## Differential effects of AGS3 expression on D<sub>2L</sub> dopamine receptor-mediated adenylyl cyclase signaling

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**Online Resource 1** Selective activation of recombinant AC1 by the calcium ionophore, A23187 and recombinant AC2 by the phorbol ester, PMA. **A.** Cell lines expressing recombinant AC1 or **B.** recombinant AC2 were incubated under the indicated conditions and cAMP accumulation was measured using a Cisbio homogenous time-resolved fluorescent (HTRF) cAMP assay kit according to the manufacturer's protocol. Data represent the mean  $\pm$  S.E.M. of three experiments. \*\* *p* < 0.01, compared to the basal condition within each cell line (one way ANOVA, followed by Dunnett's post hoc test).



**Online Resource 2** D<sub>2L</sub>DR expression levels in cellular models used for functional studies. [<sup>3</sup>H]-Methylspiperone saturation binding assays and analysis was performed (as described in Przybyla and Watts, 2010) to measure the  $D_{2L}DR$  expression level in each cell line. Data represent the mean  $\pm$  S.E.M. of four independent experiments

Cell Line	<b>B</b> <sub>max</sub> (pmol/mg protein)	$K_{\mathbf{d}}$ ( <i>nM</i> )
HEK-AC1-D <sub>2L</sub>	2.5±0.10	$0.10\pm0.02$
HEK-AC1-D <sub>2L</sub> -AGS3-Venus	4.0±0.27	0.11±0.02
HEK-AC2-D <sub>2L</sub>	9.1±2.5	0.11±0.01
HEK-AC2-D <sub>2L</sub> -AGS3-Venus	15±3.0	$0.17 \pm 0.04$
HEK-AC2-D <sub>2L</sub> -AGS3-Q/A-Venus	8.9±2.1	$0.10\pm0.01$

Online Resource 3 AGS3-Venus expression by confocal microscopy. Representative fields were imaged for cell lines expressing AGS3 fused to the Venus fluorescent protein at 20X magnification using a Nikon A1 confocal system when excited with the 488 nm laser and filtered for detection of Venus fluorescence (Emission: 525/50)

## AC1-D<sub>2L</sub>-AGS3-Venus

## AC2-D<sub>2L</sub>-AGS3-Venus

AC2-D<sub>2L</sub>-AGS3-Q/A-Venus



