0008)
(library name:CNhs10384)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch15)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch15.srf
(DRA run alias:DRR000952)
(DRA run submitted id:hori-0013_Run_0008)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.200ng.2nd_20100527.Riken4.fc1.ch15.filtered.sb.bed.gz
(DRA analysis alias:DRZ000015)
(DRA analysis submitted id:hori-0013_Analysis_0008)
(basename:THP1.LqDynal.200ng.2nd_20100527.Riken4.fc1.ch15.filtered.sb)
(filtered reads:2681206)
(mapped reads on rDNA:216646)
(mapped reads on the genome assembly:1451840)
bedToCtss
FileITHP1.LqDynal.200ng.2nd_20100527.Riken4.fc1.ch15.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.200ng.2nd_20100527.Riken4.fc1.ch15.filtered.sb.[fwdlrev].bw
LQbCAGE
(starting RNA:500ng)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.500ng.1st
(DRA experiments alias:DRX000603)
(:hori-0013_Experiment_0009)
(library name:CNhs10380)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch2)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch2.srf
(DRA run alias:DRR000953)
(DRA run submitted id:hori-0013_Run_0009)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.500ng.1st_20100527.Riken4.fc1.ch2.filtered.sb.bed.gz
(DRA analysis alias:DRZ000016)
(DRA analysis submitted id:hori-0013_Analysis_0009)
(basename:THP1.LqDynal.500ng.1st_20100527.Riken4.fc1.ch2.filtered.sb)
(filtered reads:5716054)
(mapped reads on rDNA:618023)
(mapped reads on the genome assembly:2986610)
bedToCtss
FileITHP1.LqDynal.500ng.1st_20100527.Riken4.fc1.ch2.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.500ng.1st_20100527.Riken4.fc1.ch2.filtered.sb.[fwdlrev].bw
LQbCAGE
(starting RNA:500ng)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.500ng.2nd
(DRA experiments alias:DRX000604)
(:hori-0013_Experiment_0010)
(library name:CNhs10381)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch24)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch24.srf
(DRA run alias:DRR000954)
(DRA run submitted id:hori-0013_Run_0010)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.500ng.2nd_20100527.Riken4.fc1.ch24.filtered.sb.bed.gz
(DRA analysis alias:DRZ000017)
(DRA analysis submitted id:hori-0013_Analysis_0010)
(basename:THP1.LqDynal.500ng.2nd_20100527.Riken4.fc1.ch24.filtered.sb)
(filtered reads:5456460)
(mapped reads on rDNA:560686)
(mapped reads on the genome assembly:2935502)
bedToCtss
FileITHP1.LqDynal.500ng.2nd_20100527.Riken4.fc1.ch24.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.500ng.2nd_20100527.Riken4.fc1.ch24.filtered.sb.[fwdlrev].bw
LQbCAGE
(starting RNA:1μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.1000ng.1st
(DRA experiments alias:DRX000598)
(:hori-0013_Experiment_0004)
(library name:CNhs10378)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch8)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch8.srf
(DRA run alias:DRR000948)
(DRA run submitted id:hori-0013_Run_0004)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.1000ng.1st_20100527.Riken4.fc1.ch8.filtered.sb.bed.gz
(DRA analysis alias:DRZ000011)
(DRA analysis submitted id:hori-0013_Analysis_0004)
(basename:THP1.LqDynal.1000ng.1st_20100527.Riken4.fc1.ch8.filtered.sb)
(filtered reads:13678225)
(mapped reads on rDNA:1566563)
(mapped reads on the genome assembly:7433452)
bedToCtss
FileITHP1.LqDynal.1000ng.1st_20100527.Riken4.fc1.ch8.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.1000ng.1st_20100527.Riken4.fc1.ch8.filtered.sb.[fwdlrev].bw
LQbCAGE
(starting RNA:1μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.LqDynal.1000ng.2nd
(DRA experiments alias:DRX000599)
(:hori-0013_Experiment_0005)
(library name:CNhs10379)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100527.Riken4.fc1.ch5)
FileIRIKEN04.2010-05-27T14_02_22.0244193441010.fc1.ch5.srf
(DRA run alias:DRR000949)
(DRA run submitted id:hori-0013_Run_0005)
(original path:/sequencedata/heliscope/srf/2010_05_27_R4_30Q_550FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.LqDynal.1000ng.2nd_20100527.Riken4.fc1.ch5.filtered.sb.bed.gz
(DRA analysis alias:DRZ000012)
(DRA analysis submitted id:hori-0013_Analysis_0005)
(basename:THP1.LqDynal.1000ng.2nd_20100527.Riken4.fc1.ch5.filtered.sb)
(filtered reads:11873336)
(mapped reads on rDNA:1338237)
(mapped reads on the genome assembly:6501153)
bedToCtss
FileITHP1.LqDynal.1000ng.2nd_20100527.Riken4.fc1.ch5.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.LqDynal.1000ng.2nd_20100527.Riken4.fc1.ch5.filtered.sb.[fwdlrev].bw
TRIzol_RNA_extraction
SourceTHP-1
(sample_cell_line:TIB-202 THP-1)
(sample_cell_type:monocyte)
(sample_developmental_stage:1 year infant)
(sample_sex:male)
(DRA sample alias:DRS000625)
(DRA sample submitted id:hori-0013_Sample_0002)
TRIzol_RNA_extraction
Sample2855-61I3
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:0.2 mg/mL)
SampleTHP1.ActinomycinD_conc2.1st
(DRA experiments alias:DRX000610)
(:hori-0013_Experiment_0016)
(library name:CNhs11667)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch17)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch17.srf
(DRA run alias:DRR000960)
(DRA run submitted id:hori-0013_Run_0016)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.ActinomycinD_conc2.1st.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000023)
(DRA analysis submitted id:hori-0013_Analysis_0016)
(basename:THP1.ActinomycinD_conc2.1st.filtered.align_sb)
(filtered reads:10271874)
(mapped reads on rDNA:785218)
(mapped reads on the genome assembly:5653262)
bedToCtss
FileITHP1.ActinomycinD_conc2.1st.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.ActinomycinD_conc2.1st.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:0.1 mg/mL)
SampleTHP1.ActinomycinD_conc1.2nd
(DRA experiments alias:DRX000609)
(:hori-0013_Experiment_0015)
(library name:CNhs11666)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch20)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch20.srf
(DRA run alias:DRR000959)
(DRA run submitted id:hori-0013_Run_0015)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.ActinomycinD_conc1.2nd.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000022)
(DRA analysis submitted id:hori-0013_Analysis_0015)
(basename:THP1.ActinomycinD_conc1.2nd.filtered.align_sb)
(filtered reads:10850048)
(mapped reads on rDNA:683569)
(mapped reads on the genome assembly:6071239)
bedToCtss
FileITHP1.ActinomycinD_conc1.2nd.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.ActinomycinD_conc1.2nd.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:0.2 mg/mL)
SampleTHP1.ActinomycinD_conc2.2nd
(DRA experiments alias:DRX000611)
(:hori-0013_Experiment_0017)
(library name:CNhs11669)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch11)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch11.srf
(DRA run alias:DRR000961)
(DRA run submitted id:hori-0013_Run_0017)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.ActinomycinD_conc2.2nd.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000024)
(DRA analysis submitted id:hori-0013_Analysis_0017)
(basename:THP1.ActinomycinD_conc2.2nd.filtered.align_sb)
(filtered reads:7889223)
(mapped reads on rDNA:532781)
(mapped reads on the genome assembly:4193727)
bedToCtss
FileITHP1.ActinomycinD_conc2.2nd.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.ActinomycinD_conc2.2nd.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.Std.5000ng.4th
(DRA experiments alias:DRX000614)
(:hori-0013_Experiment_0020)
(library name:CNhs11678)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch9)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch9.srf
(DRA run alias:DRR000964)
(DRA run submitted id:hori-0013_Run_0020)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.Std.4th.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000027)
(DRA analysis submitted id:hori-0013_Analysis_0020)
(basename:THP1.Std.4th.filtered.align_sb)
(filtered reads:24689530)
(mapped reads on rDNA:1257893)
(mapped reads on the genome assembly:13817733)
bedToCtss
FileITHP1.Std.4th.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.Std.4th.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.Std.5000ng.5th
(DRA experiments alias:DRX000615)
(:hori-0013_Experiment_0021)
(library name:CNhs11679)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch6)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch6.srf
(DRA run alias:DRR000965)
(DRA run submitted id:hori-0013_Run_0021)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.Std.5th.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000028)
(DRA analysis submitted id:hori-0013_Analysis_0021)
(basename:THP1.Std.5th.filtered.align_sb)
(filtered reads:24295949)
(mapped reads on rDNA:1363538)
(mapped reads on the genome assembly:13349818)
bedToCtss
FileITHP1.Std.5th.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.Std.5th.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:0.4 mg/mL)
SampleTHP1.ActinomycinD_conc3.1st
(DRA experiments alias:DRX000612)
(:hori-0013_Experiment_0018)
(library name:CNhs11670)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch8)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch8. s r f
(DRA run alias:DRR000962)
(DRA run submitted id:hori-0013_Run_0018)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.ActinomycinD_conc3.1st.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000025)
(DRA analysis submitted id:hori-0013_Analysis_0018)
(basename:THP1.ActinomycinD_conc3.1st.filtered.align_sb)
(filtered reads:7401073)
(mapped reads on rDNA:583838)
(mapped reads on the genome assembly:3739668)
bedToCtss
FileITHP1.ActinomycinD_conc3.1st.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.ActinomycinD_conc3.1st.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:0.4 mg/mL)
SampleTHP1.ActinomycinD_conc3.2nd
(DRA experiments alias:DRX000613)
(:hori-0013_Experiment_0019)
(library name:CNhs11674)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch21)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch21.srf
(DRA run alias:DRR000963)
(DRA run submitted id:hori-0013_Run_0019)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.ActinomycinD_conc3.2nd.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000026)
(DRA analysis submitted id:hori-0013_Analysis_0019)
(basename:THP1.ActinomycinD_conc3.2nd.filtered.align_sb)
(filtered reads:7253698)
(mapped reads on rDNA:603855)
(mapped reads on the genome assembly:3772108)
bedToCtss
FileITHP1.ActinomycinD_conc3.2nd.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.ActinomycinD_conc3.2nd.filtered.align_sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:0.1 mg/mL)
SampleTHP1.ActinomycinD_conc1.1st
(DRA experiments alias:DRX000608)
(:hori-0013_Experiment_0014)
(library name:CNhs11665)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20101202.Riken2.fc2.ch23)
FileIRIKEN02.2010-12-02T11_26_57.0239502531008.fc2.ch23.srf
(DRA run alias:DRR000958)
(DRA run submitted id:hori-0013_Run_0014)
(original path:/sequencedata/heliscope/srf/2010_12_02_R2_30Q_1100FOV/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.ActinomycinD_conc1.1st.filtered.align_sb.bed.gz
(DRA analysis alias:DRZ000021)
(DRA analysis submitted id:hori-0013_Analysis_0014)
(basename:THP1.ActinomycinD_conc1.1st.filtered.align_sb)
(filtered reads:10145860)
(mapped reads on rDNA:714844)
(mapped reads on the genome assembly:5641921)
bedToCtss
FileITHP1.ActinomycinD_conc1.1st.filtered.align_sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.ActinomycinD_conc1.1st.filtered.align_sb.[fwdlrev].bw
TRIzol_RNA_extraction
Sample2855-61D3
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.Std.5000ng.3rd
(DRA experiments alias:DRX000607)
(:hori-0013_Experiment_0013)
(library name:CNhs10349)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100614.Riken2.fc2.ch22)
FileIRIKEN02.2010-06-14T14_52_33.0244193431010.fc2.ch22.srf
(DRA run alias:DRR000957)
(DRA run submitted id:hori-0013_Run_0013)
(original path:/sequencedata/heliscope/srf/2010_06_14_R2_30Q/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.Std.5000ng.3rd_20100614.Riken2.fc2.ch22.filtered.sb.bed.gz
(DRA analysis alias:DRZ000020)
(DRA analysis submitted id:hori-0013_Analysis_0013)
(basename:THP1.Std.5000ng.3rd_20100614.Riken2.fc2.ch22.filtered.sb)
(filtered reads:27976845)
(mapped reads on rDNA:1944242)
(mapped reads on the genome assembly:16312956)
bedToCtss
FileITHP1.Std.5000ng.3rd_20100614.Riken2.fc2.ch22.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.Std.5000ng.3rd_20100614.Riken2.fc2.ch22.filtered.sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.Std.5000ng.1st
(DRA experiments alias:DRX000605)
(:hori-0013_Experiment_0011)
(library name:CNhs10333)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100614.Riken2.fc1.ch23)
FileIRIKEN02.2010-06-14T14_52_33.0244193431015.fc1.ch23.srf
(DRA run alias:DRR000955)
(DRA run submitted id:hori-0013_Run_0011)
(original path:/sequencedata/heliscope/srf/2010_06_14_R2_30Q/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.Std.5000ng.1st_20100614.Riken2.fc1.ch23.filtered.sb.bed.gz
(DRA analysis alias:DRZ000018)
(DRA analysis submitted id:hori-0013_Analysis_0011)
(basename:THP1.Std.5000ng.1st_20100614.Riken2.fc1.ch23.filtered.sb)
(filtered reads:19338385)
(mapped reads on rDNA:1760065)
(mapped reads on the genome assembly:10865424)
bedToCtss
FileITHP1.Std.5000ng.1st_20100614.Riken2.fc1.ch23.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.Std.5000ng.1st_20100614.Riken2.fc1.ch23.filtered.sb.[fwdlrev].bw
hCAGE
(starting RNA:5μg)
(ActinomycinD in RT reaction:-)
SampleTHP1.Std.5000ng.2nd
(DRA experiments alias:DRX000606)
(:hori-0013_Experiment_0012)
(library name:CNhs10341)
OP-HELISCOPE-sequencing-v1.0
(flow cell:20100614.Riken2.fc1.ch24)
FileIRIKEN02.2010-06-14T14_52_33.0244193431015.fc1.ch24.sr f
(DRA run alias:DRR000956)
(DRA run submitted id:hori-0013_Run_0012)
(original path:/sequencedata/heliscope/srf/2010_06_14_R2_30Q/FullRun/srf)
filterSMS_indexDPgenomic
(genome assembly:hg18)
(rDNA accession:u13369)
FileITHP1.Std.5000ng.2nd_20100614.Riken2.fc1.ch24.filtered.sb.bed.gz
(DRA analysis alias:DRZ000019)
(DRA analysis submitted id:hori-0013_Analysis_0012)
(basename:THP1.Std.5000ng.2nd_20100614.Riken2.fc1.ch24.filtered.sb)
(filtered reads:23625075)
(mapped reads on rDNA:1090712)
(mapped reads on the genome assembly:14446044)
bedToCtss
FileITHP1.Std.5000ng.2nd_20100614.Riken2.fc1.ch24.filtered.sb.ctss.bed.gz
bigBedToBigWig
FileITHP1.Std.5000ng.2nd_20100614.Riken2.fc1.ch24.filtered.sb.[fwdlrev].bw