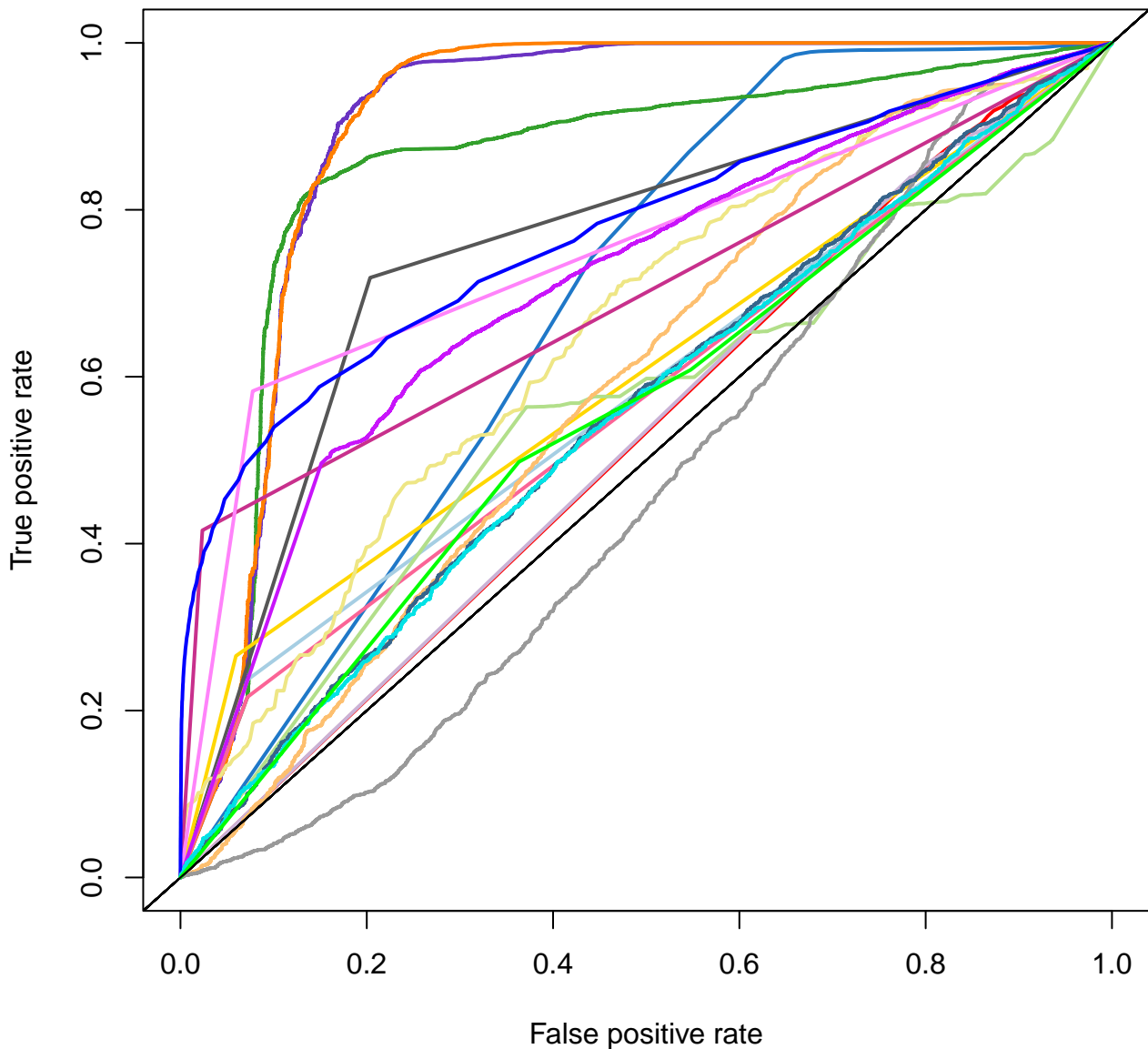























Single Feature Logistic Regression AUCs



	Homology Length
	Insertion Length
	SPAN
	gnomAD distance (5')
	gnomAD distance (3')
	Deletion
	Duplication
	Inversion
	Translocation
	Homology GC Content
	Insertion GC Content
	Distance to LINEs
	Distance to SINEs
	Sample SV Count
	Whole Gene Impact
	Exon Impact
	Nearest SV Distance
	SV Count in 5Mbp
	Replication time (5')
	Replication time (3')
	tp53 mutation status