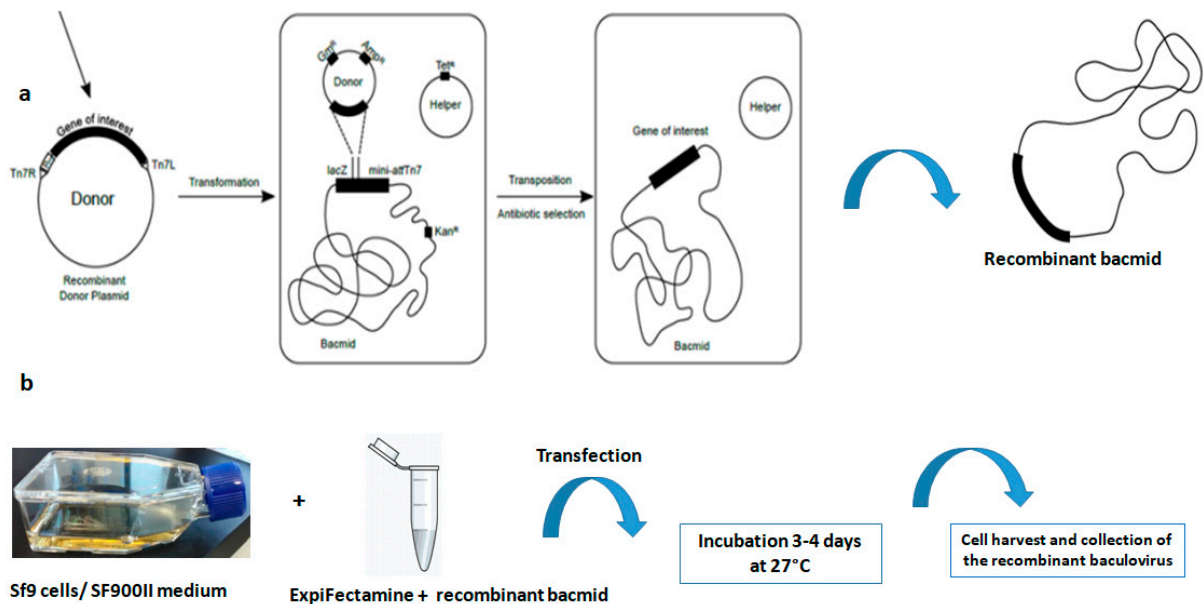
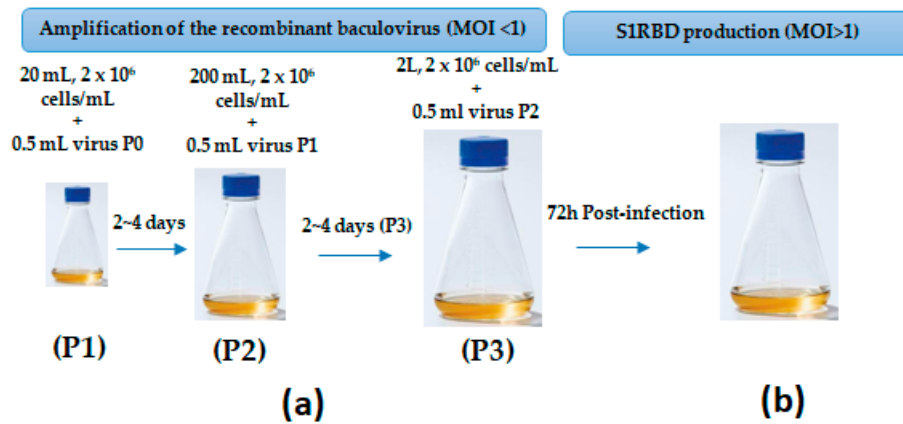


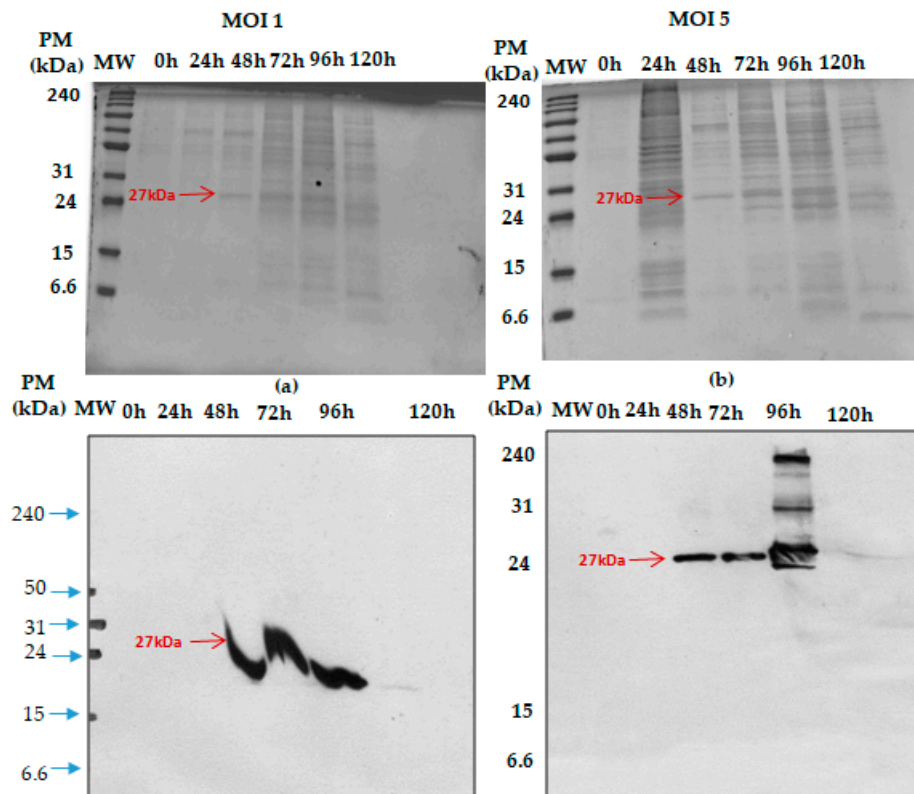
**Figure S1.** Map of the recombinant pFASTBac-S1RBD vector.



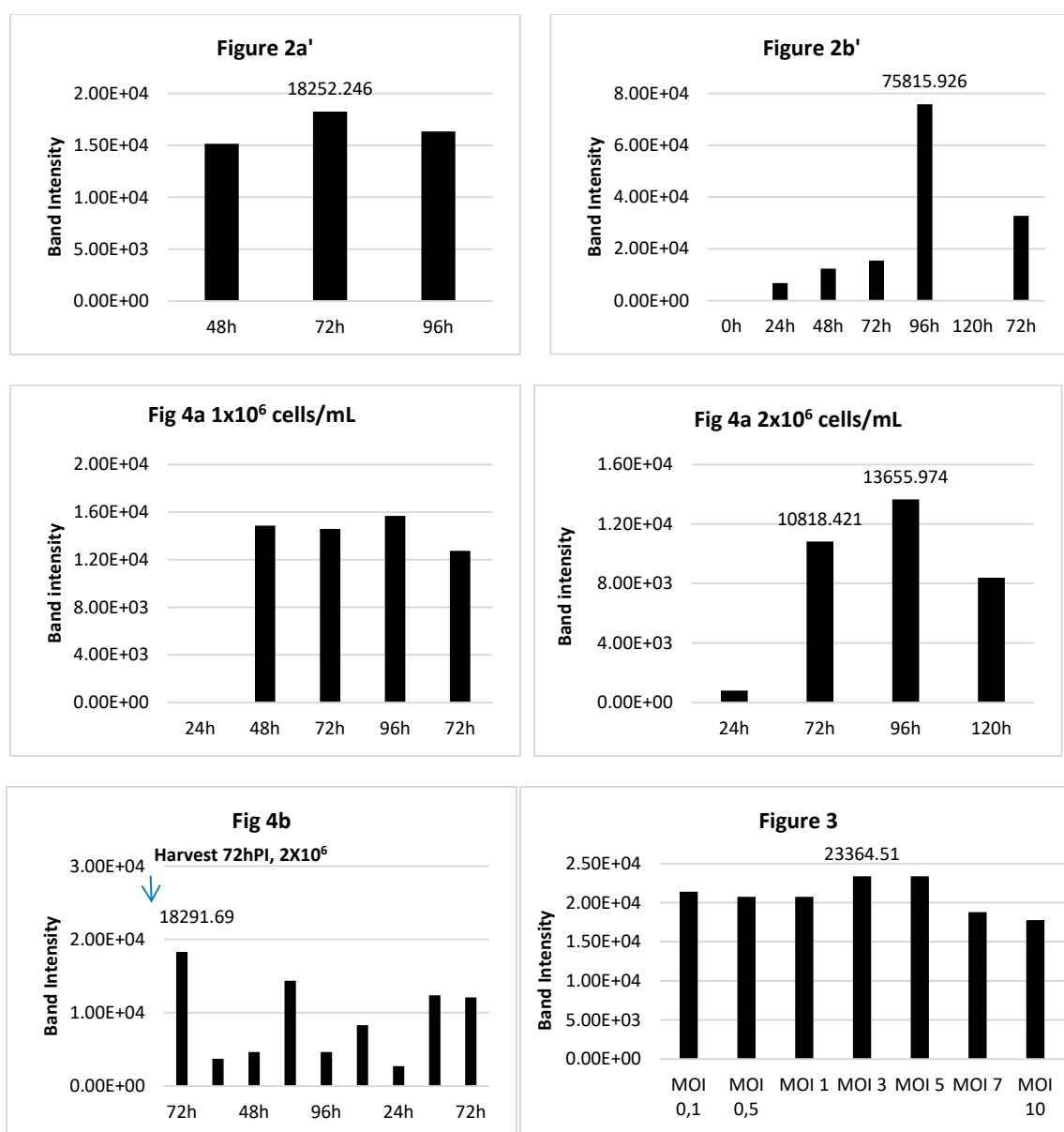
**Figure S2.** Schematic representation of the preparation of the recombinant bacmid (a) and transfection of Sf9 cells (b).



**Figure S3.** Schematic representation of the amplification process of recombinant baculovirus (a) and S1RBD protein (b).



**Figure S4.** S1RBD expression in Sf9 cells. Total proteins from the culture supernatant were analysed at different time points post infection (tpi) with MOI 1 (a) and MOI 5 (b). The assays were analysed by Coomassie-stained SDS-PAGE 12% and Western blot. The cell density at infection was  $2 \times 10^6$  cells/mL in baffled shake-flask. Equal amounts of total proteins (10  $\mu$ g) were loaded onto the SDS-PAGE gel.



**Figure S5.** Examples of band density evaluation using the software “image J”

**Table S1.** Supplementary data related to downstream process

Culture Conditions	Purification method	Volume <sup>a</sup> (mL)	Total quantity of S1RBD (mg/L of crude supernatant) <sup>b</sup>
Shake-flasks	Ni-NTA affinity column Followed by gel filtration	150	4.45±0.15
	Ni-NTA column + Superdex increase column	30	1.69±0.4
7L-bioreactor	NTA column Followed by gel filtration	400	70±8

<sup>a</sup>Volume after buffer exchange and concentration (after TFF)

<sup>b</sup>The results are expressed as mean ± standard deviation (n = 3)