Figure S4. Relationship between tandem duplication within each transcription factor (TF) family and percentage of the TF family in triads. a) and b) show TF families with >30 genes in triads. c), d) and e) show all TF families. a) and d) Define tandem duplicates as up to ±2 genes away from each other (i.e. with 0 or 1 gene between duplicated TFs). b) and e) Define tandem duplicates as ±1 genes away from each other (i.e. with 0 genes between duplicated TFs). c) Defines tandem duplicates as up to ±3 genes away from each other (i.e. with 0, 1 or 2 genes between duplicated TFs).