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penn state u

statistical genetics other comparative effectiveness research

clinical trials biomarkers cancer applications epidemiologic methods longitudinal/survival models

prediction/prognostic modeling

next generation sequencin high dimensional data

causal inference categorical data genome wide association studies infectious disease models meta-analysis multiple testing

u of alabama

u of iowa

texas a&m u

nichd

u of missouri

survival analysis

time series
longitudinal data
genome wide association studies
functional data analysis
biopharmaceutical research causal inference
high dimensional data
infectious disease models
next generation sequencing

nonparametric methods

longitudinal/survival models bayesian methods diagnostic/screening tes statistical genetics

survival analysis longitudinal data

semi-parametric/non-parametric models longitudinal/survival models missing data Imaging

health services research
epidemiologic methods
clustered data methods
other cancer applications
adaptive design
surrogate markers causal inference model selection environmental/ecological high dimensional data

u of minnesota

statistical genetics

otherimaging graphical models demography/population studies

bayesian methods dynamic treatment regimens genome wide association studies

meta-analysis

survival analysis

spatial/temporal modeling

prediction/prognostic modeling

longitudinal data personalized medicine spatial/temporal modeling

statistical genetics

model selection high dimensional data epidemiologic methods data mining/massive data survey research data^{Clustered} data methods clinical trials spatial/temporal modeling

u of north carolina

bayesian methods
adaptive design imaging
cancer applicationsgraphical models
categorical data machine learning causal inference genome wide association studies next generation sequencing personalized medicine semi-parametric/non-parametric models survival analysis

harvard u

biologics adaptive design biomarkers causal inference epidemiologic methods genomics survival analysis

other comparative effectiveness research

north carolina state u

Year: 2013

u of pittsburg

nonparametric methods

machine learning

hierarchical models epidemiological study design missing datacompeting risks

comparative effectiveness research diagnostic/screening tests genome wide association studies

high dimensional data longitudinal/survival models

survival analysis

meta-analysis

roc analysis

statistical genetics

spatial/temporal modeling

longitudinal data

adaptive design time series

random effects model selection dynamic treatment regimens constrained estimation bayesian methods computational methods functional data analysis measurement error

prediction/prognostic modeling survival analysis

toxicology/dose-response spatial/temporal modeling spatial/temporal modeling other environmental/ecological dynamic treatment regimens bayesian methods biomarkers

imaging

u of texas

clinical trials causal inference agreement

emory u

longitudinal/survival models

othergenomics

adaptive design

bayesian methods cancer applications epidemiologic methods epidemiological study design next generation sequencing

survey research data

prediction/prognostic modeling next generation sequencing causal inference

u of pennsylvania

other biomarkers appreached bayesian methods epidemiological study design genomics imaging time series longitudinal data

harvard sph

statistical genetics

time series other meta-analysis infectious disease models causal inference

bayesian methods genome wide association studies high dimensional data measurement error model selection prediction/prognostic modeling survival analysis

food and drug administration

johns hopkins u

u of florida

johns hopkins bloomberg sph

diagnostic/screening tests other clinical trials biomarkers adaptive design categorical data data mining/massive data medical devices

next generation sequencing longitudinal data genomics epidemiologic methods functional data analysis missing data statistical graphics personalized medicine statistical genetics

u of wisconsin

emi-parametric/non-parametric models next generation sequencing missing data environmental/ecological data mining/massive data functional data analysis imaging

survey research data

next generation sequencing multivariate survival roc analysis imaging

functional dată analysis bayesian methods environmental/ecological genomics high dimensional data medical devices personalized medicine

duke u

u of south carolina

columbia u

national cancer institute

penn state u

statistical genetics roc analysis high dimensional data environmental/ecological bayesian methods causal inference functional data analysis spatial/temporal modeling time series

toxicology/dose-response random effects other model selection bayesian methods survival analysis

prediction/prognostic modeling multiple testing generalized linear models competing risks data mining/massive data next generation sequencing statistical genetics survey research data

personalized medicine other
epidemiologic methods
biomarkers
genome wide association studies
health policy applications prediction/prognostic modeling statistical genetics

spatial/temporal modeling prediction/prognostic modeling multiple testing survival analysis graphical models

clustered data methods
causal inference
genome wide association studies
genomics other
infectious disease models
next generation sequencing
personalized medicine
statistical genetics

u of alabama

u of iowa

texas a&m u

nichd

u of missouri

bayesian methods

model selection bayesian methods survival analysis

roc analysis longitudinal data bayesian methods

agreement
clustered data methods
epidemiological study design
meta-analysis multivariate survival statistical genetics

survival analysis nonlinear models longitudinal data high dimensional data model selection multiple testing