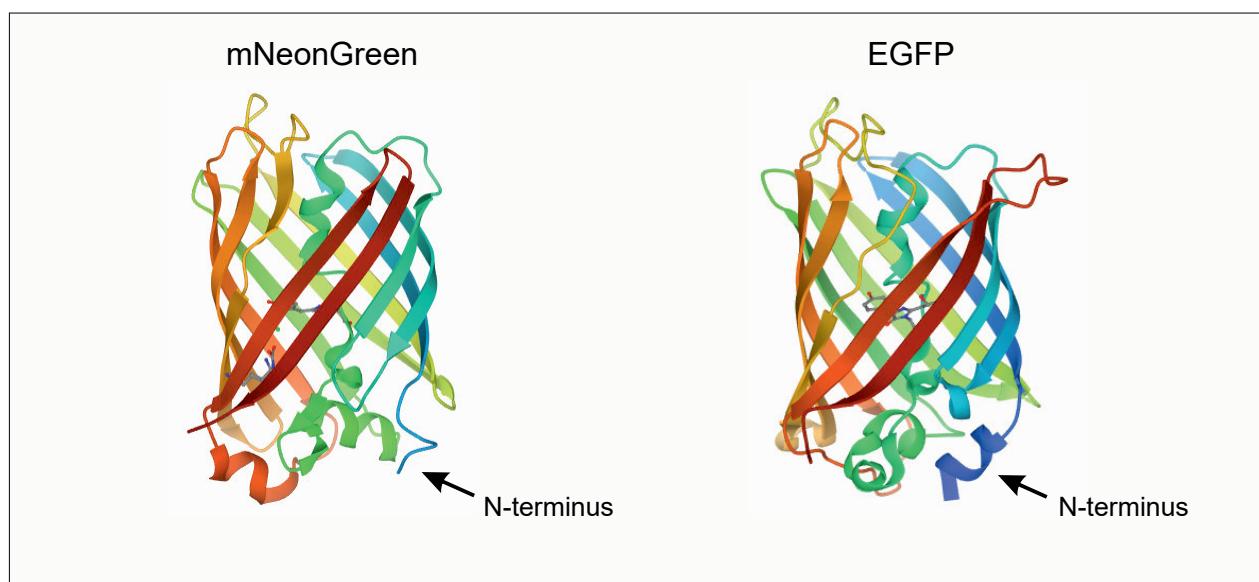


A

CLUSTAL O (1.2.4) multiple sequence alignment

mNeonGreen	MVSKGEEDNMASLPA T HELHIFGSINGVDFDMVGQGTGNPNDGYEELNLKSTKGD L QFSP 60
EGFP	MVSKGEELFTGVVPIL--VELD G DVN G HKE S VSG E E GDATY G KLT L KFICTTG K L P V P W 58
	***** . :* . : . * . :* . * : * : . * * : * : . * . :* .
mNeonGreen	WI L VPHIGYG F H Q YLP P PDGMSPF Q AA-MVD G S G Y Q VHRT M Q F ED G ASLT V N Y R Y TY E S 119
EGFP	PT L VT T LT T Y G V Q C F SR Y PD H M K Q H FFF K SAMPE G Y Q ERT I FF K D D G N Y K TRA E V K F E G 118
	** : ** . : * ** * . : . . ** . ** : * : * . . . :** .
mNeonGreen	HIKG E A Q V K GTGF P ADGPVMT N S LTAAD W CRSKKT Y PN---DKT I I S T F KWSYTTGN G K 175
EGFP	TL V N R IEL K GI D F KED G N I L G H K LE Y NY N SH--NV Y IMAD K Q K NG I KAN F KIRHN I E D G 176
	: . . :** . * ** : : . * .. : . * . : * . :** . . :*
mNeonGreen	R Y R S T A R T T Y TF A K P MA A N-Y L K N Q P MY V FR K TEL---KHS K TE L N F K E W Q K A F T D V M G 230
EGFP	V---QLADH Y QQNT P IGDGPV L LPDN H YLSTQS A LSKD P NEKRDH M V L E F V T A A G I T L G 233
	* . * : . * : * : : * : . : . : * : . * . :*
mNeonGreen	M DELY K
EGFP	M DELY K

B

C

Construct	Promoter	palmNeonGreen	palmGFP
	Vector	CMV	
		CSCGW2 lentivirus	pLNCX retrovirus
EVs characterisation	EVs purification method	Size exclusion chromatography	Sucrose gradient centrifugation
	EVs size	40 to 240 nm; mean 118.6 +\-.1.4 nm	Various (not provided in details); EVs detected in 0.22 µm and 0.8 µm - filtered sample
	EVs markers	Alix, Flotillin-2, CD9, CD63, DRα	Alix
EVs uptake	EVs donor cells	MelJuSo	HEK 293T; primary glioblastoma cells
	EV recipient cells	Huh7	HEK 293T; primary glioblastoma cells
Fluorescent reporter	Organism	<i>Aequorea Victoria</i>	<i>Branchiostoma lanceolatum</i>
	Excitation	506 nm	488 nm
	Emission	517 nm	509 nm
	Brightness	92.8	33.54
	Maturation	25 minutes	10 minutes
	Sequence identity	20-25%	

Supplementary Figure S2. Comparison of mNeonGreen and EGFP reporters. A. Amino acid sequence alignment of mNeonGreen (GenBank accession number AGG56535.1) and EGFP (Addgene expression vector 14757). B. X-ray structure of mNeonGreen (RCSB PDB accession number 5LTR) and EGFP (RCSB PDB accession number 2YGD). C. The properties of mNeonGreen and EGFP and the comparison of methodological details between this study using palmNG and the report by Lai *et al.* [20] using palmGFP.