



| Key | Table of mutations |
|---|---|
| 1-2 Calcium ion coordinating residue (numbers denote calcium ion coordinating site) | E-cadherin (Ozawa et al., 1990) D134K, D134A |
| between these domains, triangle denotes calcium ion coordinating carbonyl | E-cadherin (Pertz et al., 1999) W2A, A80I |
| Glycosylated residue | E-cadherin (Bex et al., 1998) T397, N161S, E162D, D216A, E390Q, T316I, V319N, A438T, R444Q, A463T |
| Mutation abolishes adhesion | N-cadherin (Tamura et al., 1998) W2A, W2I(partial), W2Y, V3Y, A78M, A80M |
| Mutation does not abolish adhesion | N-cadherin (Kitagawa et al., 2000) W2A, G15A, D27A, D29A, Y36A, T39A, D44A, F51A, G58A, Y74A(partial), H79A, A80I, V81A, D82A, N84A, E89A, Y147A, F162A |
| Mutation seen in tumors | |
| Disulphide bond | |
| Conserved Pro at start of each domain | |
| Strand dimer interaction W2 donor | |
| Strand dimer interaction W2 acceptor | |
| Cis dimer interaction | |