# Statistical Modeling of Genomic and Biomedical Data

# **Overall Research Objectives:**

- Developing statistical methods for inference-based analysis of biological data
- Collaborating with scientific teams to answer research questions using statistical methodology

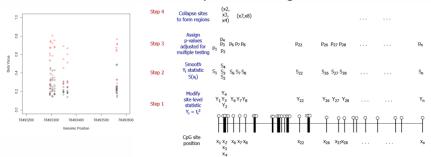
#### **Statistical Genomics Topics:**

- Detecting Differentially Methylated Regions
- Meta-Analysis of Gene Expression Data

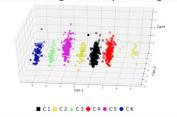
#### **Statistical Modeling of Biomedical Data Topics:**

- Modeling Sleep in the Fruit Fly
- Statistical Methods for Explainability in Unsupervised Learning Applications of Biomedical Data

### Differential methylation testing



#### Visualizing and explaining TBI clusters via statistical analysis



Hematocrit level (highest)		C1	C2	C3	C4	C5	C6		Hematocrit level (lowest)
39.4 ± 5.2	C1	*						C1	32.8 ± 6.7
40.1 ± 4.9	C2		*					C2	33.9 ± 7.4
41.3 ± 4.0	C3			*				C3	36.7 ± 5.1
41.4 ± 4.3	C4							C4	34.9 ± 6.6
40.7 ± 5.3	C5					*		C5	35.6 ± 6.2
42.0 ± 3.4	C6						*	C6	38.3 ± 5.9
		C1	C2	C3	C4	C5	C6		

#### **PoC**

• Gayla Olbricht, Professor

• E-Mail: olbrichtg@mst.edu

• Phone: 1-573-341-4913

# **Funding**

 Funding agencies: NIH, Leonard Wood Institute, Missouri Department of Higher Education **Statistical Keywords**: Cluster analysis, functional data analysis, imputation, meta-analysis, multivariate statistics, multiple testing, regression

**Biomedical Data Keywords:** Traumatic brain injury (TBI), sleep, cancer, mesenchymal stem cells

#### **Publications:**

Link to my publications

