Mix: 0.1, longmean: 1.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
IETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.1, longmean: 2, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low_
TEROZYGOSITY_STATS.mean_mid_
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

val

refuges

Mix: 0.1, longmean: 2.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.1, longmean: 3, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
FIT_DISTRIBUTION_WEIBULL.shape
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercept
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low_
TEROZYGOSITY_STATS.mean_mid_
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.05, longmean: 1.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
IETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.05, longmean: 2, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept

ALLELE_STATS.pval

ALLELE_STATS.rsq

ALLELE_STATS.slope

ALLELE_STATS.var_high_lat

ALLELE_STATS.var_low_lat

ALLELE_STATS.var_mid_lat

FIT_DISTRIBUTION_GAMMA.scale

FIT_DISTRIBUTION_GAMMA.shape

FIT_DISTRIBUTION_WEIBULL.scale

IT_DISTRIBUTION_WEIBULL.shap

FST_STATS.intercept

FST_STATS.pval

FST_STATS.rsq

FST_STATS.slope

HETEROZYGOSITY_STATS.intercep

TEROZYGOSITY_STATS.mean_high

TEROZYGOSITY_STATS.mean_low_

TEROZYGOSITY_STATS.mean_mid_

HETEROZYGOSITY_STATS.pval

HETEROZYGOSITY_STATS.rsq

HETEROZYGOSITY_STATS.slope

TWO_REG_MODEL.breakpoint

TWO_REG_MODEL.diffLL

val

refuges

Mix: 0.05, longmean: 2.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.05, longmean: 3, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
FIT_DISTRIBUTION_WEIBULL.shape
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercept
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low_
TEROZYGOSITY_STATS.mean_mid_
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.01, longmean: 1.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
IETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.01, longmean: 2, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept

ALLELE_STATS.pval

ALLELE_STATS.rsq

ALLELE_STATS.slope

ALLELE_STATS.var_high_lat

ALLELE_STATS.var_low_lat

ALLELE_STATS.var_mid_lat

FIT_DISTRIBUTION_GAMMA.scale

FIT_DISTRIBUTION_GAMMA.shape

FIT_DISTRIBUTION_WEIBULL.scale

FIT_DISTRIBUTION_WEIBULL.shape

FST_STATS.intercept

FST_STATS.pval

FST_STATS.rsq

FST_STATS.slope

HETEROZYGOSITY_STATS.intercept

HETEROZYGOSITY_STATS.mean_high

HETEROZYGOSITY_STATS.mean_low

HETEROZYGOSITY_STATS.mean_mid

HETEROZYGOSITY_STATS.pval

HETEROZYGOSITY_STATS.rsq

HETEROZYGOSITY_STATS.slope

TWO_REG_MODEL.breakpoint

TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.01, longmean: 2.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.01, longmean: 3, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
FIT_DISTRIBUTION_WEIBULL.shape
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercept
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low_
TEROZYGOSITY_STATS.mean_mid_
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.001, longmean: 1.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

val

refuges

Mix: 0.001, longmean: 2, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept

ALLELE_STATS.pval

ALLELE_STATS.rsq

ALLELE_STATS.slope

ALLELE_STATS.var_high_lat

ALLELE_STATS.var_low_lat

ALLELE_STATS.var_mid_lat

FIT_DISTRIBUTION_GAMMA.scale

FIT_DISTRIBUTION_GAMMA.shape

FIT_DISTRIBUTION_WEIBULL.scale

IT_DISTRIBUTION_WEIBULL.shape

FST_STATS.intercept

FST_STATS.pval

FST_STATS.rsq

FST_STATS.slope

HETEROZYGOSITY_STATS.intercep

TEROZYGOSITY_STATS.mean_high

TEROZYGOSITY_STATS.mean_low_

TEROZYGOSITY_STATS.mean_mid_

HETEROZYGOSITY_STATS.pval

HETEROZYGOSITY_STATS.rsq

HETEROZYGOSITY_STATS.slope

TWO_REG_MODEL.breakpoint

TWO_REG_MODEL.diffLL

val

refuges

Mix: 0.001, longmean: 2.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.001, longmean: 3, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
FIT_DISTRIBUTION_WEIBULL.shape
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercept
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low_
TEROZYGOSITY_STATS.mean_mid_
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.15, longmean: 1.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.15, longmean: 2, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept

ALLELE_STATS.pval

ALLELE_STATS.rsq

ALLELE_STATS.slope

ALLELE_STATS.var_high_lat

ALLELE_STATS.var_low_lat

ALLELE_STATS.var_mid_lat

FIT_DISTRIBUTION_GAMMA.scale

FIT_DISTRIBUTION_GAMMA.shape

FIT_DISTRIBUTION_WEIBULL.scale

IT_DISTRIBUTION_WEIBULL.shap

FST_STATS.intercept

FST_STATS.pval

FST_STATS.rsq

FST_STATS.slope

HETEROZYGOSITY_STATS.intercep

TEROZYGOSITY_STATS.mean_high

TEROZYGOSITY_STATS.mean_low_

TEROZYGOSITY_STATS.mean_mid_

HETEROZYGOSITY_STATS.pval

HETEROZYGOSITY_STATS.rsq

HETEROZYGOSITY_STATS.slope

TWO_REG_MODEL.breakpoint

TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.15, longmean: 2.5, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
IT_DISTRIBUTION_WEIBULL.shap
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercep
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low
TEROZYGOSITY_STATS.mean_mid
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges

Mix: 0.15, longmean: 3, shortscale: 0.5, Glac front: 20, Marginal decrease: 1

ALLELE_STATS.intercept
ALLELE_STATS.pval
ALLELE_STATS.rsq
ALLELE_STATS.slope
ALLELE_STATS.var_high_lat
ALLELE_STATS.var_low_lat
ALLELE_STATS.var_mid_lat
FIT_DISTRIBUTION_GAMMA.scale
FIT_DISTRIBUTION_GAMMA.shape
FIT_DISTRIBUTION_WEIBULL.scale
FIT_DISTRIBUTION_WEIBULL.shape
FST_STATS.intercept
FST_STATS.pval
FST_STATS.rsq
FST_STATS.slope
HETEROZYGOSITY_STATS.intercept
TEROZYGOSITY_STATS.mean_high
TEROZYGOSITY_STATS.mean_low_
TEROZYGOSITY_STATS.mean_mid_
HETEROZYGOSITY_STATS.pval
HETEROZYGOSITY_STATS.rsq
HETEROZYGOSITY_STATS.slope
TWO_REG_MODEL.breakpoint
TWO_REG_MODEL.diffLL

value

refuges