Test File for Attached File Endpoint

File Name	Sample ID	File Type	Checksum	Description
file_001.bam	SMPL001	BAM	abc12345	BAM file for whole genome sequ
file_002.vcf	SMPL002	VCF	def67890	VCF file with called variants
file_003.fastq.gz	SMPL003	FASTQ	ghe98765	Raw sequencing reads