



## A split-luciferase reporter recognizing GFP and

## 2 mCherry tags to facilitate studies of protein-protein

## 3 interactions

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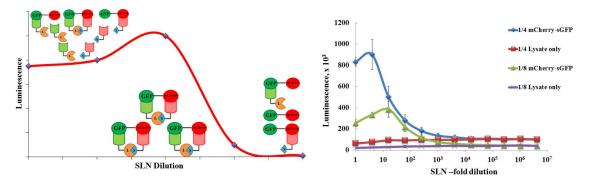


Figure S1 (Supplementary info): **A** – Typical 'hook' effect explained when the SLN is serially diluted instead of the analyte. At highest concentration of the SLN, the signal stays higher than background but doesn't reach maxima. Here, the SLNs are competing against each other for binding to their target where the two parts of split-Luc may end up on different protein targets. The signal reaches to maximum when the ratio between SLN and protein pair reaches 1:1 but drops down with further dilution due to lack of SLNs present for the reaction. **B** – Graphical representation of luminescence signal observed with serial dilution of SLN at a given concentration of analyte. Here, for each concentration analyte, we have corresponding concentration of non-expressing LTE that acts as background. It is evident that, at any given concentration of analyte, the signal is always higher than the background, confirming that we see limited split-Luc self-association and the complementation is driven by the presence of analyte.

Table S1: Protein sequences of the multiple protein used in construction of the universal split-Luc system.

Name of the Gene	Sequence		
Anti – GFP	MQVQLVESGGALVQPGGSLRLSCAASGFPVNRYSMRWYRQAPGKERE		
	WVAGMSSAGDRSSYEDSVKGRFTISRDDARNTVYLQMNSLKPEDTAVYY		
	CNVNVGFEYWGQGTQVTVSS		
Anti – mCherry	MAQVQLVESGGGLVQAGGSLRLSCATSGFTFSDYAMGWFRQAPGKERE		
	FVAAISWSGHVTDYADSVKGRFTISRDNVKNTVYLQMNSLKPEDTAVYS		
	CAAAKSGTWWYQRSENDFGSWGQGTQVTVSKEAI		
Split-Luc Large bit	VFTLEDFVGDWEQTAAYNLDQVLEQGGVSSLLQNLAVSVTPIQRIVRSG		
	ENALKIDIHVIIPYEGLSADQMAQIEEVFKVVYPVDDHHFKVILPYGTLVI		
	DGVTPNMLNYFGRPYEGIAVFDGKKITVTGTLWNGNKIIDERLITPDGSM		
	LFRVTINS		
Split-Luc Large bit	VTGYRLFEEIL		
Flexible Linker	GSSGGGGSGGGSSG		
Rigid Linker	EAAAKEAAAK		
Flexible-Rigid linker	GSSGGGGSGGGSSGEAAAKEAAAKGSSGGGGSGGGSSGEAAAKEAA		
	AKGSSGGGGGGSSG		

Table S2: Accession numbers for the multiple proteins used as targets to test the universal split-Luc system

Name of Gene	Accession number	
eGFP	DQ389577.1	
sfGFP	ASL68970	
mCherry	AKH87425	
Foldon	4NCV_A	
cMyc (b/HLH/Zip)	P01106	
HDAC3	O15379	
SOX9	P48436	
MyD88	Q99836	
Cav1	Q03135	
MaxZ (b/HLH/Zip)	P61244	

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