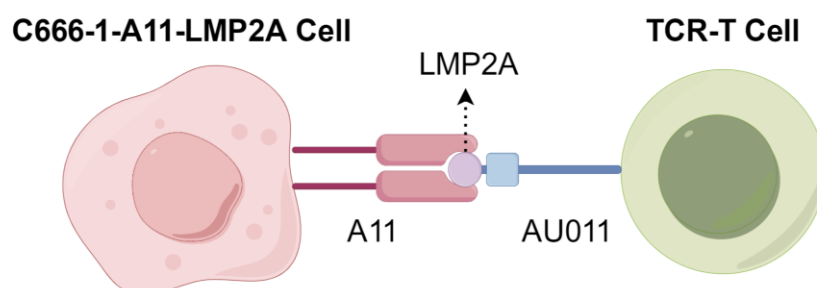
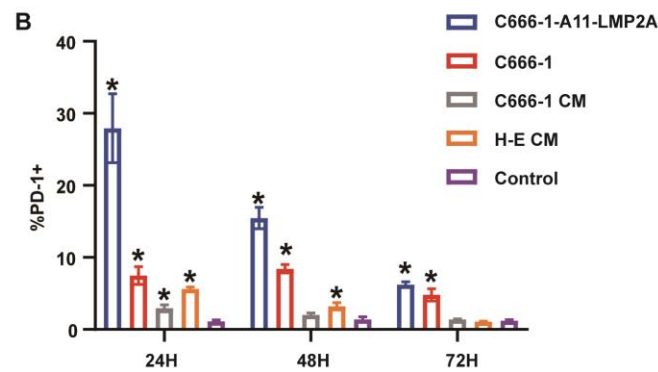
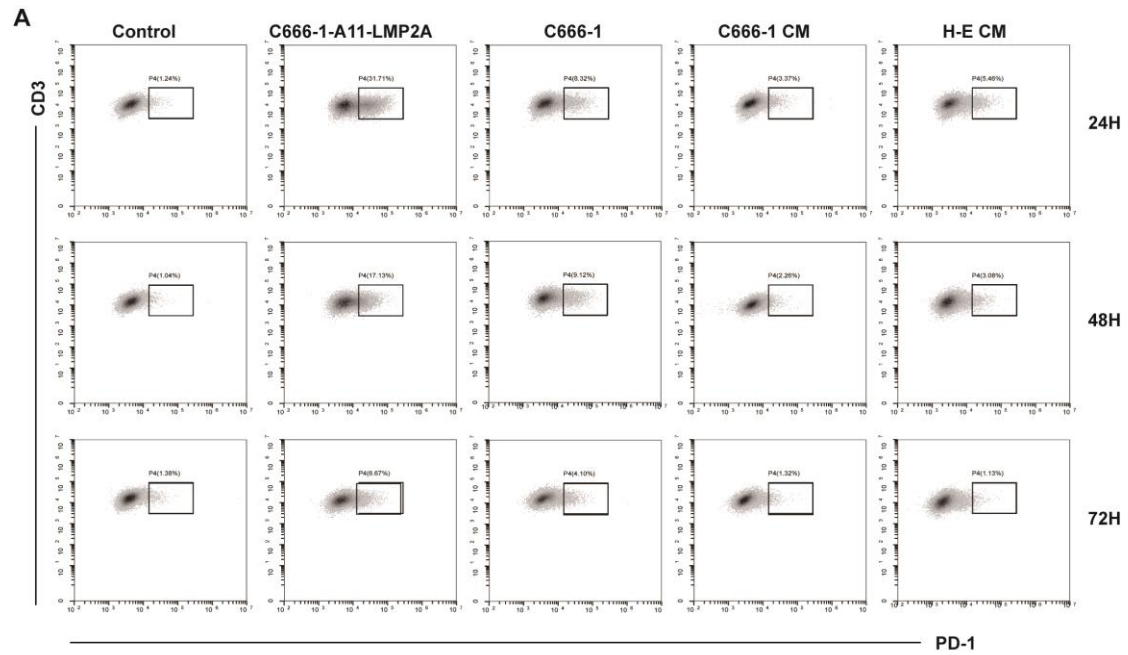


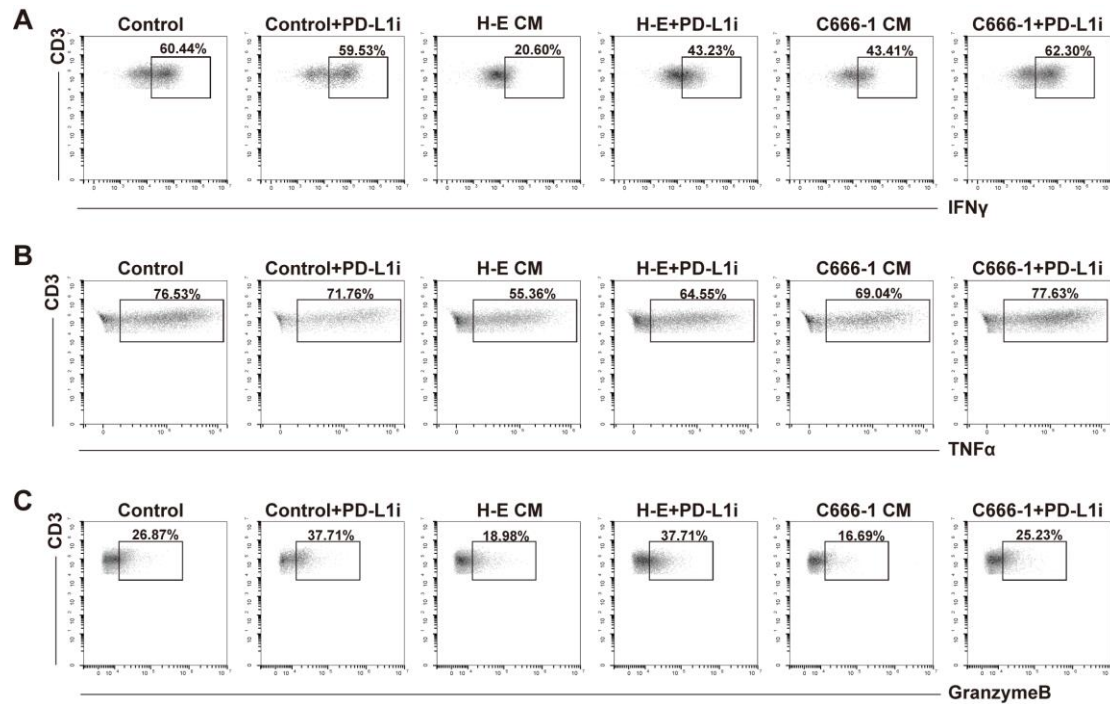
Supplementary Fig. 1. PD-L1 expression in HUVECs treated with supernatants from EBV- and EBV+ tumor cells for 48h using WB detection.



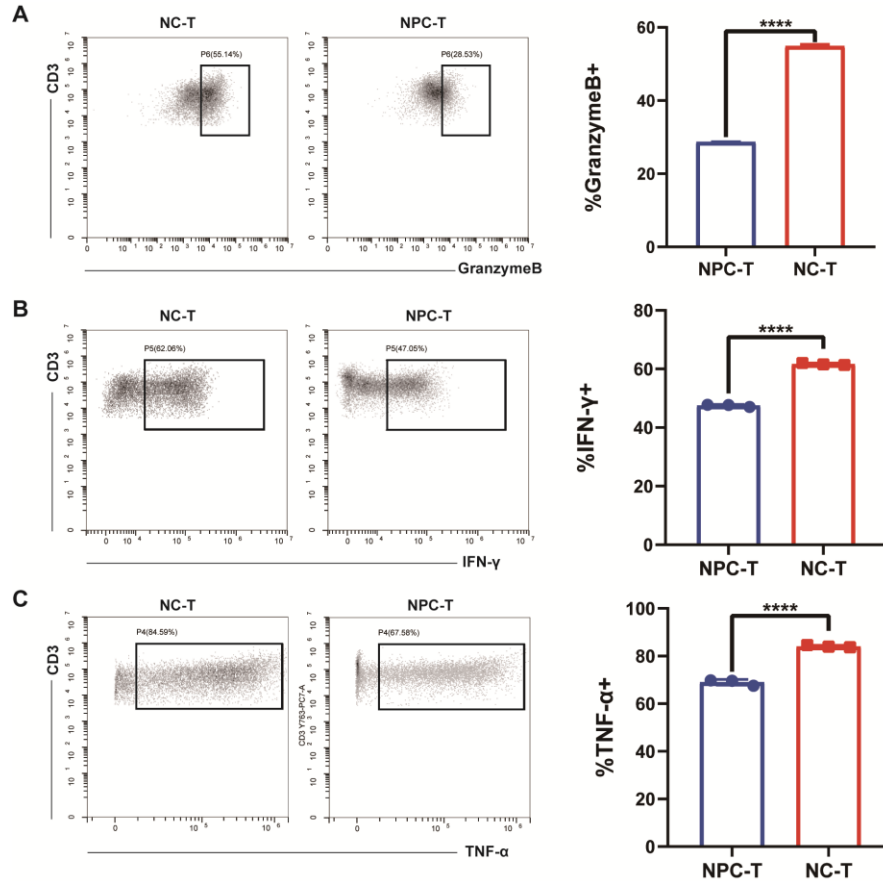
Supplementary Fig. 2. Schematic diagram depicting TCR-T cells recognising C666-1-A11-LMP2A tumour cells.



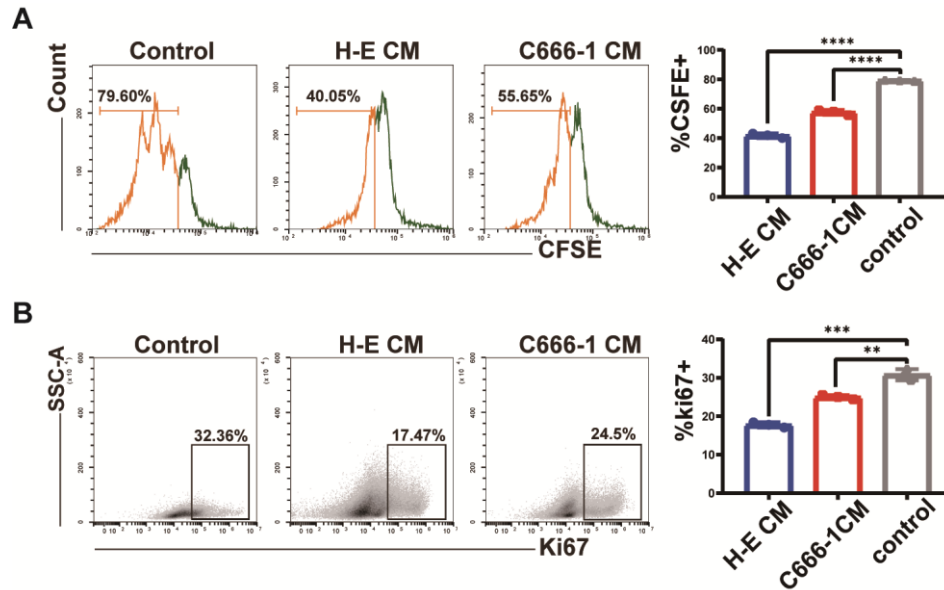
Supplementary Fig. 3. Expression of PD-1 on HUVECs after co-culture with C666-1-A11-LMP2A, C666-1, C666-1 CM, and H-E CM for 24h, 48h, and 72h, respectively. a) Representative images of Flow cytometry. b) Statistical plot of flow cytometry results. Data are mean \pm SD, $n=3$, two-tailed Student's t test. $*P < 0.05$.



Supplementary Fig. 4. Representative images of flow cytometry of cytokines in TCR-T cells after different treatments. **a)** Representative images of flow cytometry of IFN- γ in TCR-T cells after different treatments. **b)** Representative images of flow cytometry of TNF- α in TCR-T cells after different treatments. **c)** Representative images of flow cytometry of GranzymeB in TCR-T cells after different treatments. *Control* indicated TCR-T co-incubated with control HUVECs, *H-E* indicated TCR-T co-incubated with HK1-EBV CM-treated HUVECs, C666-1 indicated TCR-T co-incubated with C666-1 CM-treated HUVECs, and PD-L1i indicated incorporation of PD-L1 inhibitor.



Supplementary Fig. 5. Representative images of flow cytometry and statistical plots of cytokines in NC-T cells and NPC-T cells after co-incubation with CM-HUVECs. a) Representative images of flow cytometry and statistical plot of GranzymeB in NC-T cells and NPC-T cells after co-incubation with CM-HUVECs. **b)** Representative images of flow cytometry and statistical plot of IFN-γ in NC-T cells and NPC-T cells after co-incubation with CM-HUVECs. **c)** Representative images of flow cytometry and statistical plot of TNF-α in NC-T cells and NPC-T cells after co-incubation with CM-HUVECs. Data are mean ± SD, n=3, two-tailed Student's t test, **** $P < 0.0001$.



Supplementary Fig. 6. Effect of CM-treated versus untreated HUVECs on the proliferative capacity of TCR-T cells. *a*, *b*) CSFE staining *a* and Ki67 flow cytometry *b* results suggested that CM-treated HUVECs inhibited the proliferative capacity of TCR-T cells. *Control* indicated untreated HUVECs. *H-E CM* indicated H-E CM-treated HUVECs. *C666-1 CM* indicated C666-1 CM-treated HUVECs. Data are mean \pm SD, $n=3$, two-tailed Student's *t* test. $**P < 0.01$, $***P < 0.001$, $****P < 0.0001$.

Supplementary Table 1. The AUCs of classification models for predicting OS and PFS of patients with NPC.

OS		PFS	
Classification models	AUC	Classification models	AUC
TEC-PD-L1	0.7733 (95%CI [0.7055-0.8412])	TEC-PD-L1	0.6981 (95%CI [0.6305-0.7657])
TC-PD-L1	0.5927 (95%CI [0.5150-0.6705])	TC-PD-L1	0.5695 (95%CI [0.4974-0.6417])
Dual-scoring	0.7123 (95%CI [0.6445-0.7800])	Dual-scoring	0.6448 (95%CI [0.5739-0.7156])