Dilsher Singh Dhillon

Statistician

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**** 9796762387

♦ Houston, TX

Skills

Statistical Programming in SAS

Python

SQL

Statistics

Survival Analysis

Analysis of Categorical Data

Design of Experiments

Unsupervised Learning

Analysis of correlated data

Mathematical Statistics

Luminex bead based assays

ELISA

Tissue culture

Bacterial Cloning techniques

Human Biospecimen Collection (CLIA)

Gel Electrophoresis

Immunoflourescence

Education

Texas A&M University MS Statistics 2017 MS Biotechnology 2012

Employment

Baylor College of Medicine

Houston, TX Jan. 2018 to Current

Identifying risk factors for poor outcomes in co-infected TB-HIV patients

- Data extraction o Worked with monitoring and evaluation officers in Africa to identify key variables to extract o Extracted longitudinal data for -25000 subjects from 7 clinics in Sub-Saharan Africa during the time period 2006-2018
- o Cleaned, processed, run quality checks and check completeness of data to convert it into a usable format for analysis

- o Consulted subject matter experts to identify a priori list of variables that are hypothesized to be related to worse TB outcomes
 o Developed cox regression models (multiple models categorizing subjects according to Anti-retroviral therapy use) to identify risk factors associated with TB outcomes in HIV infected
 - o Presented findings of the project in the network wide meeting held in Johannesburg, South Africa (November 2018)

- Automated a pipeline in Python for merging of datasets from clinical research forms(CRFs) entered in eSwatini
 Run quality checks monthly on data extracted from the database
 Worked with database management team to remove redundancies across datasets reducing the size and number of variables to be extracted

- Studying underlying epigenetic mechanism of TB infection

 Processed methylation data from EPIC arrays consisting of -850,000 individual probes using the minfi package from Bioconductor

 Produced exploratory data analysis reports consisting of heatmaps, violin plots

University of Texas MD Anderson Cancer Center

Apr. 2016 to Jan. 2018

Promoted to a supervisory capacity to oversee assay development of novel biomarkers and testing of in-house and commercial assays in clinical samples for validation Manage CLIA facility for sample collection for validation of blood based biomarkers

Formulate final biomarker panel of autoantibodies in response to aberrantly expressed proteins in plasma and serum/plasma of lung cancer subjects

Maintaining a bio-repository of multiple pre-diagnostic, diagnostic and diseased subject samples to be used for the final validation of the down selected biomarker panel

Houston Jan. 2013 to Mar. 2016

Translated two ELISA assays for proteins differentially expressed in lung cancer into a bead based assay using the Luminex MagPix platform and achieved higher sensitivity and better inter-assay precision through development and optimization

Executed an expanded panel of 34 protein biomarkers for lung cancer assembled into in-house as well as MilliPlex designed panels on various pre-diagnostic and diagnostic lung cancer subjects

Contribute to down selecting a panel of proteins, miRNAs and autoantibodies to develop a biomarker panel in multiple diagnostic, pre-diagnostic subjects and patients from the initial expansion set of biomarkers

Projects

Forecasting Red Light Camera Violations in Chicago

May 2017 to Oct. 2017

Develop a regression model to predict red light camera violations in the Chicago

- Develop a regression model to predict red light camera violations in the Chicago from the beginning of 2014

 Expanded the dataset to impute missing camera IDs using intersection names and included dummy variables for all Chicago neighborhoods based on the co-ordinates of each Camera ID in order to use these neighborhoods as potential predictors

 Using the "moad" package in R, downloaded weather data for the Chicago Midway Airport and the Chicago O'Hare Airport which includes potential predictors such as snowfall, precipitation, wind speed, visibility in addition to temperature conditions on a daily basis from the start of 2014

Testing Efficacy of Horn Fly Counts in Cattle

Jan. 2017 to Apr. 2017

- Cleaned and pre-processed dataset obtained from client for horn fly counts over 26 weeks in cattle across 15 counties in Texas
 Developed a repeated measures mixed effects model to analyze outcomes of pesticides treatment under the assumption that the counts are correlated with each other
 Prepared a client report summarizing research goals, exploratory data analysis, statistical methods and key findings of the analysis. In addition, provided suggestions on how to improve experimental design in order to have greater power to detect differences

Awards

Texas A&M University · Regents Fellowship

Aug. 2010

International Student Services, Texas A&M University - International Education Scholarship

Sept. 2011

Custom Section 1

Guida F, Sun N, Bantis LE, Muller DC, Li P, Taguchi A, **Dhillon D**, Kundnani DL, Patel NJ, Yan Q, Byrnes G, Moons KGM, Tjønneland A, Panico S, Agnoli C, Vineis P, Palli D, Bueno-de-Mesquita B, Peeters PH, Agudo A, Huerta JM, Dorronsoro M, Barranco MR, Ardanaz E, Travis RC, Byrne KS, Boeing H, Steffen A, Kaaks R, Hüsing A, Trichopoulou A, Lagiou P, La Vecchia C, Severi G, Boutron-Ruauth MC, Sandanger TM, Weiderpass E, Nost TH, Tsilidis K, Ribiol E, Carankvis K, Johansson M, Goodman GE, Feng Z, Brennan P, Johansson M, Hanash SM Assessment of Lung Cancer Risk on the Basis of a Biomarker Panel of Circulating Proteins (JAMA Oncology 2018)

Shiels MS, Kirk GD, Drummond MB, Dhillon D, Hanash SM, Taquchi A, Engels EA HIV Infection and Circulating Levels of Prosurfactant Protein B and Surfactant Protein D (Journal of Infectious

Çeliktas M, Tanaka I, Tripathi SC, Fahrmann JF, Aguilar-Bonavides C, Villalobos P, Delgado O, Dhillon D, Dennison JB, Ostrin EJ, Wang H, Behrens C, Do KA, Gazdar AF, Hanash SM, Taguchi A Role of CPS1 in Cell Growth, Metabolism and Prognosis in LKB1-inactivated Lung Adenocarcinoma (JNCI 2017)

Capello M, Bantis LE, Scelo G, Zhao Y, Li P, Dhillon DS, Patel NJ, Kundnani DL, Wang H, Abbruzzese JL, Maitra A, Tempero MA, Brand R, Firpo MA, Mulvihill SJ, Katz MH, Brennan P, Feng Z, Taguchi A, Hanash SM Sequential Validation of Blood-Based Protein Biomarker Candidates for Early-Stage Pancreatic Cancer (JNCI 2017)