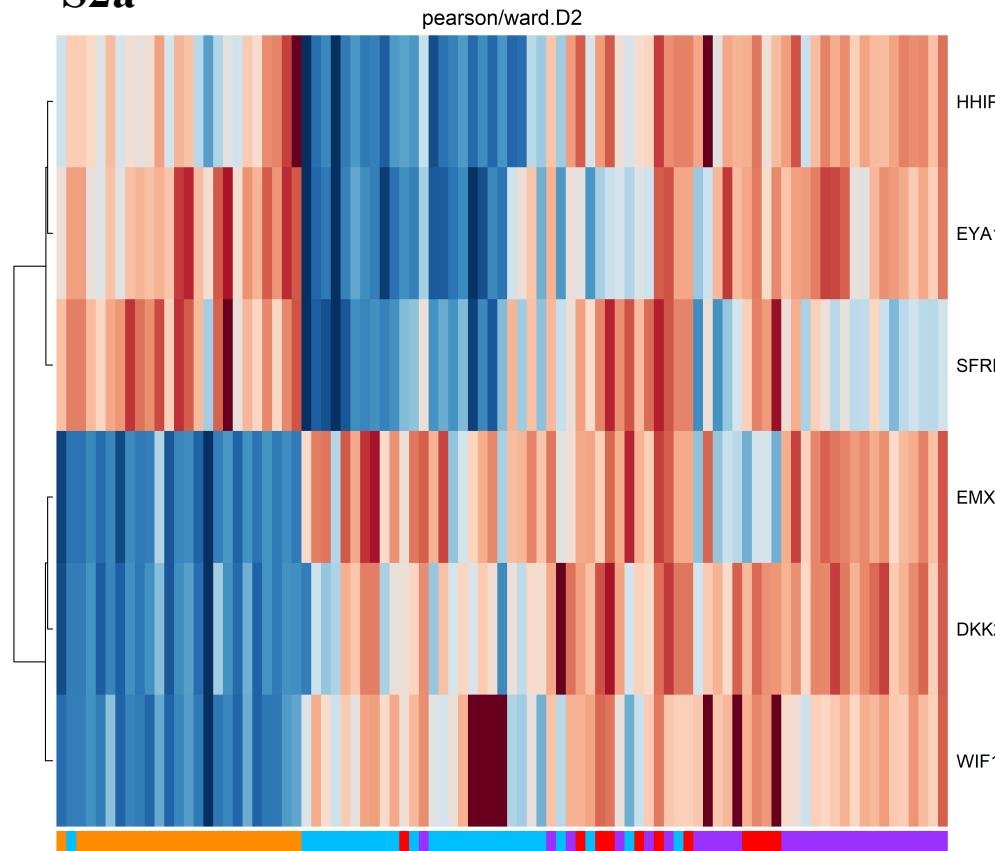
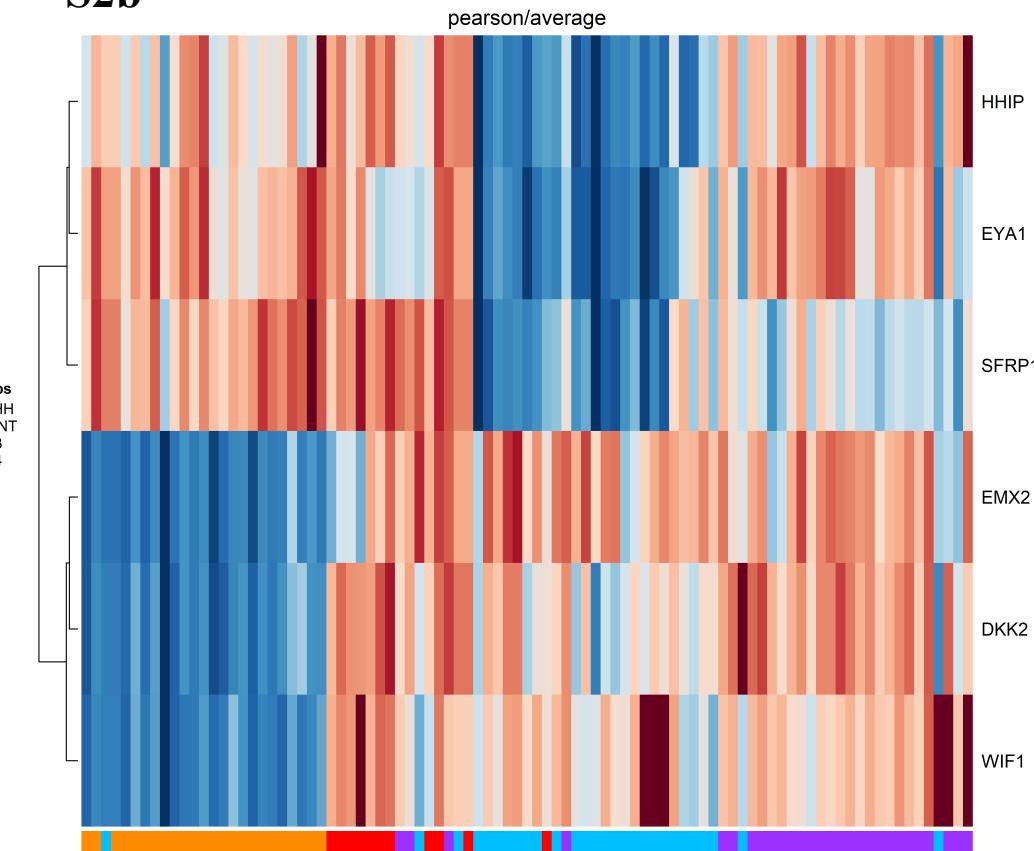


**S2a****S2b**

**Fig. S2.** Comparison of clustering algorithms in our study ( $n=92$ ) with 6 genes *HHIP*, *EYA1*, *SFRP1*, *EMX2*, *DKK2*, *WIFI1*. **(a)** Ward.D2 algorithms **(b)** Average-linkage algorithms.