

Supplemental Table 3: Characterization of SFF colonies following targeted disruption of both *IFNAR1* and *IFNAR2*.

Cell line (Sex)	Targeted Genes	No. of Colonies Isolated	Biallelic Disruption of <i>IFNAR1</i> (%)	Biallelic Disruption of <i>IFNAR2</i> (%)	Biallelic Disruption of Both Genes (%)
SFF3 (F)	<i>IFNAR1</i> and <i>IFNAR2</i>	268	21 (7.8)	--	8 (3.0)
SFF5 (M)	<i>IFNAR1</i> and <i>IFNAR2</i>	139	--	31 (22.3)	18 (12.9)