

Hitting BRCA1/2 Mutant Cells in their Achilles Heel

1. The BRCA2 gene is located on chromosome 13 and contains approximately 27 exons, 1000 bases and 3418 amino acids.
2. In Dec 1995, a group of scientists found that **one missing coding "base"** in the gene sequence increases the risk of cancer (both ovarian and breast).
 - This deletion causes a frameshift, which leads to a stop codon in its place and thus the **truncation** of the protein.
3. Most people have two normal BRCA2 genes, but carriers of mutations have **one normal and one fault gene**.
 - This means that these individuals have an 85% lifetime risk of breast and 30% risk of ovarian cancer.
 - Therefore, cancers develop from cell that **lost the final normal BRCA2 copy**.
 - The mere presence of the mutant allele increases the risk of further mutations of the normal copy.

What do the normal BRCA1 and 2 proteins do in the cell?



1. THE BRCA2 gene contains 30-40 repeat motifs – the so-called BRC repeats – which are encoded in exon 11 and are conserved between several mammalian species, thus suggesting that they have an essential function.
2. In fact, the BRC repeats have been shown to **mediate the binding of BRCA2 to RAD51**, which is a mammalian protein that is essential for DNA repair and genetic recombination.
3. Evidence supporting importance of this protein in gene repair:
 - Knockout experiments have shown that a lack of this gene causes chromosomal breaks or chromosomal aberrations, for example, the formation of a tri-radial chromosome.
 - Differential mutational processes often generate different combinations of mutation types, and are termed '**signatures**'.
 - o Analysis of 21 breast cancer whole-genome sequences found that breast cancer contains **highly variable chromosomal structural instability**.
 - o This means that the BRCA2 genome is unstable, and is possibly due to faulty DNA repair mechanisms.
 - Breast cancers develop through the **acquisition of mutations** over time.
 - o An inherited mutant copy increases the risk of developing serial mutations that are needed to develop cancer.