## **Supplementary Table 1**. Domains for construction of three-finger ZFPs

Domain	Source	Amino acid sequences or Reference	Target subsite(s)
CSNR1	Human	YKCKQCGKAFGCPSNLRRHGRTH	GAC
HSNK	Human	YKCKECGKAFNHSSNFNKHHRIH	GAC
ISNR	Human	YRCKYCDRSFSISSNLQRHVRNIH	GAT
QSHR2	Human	YKCGQCGKFYSQVSHLTRHQKIH	GGA
QSHR3	Human	YACHLCGKAFTQCSHLRRHEKTH	GGA
QSHT	Human	YKCEECGKAFRQSSHLTTHKIIH	MGA
QSNR1	Human	FECKDCGKAFIQKSNLIRHQRTH	GAA
QSNR3	Human	YECEKCGKAFNQSSNLTRHKKSH	GAA
QSNV2	Human	YVCSKCGKAFTQSSNLTVHQKIH	MAA
QSSR1	Human	YKCPDCGKSFSQSSSLIRHQRTH	GYA
QTHQ	Human	YECHDCGKSFRQSTHLTQHRRIH	HGA
RDHT	Human	FQCKTCQRKFSRSDHLKTHTRTH	NGG
RDER1	Human	YVCDVEGCTWKFARSDELNRHKKRH	GCG
RSHR	Human	YKCMECGKAFNRRSHLTRHQRIH	GGG
RSNR	Human	YICRKCGRGFSRKSNLIRHQRTH	GAG
VSNV	Human	YECDHCGKAFSVSSNLNVHRRIH	AAT
VSSR	Human	YTCKQCGKAFSVSSSLRRHETTH	GTW
VSTR	Human	YECNYCGKTFSVSSTLIRHQRIH	GCW
DGAR	Phage display	18	GTC
DGHR	Phage display	18	GGC
DGNR	Phage display	18	GAC
DGNV	Phage display	19	AAC
DRDR	Phage display	18	GCC
GRER	Phage display	19	GCC
NDTR	Phage display	20	GTT
QAHR	Phage display	18	GGA
QGDR	Phage display	18	GCH
QGTR	Phage display	6	ACA
QSDR	Phage display	21	GCT
QSNR	Phage display	18	GAA
QSSR	Phage display	18	GTA
RDKR	Phage display	19	GGG

RDNR	Phage display	18	GAG
RDTN	Phage display	19	AAG
TDKR	Phage display	18	GGK
TGNR	Phage display	18	GAW