

Term

- # Amplification of PVT1 contributes to the pathophysiology of ovarian and breast cancer
- Aneuploidy, TP53 mutation, and amplification of MYC correlate with increased intratumor heterogeneity and poor prognosis of breast cancer patients
- Breast cancer
- Breast implant-associated anaplastic large cell lymphoma
- Genes from the most frequent genomic gains and amplifications in a panel of patients with lymph node negative breast cancer (NNBC)
- Genes up-regulated in MCF7 cells (breast cancer) after stimulation with EGF [GeneID=1950]
- Human Breast Adler06 57genes
- Human Breast Pawitan05 64genes prognosisGenes
- Mouse Breast Fleming09 114genes
- Human Breast Horlings10 271genes
- Reoplasm of the breast
- Tumours of the breast
- Genes up-regulated in ME-A cells (breast cancer) undergoing apoptosis upon serum starvation (5% to 0% FCS) for 22 hr
- Examples Genes within amplicon 8q23-q24 identified in a copy number alterations study of 191 breast tumor samples