**Human Population Genetics data – Dr. Schrodi Lab**

**1, Personal Medical Research Project (PMRP, 18,604 samples)**

* 1. PLINK FORMAT (PHASE1 and PHASE2 MERGED DATA)

/mnt/sas0/AD/sguo234/pmrp/MCRI20000\_R2.bed

/mnt/sas0/AD/sguo234/pmrp/MCRI20000\_R2.bim

/mnt/sas0/AD/sguo234/pmrp/MCRI20000\_R2.fam

/mnt/sas0/AD/sguo234/pmrp/MCRI20000\_R3.pgen

/mnt/sas0/AD/sguo234/pmrp/MCRI20000\_R3.psam

/mnt/sas0/AD/sguo234/pmrp/MCRI20000\_R3.pvar

* 1. VCF FORMAT (non-imputation and non-phasing)

/mnt/sas0/AD/sguo234/pmrp/MCRI20000.vcf.gz

/mnt/sas0/AD/sguo234/pmrp/MCRI20000.vcf.gz.tbi

* 1. IMPUTED and PHASED PMRP DATA

/mnt/sas0/AD/sguo234/pmrp/phased/

**2. Exome-Sequencing data to PMI (Personal Medicine Imitate)**

2.1 INDIVIDUAL VCF FILES

/mnt/sas0/AD/sguo234/exome/ 226764-a.filtered.vcf.gz

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/mnt/sas0/AD/sguo234/exome/ 282139-a.filtered.vcf.gz

/mnt/sas0/AD/sguo234/exome/ 296044-a.filtered.vcf.gz

2.2 MERGED VCF

/mnt/sas0/AD/sguo234/exome/Final.vcf.gz

/mnt/sas0/AD/sguo234/exome/final.rs.vcf.gz

/mnt/sas0/AD/sguo234/exome/exome.rename.vcf.gz.gz

**3. Cytokine RNA-SEQ Dataset**

3.1 Saved in DataLocker since DeepThought don’t have enough space to store it.

**4. dbSNP153 in hg19**

Deepthought don’t have enough space, hence, dbSNP153 in hg38 was not generated.

/mnt/sas0/AD/sguo234/db/dbSNP153/dbSNP153.hg19.fam

/mnt/sas0/AD/sguo234/db/dbSNP153/dbSNP153.hg19.fam

/mnt/sas0/AD/sguo234/db/dbSNP153/dbSNP153.hg19.fam

/mnt/sas0/AD/sguo234/db/dbSNP153/ dbSNP153.norm.hg19.vcf.gz

/mnt/sas0/AD/sguo234/db/dbSNP153/ dbSNP153.norm.hg19.vcf.gz.tbi

**5. 1K Northeast Asians Genome Project (**1,781 Northeast Asians Genome)

/mnt/sas0/AD/sguo234/db/nard/NARD\_MAF.hg38.vcf.gz

/mnt/sas0/AD/sguo234/db/nard/NARD\_MAF.hg38.zip

**6. ANNOVAR annotation to dbSNP153 for Loss-of-function prediction**

7.