

Figure S1: Validation of DPP8 or DPP9 knockdown in MCF-7 cells. (A) pTCEBAC with constitutive mCherry and inducible ECFP fluorescence and shDPP8. **(B)** pTREBAV with constitutive mVenus and inducible dsRed fluorescence and shDPP9. **(C-D)** FACS of MCF-7 cells \pm Dox with pTCEBAC + shDPP8 **(C)** or pTREBAV + shDPP9 **(D)**. **(E-F)** Relative *DPP8* **(E)** and *DPP9* **(F)** mRNA expression normalized to *ACTB*, in *shDPP8*- or *shDPP9*-transduced MCF-7 cells \pm Dox by qRT-PCR (n=3). **(G-H)** DPP8 **(G)** and DPP9 **(H)** with TUBA protein levels of *shDPP8*- or *shDPP9*-transduced MCF-7 cells \pm Dox by Western Blot. **(I)** Relative reduction of inhibitable DPP activity in *shDPP8*- or *shDPP9*-transduced MCF-7 cells (n=3). Bar charts show all data points with mean \pm SEM and *p*-value calculated by paired-sample t-test.

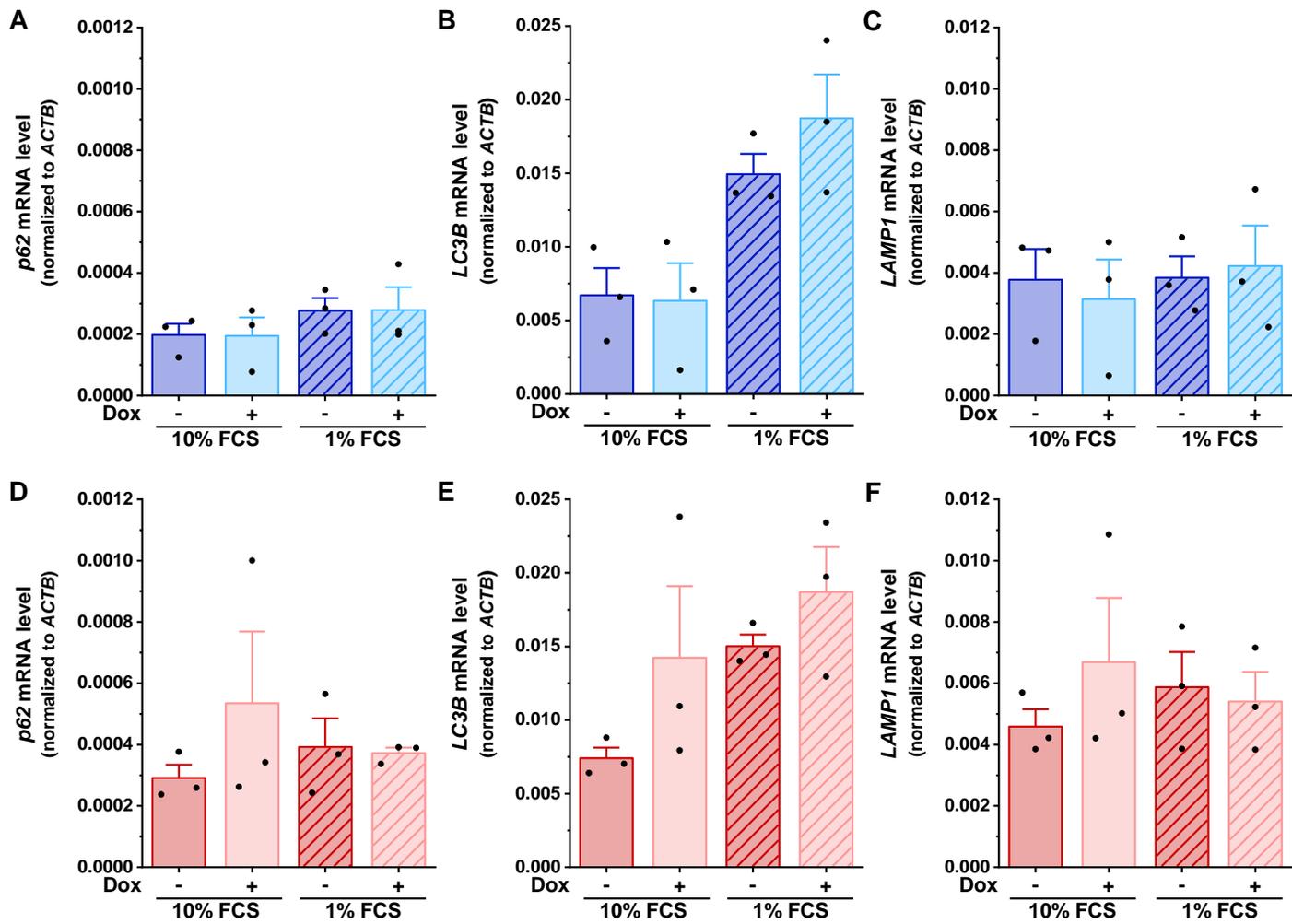


Figure S2: Expression of autophagosomal and lysosomal marker upon DPP8 or DPP9 knockdown in MCF-7 cells. (A-C) Relative *p62/SQSTM1* (A), *LC3B* (B), and *LAMP1* (C) mRNA expression normalized to *ACTB* in *shDPP8*-transduced MCF-7 cells \pm Dox and 10% or 1% FCS by qRT-PCR (n=3). (D-F) Relative *p62/SQSTM1* (D), *LC3B* (E), and *LAMP1* (F) mRNA expression normalized to *ACTB* in *shDPP9*-transduced MCF-7 cells \pm Dox and 10% or 1% FCS by qRT-PCR (n=3). Bar charts show all data points with mean + SEM.