

Figure S2.1. Response plot• for combined model

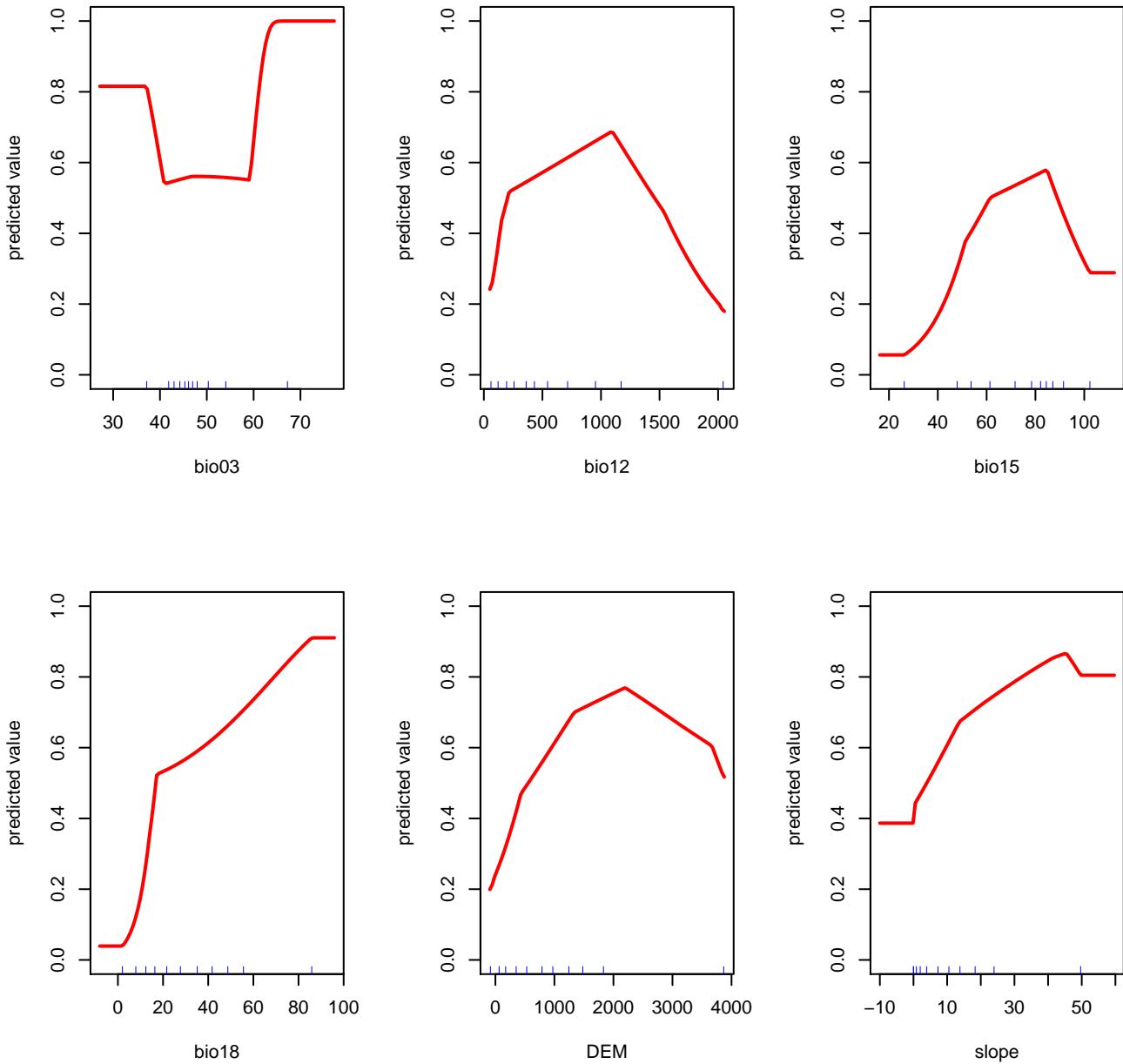


Figure S2.2. Response plots for hibernacula model

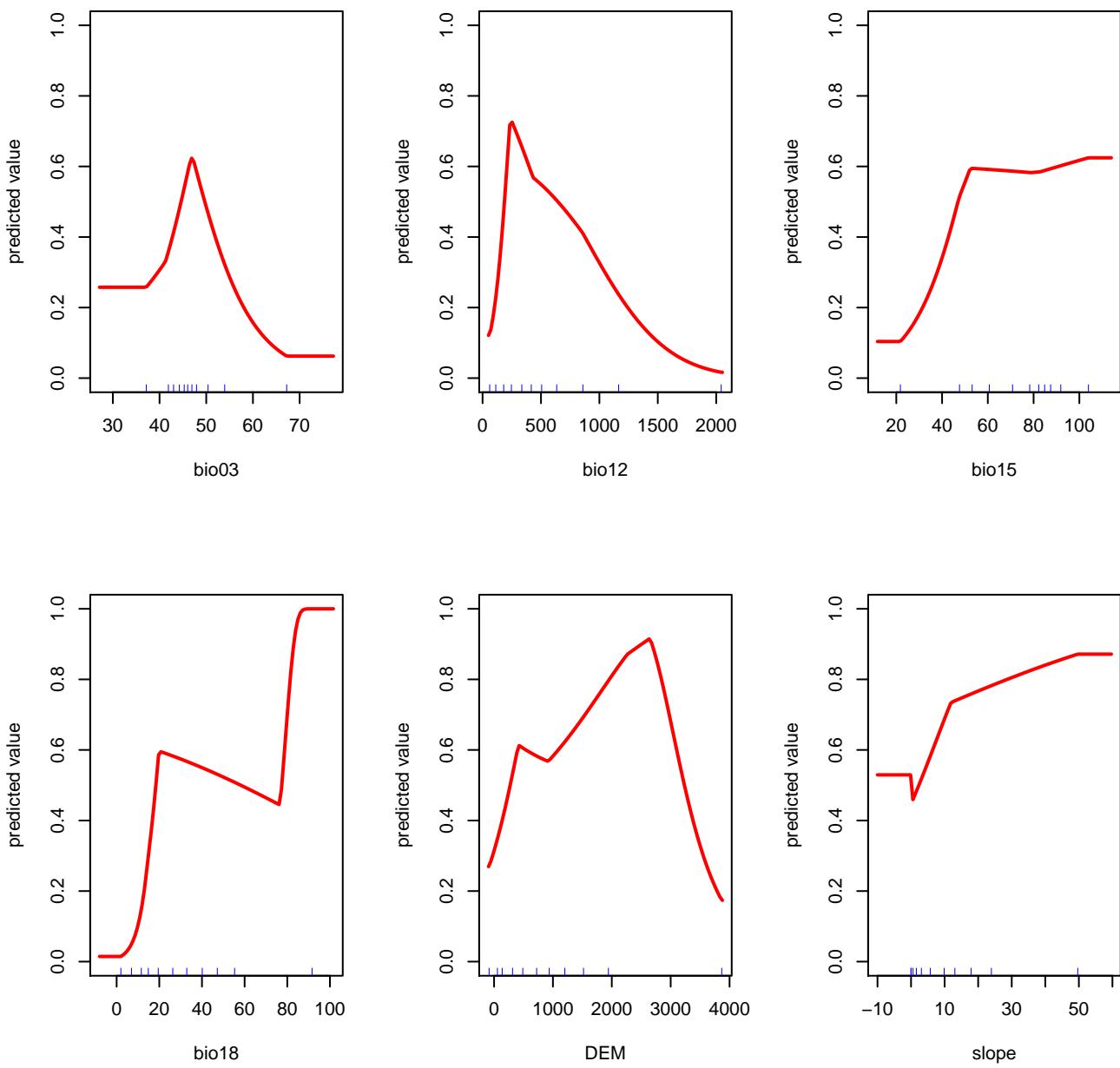
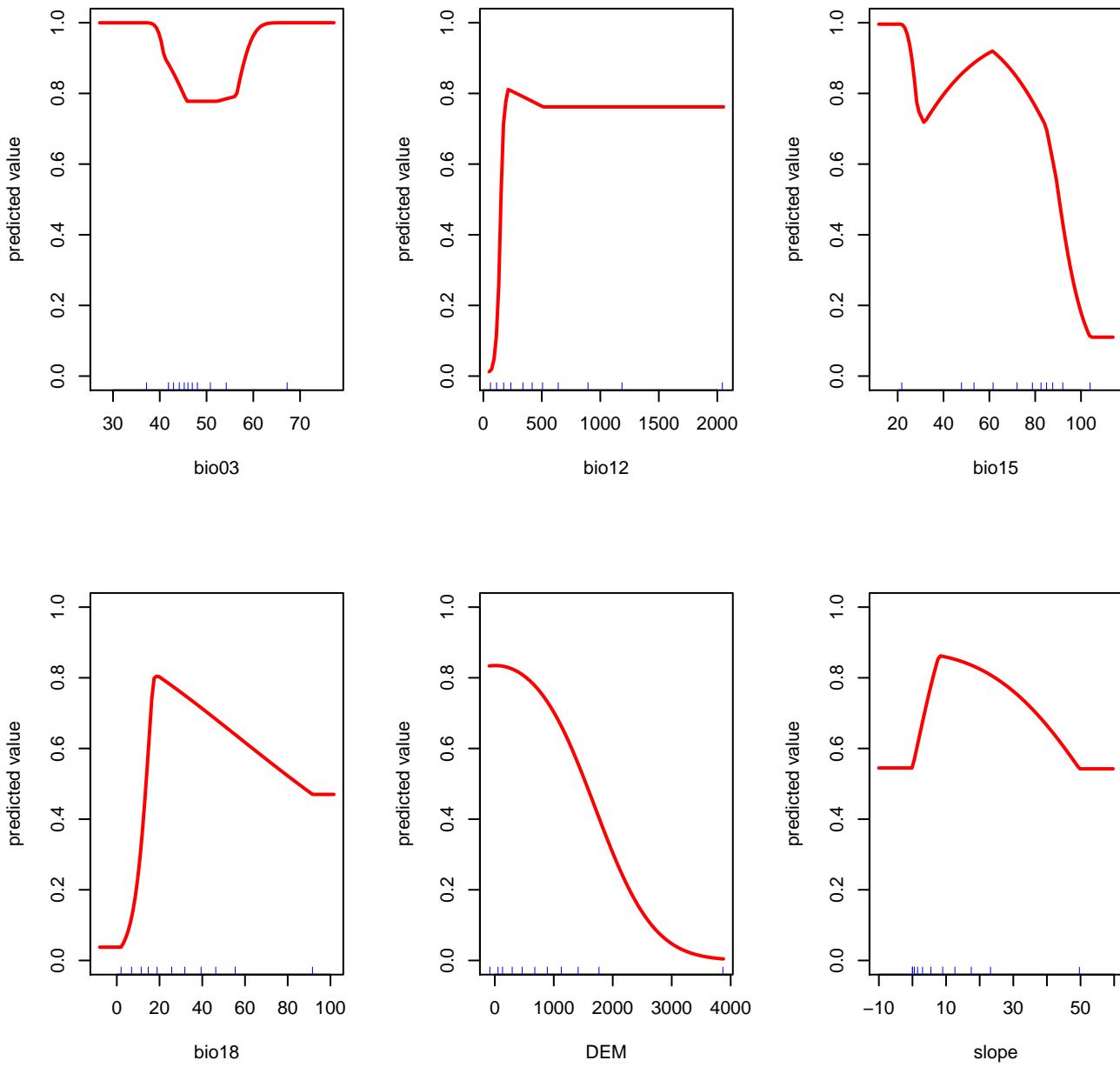
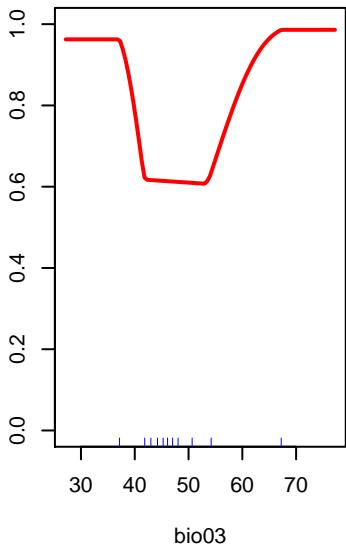


Table S2.3. Response plots for maternity models



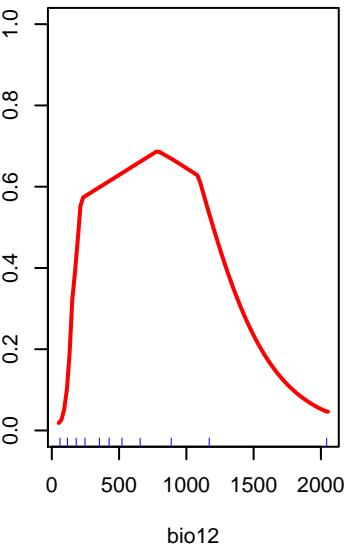
Væl|<sup>^</sup>AUGÈ EU<sup>^</sup>•] [ }•<sup>^</sup>Á|[ C Á| Á a<sup>^</sup>•á{ Á|[ á<sup>^</sup>|

predicted value



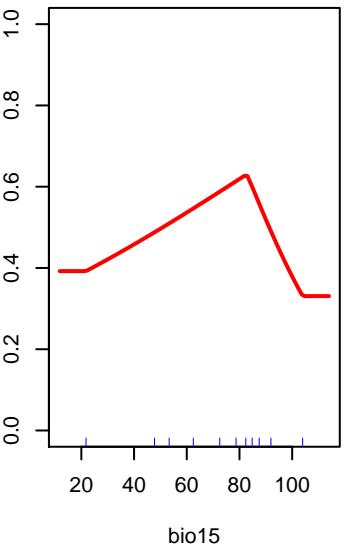
bio03

predicted value



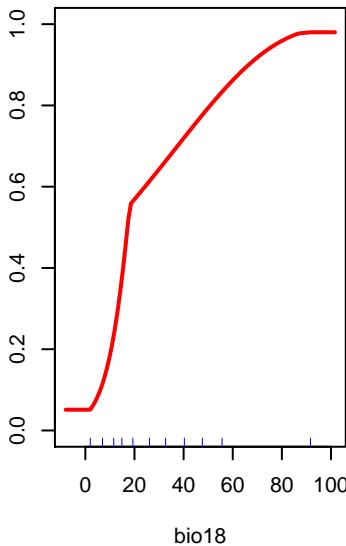
bio12

predicted value



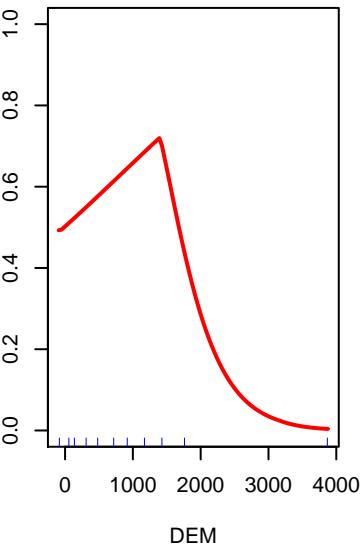
bio15

predicted value



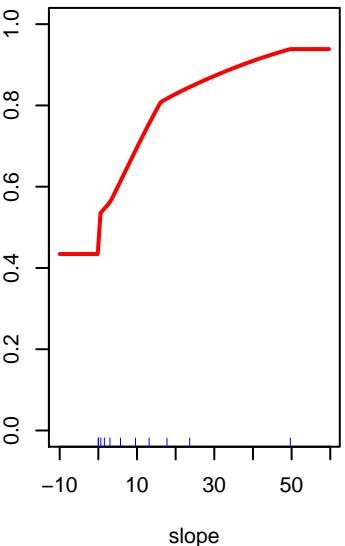
bio18

predicted value



DEM

predicted value



slope

Figure S2.5. Response plot for ecoregion 1

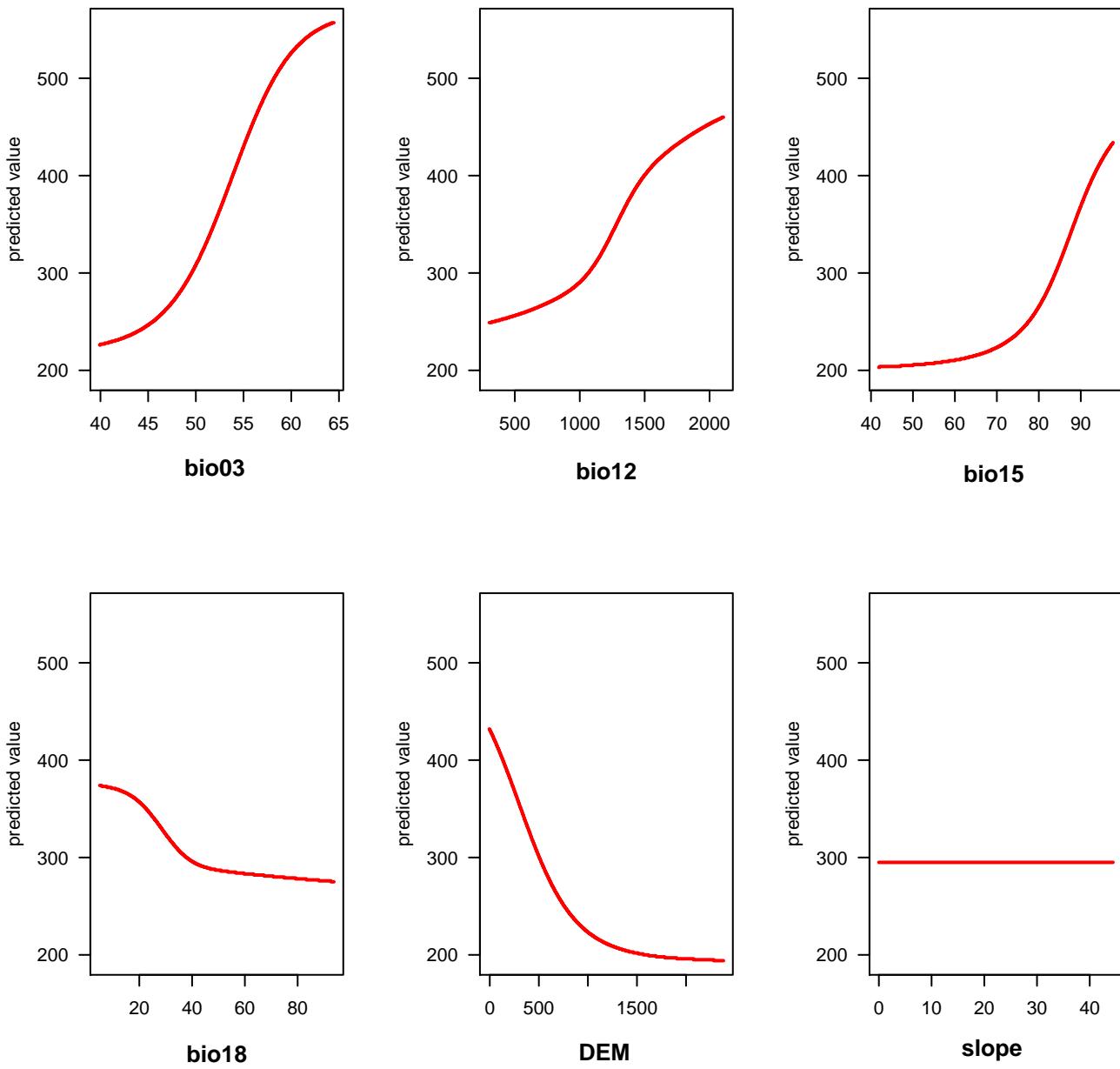


Figure S2.6. Response plots for ecoregion 4

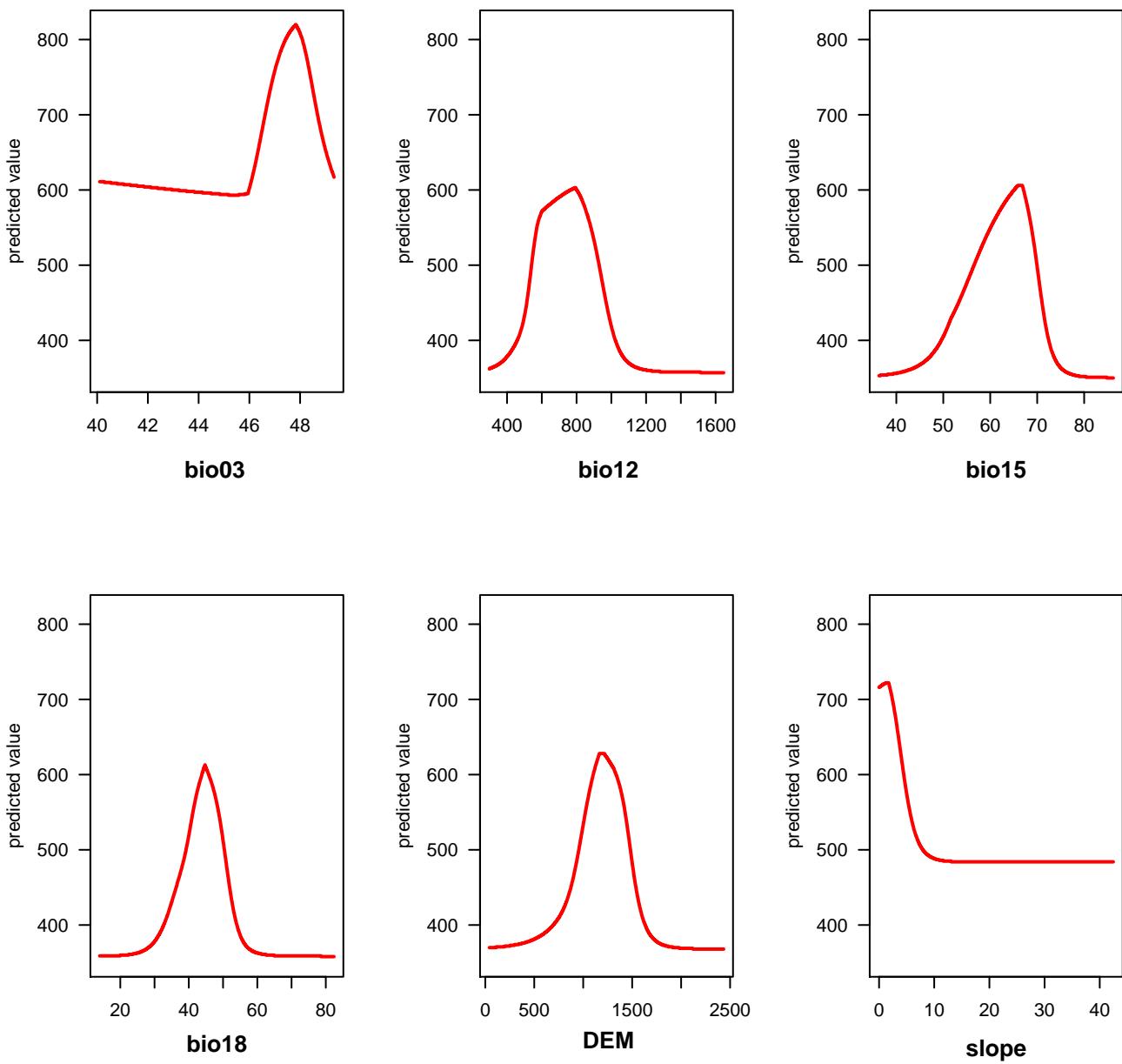


Table S2.7 Response plots for ecoregion 5

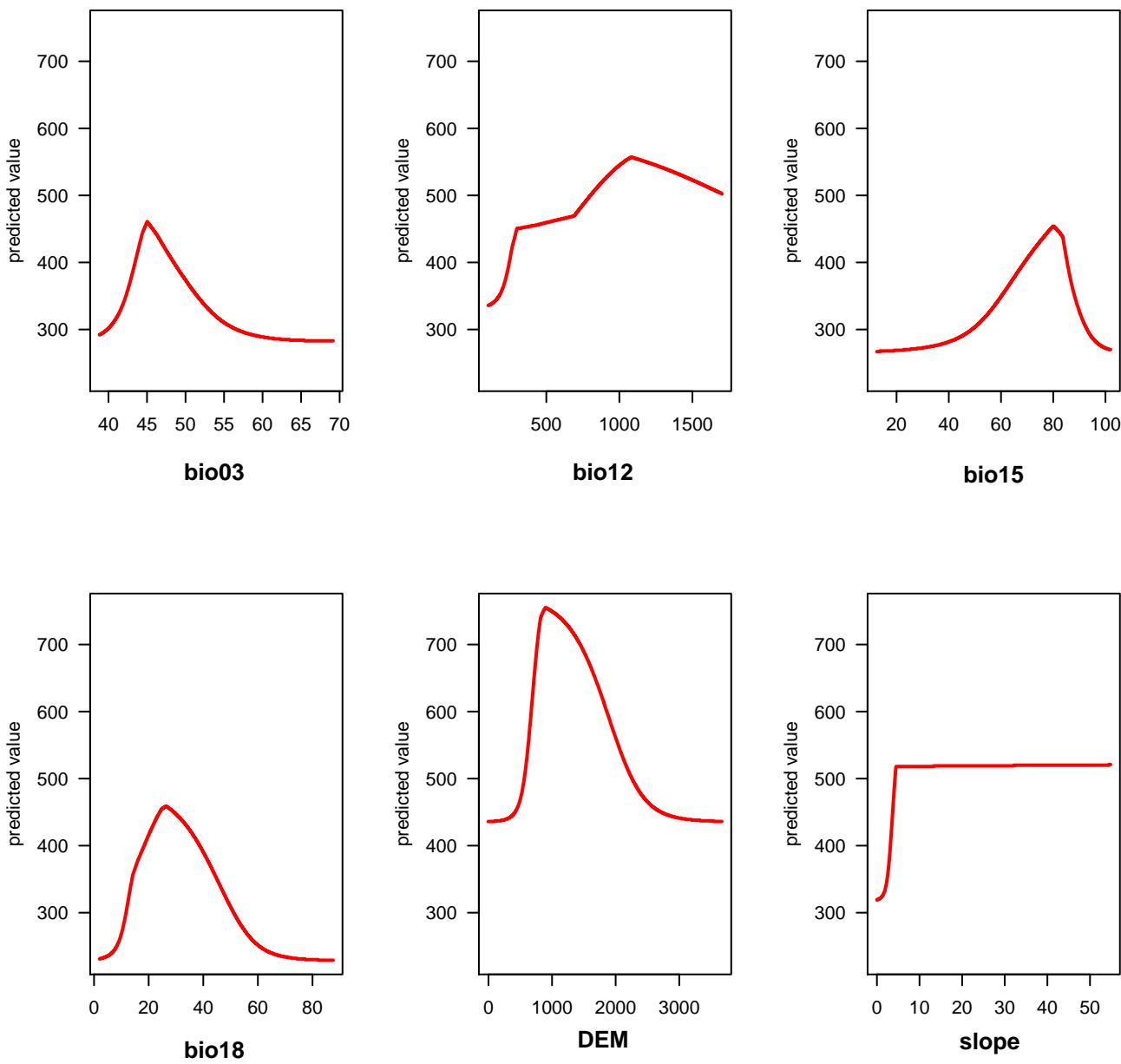


Figure S2.8. Response plots for ecoregion 6

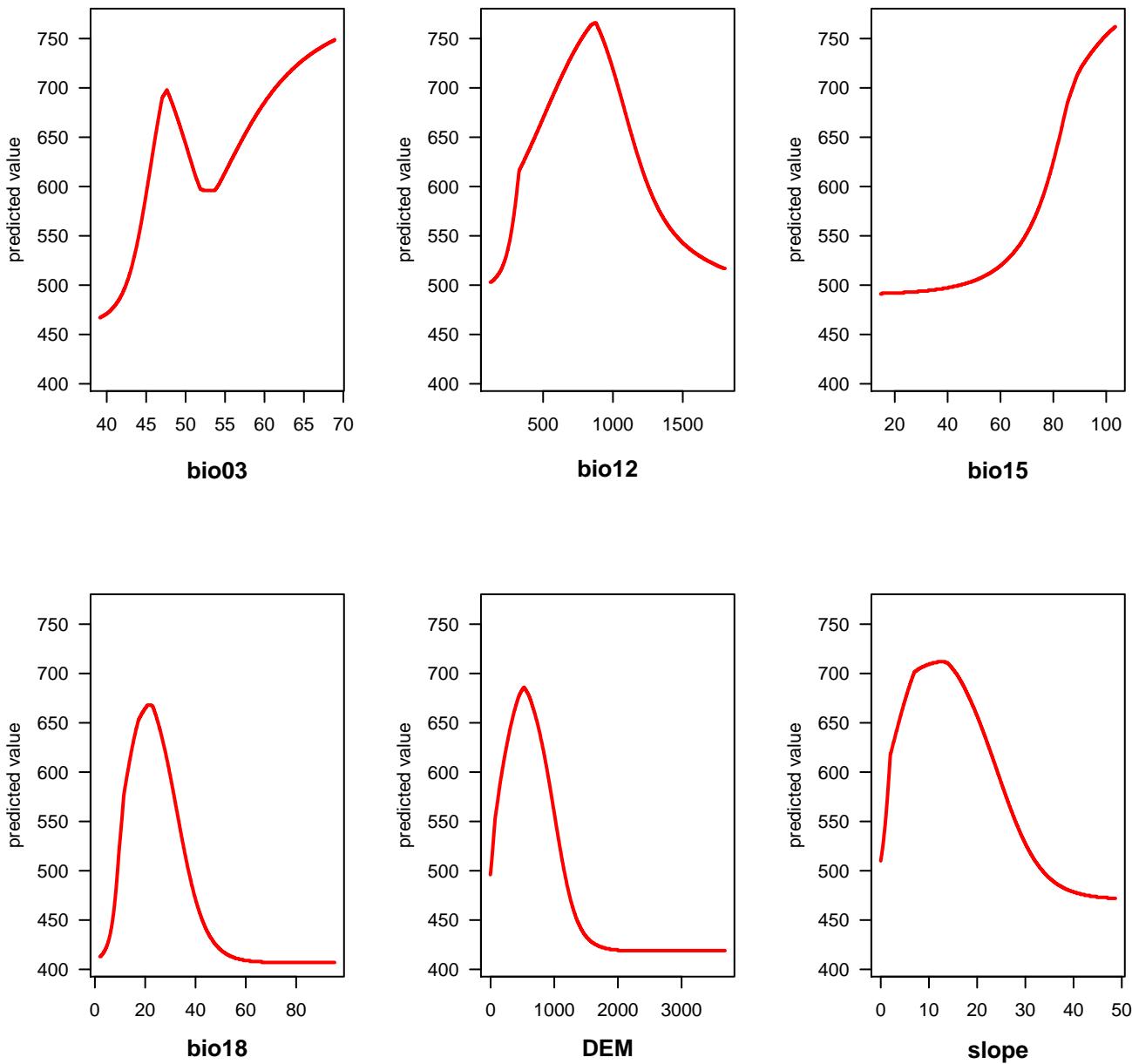


Figure S2.9. Response plots for ecoregion 8

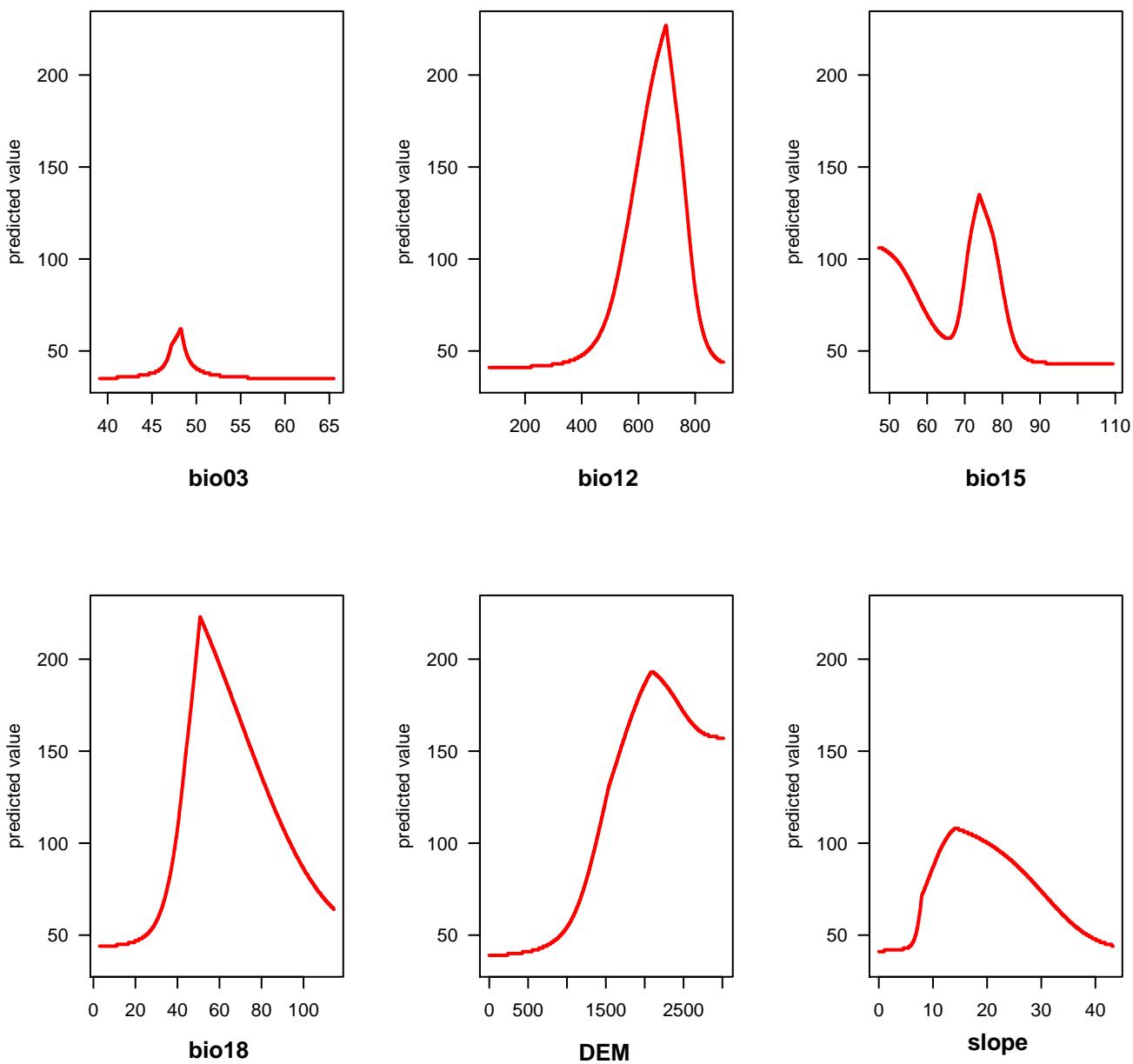


Figure S2.10. Response plots for ecoregion 13

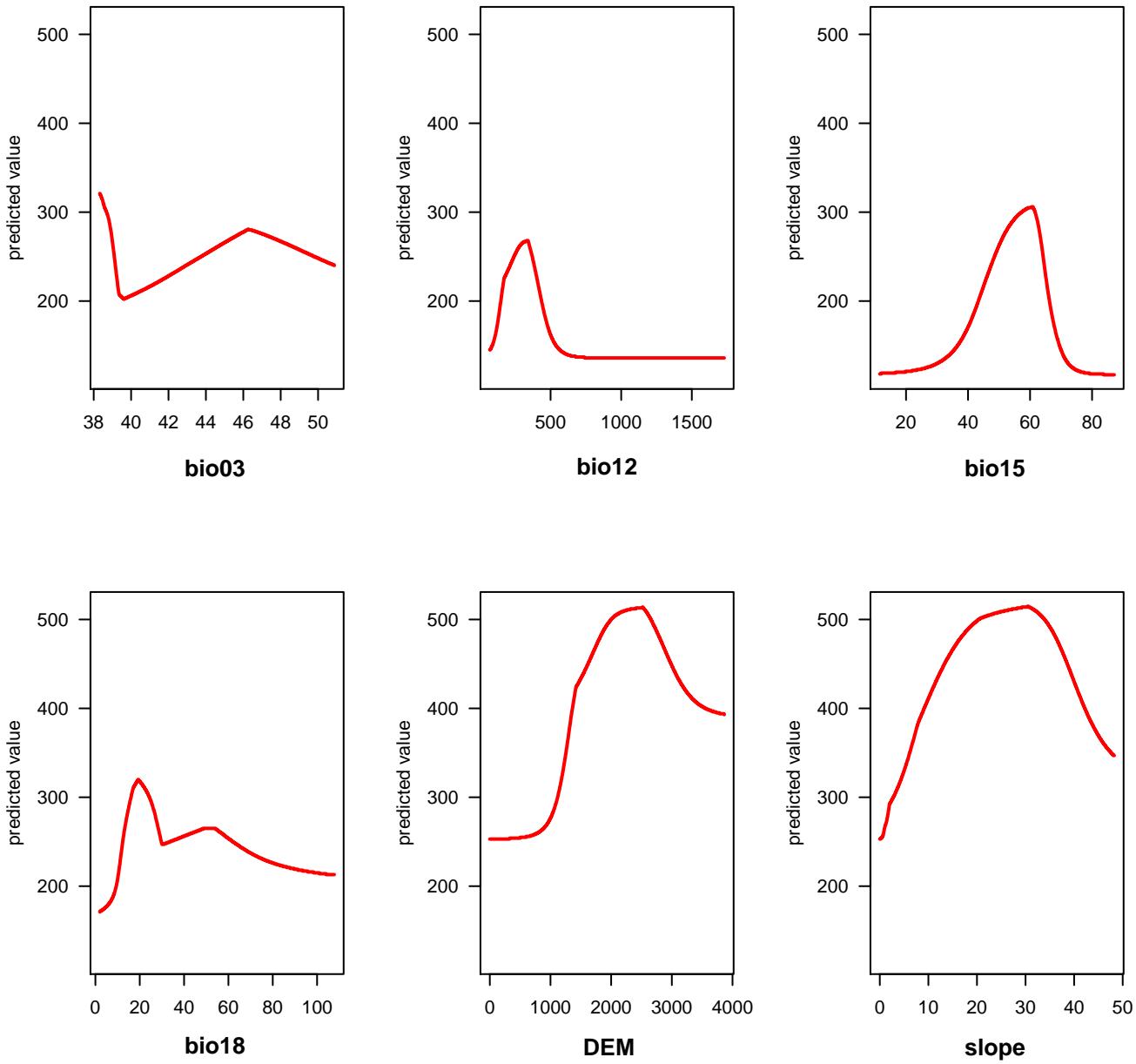


Figure S2.11. Response plots for ecoregion 14

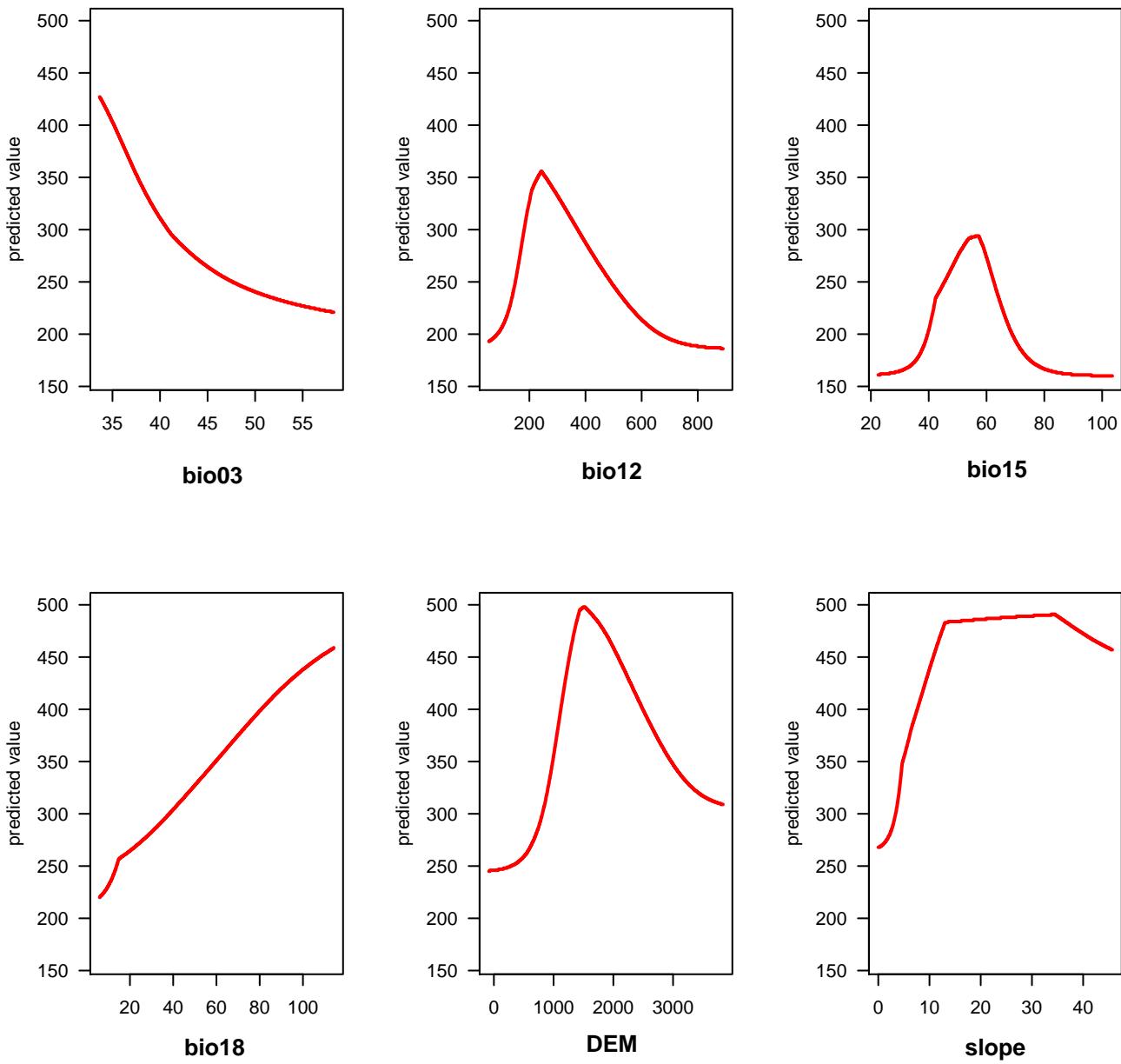


Figure S2.12. Response plots for ecoregion 78

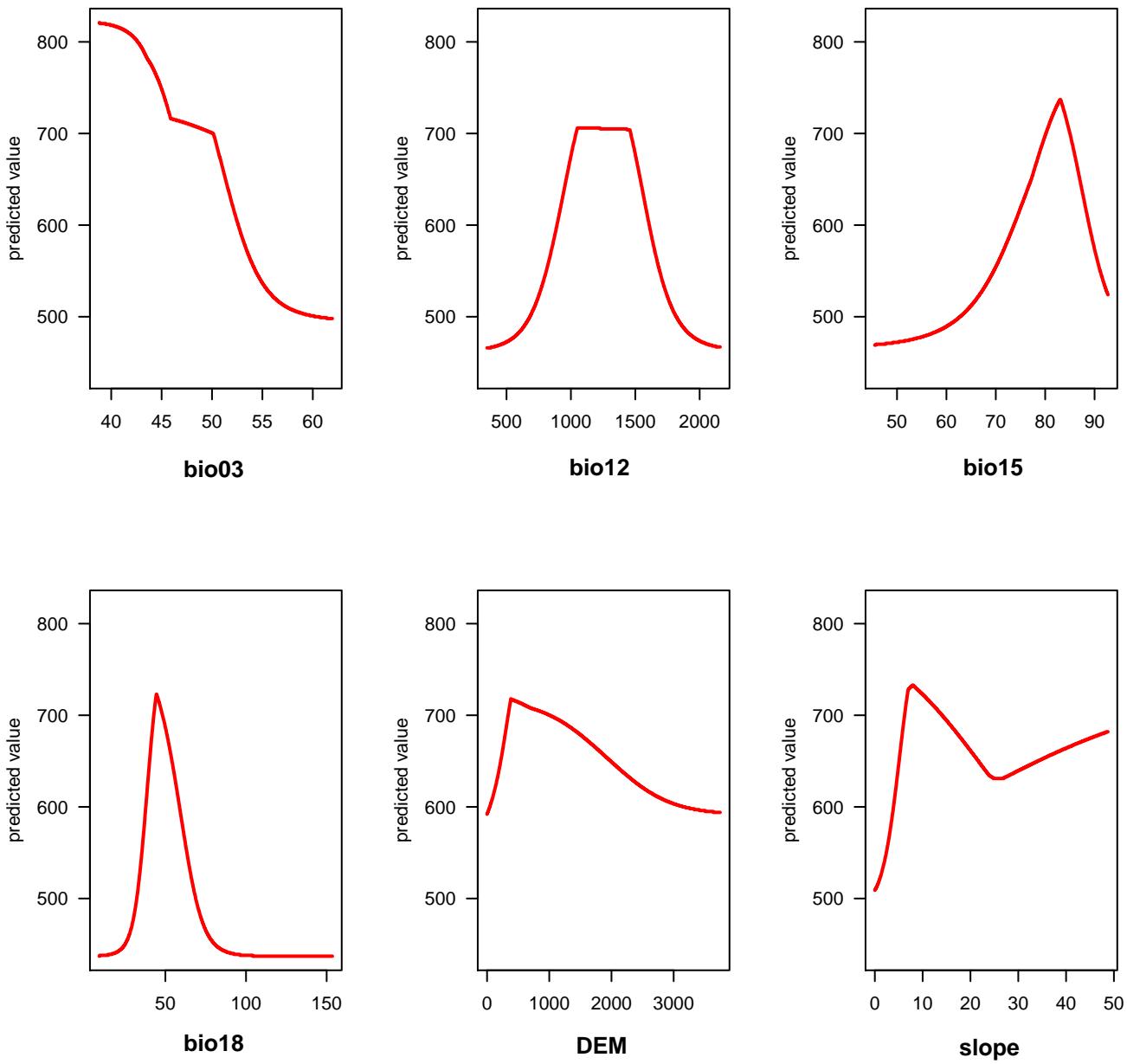


Figure S2.13. Response plots for ecoregion 85

