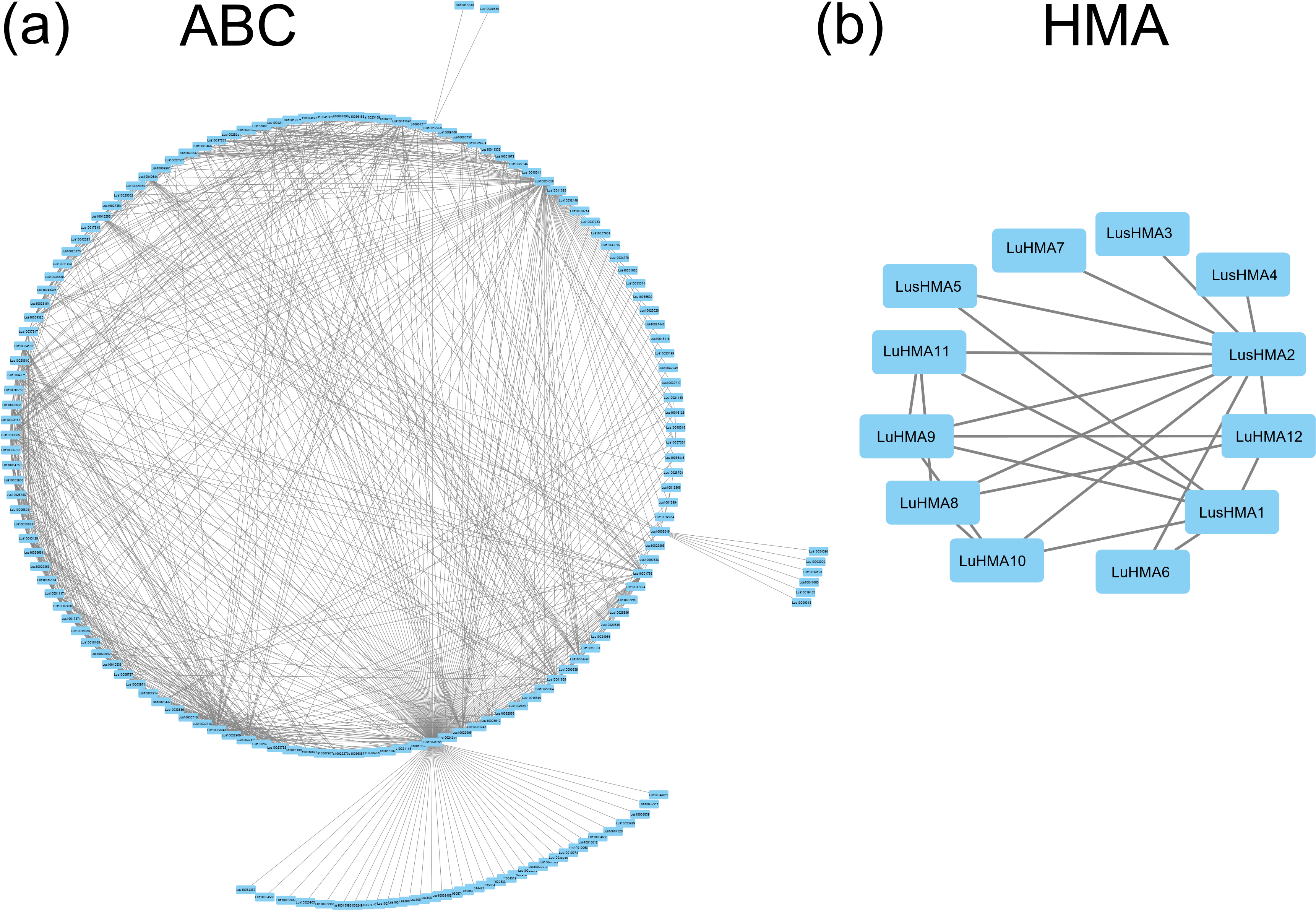


**Figure S1.** Multidimensional scaling (MDS) plot of nine different organs displaying the relative similarities between the biological replicates based on the log fold change values.



**Figure S2.** Interaction network of ABC transporter genes (a) and HMA genes (b).



**Figure S3.** Schematic representations of the two most stable genes among nine Cd candidate genes based on their conserved gene structure (a), gene expression (b), non-synonymous/synonymous substitution rates (c), and Pearson correlation coefficient (PCC) (d).