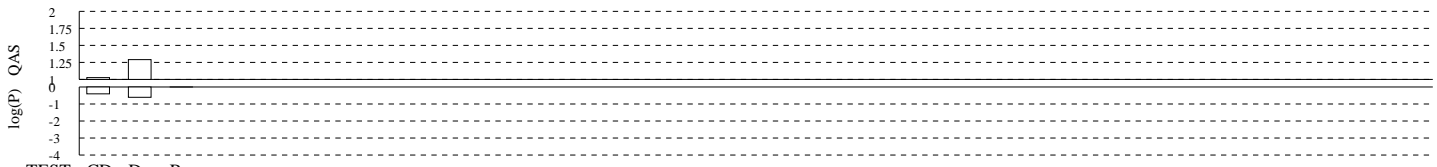


TRAX PAGE for rs11884476

Project: ARG Group A European Americans				Race	Freq 1		Freq 2
Marker:	SNP_A-8393199	Maf: 0.06052		Caucasian	917	0.93948	0.06052
SNP rs ID:	rs11884476			Total	917		
Allele 1:	C (0.93948)	Allele 2:	G (0.06052)				
Chromosome:	2	Coordinates:	206,318,593	Gene region: PARD3B			

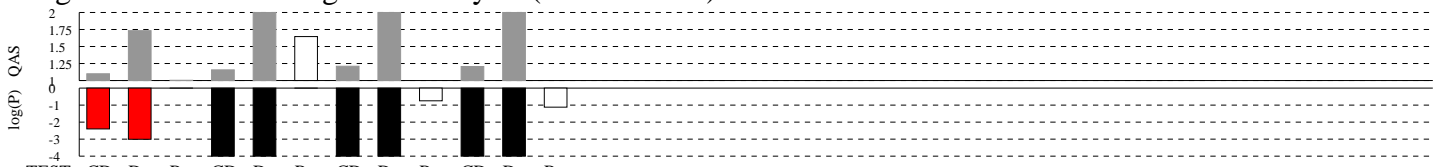
Caucasian	
SC	699
OSN	81
HREU	121
SP	16
Total	917

Infection Tests



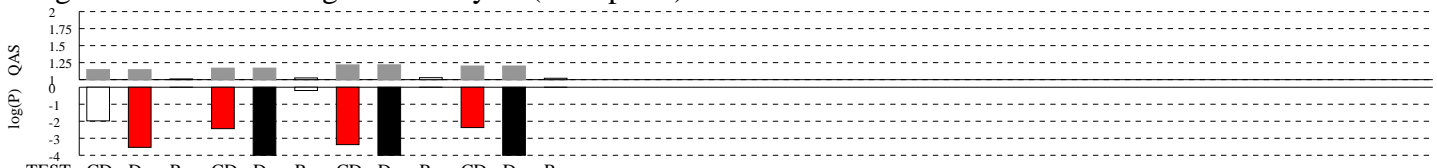
TEST: CD D R
 GRP: H H H
 POP: CA CA CA

Progression disease categorical analysis (dichotomous)



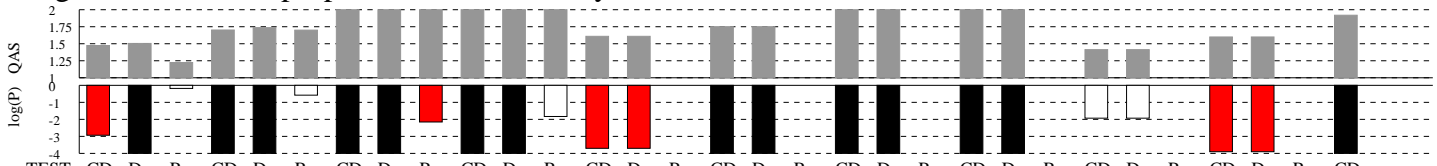
TEST: CD D R CD D R CD D R CD D R
 GRP: CD4 CD4 CD4 93 93 93 87 87 87 DTH DTH DTH
 POP: CA CA CA CA CA CA CA CA CA CA CA CA

Progression disease categorical analysis (multipoint)



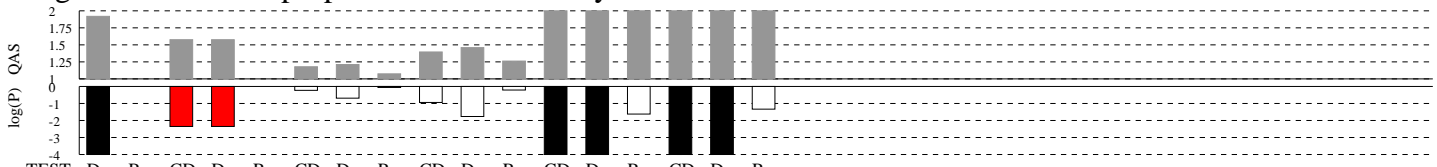
TEST: CD D R CD D R CD D R CD D R
 GRP: CD4 CD4 CD4 93 93 93 87 87 87 DTH DTH DTH
 POP: CA CA CA CA CA CA CA CA CA CA CA CA

Progression survival proportional hazard analysis



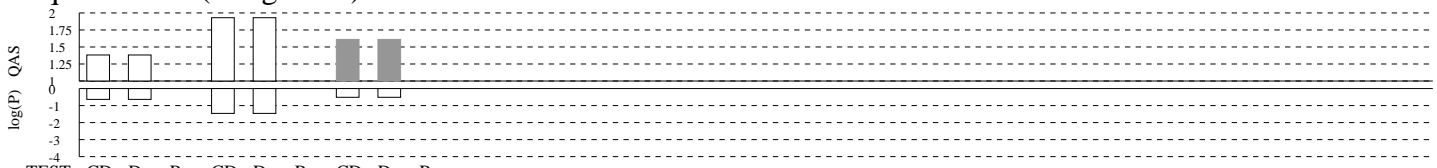
TEST: CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R CD D R
 GRP: CD4 CD4 CD4 93 93 93 87 87 87 DTH DTH DTH CD4 CD4 CD4 93 93 93 87 87 87 DTH DTH DTH CD4 CD4 CD4 93 93 93 87
 POP: CA CA CA CA CA CA CA CA CA CA CA CA GA GA GA GA GA GA GA GA GA GA MA MA MA MA MA MA MA MA MA MA MA MA MA MA MA MA

Progression survival proportional hazard analysis



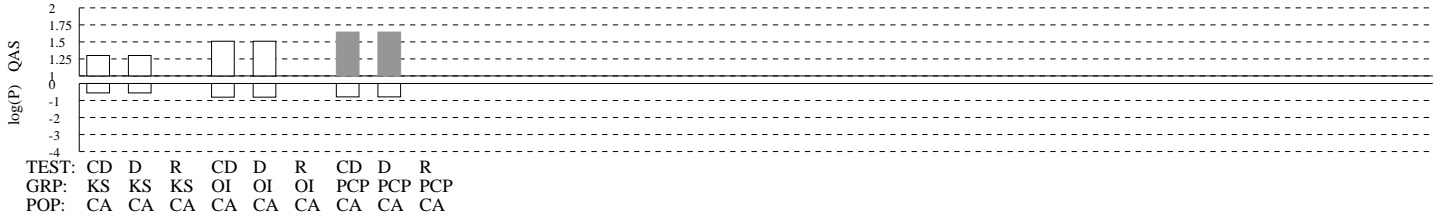
TEST: D R CD D R CD D R CD D R CD D R CD D R CD D R
 GRP: 87 87 DTH DTH DTH CD4 CD4 CD4 93 93 93 87 87 87 DTH DTH DTH
 POP: MA MA MA MA MA HE HE HE HE HE HE HE HE HE HE HE HE HE HE HE HE

Sequelae tests (categorical) for the first infection

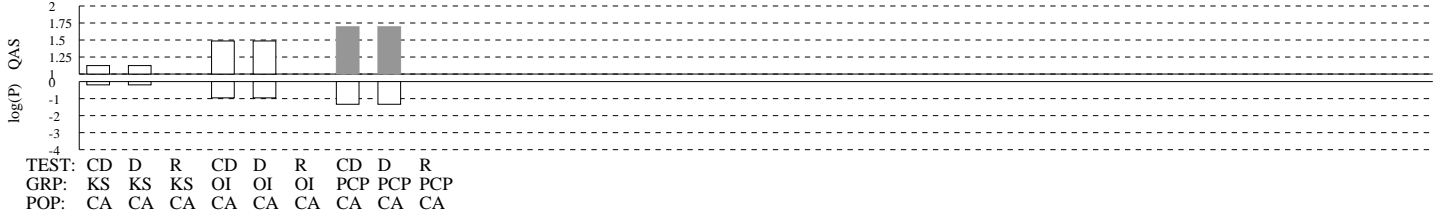


TEST: CD D R CD D R CD D R
 GRP: KS KS KS OI OI OI PCP PCP PCP
 POP: CA CA CA CA CA CA CA CA CA

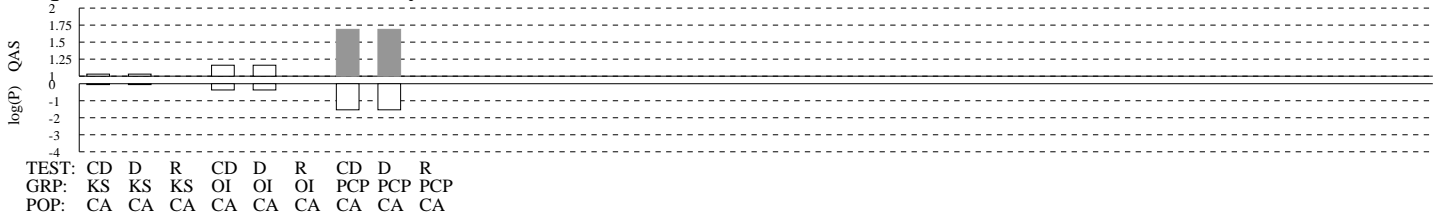
Sequela tests (categorical) for any infection order



Sequela tests (survival) for the first infection



Sequela tests (survival) for any infection order



Black: $QAS > 2$ or $-\log(P) < -4$

Red: $-4 < -\log(P) < -2$

Gray: $1/QAS$

White: QAS or $-\log(P)$