number of orthogroups that are exactly the same between method and baseline TarAar CruTar CruAar ChiTar proportionOfExactOG 0.00 species pair ChiCru 0.25 0.50 ChiAar 0.75 1.00 AthTar AthCru AthChi AthAar BR OFb OFd OĖm SPd SPm ÓИ method

number of orthogroups that are exactly the same between method and baseline 0.281 0.271 0.269 0.269 0.334 0.334 0.232 TarAar 0.316 0.352 CruTar · 0.316 0.316 0.316 0.352 0.269 0.31 0.322 0.321 0.321 0.35 0.351 0.286 CruAar 0.289 0.287 0.283 0.345 0.346 0.233 ChiTar 0.285 proportionOfExactOG 0.00 species pair ChiCru 0.421 0.415 0.413 0.414 0.501 0.502 0.337 0.25 0.50 0.286 0.293 0.291 0.291 0.35 0.351 0.248 ChiAar 0.75 1.00 0.322 0.324 0.323 0.323 0.352 0.352 0.276 AthTar 0.5 0.5 AthCru -0.497 0.499 0.543 0.543 0.417 AthChi 0.439 0.45 0.45 0.451 0.487 0.487 0.373 0.318 0.333 0.333 0.333 0.354 0.355 0.296 AthAar BR OFb OFd OĖm SPd SPm ÓИ method

mean value, diploid – JI – between method and baseline 0.377 0.356 0.355 0.355 0.43 0.43 0.332 TarAar -CruTar -0.407 0.402 0.401 0.401 0.438 0.439 0.374 0.397 0.4 0.4 0.399 0.432 0.432 0.38 CruAar 0.38 0.368 0.333 ChiTar -0.365 0.366 0.432 0.433 meanJI 0.00 species pair ChiCru -0.508 0.493 0.491 0.491 0.581 0.582 0.442 0.25 0.50 ChiAar -0.34 0.373 0.368 0.367 0.367 0.432 0.433 0.75 1.00 0.385 AthTar -0.416 0.413 0.413 0.412 0.44 0.44 AthCru -0.583 0.579 0.58 0.579 0.618 0.618 0.529 AthChi -0.534 0.536 0.536 0.536 0.57 0.57 0.49 0.392 AthAar -0.406 0.413 0.413 0.412 0.436 0.436

BR OFd OFb OFm SPd SPm ÓИ

method