## Genus-level accuracy varKoder genus 1.0 0.9 result correct ambiguous inconclusive incorrect 0.1 0.0 Skmer genus 1.0 0.9 0.8 Fraction of samples 0.7 result 0.6 0.5 correct 0.4 incorrect 0.3 0.2 0.1 0.0 Base pairs in query images ITS genus rbcL genus matK genus 1.0 0.9 Fraction of samples result correct inconclusive incorrect 0.1 Base pairs in Base/pairs in Base/pairs in query images