Figure S1 Using unsupervised clustering method, (A) four clusters captured the best segmentation, (B) all samples were assigned into these four clusters.
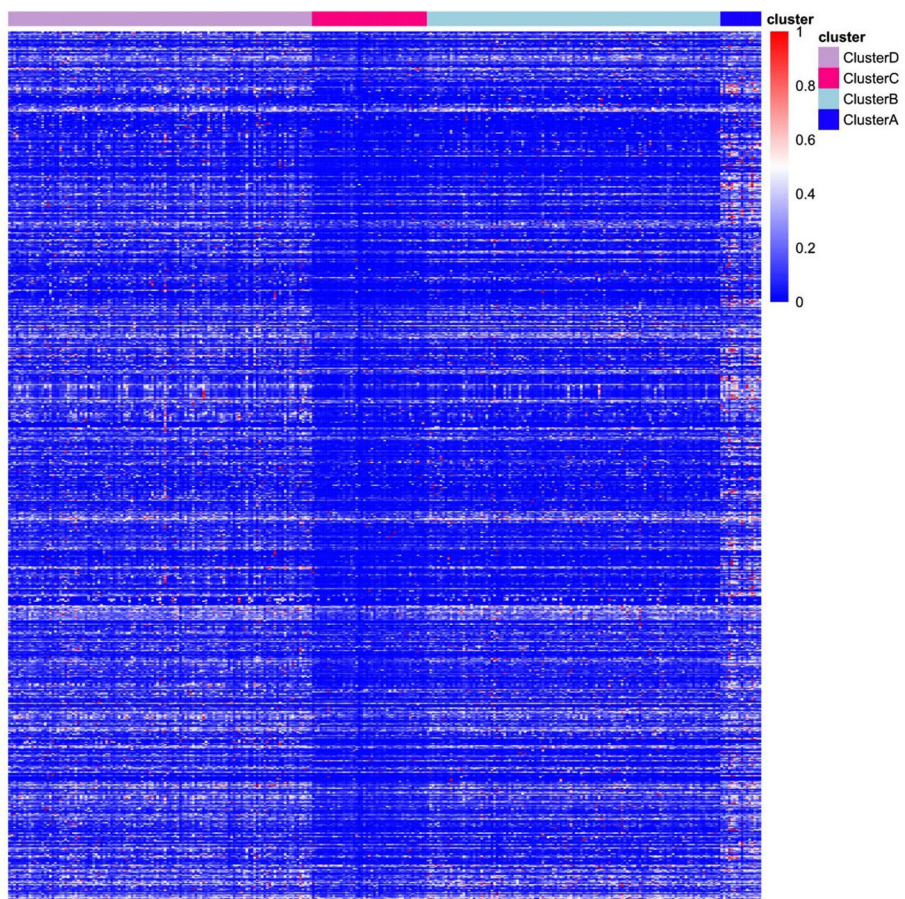

Figure S2 Expression level of immune genes divided into four clusters.

https://dx.doi.org/10.21037/tau-21-887