

Figure 1A

Enzastaurin Phase 3 Design – Prelude Trial

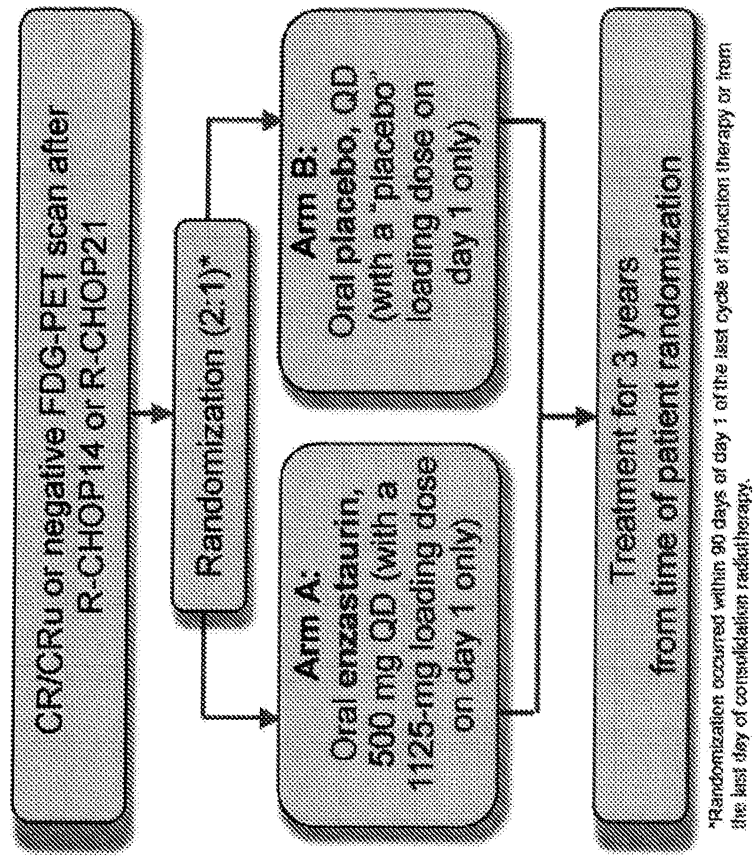


Figure 1B. Kaplan-Meier Plot Time to Overall Survival

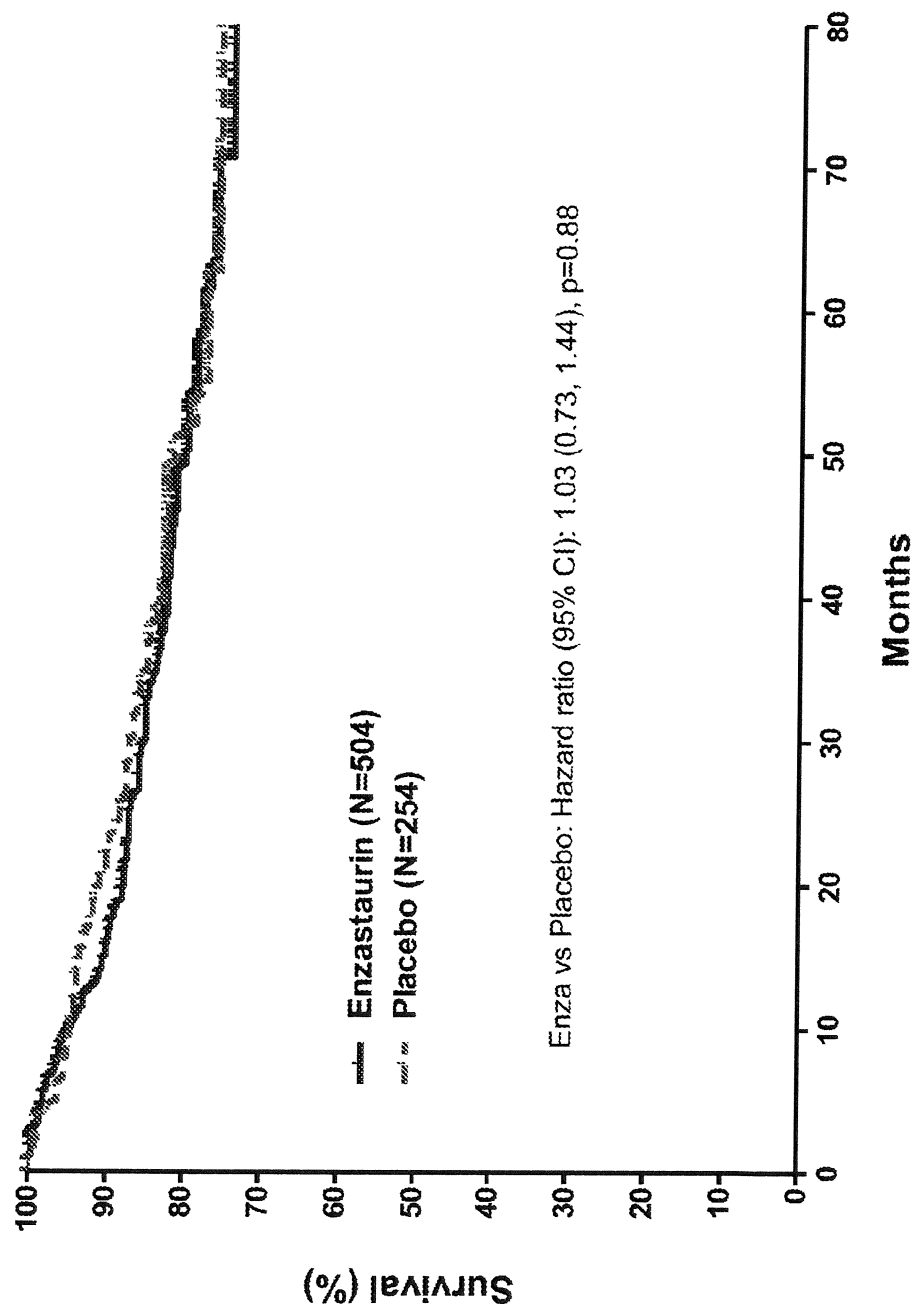


Figure 1C. Kaplan-Meier Plot Time to Overall Survival

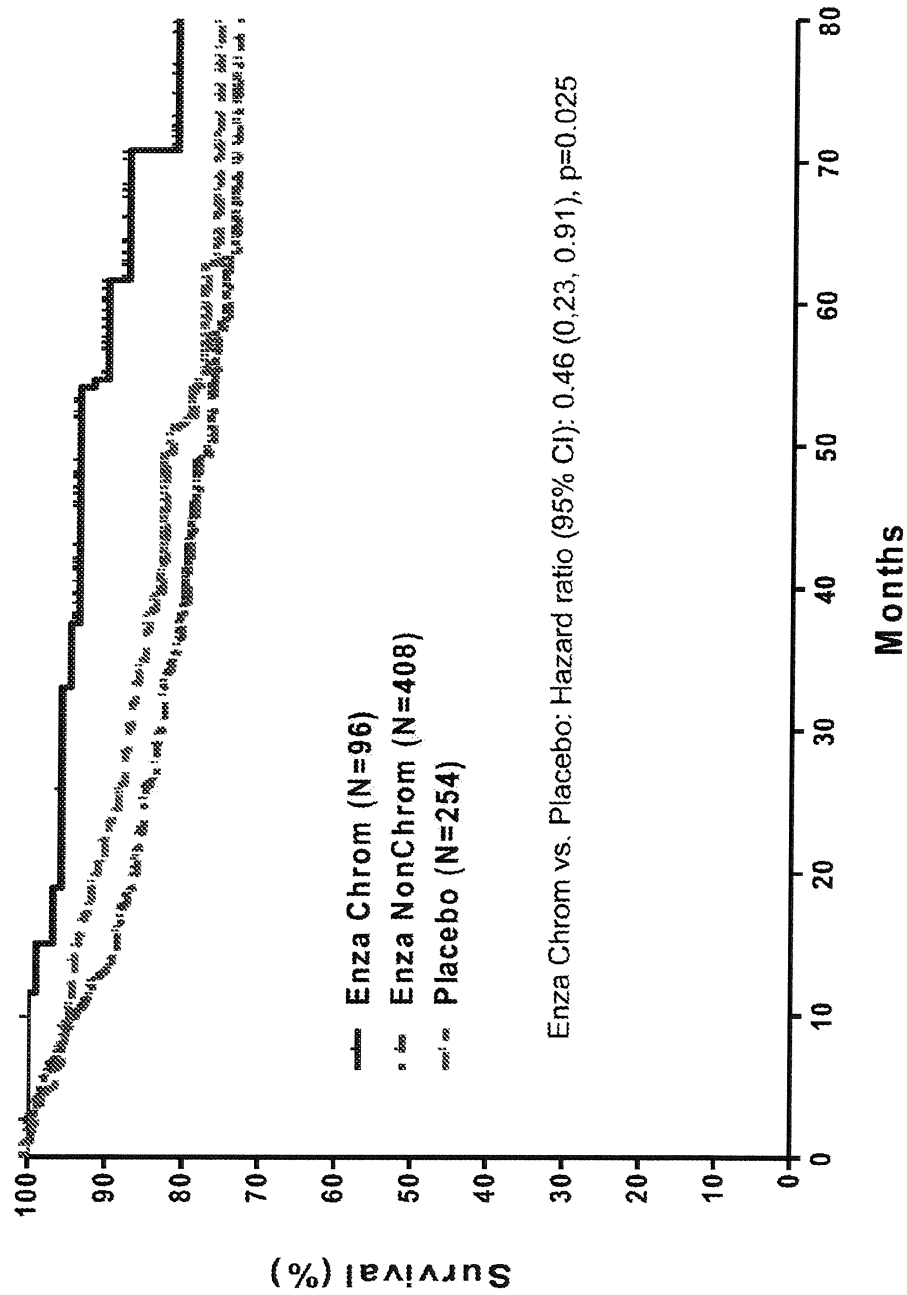


Figure 2A. Kaplan-Meier Time to Event Plot
Overall Survival in Enza Treated Patients Carrying Different Genotypes

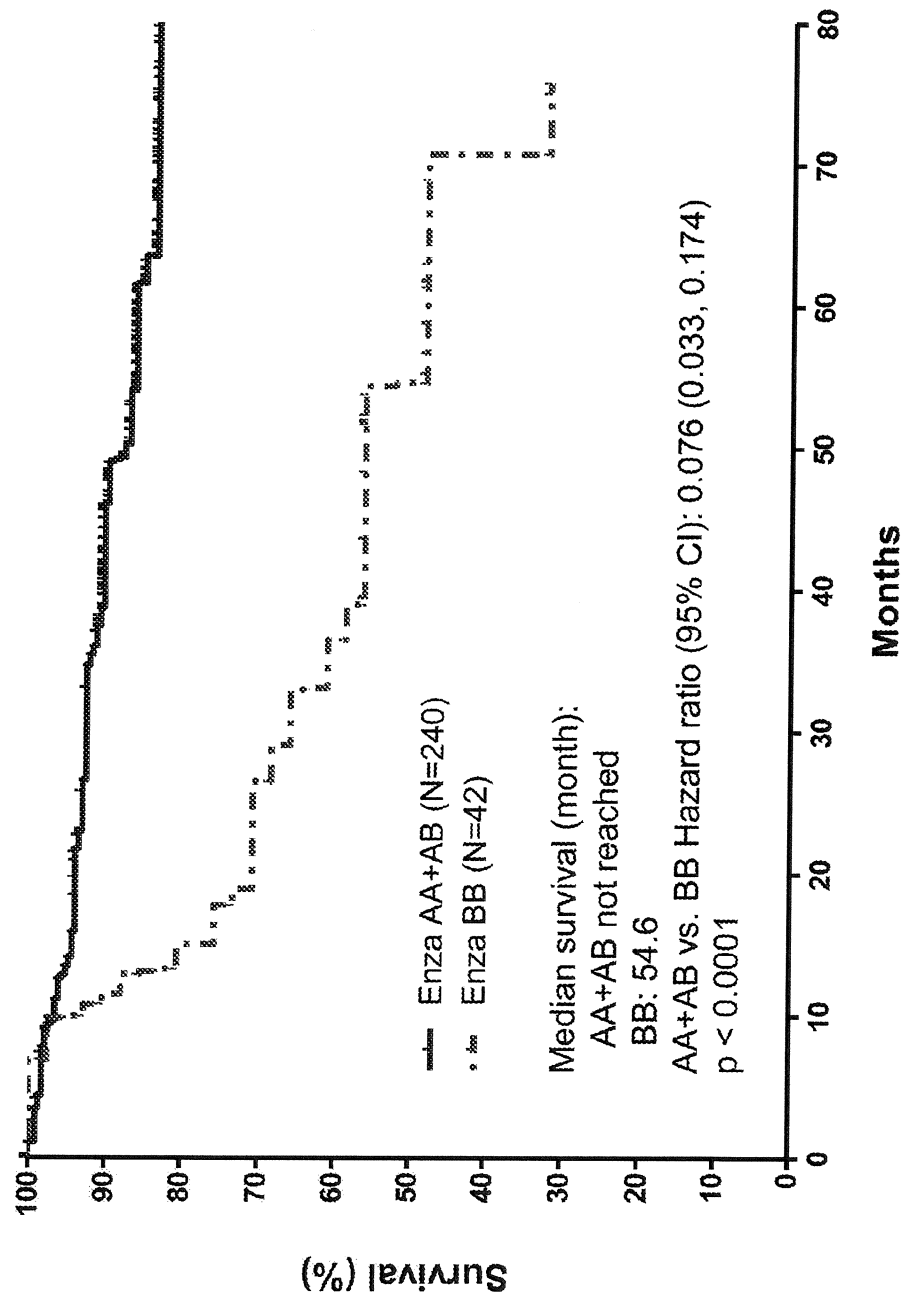


Figure 2B. Kaplan-Meier Time to Event Plot
Overall Survival in Placebo treated Patients Carrying Different Genotypes

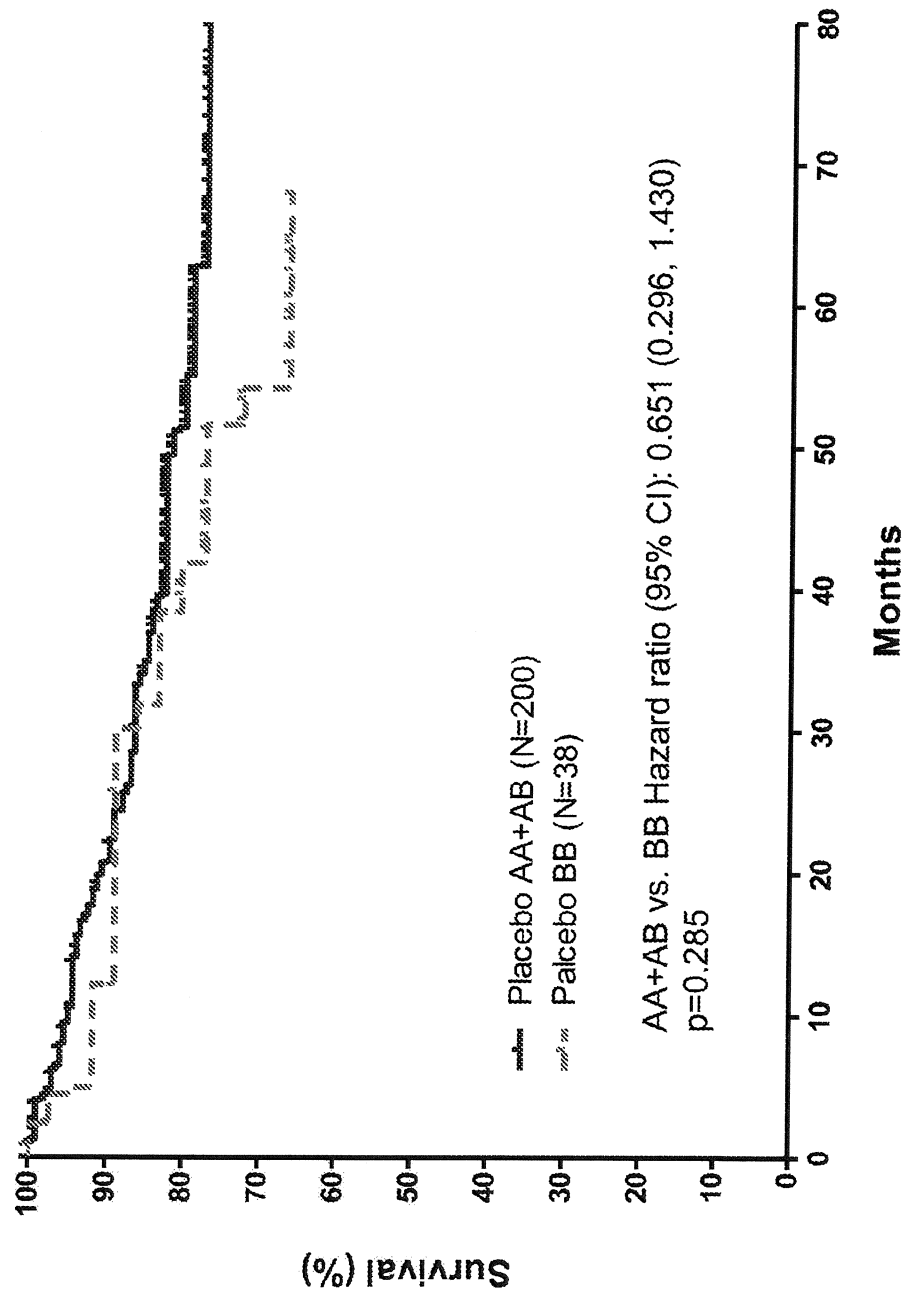
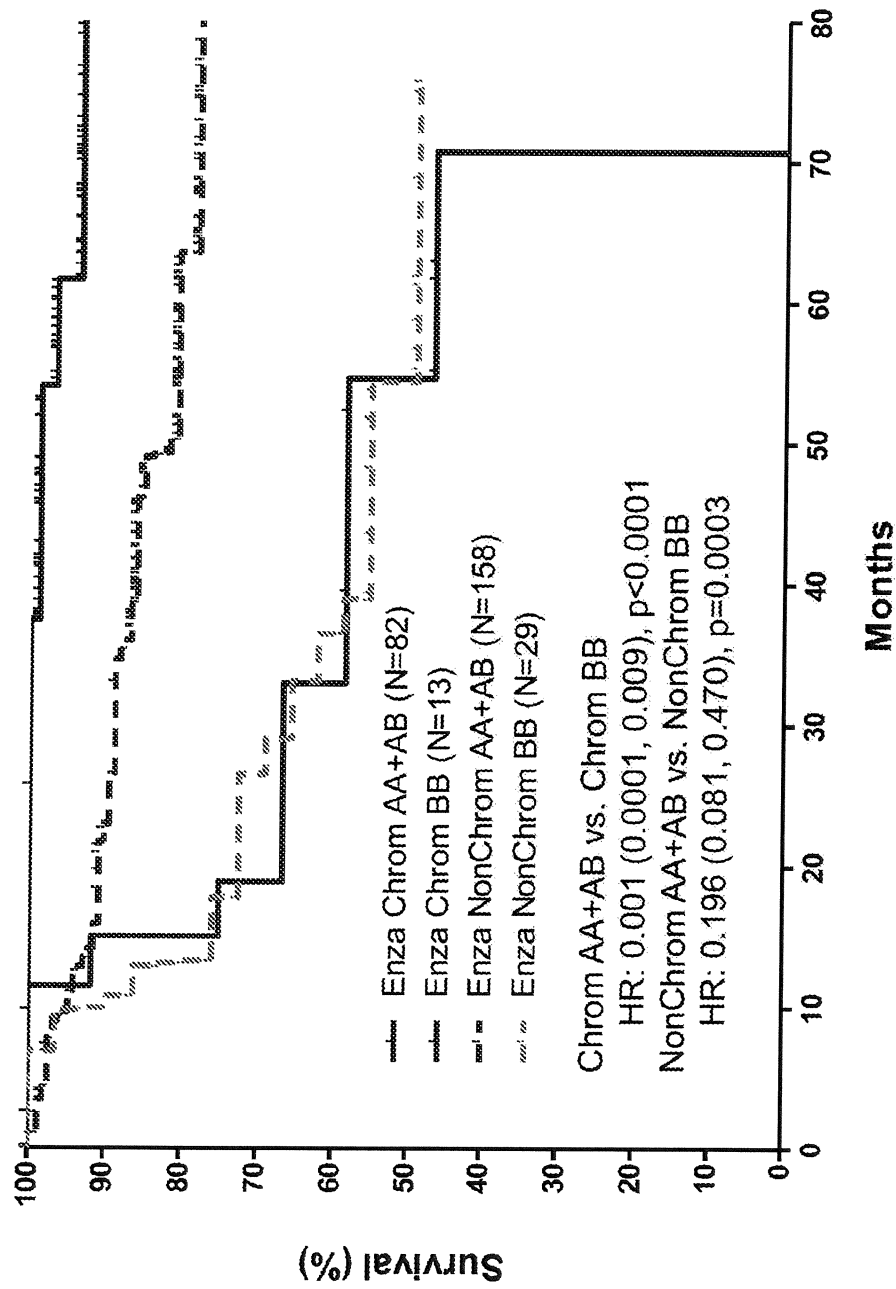


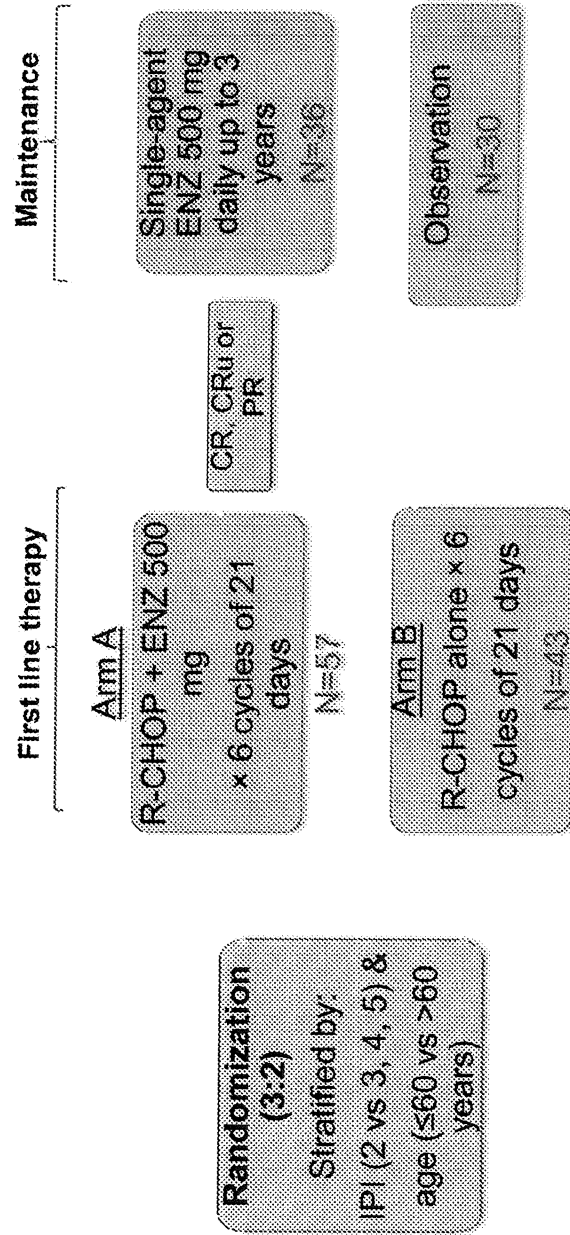
Figure 2C. Kaplan-Meier Plot Time to Overall Survival
Interaction between Chromaturia and rs309605



Note: No multiplicity adjustment for between subgroup comparison.
HR=hazard ratio; p=Logrank test p-value.

Figure 3A

Phase II 1st Line RCHOP + Enzastaurin vs. RCHOP in intermediate & high risk DLBCL



- Complete restaging was performed at the end of Cycle 6, and responses were categorized according to the International Working Group (IWG) criteria, 1999
- Maintenance therapy was continued in Arm A with response evaluated every 8 weeks

Figure 3B. S028 First Line
Kaplan-Meier Time to Event Plot Overall Survival

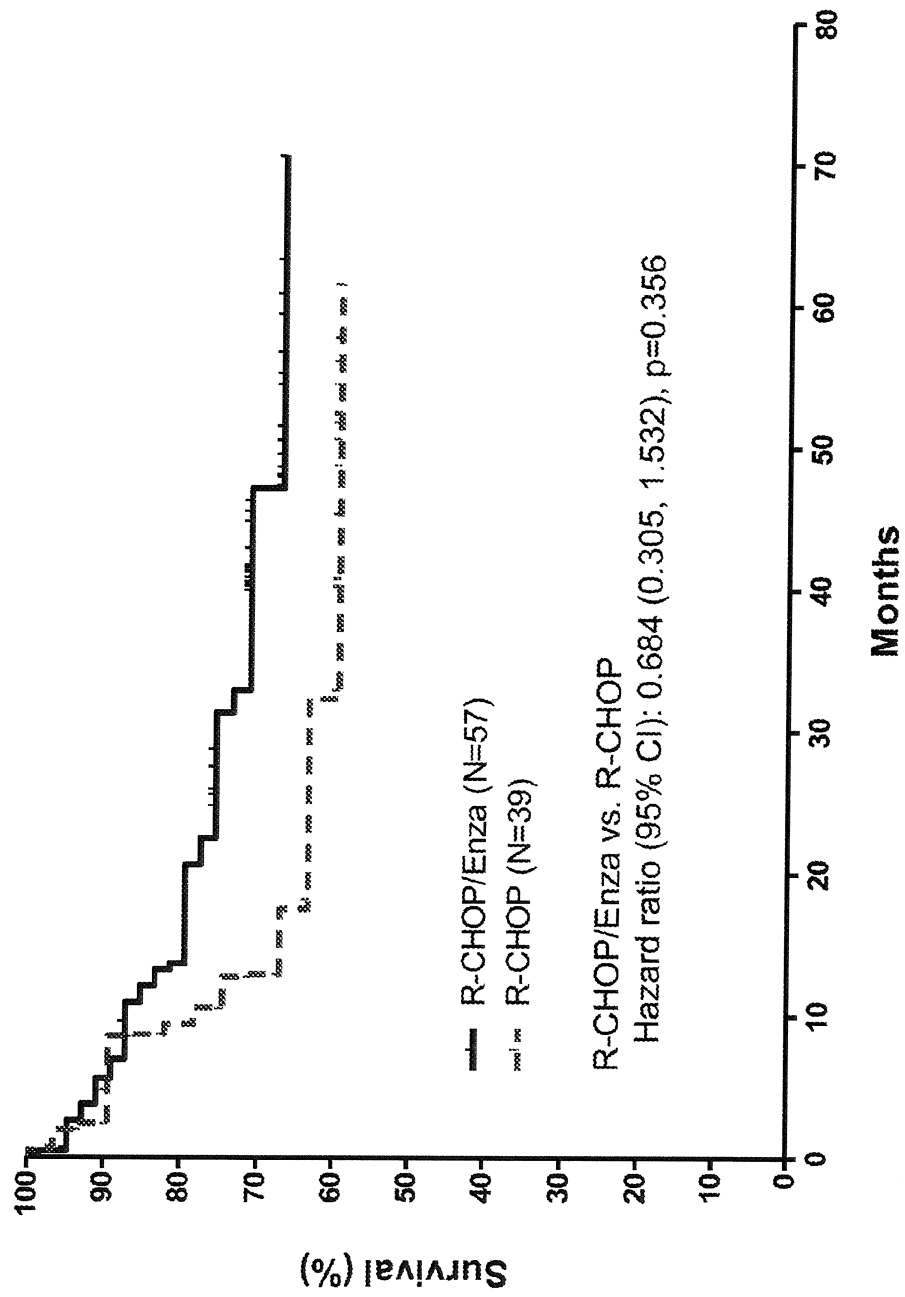


Figure 3C. S028 First Line
Overall Survival in Enza + RCHOP Treated Patients Carrying Different Genotypes

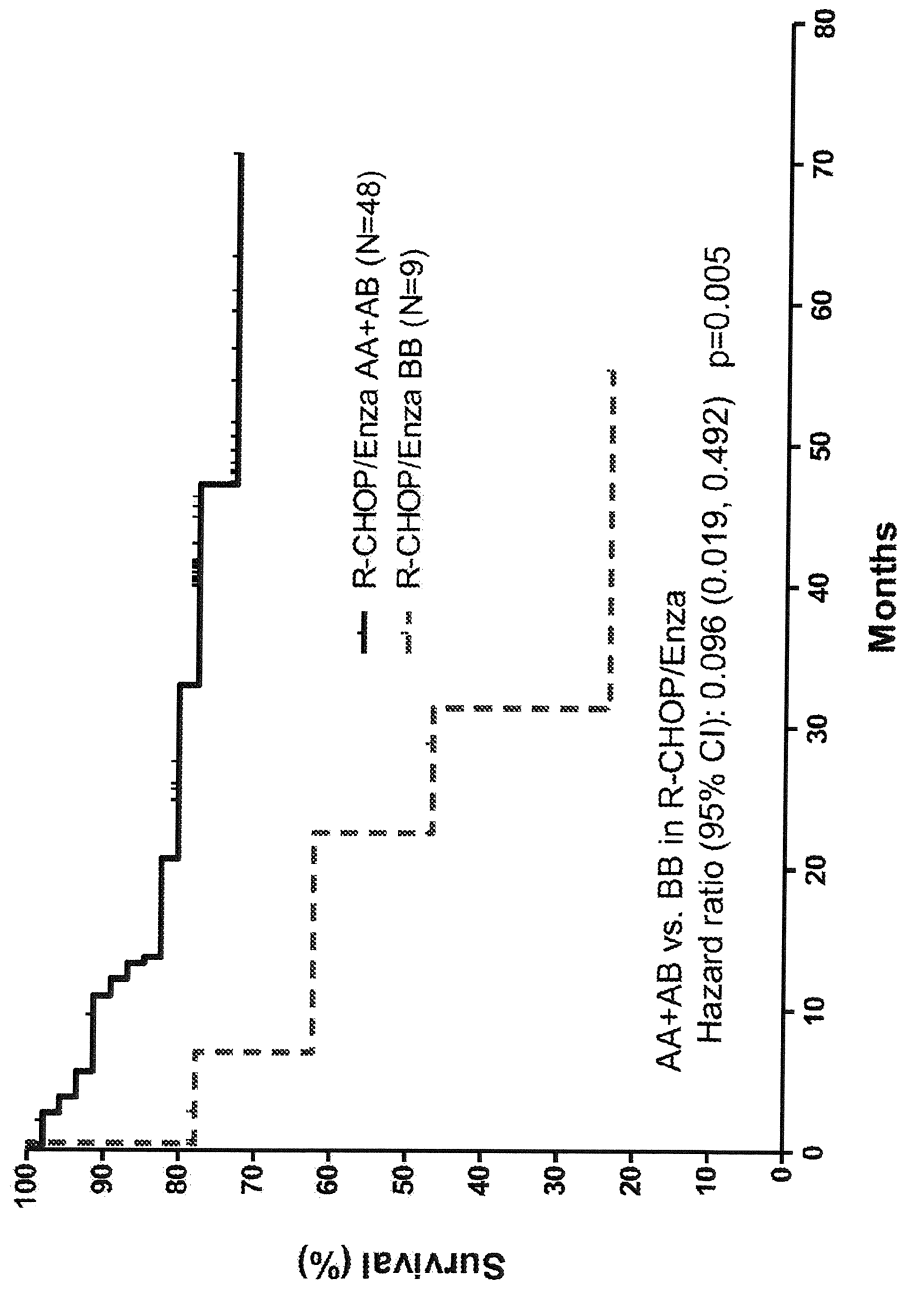


Figure 3D. S028 First Line
Overall Survival in R-CHOP Treated Patients Carrying Different
Genotypes

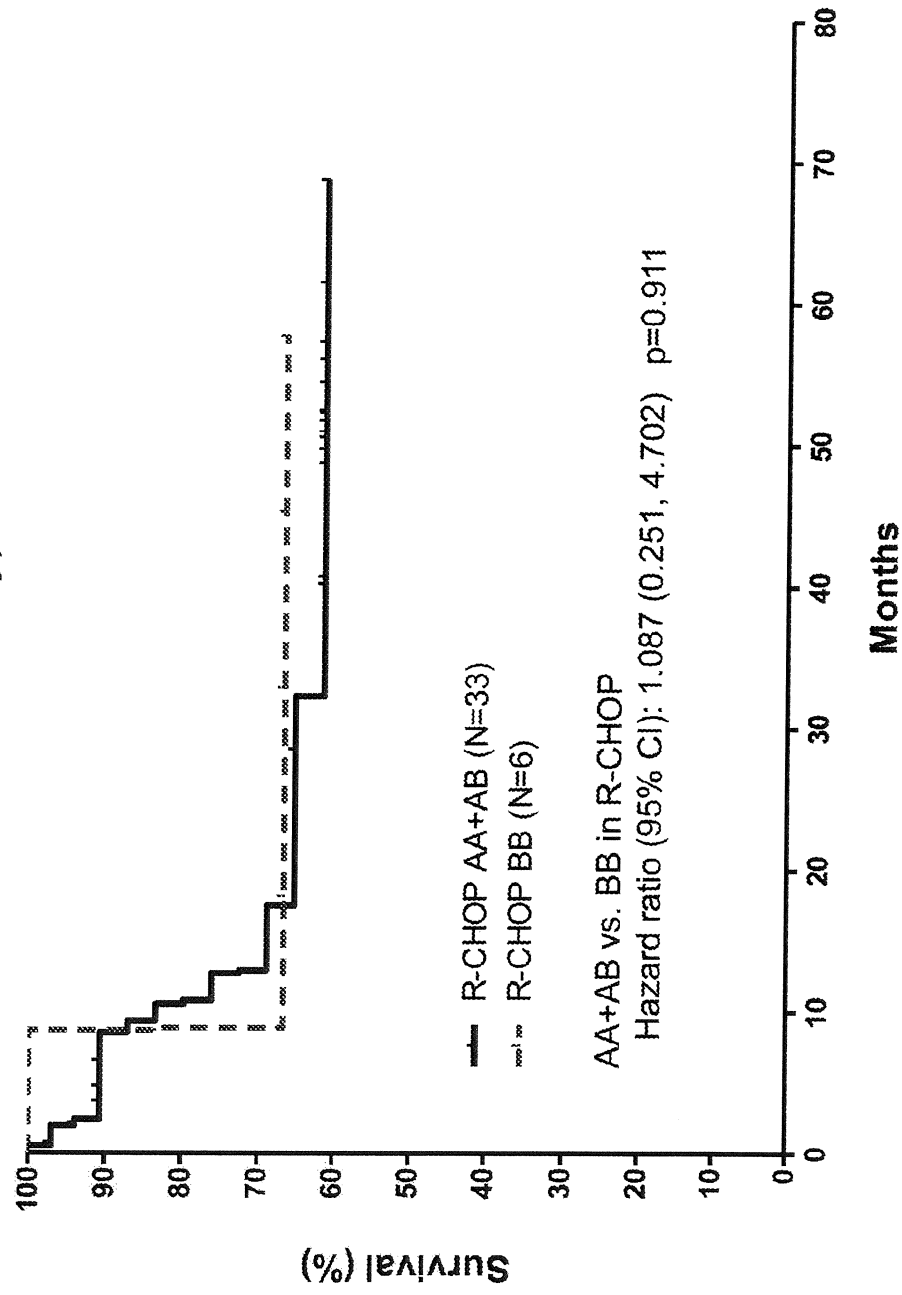


Figure 3E: Study S028 Kaplan-Meier Plot - Overall Survival
In Patients IPI Score > 2 and Genotype AA+AB

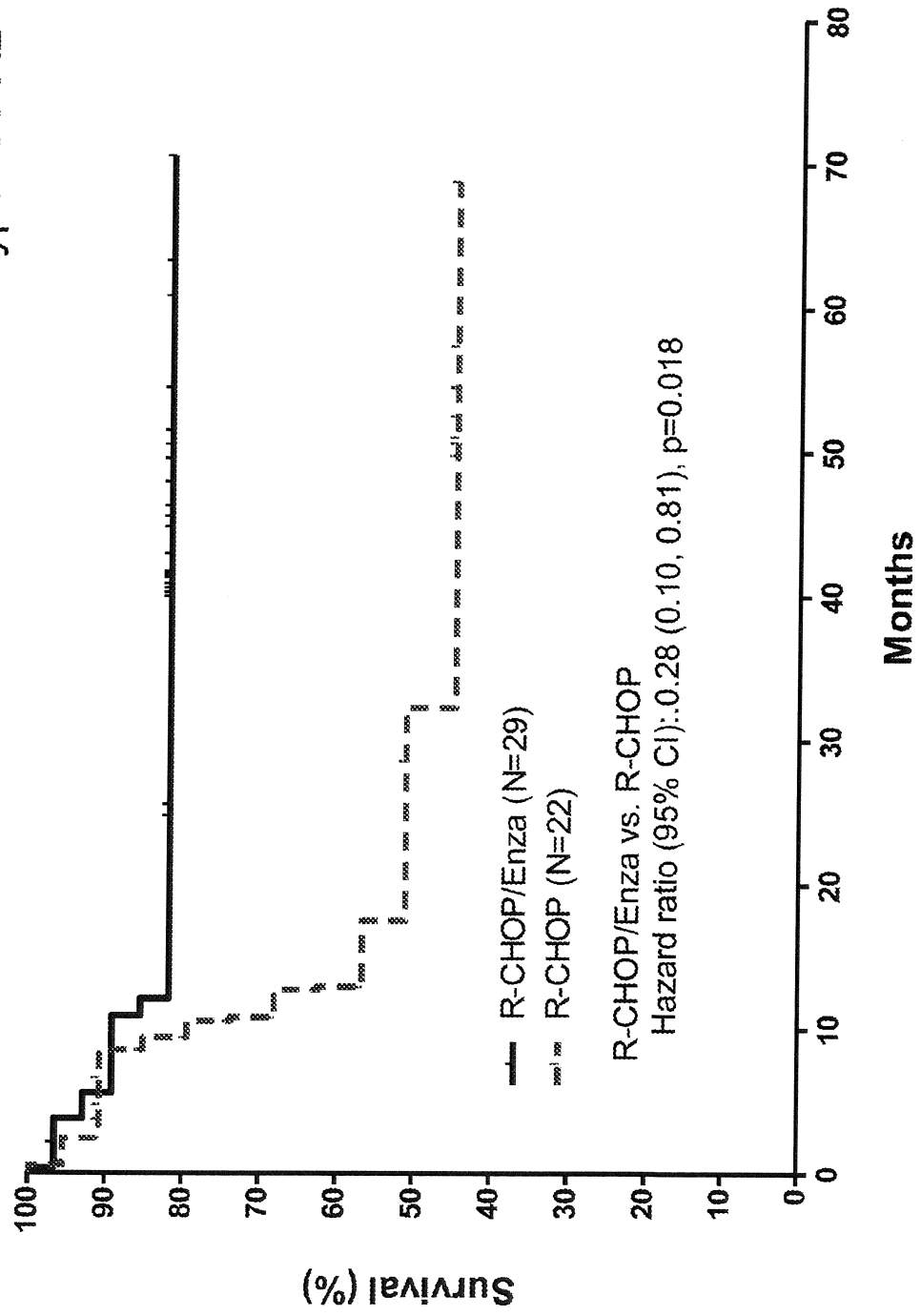


Figure 3F: Study S028 Kaplan-Meier Plot - PFS
In Patients IPI Score > 2 and Genotype AA+AB

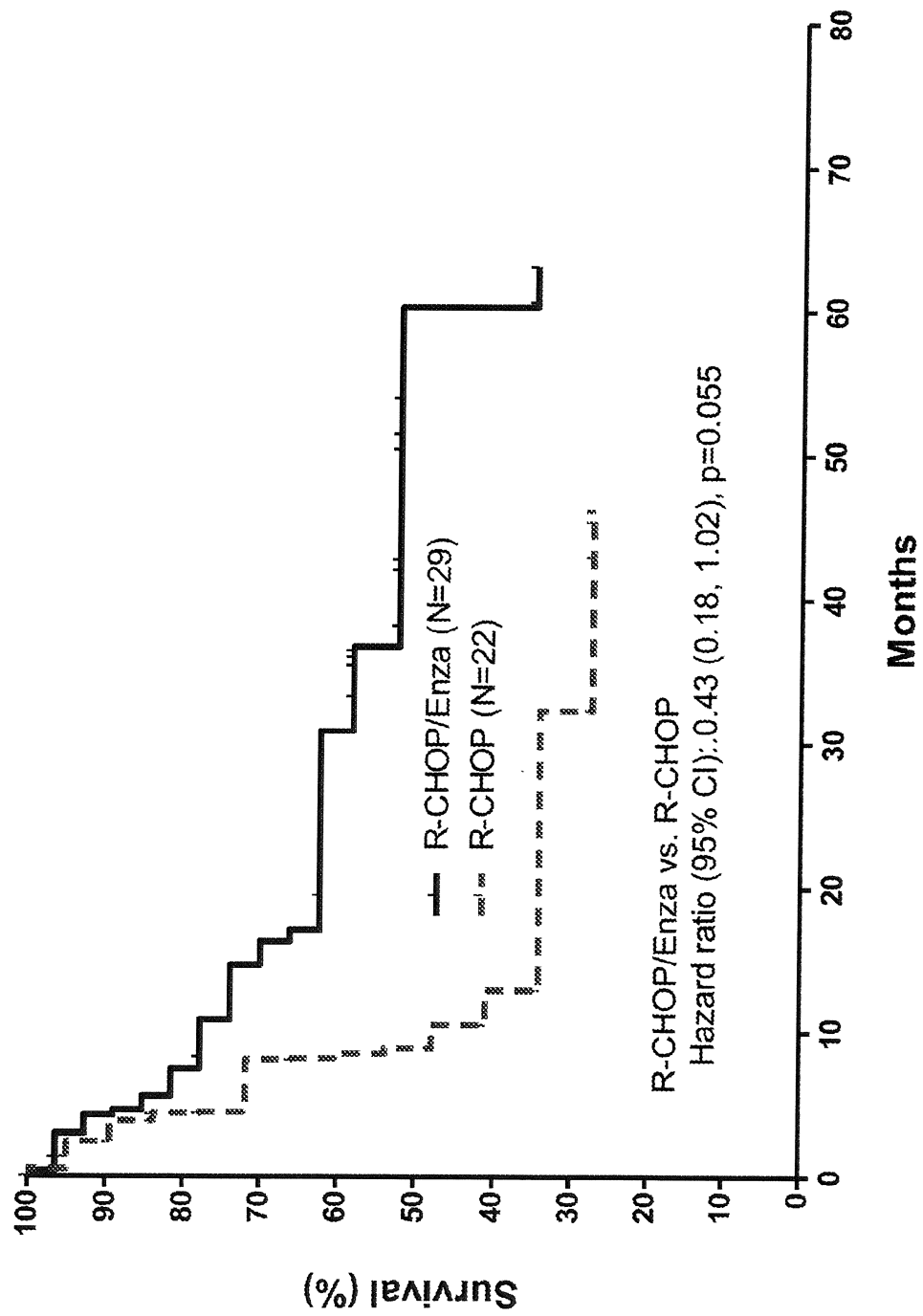


Figure 4A: S008 GBM Kaplan-Meier Plot Time to Overall Survival

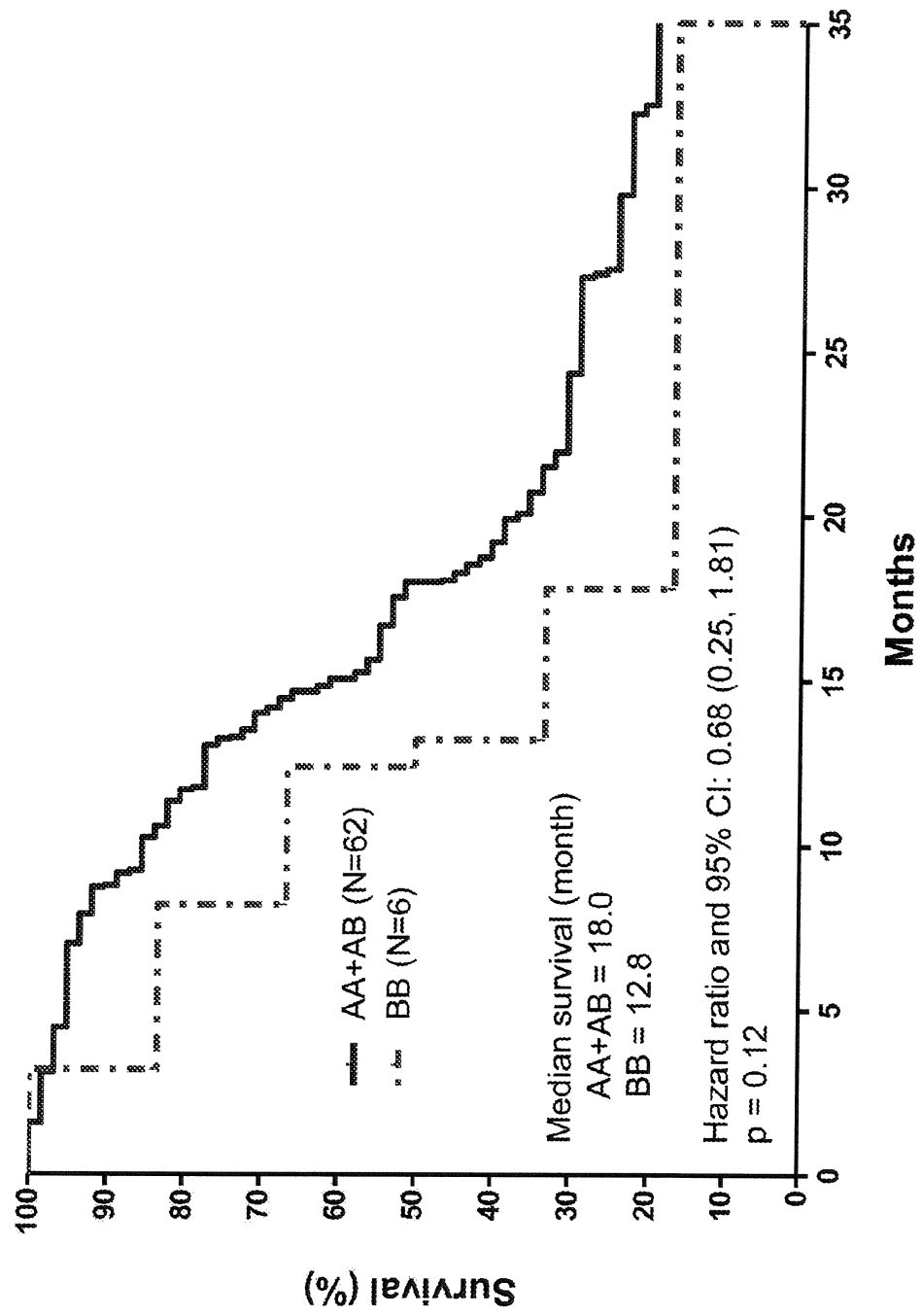
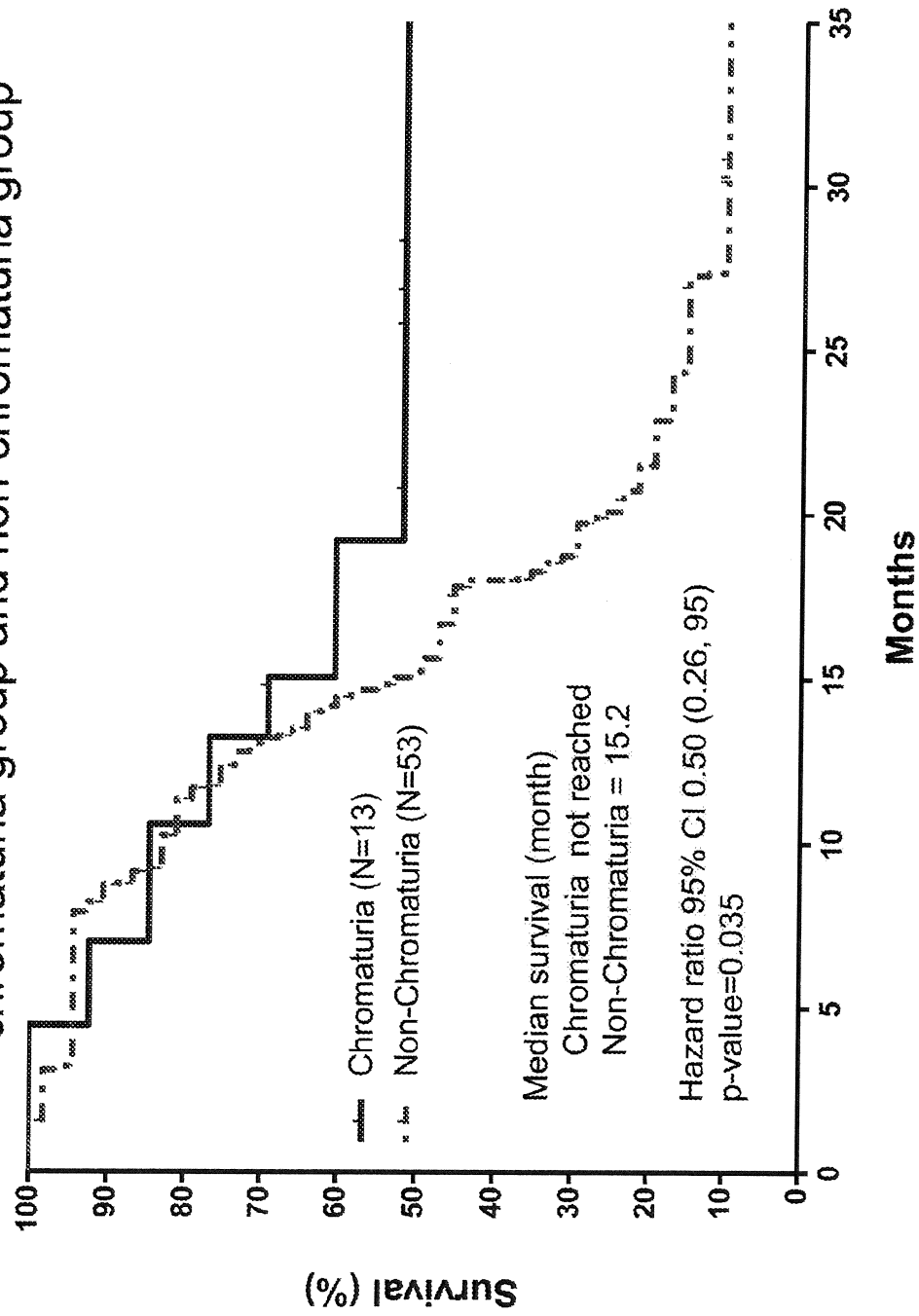


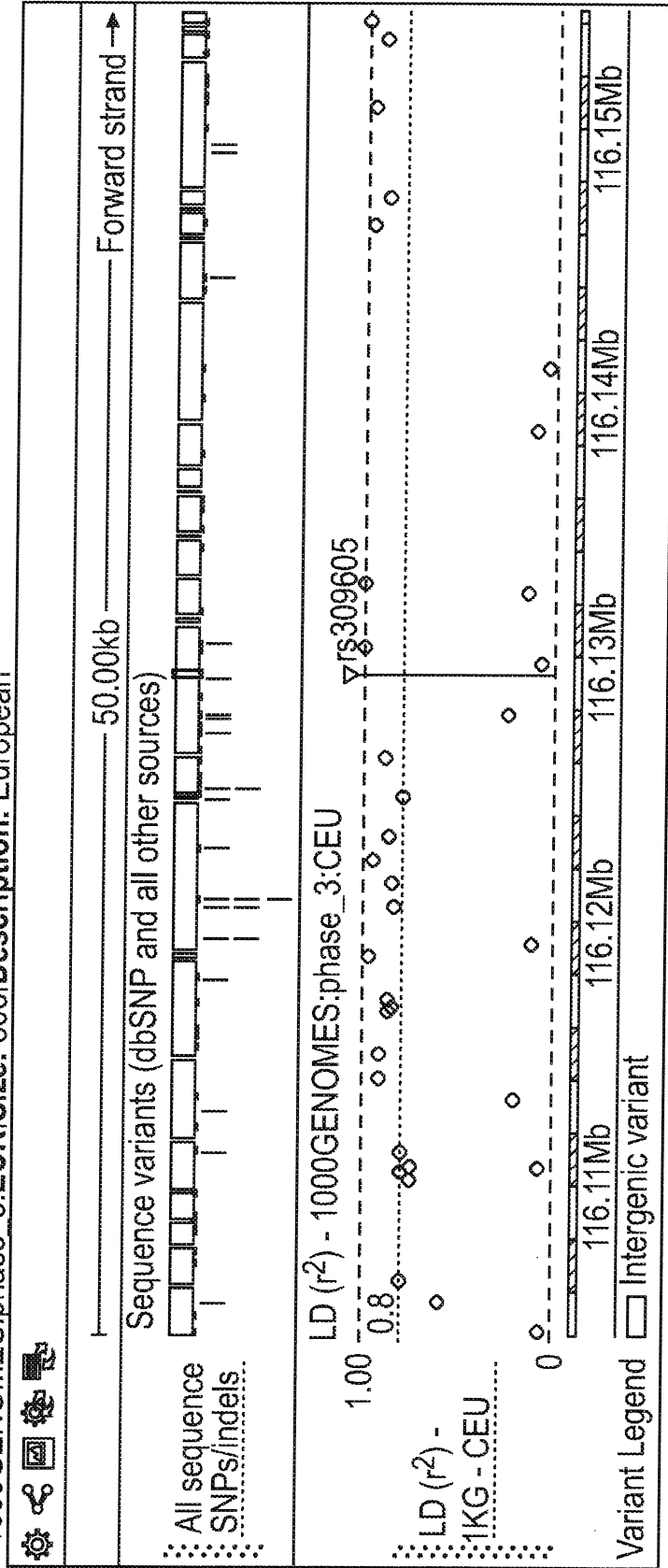
Figure 4B: Kaplan-Meier plot showing time to overall survival in the chromaturia group and non-chromaturia group



Chromaturia was coded "Urine discoloration".

FIG. 5A
1000GENOMES:phase_3:CEU
Size: 99

Description: Utah residents with Northern and Western European ancestry
Super-Population: 1000GENOMES:phase_3:EUR
Size: 503
Description: European





NC_000008.11:115M..115M(74Kbp) Find:		<=> -	+ ATG	Tools ▾ $\frac{x}{x}$	* Tracks ▾	? *
j114,800 K	j114,900 K	j115 M	j115,200 K	j115,300 K	j115,400 K	
Genes, NCBI Homo sapiens Annotation Release 107, 2015-03-13						
dbSNP Build 146 (Homo sapiens Annotation Release 107) all data						
Suspect variations, dbSNP Build 146 (Homo sapiens Annotation Release 107)						
Somatic alleles, dbSNP Build 146 (Homo sapiens Annotation Release 107)						
dbSNP Build 146 (Homo sapiens Annotation Release 107) GMAF>=0.01						
ClinVar Short Variations based on dbSNP Build 146 (Homo sapiens An...						
Cited Variants, dbSNP Build 146 (Homo sapiens Annotation Release 1...						
[1]	[2][2][1]	[1]	[1]	[3]	[1]	[1]
GWAS Results based on dbSNP Build 146 (Homo sapiens Annotation Rel...						