

FANCF sgRNA2																		PAM			GUIDE-seq read counts (frequency)								
	1								10							20		23	sgRNA		sgRNA + dRNA1		dRNA1						
	G	C	T	G	C	A	G	A	A	G	G	G	A	T	T	C	C	A	T	G	N	G	G						
OT1	A	6964	(35.5)	3495	(10.2)	0	(0.0)
ON	5502	(28.0)	13159	(38.5)	0	(0.0)
OT2	.	A	C	C	2744	(14.0)	6366	(18.6)	0	(0.0)
OT3	.	G	.	A	C	1616	(8.2)	4355	(12.7)	0	(0.0)
OT6	G	.	.	G	.	A	549	(2.8)	1481	(4.3)	0	(0.0)
OT5	A	A	.	.	T	543	(2.8)	1207	(3.5)	0	(0.0)
OT11	C	A	A	C	A	.	.	.	404	(2.1)	486	(1.4)	0	(0.0)
OT4	A	.	.	A	G	258	(1.3)	1532	(4.5)	0	(0.0)
OT8	.	.	C	A	T	A	.	367	(1.9)	106	(0.3)	0	(0.0)
OT7	T	G	.	.	A	T	255	(1.3)	633	(1.9)	0	(0.0)
OT9	T	.	.	.	C	A	122	(0.6)	499	(1.5)	0	(0.0)
OT10	T	.	.	.	A	T	112	(0.6)	285	(0.8)	0	(0.0)
OT20	A	A	G	110	(0.6)	46	(0.1)	0	(0.0)
	C	A	G	.	T	A	.	.	.	55	(0.3)	0	(0.0)	0	(0.0)
	A	A	.	.	.	T	16	(0.1)	179	(0.5)	0	(0.0)
OT17	A	T	.	.	G	A	3	(0.0)	0	(0.0)	0	(0.0)
OT16	C	A	.	.	G	C	0	(0.0)	165	(0.5)	0	(0.0)
	A	T	C	.	.	.	T	0	(0.0)	71	(0.2)	0	(0.0)
OT25	.	G	.	A	G	0	(0.0)	54	(0.2)	0	(0.0)
OT12	T	G	C	T	.	.	.	0	(0.0)	44	(0.1)	0	(0.0)
	C	A	T	.	.	.	A	.	.	.	0	(0.0)	30	(0.1)	0	(0.0)