

Figure S1: CLUSTALX alignment of 450bp of cytochrome oxidase I DNA barcodes obtained from infected honey bees (samples 19-24,26-31,34,35) and bumble bees (samples 33,36). Bidirectional Sanger sequence indicates that only two positions varied (88, 288) in a single sample each. All samples had less than 0.22% divergence (i.e. 1bp).

