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AccessRequest#13214-7

Project,Study,Consent	Status	Expiration	
#3253: Statisticalanalysisandmethodsforexpressionanalysis Genotype-TissueExpression(GTEx)CommonFundProject (phs000424.v6.p1) ExchangeArea (phs000424.v6.p1.c999),NHGRI	Dataaccess GRANTED	2018-02- 09	Publicftp

HowwillIbeabletodownloadtherequesteddata?

typeandGenotypefiles SRAdata(readsandreferencealignments)	SRAsubmittedfiles Provisional
gtex	121Tb
exchange	121Tb
☐ GTEx_phs000424	121Tb
□ exchange	1793Gb
analysis_releases	1646Gb
☐ GTEx_Analysis_2015-01-12	723Gb
☐ GTEX_Analysis_2015-01-12 ☐ GTEX_Analysis_2016-01-15_v7	922Gb
☐ genotypes	907Gb
■ WES	3882Mb
□ WGS	903Gb
☐ GTEx_Analysis_2016-01-15_v7_WholeGe	<u>-</u>
Dominat calls	450Gb
variant_calls	452Gb
GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_635Ind_AllVar_C	· ·
	159Gb
☐ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_635Ind_AllVar_C	•
	2663Kb
☑ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_635Ind_PASS_AE	•
	2687Mb
☑ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_635Ind_PASS_AE	302_GQ20_HETX_MISS15_PLINI
	2381Kb
☑ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_635Ind_PASS_AB	302_GQ20_HETX_MISS15_PLINF
	119Gb
☑ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_635Ind_PASS_AF	302_GQ20_HETX_MISS15_PLINF
	2638Kb
☐ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_652Ind_GATK_H	laplotypeCaller.vcf.gz
	162Gb
☐ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_652Ind_GATK_H	laplotypeCaller.ycf.qz.tbi
	2667Kb
☐ GTEx_Analysis_2016-01-	
15_v7_WholeGenomeSeq_652Indiv_GATK_	HaplotypeCaller.siteOnly.ven \
zs_t/_moledenomeseq_oszmanv_d/m_	6008Mb

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□ GTEx_Analysis_2016-01-
        15_v7_WholeGenomeSeq_652Indiv_GATK_HaplotypeCaller.siteOnly.vep.\(\text{v}\)

☑ GTEx_Analysis_2016-01-

        15_v7_WholeGenomeSeq_PhasedVCF_README.txt
                                                                    1Kb

☑ GTEx_Analysis_2016-01-15_v7_WholeGenomeSeq_README.txt

                                                                   13Kb
         GTEx_Analysis_2016-01-15_v7_WholeGenomeSeq_support_files
                                                                3083Mb

☑ GTEx_Analysis_2016-01-

          15_v7_WGS_652ind_VarID_Lookup_Table.txt
                                                                3083Mb

☑ GTEx_Analysis_2016-01-

          15_v7_WholeGenomeSeq_635Ind_PostVarQC_20genotPCs.txt
                                                                 123Kb

☑ GTEx_Analysis_2016-01-

          15_v7_WholeGenomeSeq_GenomeSTRiP_LCNVs.txt
                                                                    6Kb

☑ GTEx_Analysis_2016-01-

          15_v7_WholeGenomeSeq_QC_README.txt
                                                                    2Kb

☑ GTEx_Analysis_2016-01-15_v7_WholeGenomeSeq_QC_TABLE.tsv

                                                                 162Kb

✓ GTEx_Analysis_2016-01-

          15_v7_WholeGenomeSeq_VarCount_Summary.txt
                                                                  248b

✓ GTEx_Analysis_2016-01-

          15_v7_WholeGenomeSeq_flagged_samples.txt
                                                                    1Kb
   reference_files
                                                                 997Mb
     Homo_sapiens_assembly19.fasta.gz
                                                                 818Mb
     gencode.v19.genes.v7.patched_contigs.gtf
                                                                 141Mb
     gencode.v19.transcripts.patched_contigs.gtf.gz
                                                                  37Mb
   ■ rna-seq
                                                                  14Gb
     GTEx_Analysis_2016-01-15_v7_RNA-seq_RNA-
     SeQCv1.1.8_exon_reads.txt.gz
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     GTEx_Analysis_2016-01-15_v7_RSEMv1.2.22_transcript_isopct.txt.gz
                                                                2172Mb
     GTEx_Analysis_2016-01-15_v7_RSEMv1.2.22_transcript_reads.txt.gz
                                                                2669Mb
     GTEx_Analysis_2016-01-15_v7_RSEMv1.2.22_transcript_tpm.txt.gz
                                                                2003Mb
     GTEx_Analysis_v7_RNA-seq_RNA-SeQCv1.1.8_gene_reads.gct.gz
                                                                 495Mb
     GTEx_Analysis_v7_RNA-seq_RNA-SeQCv1.1.8_gene_rpkm.gct.gz
                                                                2658Mb
   sample_annotations
                                                                  11Mb
     GTEx_Analysis_2016-01-15_v7_Annotations_SampleAttributesDD.xlsx
                                                                  16Kb

☑ GTEx_Analysis_2016-01-

     15_v7_Annotations_SubjectPhenotypesDD.xlsx
                                                                  33Kb
     GTEx_Analysis_2016-01-15_v7_SampleAttributesDS.txt
                                                                  10Mb
     GTEx_Analysis_2016-01-15_v7_SubjectPhenotypesDS.txt
                                                                 586Kb
dbGaP_releases
                                                                  16Gb
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GTEx_Data_2016-01-15_v7	16Gb		
🗷 rna-seq	16Gb		
GTEx_Data_2016-01-15_v7_RNA-seq_RNA-			
SeQCv1.1.8_exon_reads.txt.gz			
	5785Mb		
☑ GTEx_Data_2016-01-15_v7_RNA-seq_RNA-			
SeQCv1.1.8_gene_reads.gct.gz			
G CTF D . 2016 01 15 7 DNA DNA	539Mb		
☑ GTEx_Data_2016-01-15_v7_RNA-seq_RNA- Scott 1.2 compared to the control of the control o			
SeQCv1.1.8_gene_rpkm.gct.gz	280CM		
CTEV Data 2016 01 15 v7 DNA cog DNA CoOCv1 1 9	2896Mb		
☑ GTEx_Data_2016-01-15_v7_RNA-seq_RNA-SeQCv1.1.8	4379Kb		
GTEx_Data_2016-01-15_v7_RSEMv1.2.22_transcript_isc			
GTEX_Data_2010-01-13_V7_K3EMV1.2.22_transcript_isc	2363Mb		
GTEx_Data_2016-01-15_v7_RSEMv1.2.22_transcript_real			
a diez_bata_zoto of is_v/_tozmvi.z.zz_transcript_tet	2901Mb		
GTEx_Data_2016-01-15_v7_RSEMv1.2.22_transcript_tp			
	2179Mb		
■ subgroups	130Gb		
□ ase	5495Mb		
□ v6_data	5495Mb		
■ eqtl	125Gb		
GTExMetasoftRelease_v6p.output.gz	1353Mb		
GTExMetasoftRelease_v6p.output.gz.aspx	64b		
■ Metasoft_20_tissues_UNC	1474Mb		
GTEx.Metasoft6p.20tissues.bestSNP.tab.gz	4455Kb		
GTEx.Metasoft6p.20tissues.input.tab.gz	646Mb		
GTEx.Metasoft6p.20tissues.output.tab.gz	823Mb		
☐ GTEx.Metasoft6p.20tissues.tissue_list	348b		
README.md	3Kb 1400Mb		
Metasoft_Results_v6p.tarfastqtl_v6_v6p_pairs_FOR_QC_ONLY.tar.gz	42Mb		
□ v6_fastQTL_FOR_QC_ONLY.tar.gz	3150Mb		
□ v6_v6p_annotations.tar.gz	65Mb		
□ v6p_All_Tissues_covariates_FOR_QC_ONLY.tar.gz	1699Kb		
v6p_All_Tissues_exon_read_counts_FOR_QC_ONLY.txt.gz	3831Mb		
□ v6p_All_Tissues_gene_rpkm_FOR_QC_ONLY.gct.gz	1845Mb		
□ v6p_All_Tissues_read_counts_FOR_QC_ONLY.gct.gz	352Mb		
□ v6p_Metasoft_Results.tar	1400Mb		
v6p_fastQTL_FOR_QC_ONLY.tar.gz	3103Mb		
v6p_fastQTL_allpairs_FOR_QC_ONLY.tar	107Gb		
<pre>sra_source_files</pre>	3 <i>7</i> Tb		
sra_source_files2	82Tb		
Create download request			
Circuit download request			
Get manifest(CSVformat).			

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