

Response Prediction to Neoadjuvant Chemotherapy: Comparison between Pre-Therapeutic Gene Expression Profiles and In Vitro Chemosensitivity Assay.

Singer CF, Klinglmüller F, Stratmann R, Staudigl C, **Fink-Retter A**, Gschwantler D, Helmy S, Pfeiler G, Dressler AC, Sartori C, Bilban M.

PLoS One. 2013 Jun 24;8(6)

Clinical implications of genetic testing for BRCA1 and BRCA2 mutations in Austria.

Singer C, Muhr D, Rappaport C, Tea MK, Gschwantler-Kaulich D, **Fink-Retter A**, Pfeiler G, Berger A, Sun P, Narod S.

Clin Genet. 2013 Jun 17.

Tuberin and p27 expression in breast cancer patients with or without BRCA germline mutations.

Dressler AC, Hudelist G, **Fink-Retter A**, Gschwantler-Kaulich D, Pfeiler G, Rosner M, Hengstschläger M, Singer CF.

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Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk.

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A nonsynonymous polymorphism in IRS1 modifies risk of developing breast and ovarian cancers in BRCA1 and ovarian cancer in BRCA2 mutation carriers.

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Common variants at the 19p13.1 and ZNF365 loci are associated with ER subtypes of breast cancer and ovarian cancer risk in BRCA1 and BRCA2 mutation carriers.

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la Hoya M, Muranen TA, Nevanlinna H, Tischkowitz MD, Spurdle AB, Neuhausen SL, Ding YC, Lindor NM, Fredericksen Z, Pankratz VS, Peterlongo P, Manoukian S, Peissel B, Zaffaroni D, Barile M, Bernard L, Viel A, Giannini G, Varesco L, Radice P, Greene MH, Mai PL, Easton DF, Chenevix-Trench G; kConFab investigators, Offit K, Simard J; Consortium of Investigators of Modifiers of BRCA1/2.

Cancer Epidemiol Biomarkers Prev. 2012 Apr;21(4):645-57.

Common variants at 12p11, 12q24, 9p21, 9q31.2 and in ZNF365 are associated with breast cancer risk for BRCA1 and/or BRCA2 mutation carriers.

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Ovarian cancer susceptibility alleles and risk of ovarian cancer in BRCA1 and BRCA2 mutation carriers.

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Presence of intratumoral stem cells in breast cancer patients with or without BRCA germline mutations.

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