

# Package ‘JKGWAS’

March 20, 2021

**Type** Package

**Title** Contains functions used for HORT545 class assignments.

**Version** 0.1.0

**Author** Lindsey Kornowske - with code from Zhiwu Zhang

**Maintainer** Lindsey Kornowske <lindsey.kornowske@wsu.edu>

**Description** Package contains functions pertaining to tasks necessary for the imputation and analysis of genomic data.

**License** MIT

**Encoding** UTF-8

**LazyData** false

**RoxygenNote** 7.1.1

## R topics documented:

JKGLM . . . . .	1
JKManhattan . . . . .	2
JKPCA . . . . .	3
JKQQ . . . . .	3
<b>Index</b>	<b>4</b>

---

JKGLM

*Compute SNP p-values for GWAS by GLM*

---

## Description

Compute p-values for the association tests between phenotype and SNPs by Generalized Linear Model fitted for the fixed effects of SNP, with option to include Principal Components and Covariates.

## Usage

JKGLM(X, y, CV = NULL, PC = NULL)

**Arguments**

X	Matrix containing genotype data organized with SNPs as columns, individuals as rows
y	Matrix containing SNP positions and chromosomes corresponding to X
CV	Optional argument to include matrix containing covariates. Default is NULL.
PC	Optional argument to include matrix containing principal components such as an object of JKPCA. Default is NULL.

**Value**

Pvals array of length ncol of X for SNPs from fitted GLM

---

JKManhattan	<i>Manhattan Plot for GWAS Visualization</i>
-------------	--

---

**Description**

Visualize the GWAS by GLM results by Manhattan plot. User can also specify QTN.

**Usage**

```
JKManhattan(Pvals, SNP, sigcutoff = NULL, QTN = NULL)
```

**Arguments**

Pvals	Input vector of Pvals such as an object returned by the JKGLM function
SNP	Input matrix containing SNP location. Must include columns Position and Chromosome
QTN	Vector of QTN positions that is provided by the user to highlight position in the Manhattan plot. If NULL, then QTN will not be identified.
sig.cutoff	Significance threshold for visualization. If NULL, then uses bonferroni correction with $\alpha = 0.05$

**Value**

Manhattan plot with user inputs.

---

JKPCA	<i>Principle Component Analysis</i>
-------	-------------------------------------

---

**Description**

Removes PCs that are linearly dependent with the given covariates and also user can specify how many PCs to choose as co-factors

**Usage**

```
JKPCA(X, CV = NULL, npc = 5)
```

**Arguments**

X	Markers data in the form n by m with n number of individuals and m number of markers
CV	Covariates matrix in the form n by t with n number of individuals and t number of co-variates
npc	Number of Principle Components (PCs) that are specified by the user

**Value**

Principle Component Analysis

---

JKQQ	<i>QQ plot</i>
------	----------------

---

**Description**

Function to generate QQ-plot

**Usage**

```
JKQQ(Pvals)
```

**Arguments**

Pvals	Input vector of Pvals such as an object returned by the JKGLM function
-------	--

**Value**

QQplot with user inputs.

# Index

JKGLM, [1](#)  
JKManhattan, [2](#)  
JKPCA, [3](#)  
JKQQ, [3](#)