Susan G. Komen

Research Grants – Fiscal Year 2015

This research grant was approved for FY2015 Research Programs funding. This grant will be funded upon the execution of grant agreements between Komen and the grantee institutions.

Molecular and clinical characterization of breast cancer heterogeneity in Israel

Investigator(s): Bella Kaufman, M.D.

Lead Organization: Chaim Sheba Medical Center

Grant Mechanism: Komen Scholars  Grant ID: SAC110019

Public Abstract:

Breast cancer is composed of different diseases which vary according to the origin of the cancer cell types as well as their location. Breast cancer originates and progresses as a result of molecular processes. Our aim in this study is to understand the molecular processes that are the basis of breast cancer development, progression, metastasis and resistance to therapy.

As cancer progresses, new mutations (genetic alterations) occur and accumulate in the cancer cells. Our working hypothesis is that early identification of such genetic signatures will shed light on tumor evolution and will lead to the development of rationally targeted and individualized therapy.

We address several clinically relevant questions and plan to detect and analyze the genetic differences underlying the following processes:

1. To understand the molecular mechanism(s) of the development of invasive cancer (IDC – invasive ductal carcinoma) from locally confounded cancer (DCIS – ductal carcinoma in-situ). We propose to investigate tumor evolution by analyzing the mutational spectrum of a certain patient's tumors at different stages of disease progression.
2. Characterization of breast cancer evolution throughout the timeline of pre-operative (neo-adjuvant) chemotherapy. We plan to identify expression signatures that predict recurrence at the biopsy stage. Overall, in addition to the contribution of this study to the improvement of breast cancer prevention and management, it also has the potential to contribute to cancer study in general. The produced data from these studies have the following potential: 1) providing the research community with a basic understanding of the tumorigenic process, and 2) detecting a set of potential biomarkers involved in these processes. Identifying the cases prone either to resist therapy, cause tumor progression, or recur, would justify more aggressive therapy while avoiding unnecessary morbidity. On the other hand, identification of low-risk patients can help reconsider the most appropriate treatment management, thus avoiding short- and long-term morbidities, as well as side effects undermining patients’ quality of life.

3. The mechanism(s) leading to the significant difference in the response rate to treatment (pathological complete response) between young and older patients receiving pre-operative chemotherapy (neo-adjuvant).