

<i>Gene symbol</i>	<i>siRNA construct</i>	Rel. rate of prion propagation		
		<i>FC</i>	<i>SD</i>	<i>t-test</i>
<i>Fn1</i>	<i>siRNA-Fn.1</i>	1.70	0.86	9.3×10^{-3}
<i>Chga</i>	<i>siRNA-Chga.2</i>	1.29	0.66	2.5×10^{-1}
<i>Galt</i>	<i>siRNA-Galt.2</i>	2.82	2.64	1.1×10^{-4}
	<i>siRNA-Galt.3</i>	1.80	0.88	1.7×10^{-3}
<i>IL11ra1</i>	<i>siRNA-IL11ra1.1</i>	2.73	1.51	1.1×10^{-6}
<i>Igsf5</i>	<i>siRNA-Igsf5.2</i>	1.41	0.67	1.2×10^{-1}
<i>Itga8</i>	<i>siRNA-Igga8.1</i>	1.05	0.53	8.5×10^{-1}
	<i>siRNA-Igga8.2</i>	2.18	1.04	7.7×10^{-5}
<i>Iqgap2</i>	<i>siRNA-Iqgap2.1</i>	1.41	0.93	1.4×10^{-1}

Supplementary Table S10: Susceptibility of CAD5 cells to RML prions after transient gene silencing of candidate genes. 1.5×10^4 CAD5 cells per well of a 96-well plate were transfected with siRNA followed by RML prion infection (10^{-5} dilution) as described in Methods. Fold changes (FC) in relative prion propagation rates $\pm$ SD were determined.