BIOINFORMATICS ASSIGNMENT 1 (Day 1 - 5)

1. Gene Name: TP53 (tumor protein p53)

2. **Function of the Gene**: This gene encodes a tumor suppressor protein containing transcriptional activation, DNA binding, and oligomerization domains. The encoded protein responds to diverse cellular stresses to regulate the expression of target genes, thereby inducing cell cycle arrest, apoptosis, senescence, DNA repair, or changes in metabolism. Mutations in this gene are associated with a variety of human cancers, including hereditary cancers such as Li-Fraumeni syndrome. Alternative splicing of this gene and the use of alternate promoters result in multiple transcript variants and isoforms. Additional isoforms have also been shown to result from the use of alternate translation initiation codons from identical transcript variants (PMIDs: 12032546, 20937277). [provided by RefSeq, Dec 2016]

3. NCBI accession number: NC_000017

4. Forward Primer: TTGTAATGCAGGGCTGAGGA5. Reverse primer: CTGCTCCCACCTCCTGTTAA

6. Features of primers:

	SEQUENCE	TEMPLATE	START	Tm	Gc%
		STRAND	LENGTH		
LEFT PRIMER	TTGTAATGCAGGGCTGAGGA	plus	10053	59.01	50
RIGHT PRIMER	CTGCTCCCACCTCCTGTTAA	minus	10260	59.02	55

7. Amplicon length and sequence:

Amplicon length: 208bp

Amplicon sequence: T

TGTAATGCAGGGCTGAGGAGTGTCCGAAGAGAATGGGCAGGTGAGCCAGAGACAGTTGTTCTTCCAGAAGCTTTGCAGTGAAAGGAATCAAAGAAATGGAGCCGTGTATCAGGTGGGGAAGGGTGGGGGCCAAGGGGGTGTCCTTCCCCATACAGAGATTGCAGGCTGAGAATGACTATATCCTTGTTAACAGGAGGTGGGAGCAG

qPCR Data analysis (DAY 5)

		Ct values			
Housekeeping genes(GAPDH)	Ct 1		Ct 2		
Untreated (control)		18.5		18.5	
Untreated (control)		17.8		17.8	
Untreated (control)		17.5		17.5	
Treated		18.3		18.3	
Treated		18.5		18.5	
Treated		18.2		18.2	

	Ct values			
Gene of interest (HER2)	Ct 1		Ct 1	
Untreated(control)		23.3		22.5
Untreated(control)		22.5		22.2
Untreated(control)		21.2		21.9
Treated		25.3		25.3
Treated		26.5		26.5
Treated		27.5		27.5

The following data are results of qPCR from cancer cell lines. HER2 stands for human epidermal growth factor. It's healthy in normal amounts, but too much may be a sign of a certain type of breast cancer. Calculate the 2 Delta Ct values for the following data and plot the values on a graph using graphpad prism.

Ans:

A	В	С	D	E	F	G	Н	I	J	K	L	M	N
Hou	seKeppin	g Gene	Ger	ne Of Int	rest		Avg ct value for HG	Avg ct value for GOI	Δct Value	ΔΔct Valu	e fold change		
	R1	R2		R1	R2								
control(untreated	18.5	18.5		23.3	22.5		18.5	22.9	4.4		1		
sample 1	18.3	18.3		25.3	25.3		18.3	25.3	7	2.6	0.16493849		
sample 2	18.5	18.5		26.5	26.5		18.5	26.5	8	3.6	0.08246924		
sample 3	18.2	18.2		27.5	27.5		18.2	27.5	9.3	4.9	0.03349292		