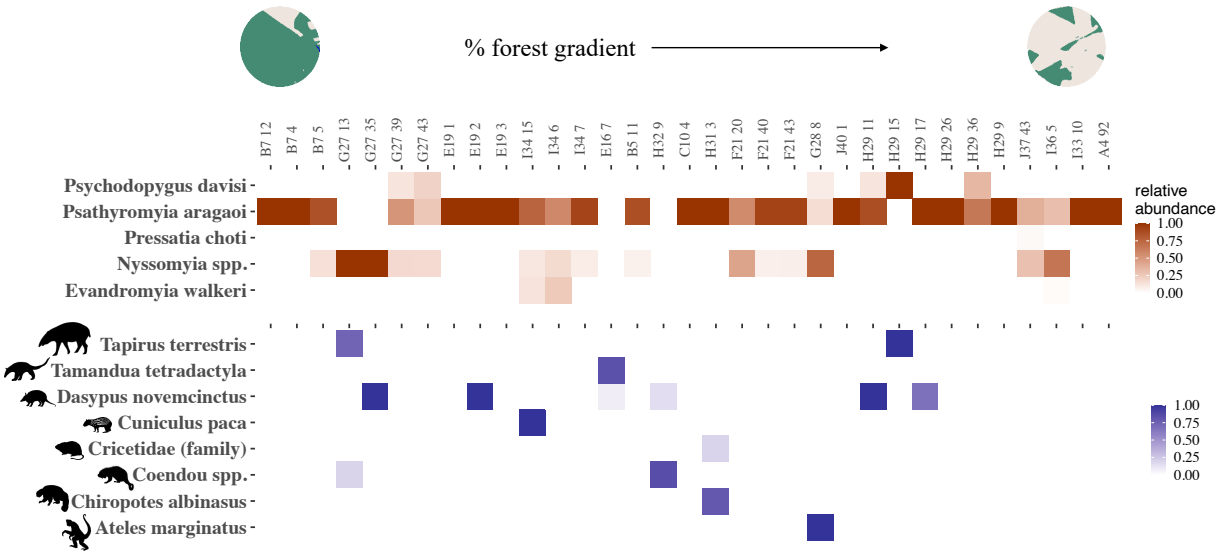Figure 6: Of 1130 samples, 42 were positive for the presence of at least one *Leishmania* species, 33 of which containing either sandfly and/or vertebrate data from DNA metabarcoding. The samples that tested positive for *Leishmania* are organized on the horizontal axis by amount of forest cover at their corresponding site. The sandfly and vertebrate taxa that were found in these samples are shown with either orange (sandfly) or purple (vertebrate) color showing the relative abundance of sequence reads for each taxon. Domesticated species found with DNA metabarcoding have been excluded.