

## shRNAmir Screening/Validation

**Gene Name:** rat DRD5      rat DRD1  
**RefSeq#**      NM\_012768      NM\_012546

shRNAmir Vector	KD %	KD %	shRNAmir sequence
Empty Vector control	0%	0%	
Scramble shRNA	25%	14%	GCT GAAATGTACTGCGCGTGGAGACGTTTTGGCCACTGACTGACGTCTCCACGCAGTACATTT
rDRD5-shmir#1 **	85%	24%	GCT GAAACCAGACGAATATGTCGAAGTTTTGGCCACTGACTGACTTCGACATTGCTGGTTT CAG
rDRD5-shmir#2	70%	32%	GCT GTACTGATGTTTACCGTCTGCAGTTTTGGCCACTGACTGACTGCAGACGAAACATCAGTA CAG
rDRD5-shmir#3	84%	41%	GCT GATCATGTGGACATAGGCAGTAGTTTTGGCCACTGACTGACTACTGCCTGTCCACATGAT CAG
rDRD5-shmir#4	36%	33%	GCT GATTAGGAGAGTCAGGAGGCCTGTTTTGGCCACTGACTGACAGGCCTCCACTCTCCTAAT CAG

\*\* will be used for viral production

Note:

shRNAmir155 sequence Scheme

GCT-target-GTTT TGGCCACTGAC TGAC -antisense- CAG

Promoter

CAG

Vector Backbone

AAV-CAG-GFP-shRNAmir155

**Figure S1.** Comparison of the efficacy of different rDRD5-shRNAmir constructs on the suppression of DRD5 and DRD1 mRNA expression. shRNAmir #1 showed the best DRD5 mRNA knockdown rate at 85% with the lowest impact on DRD1 gene expression at 24%. Validation of knockdown for each shRNA was performed in HEK 293 cells by co-transfection of the shRNA plasmid with the appropriate DRD cDNA plasmid and using the luciferase reporter assay.