		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
CADC	Gene expression	1.001	1.002	2.418308e-02	1.001	1.002	2.549198e-01
SARC	Replication timing	1.001	1.001	1.981822e-04	1.001	1.001	1.513193e-03
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
DDC A	Gene expression	1.001	1.001	2.452259e-04	1.000	1.000	4.575905e-01
BRCA	Replication timing	1.001	1.001	7.809351e-20	1.001	1.001	5.364861e-17
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
CDM	Gene expression	1.001	1.002	0.0168452408	1.000	1.001	5.841908e-01
GBM	Replication timing	1.001	1.001	2.795532e-10	1.001	1.001	4.492228e-09
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
OV	Gene expression	1.001	1.002	2.802480e-07	1.001	1.001	1.691032e-02
	Replication timing	1.001	1.001	1.280088e-16	1.001	1.001	4.708444e-12
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
LUSC	Gene expression	1.002	1.002	0.0005681217	1.001	1.002	0.009876143
	Replication timing	1.000	1.001	0.001978267	1.000	1.001	0.037026772
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Gene expression	1.004	1.005	7.530880e-10	1.003	1.004	2.772250e-05
STAD	Replication timing	1.004	1.003	2.101723e-10	1.003	1.004	7.506947e-06
	Replication timing	1.001	1.002	2.1017236-10	1.001	1.001	7.5009476-00
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Gene expression	1.003	1.004	1.936049e-07	1.001	1.002	2.762368e-02
UCEC	Replication timing	1.002	1.002	2.574777e-19	1.002	1.002	2.040572e-14
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
LINICC	Gene expression	1.001	1.001	0.007045664	1.000	1.001	0.141664710
HNSC	Replication timing	1.000	1.001	7.383168e-05	1.000	1.001	0.001125938
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
LIHC	Gene expression	1.001	1.002	2.249945e-02	1.000	1.001	8.664341e-01
LITIC	Replication timing	1.001	1.001	5.639720e-09	1.001	1.001	8.128367e-08
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
SKCM	Gene expression	1.001	1.001	1.568880e-01	1.000	1.001	0.726452759
	Replication timing	1.000	1.001	0.0009835809	1.000	1.001	0.002735191

SARC	Replication timing	1.001	1.002	0.07560758
		OR	95% CI	p-value
BRCA	Replication timing	1.001	1.001	0.0001706559
		OR	95% CI	p-value
GBM	Replication timing	1.001	1.002	0.003476638
		OR	95% CI	p-value
OV	Replication timing	1.000	1.001	0.0948398
		OR	95% CI	p-value
LUSC	Replication timing	1.001	1.002	0.02497611
		OR	95% CI	p-value
STAD	Replication timing	1.000	1.001	0.6856750
		OR	95% CI	n-value

		OR	95% CI	p-value
SARC	Replication timing	1.001	1.002	0.07560758
		OR	95% CI	p-value
BRCA	Replication timing	1.001	1.001	0.0001706559
		OR	95% CI	p-value
GBM	Replication timing	1.001	1.002	0.003476638
		OR	95% CI	n valua
OV	Replication timing	1.000	1.001	<b>p-value</b> 0.0948398
		OR	95% CI	p-value
LUSC	Replication timing	1.001	1.002	0.02497611
		OR	050/ 61	
STAD	Replication timing	1.000	<b>95% CI</b> 1.001	<b>p-value</b> 0.6856750
		OP	050/ 61	
UCEC	Replication timing	<b>OR</b> 1.001	<b>95% CI</b> 1.002	<b>p-value</b> 0.005858224
		OR	95% CI	n valua
HNSC	Replication timing	1.000	1.001	<b>p-value</b> 0.2001579
		OR	95% CI	p-value
LIHC	Replication timing	1.001	1.001	0.01162314
		OR	95% CI	p-value
SKCM	Replication timing	0.99997	1.001	0.95351903

		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Replication timing	1.001	1.001	1.300404e-07	1.001	1.001	5.327523e-0
LIHC	Gene expression	1.002	1.001	0.001619439	1.001	1.003	9.492509e-0
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
HNSC	Replication timing	1.000	1.001	0.0000727371	1.000	1.001	0.005838784
	Gene expression	1.002	1.003	9.260708e-07	1.002	1.003	0.000065499
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Replication timing	1.002	1.002	3.361996e-16	1.002	1.002	1.094692e-1
UCEC	Gene expression	1.005	1.007	3.225537e-06	1.003	1.005	1.632000e-02
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
		.,,,,				.,,,,	
STAD	Replication timing	1.000	1.002	1.843887e-09	1.003	1.007	4.243039e-0
	Gene expression	1.006	1.008	1.392866e-08	1.005	1.007	3.333984e-0
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Replication timing	1.000	1.001	0.03669563	1.000	1.000	3.756591e-0
LUSC	Gene expression	1.004	1.006	2.790550e-06	1.004	1.005	1.814436e-0
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Replication timing	1.001	1.001	4.0393716-10	1.001	1.001	1.5110256-1
OV	Replication timing	1.003	1.004	4.039371e-16	1.002	1.003	1.511823e-1
	Gene expression	1.003	1.004	4.533474e-09	1.002	<b>adj. 95% CI</b> 1.003	<b>adj. p-value</b> 2.163129e-0-
		OR	95% CI	p-value	adj. OR	ad: 05% CI	adi n value
GDIVI	Replication timing	1.001	1.001	8.254689e-08	1.001	1.001	1.418192e-0
GBM	Gene expression	1.002	1.003	0.00234432	1.001	1.003	5.218249e-02
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Replication timing	1.001	1.001	3.731477e-15	1.001	1.001	2.488881e-13
BRCA	Gene expression	1.001	1.002	1.691989e-03	1.000	1.001	2.061333e-0
		OR	95% CI	p-value	adj. OR	adj. 95% CI	adj. p-value
	Replication tilling	1.001	1.001	3.677012603	1.001	1.001	0.00722311
SARC	Replication timing	1.002	1.004	3.877042e-03	1.001	1.003	0.00922311
	Gene expression	<b>OR</b> 1.002	<b>95% CI</b> 1.004	<b>p-value</b> 1.432804e-01	<b>adj. OR</b> 1.001	<b>adj. 95% CI</b> 1.003	<b>adj. p-valu</b> 0.44579173

		OR	95% CI	p-value
SARC	Replication timing	1.001	1.001	4.534571e-02
		OR	95% CI	p-value
BRCA	Replication timing	1.000	1.001	6.411577e-11
		OR	95% CI	p-value
GBM	Replication timing	1.000	1.001	7.825698e-05
		OR	95% CI	p-value
OV	Replication timing	1.000	1.000	0.0002432668
		OR	95% CI	p-value
LUSC	Replication timing	0.9999	1.000	0.6567748223
		OR	95% CI	p-value
STAD	Replication timing	1.001	1.002	0.0001362056
		OR	95% CI	p-value
UCEC	Replication timing	1.002	1.002	2.835927e-12
		OR	95% CI	p-value
HNSC	Replication timing	1.000	1.000	0.35956082
		OR	95% CI	p-value
LIHC	Replication timing	1.001	1.001	2.491517e-11
		OR	95% CI	p-value
SKCM	Replication timing	1.000	1.000	0.1759344

1.002

1.001

1.004

1.001

0.0006287536

8.868630e-06

1.002

1.001

1.003

1.001

0.0229081173

0.0002759546

Gene expression

Replication timing

SKCM

d

Supp Table 1. Gene expression and replication timing are determinants of chromosomal translocation (p<0.001). Linear regression model of normalized translocation number on gene expression and replication timing for (a) all protein coding genes (b) non-silent genes (c) silent genes (d) intergenic regions.