

Table S4Yeast **CAR** genes, recognized functions, and human orthologs

CAR gene	recognized function	human ortholog
<i>CDC73</i>	transcription factor (Paf1 complex)	CDC73
<i>CHL1</i>	DNA helicase	DDX11
<i>CTK1</i>	CTD kinase I, catalytic subunit	CDK12
<i>CTK2</i>	CTD kinase I, cyclin-like subunit	CYCLIN K
<i>CTK3</i>	CTD kinase I, regulatory (?) subunit	?
<i>FUN12</i>	eIF5B	EIF5B
<i>GCN5</i>	HAT	KAT2A
<i>HOG1</i>	protein kinase	p38 MAP kinase
<i>HRR25</i>	casein kinase	CSNK1D
<i>HTZ1</i>	histone variant	H2AFX
<i>JEM1</i>	chaperone	DNAJA3
<i>LIA1</i>	hydroxylase	DOHH
<i>MRT4</i>	mRNA turnover	MRTO4
<i>NOT5</i>	transcription; RNA degradation	CNOT3
<i>PHO2</i>	transcription factor	PITX3
<i>PUS1</i>	pseudo-U synthase	PUS1
<i>RAI1</i>	RNase subunit; mRNA 5' PPase	DOM3Z
<i>RDS2</i>	unk	
<i>RVS161</i>	cytoskeleton; budding	BIN3
<i>SAC1</i>	inositol phosphate phosphatase	SACM1L
<i>SET2</i>	histone methyltransferase	SETD2
<i>SPT7</i>	subunit of SAGA txn complex	SUPT7L
<i>UME6</i>	transcription factor	
<i>YPL260w</i>	unk	
<i>ZUO1</i>	chaperone	DNAJC2

CAR genes from Table 1, and recognized functions from SGD. Human orthologs from NCBI Entrez Gene, via NCBI BLAST, NCBI HOMOLOGENE, ExPASy GPSDB, SMART database, GeneCards.