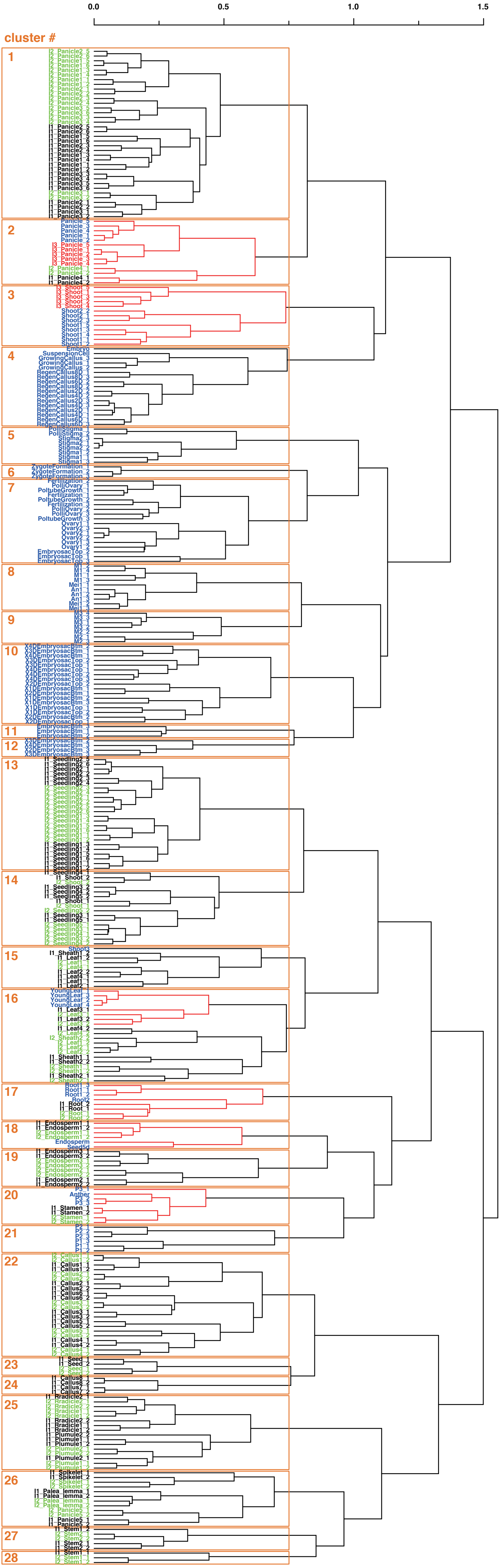Figure S2. Dendrogram of 321 arrays from *japonica* Nipponbare (blue), three *indica* varieties 9311 (red), Zhenshan97 (black), and Minghui63 (green). By using 12,097 similarly expressed genes between Nipponbare and 93-11 in both shoot and panicle, 321 arrays were classified into similar tissues. 28 clusters were produced by cutting the dendrogram at height 0.75. Six clusters with red colored branches contain both *japonica* and *indica* array data.