## **Supplementary Materials**

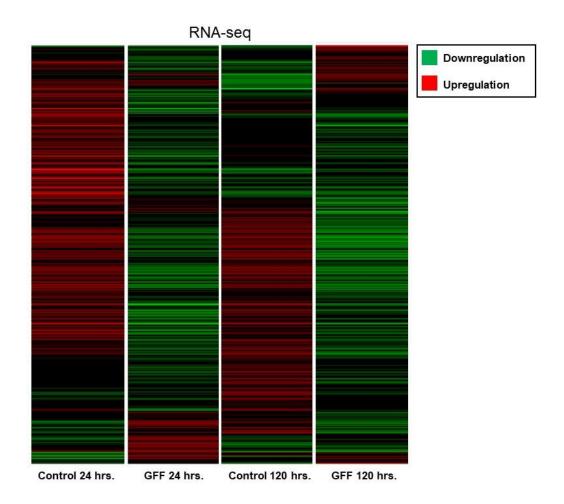


Figure S1: GFF effect on gene transcription in NHEM.

RNA-seq heat maps visually illustrate notable quantities of genes in NHEM modulated due to treatment with 10% GFF-B at both 24 and 120 hours (5 days). At 24 hours, 3,574 genes were altered by 2-fold and 533 of those genes had p-values 0.05. At 120 hours, 3,466 genes were modulated by 2-fold and 572 of those genes had p-values <0.05. Green indicates downregulation, red indicates upregulation, and black indicates no change; the intensity of the color indicates the fold change degree of the modulation, n=1.

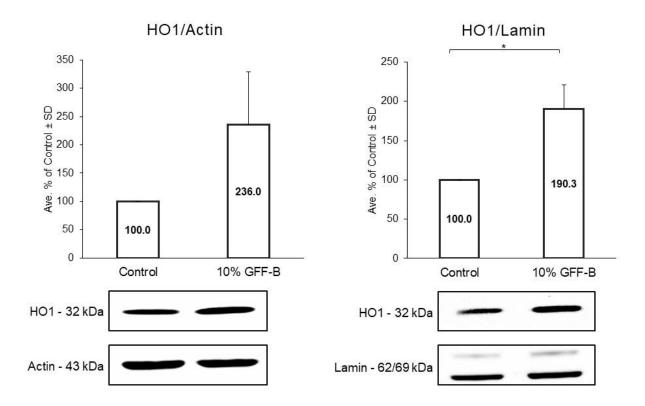


Figure S2: Cytoplasmic and nuclear HO1 expression.

HO1 expression was increased in both the cytoplasm and nucleus of NHEM treated with 10% GFF-B for 8 hours. Results are expressed as a percentage of the control. p<0.05, SD, n=3.

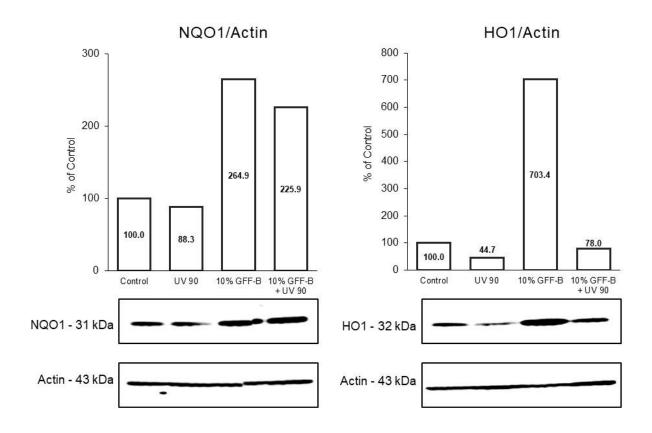


Figure S3: GFF effect on Nrf2-ARE pathway challenged by UVB.

Lightly pigmented NHEM were pretreated with 10% GFF-B for 6 days. Cells were irradiated with UVB (90 mJ/cm²) while in PBS and then immediately rescued with or without 10% GFF-B for an additional 24 hours. At harvest, cells were processed with Western blot and densitometric analyses, n=1. NQO1 expression was maintained with GFF-B treatment post-irradiation 2-fold over untreated, irradiated cells. HO1 was highly inducible with GFF-B treatment alone, yet expression was not maintained post-irradiation. Results are expressed as a percentage of the control.

Table S1 – RNA-seq antioxidant enzyme targets – 24 hours.						
Upregulation	■ Downregulation					
Gene ID	Control	GFF-B	Fold Change	p-value		
HMOX1	94.40	350.65	3.71	0.0003*		
NQO1	1623.61	3328.54	2.05	0.024*		
TXNRD1	652.28	1307.26	2.004	0.024*		

Table S2 – RNA-seq antioxidant enzyme targets – 120 hours.						
■ Upregulation	Downregulation	l				
Gene ID	Control	GFF-B	Fold Change	p-value		
HMOX1	92 51	260 48	2 82	0.002*		
	02.01	200.10	2.02	0.002		
NQO1	1327.87	3885.33	2.93	0.0002*		
NQO1	92.51 1327.87	260.48 3885.33	2.82	0.002		

**Table S1 and S2 – RNA-seq antioxidant enzyme targets.** 10% GFF-B treatment for 24 hours and 120 hours (5 days) significantly upregulated mRNA levels of antioxidant enzymes from the Nrf2-ARE pathway that are involved in ROS neutralization. Significance: Fold Change  $\geq 2$ , \*p<0.05.