

Variant_ID	Chr	Region_Start	Region_End	ContigStart	ContigEnd	Local_FragStart	Local_FragEnd	Type_OF_Inv	Length_1bp	Max_Length_of_Samples	Samples	Population	MR_Data	SeqDup_Regi	GENCODE_BAGENCODE_BAH	Main_Data1	Main_Data2	Linked_Reads	3000_Decon	Comment	Y-Chromosome	Pilot_1_DOC	Pilot_1_PI_T	Complete_De	Complete_De	Pilot_1_Rego	Phase_1_Reg	OMNI_Thic	Library	PCR	#_OMNI_MP	#_Seq_Over1	Source_of_Val	Comment_Validation	QCData	Validation_Status
CG_JV_11717921_11718340	Y	11717921	11718340	11717921	11718340	11717921	11718340	Del	419	419	1	NA20904	YR	Heterochrom chr1:11717921	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_11718967_11719807	Y	11718967	11719807	11718967	11719807	11718967	11719807	Del	780	780	1	NA21891	CEU	Heterochrom chr1:11718967	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_9968118_9970910	Y	9968118	9970910	9968118	9970910	9968118	9970910	Del	2796	2796	6	NA20994	CEU	Other chr1:9968118	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_9968118_9970910	Y	9968118	9970910	9968118	9970910	9968118	9970910	Del	2796	2796	6	NA21817	CEU	Other chr1:9968118	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_9968118_9970910	Y	9968118	9970910	9968118	9970910	9968118	9970910	Del	2796	2796	6	NA20851	CEU	Other chr1:9968118	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_9968118_9970910	Y	9968118	9970910	9968118	9970910	9968118	9970910	Del	2796	2796	6	NA21891	CEU	Other chr1:9968118	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_9968118_9970910	Y	9968118	9970910	9968118	9970910	9968118	9970910	Del	2796	2796	6	NA20901	YR	Other chr1:9968118	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_9968118_9970910	Y	9968118	9970910	9968118	9970910	9968118	9970910	Del	2796	2796	6	NA20904	YR	Other chr1:9968118	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
CG_JV_10019119_10021620	Y	10019119	10021620	10019119	10021620	10019119	10021620	Del	2500	2500	1	NA20851	CEU	Other chr16:10019119	NA	PI.Completed	CG	Complete_DeNA	Pilot_1	Repeat	No	-	-	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
BD_31	Y	7186799	7187145	7186799	7187145	7186799	7187145	Del	346	347	1	NA20996	CEU	X-Degenerate chr1:7186799	RP.BreakDanc	SLX	This_works	NA	Pilot_1	BreakDancr	No	-	+	-	-	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
BD_32	Y	1714951	1714952	1714951	1714952	1714951	1714952	Del	297	301	1	NA21819	CEU	X-Degenerate NA	NA	NA	RP.BreakDanc	SLX	This_works	NA	Pilot_1	BreakDancr	No	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
BD_33	Y	17946760	17946762	17946760	17946762	17946760	17946762	Del	269	300	1	NA21819	CEU	X-Degenerate NA	NA	NA	RP.BreakDanc	SLX	This_works	NA	Pilot_1	BreakDancr	No	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
BD_34	Y	17150952	17150953	17150952	17150953	17150952	17150953	Del	291	300	1	NA21819	CEU	X-Degenerate NA	NA	NA	RP.BreakDanc	SLX	This_works	NA	Pilot_1	BreakDancr	No	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		
BD_35	Y	17802807	17802808	17802807	17802808	17802807	17802808	Del	218	300	1	NA20910	YR	X-Degenerate NA	NA	NA	RP.BreakDanc	SLX	This_works	NA	Pilot_1	BreakDancr	No	-	+	-	-	-	0	1	NA	Failed_PCR	Non-Redundant	Unvalidated		