Figure S2

Reconstructions of the evolution of host plant usage in Tenthredinoidea, based on maximum parsimony optimisation of host plant usage on the Bayesian phylogeny.

Blasticotoma filiceti pacifica
Runaria reducta
Acordulecera sp.
Decameria similis
Perga condei
Phylacteophaga froggatti
Lophyrotoma analis
Sterictiphora furcata
Arge nigripes
Arge cyanocrocea
Gilpinia sp.
Diprion pini
Monoctenus juniperi
Corynis crassicornis
Abia sericea
Cimbex americana
Trichiosoma sp.
Athalia sp.
Athalia japonica
Hoplocampa fulvicornis
Nematinus luteus
Nematinus sp.1
Nematinus sp.2
Nematus sp.
Selandria serva
Thrinax contigua
Thrinax japonica
Strongylogaster osmundae
Strongylogaster secunda
Aneugmenus japonicus
Aneugmenus kitonis
Nipponorhynchus bimaculatus
Dolerus japonicus
Loderus genucinctus
Dolerus ephippiatus
Dolerus sp.
Stromboceros koebelei
Rocalia longipennis
Taxonus agrorum
Empria sp.
Allantus sp.
Metallus rohweri
Paracharactus japonicus
Siobla sp.
Macrophya sp.2
Macrophya apicalis
Macrophya sp.3
Macrophya sp.1
Rhogogaster sp.1
Pachyprotasis sp.
Lagidina irritans
Rhogogaster sp.2
Tenthredo sp.
Tenthredo mesomela
Aglaostigma occipitosum
Aglaostigma lichtwardti

Angiosperms
Gymnosperms
Pteridophytes