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Background: Multigene panels are routinely used to assess for predisposing germline mutations in families at high breast cancer risk. The number of variants of unknown significance thereby ...

Tumor Sequencing Is Useful to Refine the Analysis of Germline Variants in Unexplained High-Risk Breast Cancer Families
Effector memory T cells have a reduced capacity to induce GVHD but can transfer functional T-cell memory. Allogeneic haematopoietic stem-cell transplantation (SCT) is a curative therapy for ...

Graft-versus-host disease
Lledo et al. Impact of c-kit mutations, including codons 557 and/or 558, on the recurrence-free survival after curative surgery in patients with GIST. Watanabe et al. Detection of viable human ...

2011 Gastrointestinal Cancers Symposium
improvement projects through Kaizens, involvement of cross functional teams to bring cost ... Financial assistance given for providing free 100 nos hearing aid equipment at Jagatguru ...

Kirloskar Ferrous Industries Ltd.
Exploration of Pb-free polar materials for energy and electronic applications (Geneva Laurita, Chemistry and Biochemistry), STEM Faculty-Student Research Grant Morgan Baxter ’20: Magnetorotational ...

Summer Research Recipients
Our hypothesis was that matched tumor sequencing could be helpful in pinpointing genetic bases of suspected predisposition to BC in patients without pathogenic mutations in BRCA1, BRCA2, TP53, and ...

Tumor Sequencing Is Useful to Refine the Analysis of Germline Variants in Unexplained High-Risk Breast Cancer Families
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