

**Supplementary Table 1.** PCR-RFLP based mutation primers

<b>Gene</b>	<b>Forward Primer sequence</b>	<b>Reverse Primer sequence</b>	<b>Product size</b>
<i>XRCC1</i> <i>codon399</i>	5'- GGACTGTCACCGCATGC GTCGG-3'	5'- GGCTGGGACCACCTGT GTT-3'	115 bp + 34 bp (Wild)
			149 bp (Mutant)
<i>XRCC1</i> <i>codon280</i>	5'- TTGACCCCCAGTGGTGC T-3'	5'- CCCTGAAGGATCTTCCC CAGC-3'	126 bp + 62 bp (Wild)
			188 bp (Mutant)
<i>hOGG1</i> <i>codon326</i>	5'- ACTGTCACTAGTCTCAC AG-3'	5'- GGAAGGTGCTTGGGA AT-3'	200bp (wild)
			100 bp + 100 bp (mutant)

**Supplementary Table 2.** Gene specific expression primers

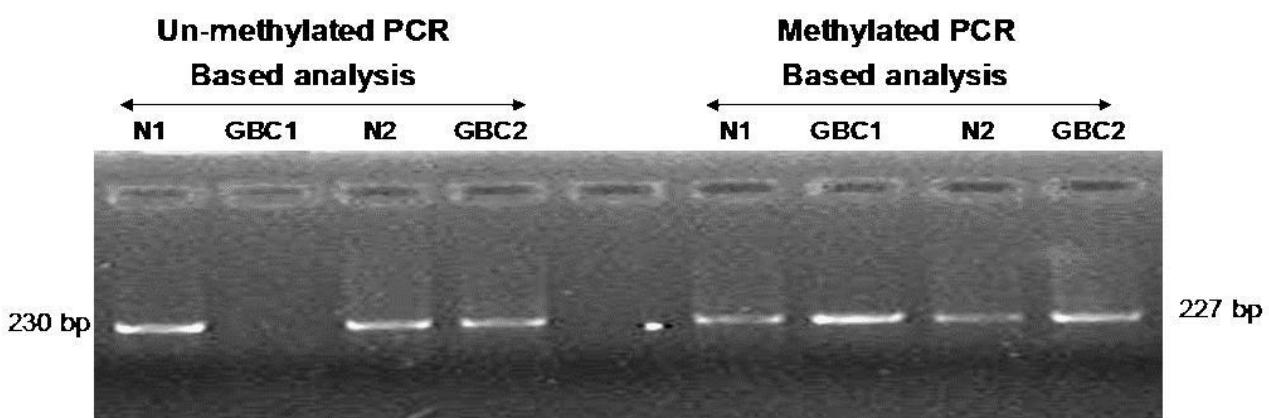
<b>Gene</b>	<b>Forward Primer sequence</b>	<b>Reverse Primer sequence</b>	<b>Product size</b>
<i>Beta-actin</i>	5'- AGATGTGGATCAGCAAG CAG -3'	5'- GCGCAAGTTAGGTTTG TCA -3'	122 bp
<i>XRCC1</i>	5'- CAGCTGTCGCCATCTGTT C-3'	5'- TGAAGGATCTTCCCCA GCTC-3'	173 bp

<i>hOGG1</i>	5'- ATCTGTTCCCTCCAACAAAC AAC-3'	5'- 'TTCCTGAGATGAGCCT CCAC-3'	161 bp
<i>PARP1</i>	5'- AAGAAATGCAGCGAGAG CAT-3'	5'- CCAGTGTGGACTTTTC CAT-3'	90 bp
<i>POL-β</i>	5'- TACATTGCTACAGTCTGT GGCAGTT-3'	5'- TGGGATGGGTCAAGGAG AACAA-3'	79 bp
<i>PCNA</i>	5'- CCATCCTCAAGAAGGTG TTGG-3'	5'- TAGGTGTCGAAGCCCT CAGA-3'	151 bp
<i>APE</i>	5'- TGGAATGTGGATGGGCT TCGAGCC-3'	5'- AAGGAGCTGACCAGTA TTGATGA-3'	169 bp

**Supplementary Table 3.** Methylation detection primers

Gene	Forward Primer sequence	Reverse Primer sequence
<i>XRCC1</i> methylated	5'- TTTTAGAAAGTAGGGTCGG ACGT-3'	5'- GCCCTACGCTAACTAAAAT ACGCA-3'
<i>XRCC1</i> unmethylated	5'- TTTTAGAAAGTAGGGTTGG	5'- TACACCCTACACTAACTAA

	ATGT-3'	AATACACA-3'
<i>hOGG1</i> methylated	5'-  AATTAGAAGAACATAGTTGT  GCGC-3'	5'-  TACATACCTCGCCCTTAC  GA-3'
<i>hOGG1</i> unmethylated	5'-  TTTAGAAGAACATAGTTGTGT  GTGT-3'	5'-  CTCCTACATACCTCACCCCTT  TACAA-3'



**Supplementary Figure 1.** Representative agarose gel showing differential methylation profile of XRCC1 promoter in GBC cases compared to controls showing hyper-methylation in GBC cases.