

Table S1 Median survival and P-values of *E. faecalis* OG1RF killing assays

Figure number	Experiment number	Strain, exposure conditions during development	Median survival (days)	P-value
1A	1	wild type, VC RNAi	8	C
		wild type, <i>skpo-1</i> RNAi	7	=0.0404
	2	wild type, VC RNAi	7	C
		wild type, <i>skpo-1</i> RNAi	6	=0.0331
	3	<i>eri-1</i> , VC RNAi	8	C
		<i>eri-1</i> , <i>skpo-1</i> RNAi	7	=0.001
	4	<i>eri-1</i> , VC RNAi	7	C
		<i>eri-1</i> , <i>skpo-1</i> RNAi	5	=0.0053
	5	<i>eri-1</i> , VC RNAi	9	C
		<i>eri-1</i> , <i>skpo-1</i> RNAi	8	=0.0037
1C	1	wild type, <i>cdc-25.1</i> RNAi	10	C
		<i>skpo-1</i> , <i>cdc-25.1</i> RNAi	6	<0.0001
	2	wild type, <i>cdc-25.1</i> RNAi	10	C
		<i>skpo-1</i> , <i>cdc-25.1</i> RNAi	8	<0.0001
	3	wild type, <i>cdc-25.1</i> RNAi	13	C
		<i>skpo-1</i> , <i>cdc-25.1</i> RNAi	8	=0.0016
	4	wild type, <i>cdc-25.1</i> RNAi	13	C
		<i>skpo-1</i> , <i>cdc-25.1</i> RNAi	8	=0.0005
	5	wild type, <i>cdc-25.1</i> RNAi	13	C
		<i>skpo-1</i> , <i>cdc-25.1</i> RNAi	7	<0.0001
4A	1	<i>wrt-2</i> , VC RNAi	5	C
		<i>wrt-2</i> , <i>skpo-1</i> RNAi	4	=0.0002
	2	<i>wrt-2</i> , VC RNAi	4	C
		<i>wrt-2</i> , <i>skpo-1</i> RNAi	3	=0.0176
	3	<i>wrt-2</i> , VC RNAi	5	C
		<i>wrt-2</i> , <i>skpo-1</i> RNAi	4	=0.0376
	4	<i>wrt-2</i> , VC RNAi	4	C
		<i>wrt-2</i> , <i>skpo-1</i> RNAi	4	=0.0058
4C	1	<i>vha-6</i> , VC RNAi	11	C
		<i>vha-6</i> , <i>skpo-1</i> RNAi	11	=0.6379
	2	<i>vha-6</i> , VC RNAi	11	C
		<i>vha-6</i> , <i>skpo-1</i> RNAi	11	=0.735
S1	1	wild type, OP50	8	C
		<i>skpo-1</i> *, OP50	3	<0.0001
	2	wild type, OP50	8	C
		<i>skpo-1</i> *, OP50	3	<0.0001

C = control, *Data only shown for GF89 *skpo-1* strain.