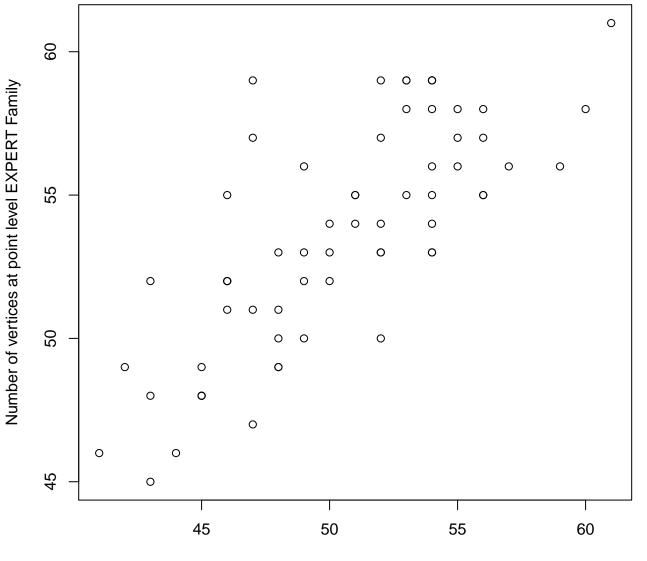
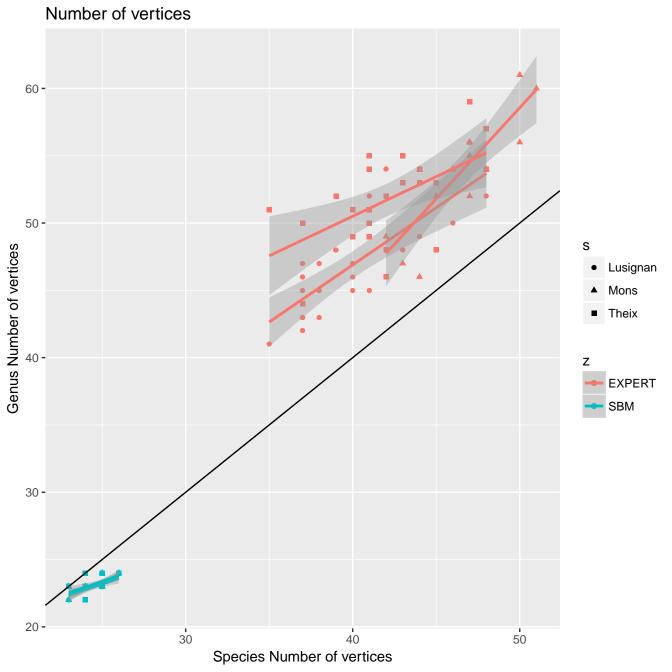
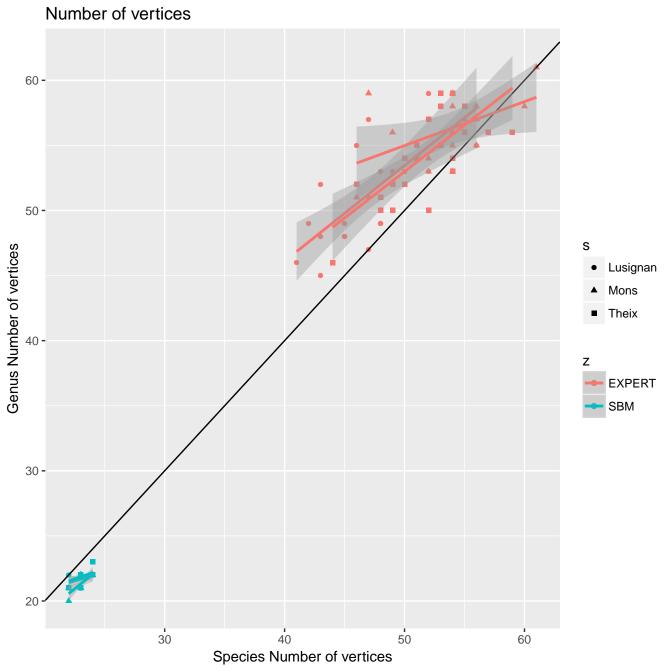


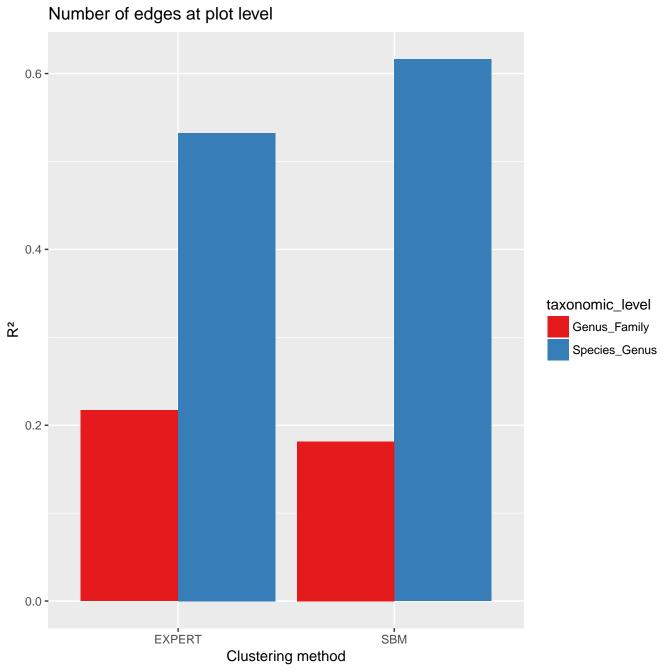
Number of vertices at point level EXPERT Species

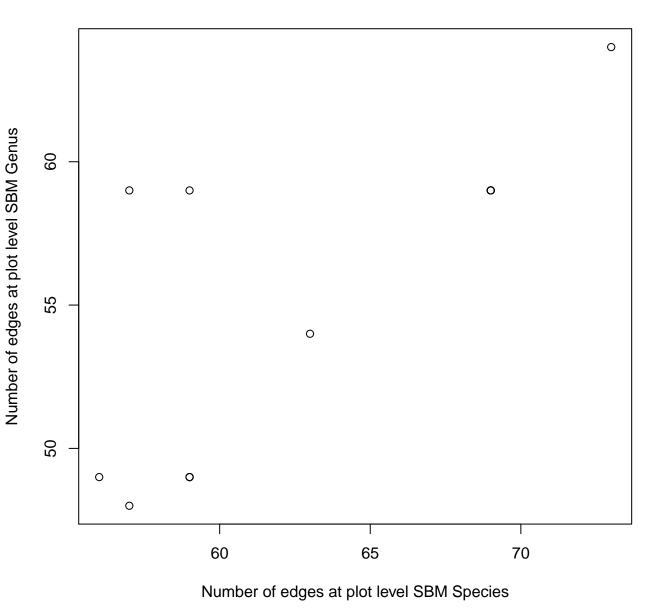


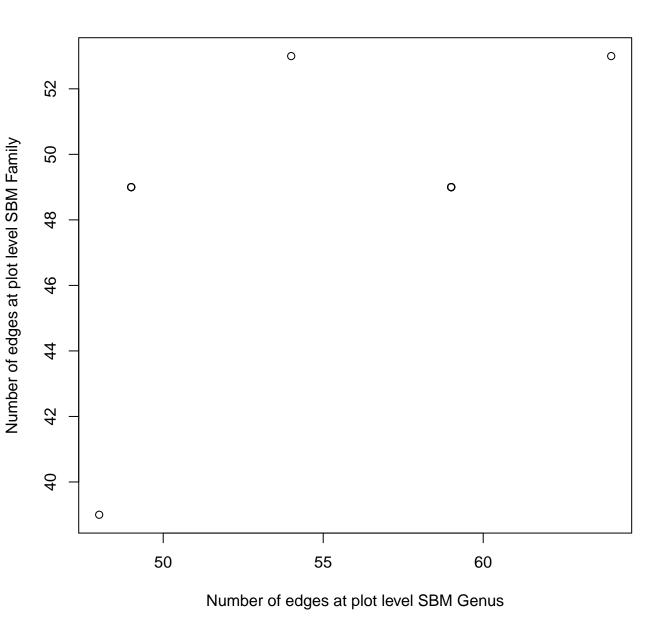
Number of vertices at point level EXPERT Genus

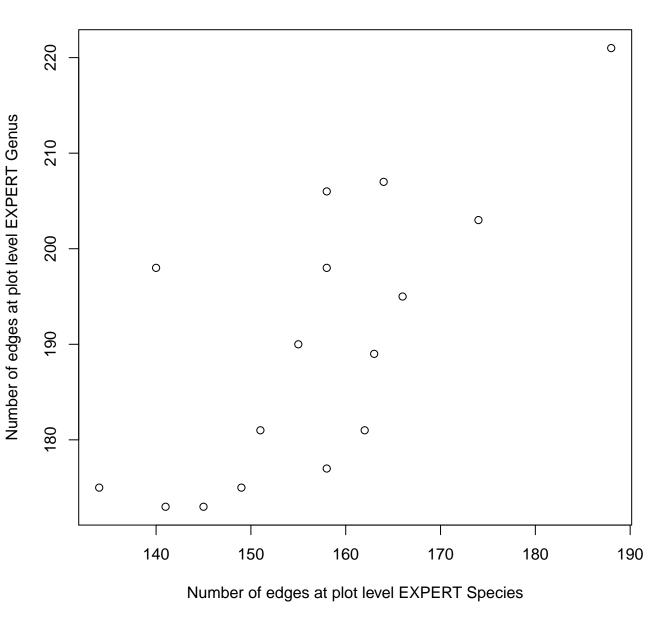


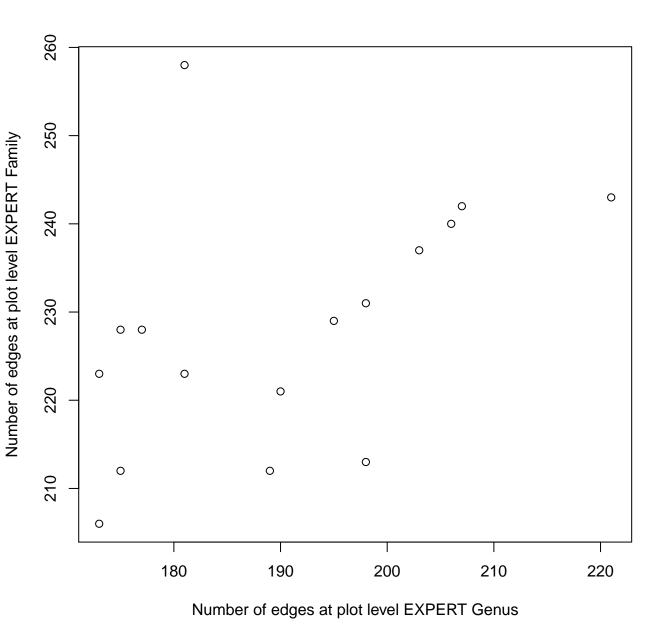


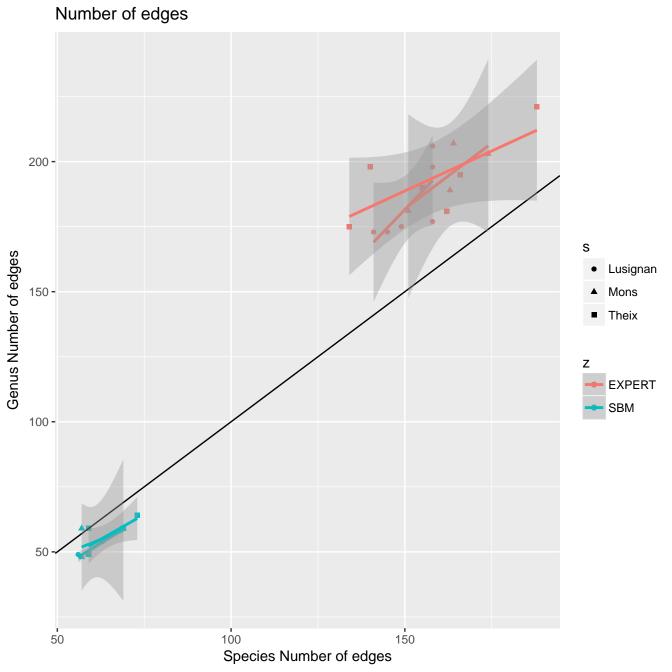


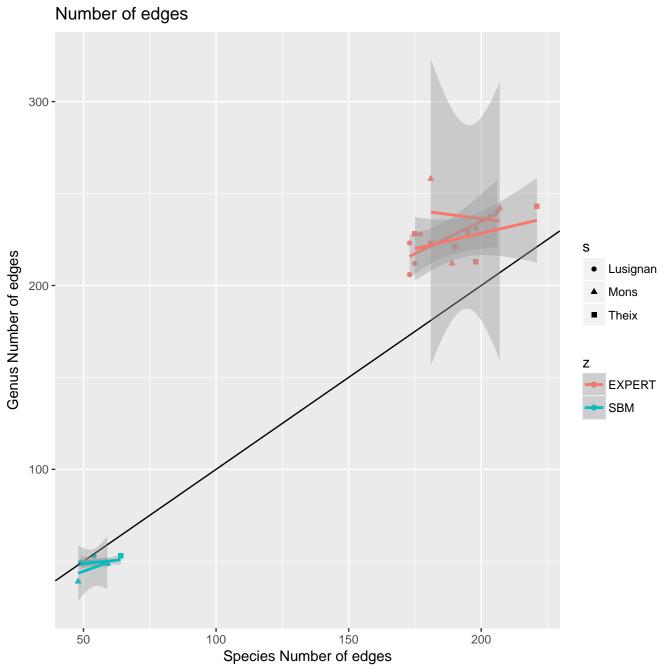


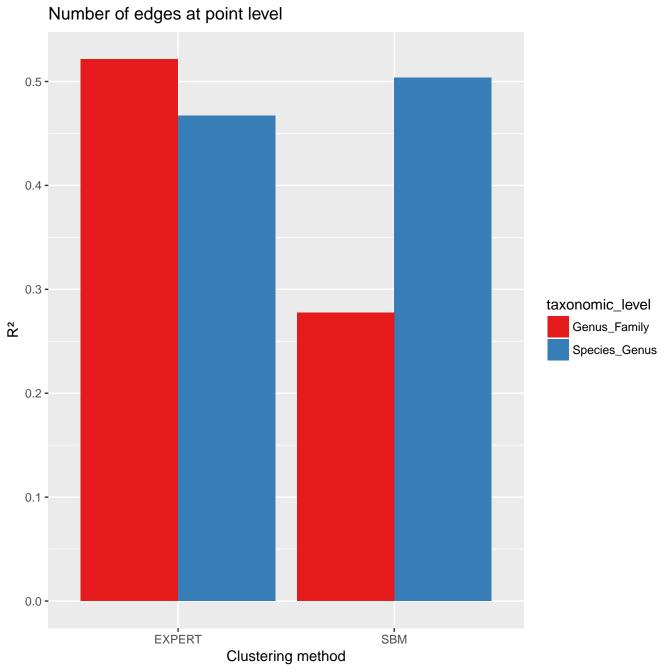


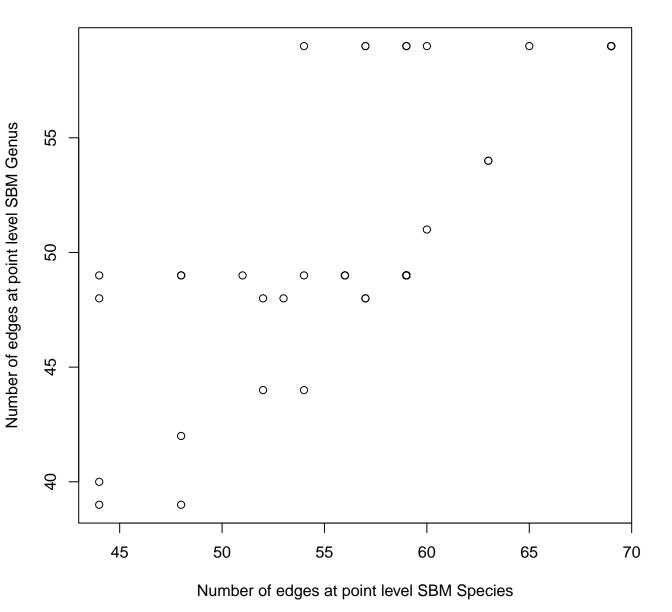


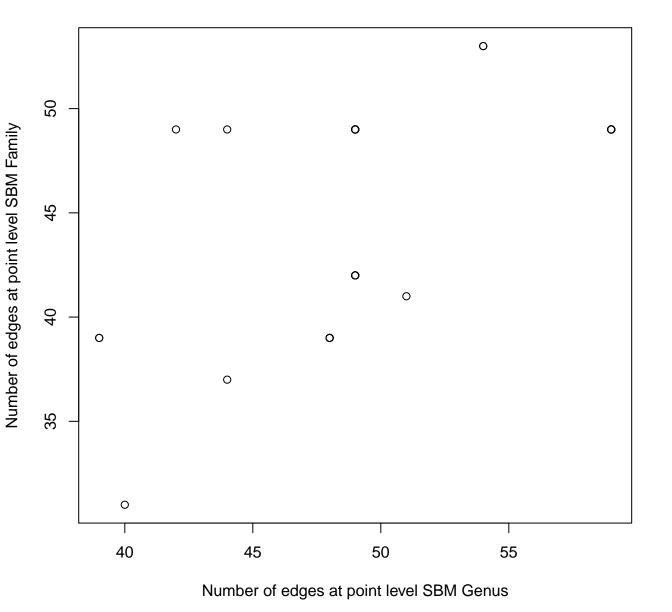


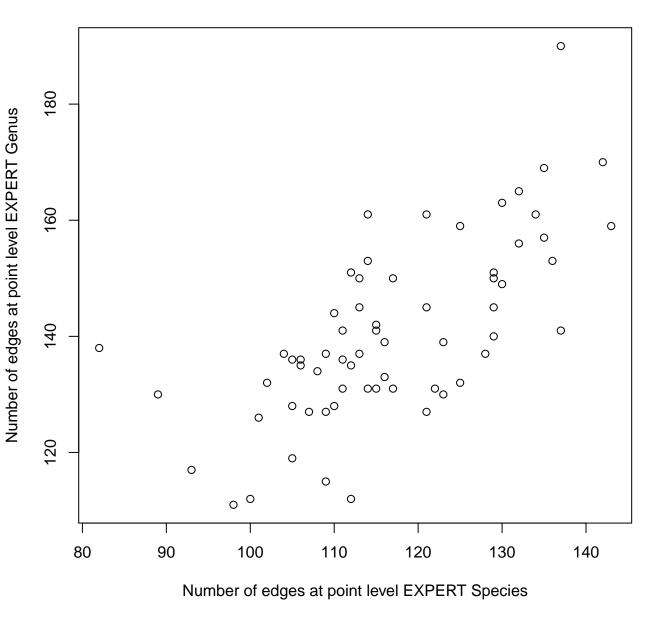


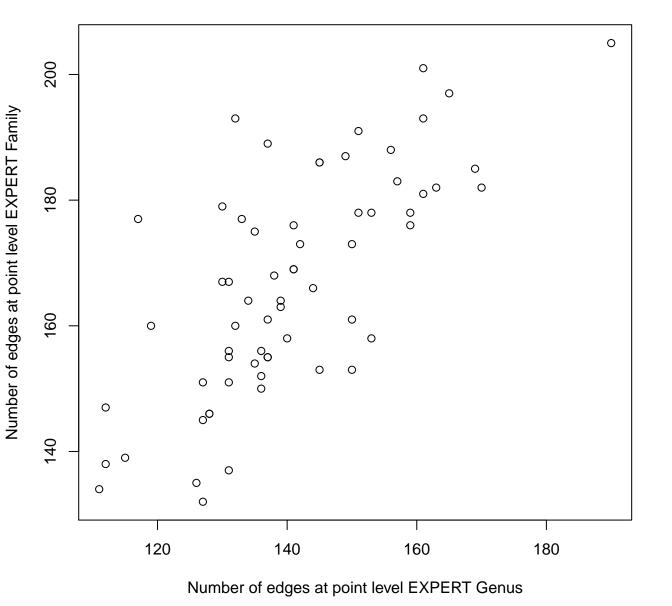


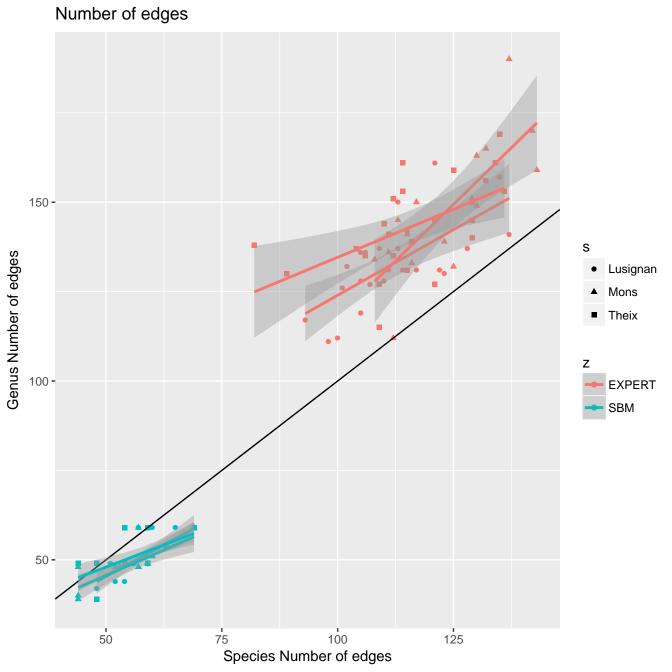


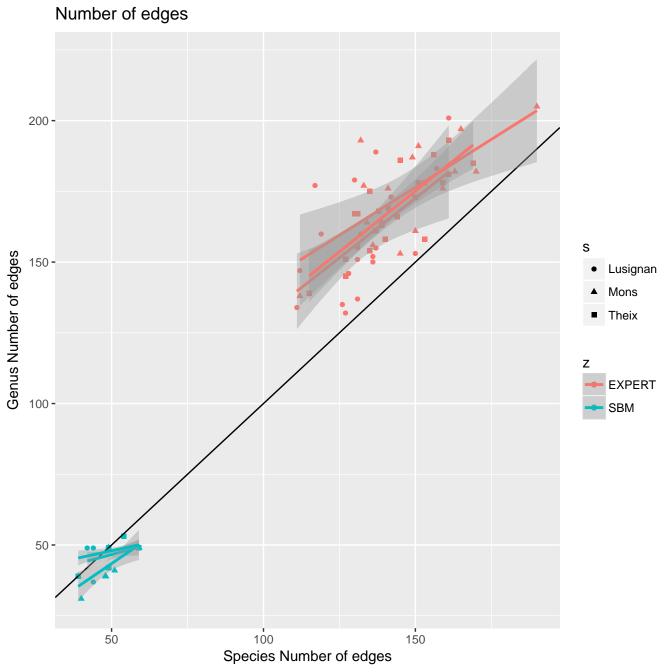




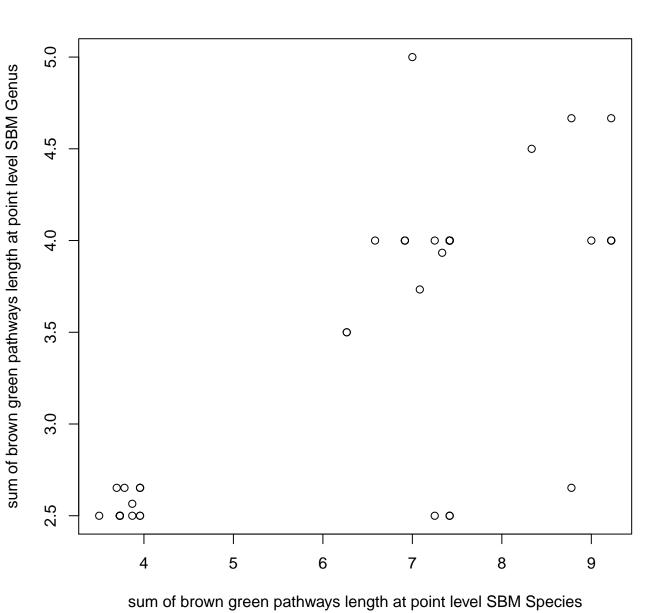


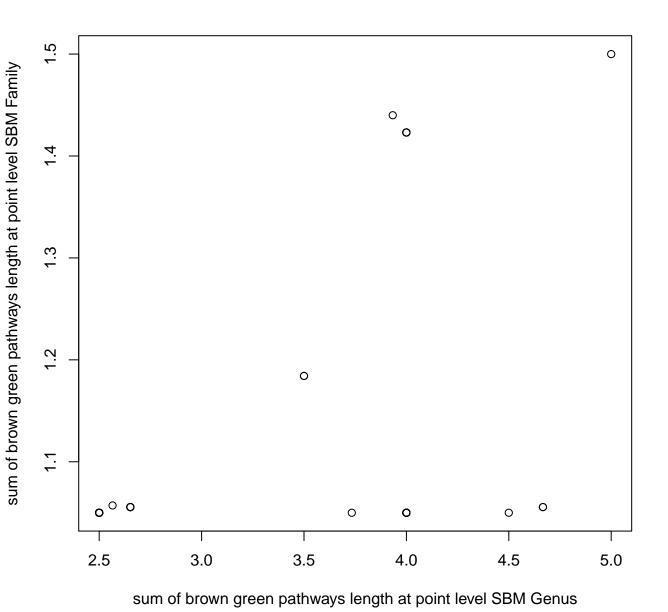


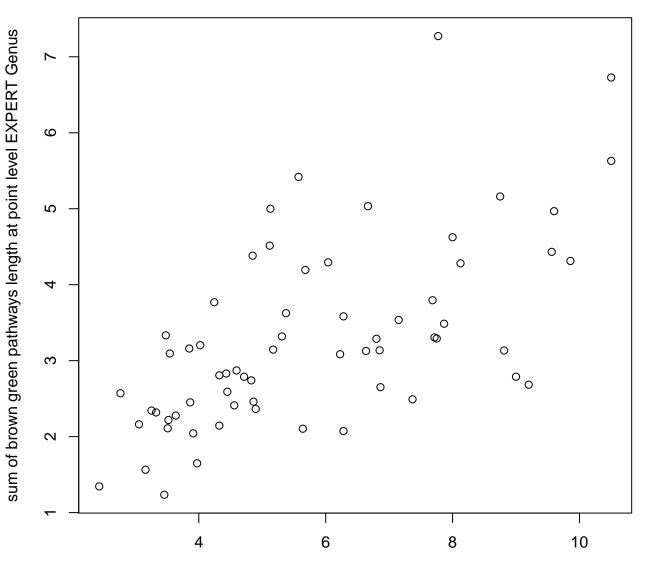




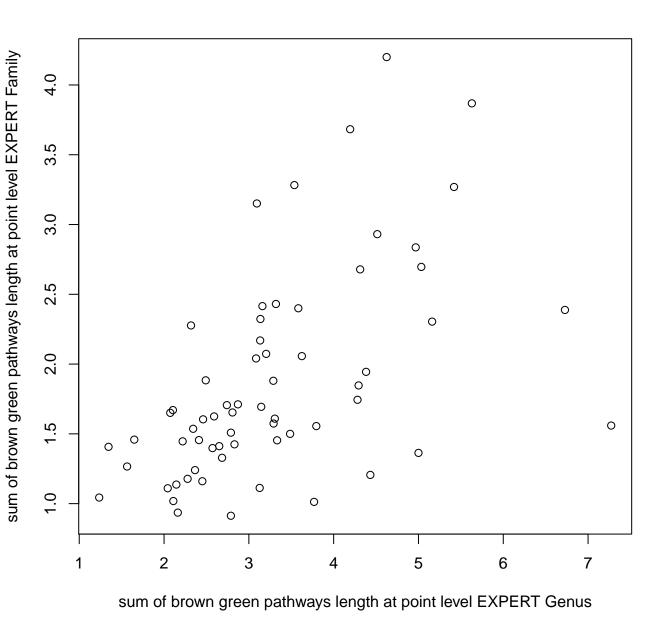
sum of brown green pathways length at point level 0.6 -0.4 taxonomic_level \mathbb{R}^2 Genus_Family Species_Genus 0.2 -0.0 -EXPERT SBM Clustering method

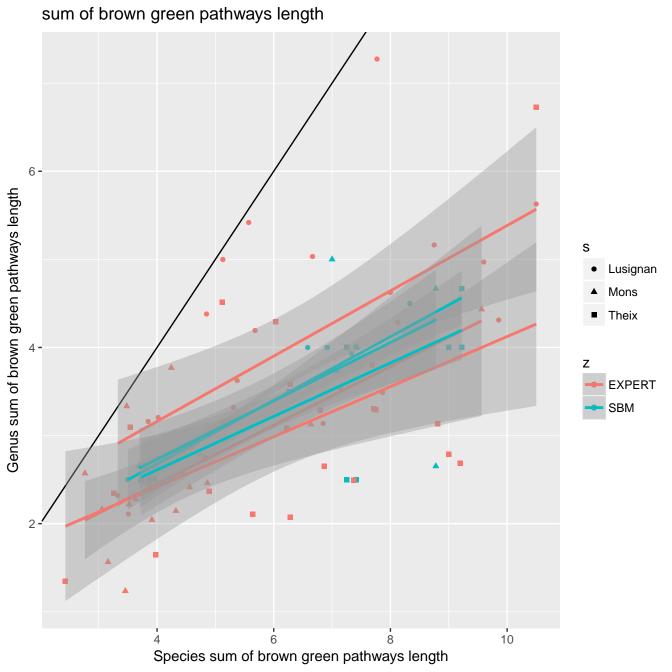


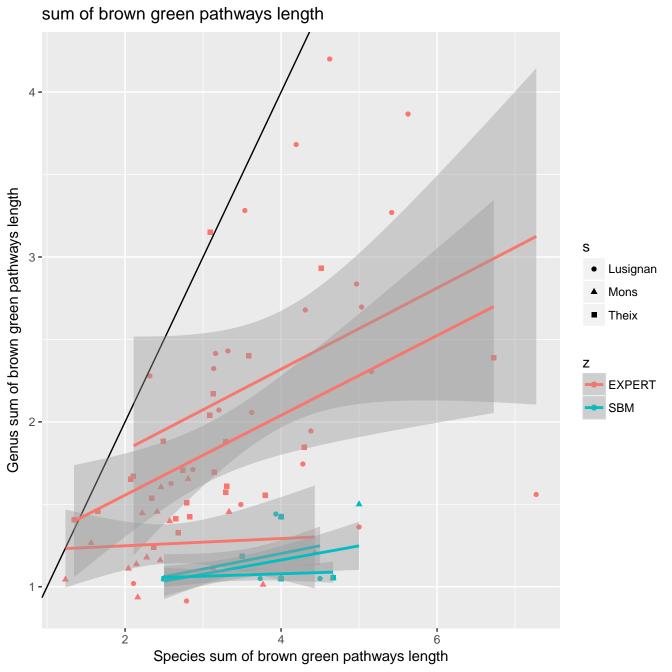




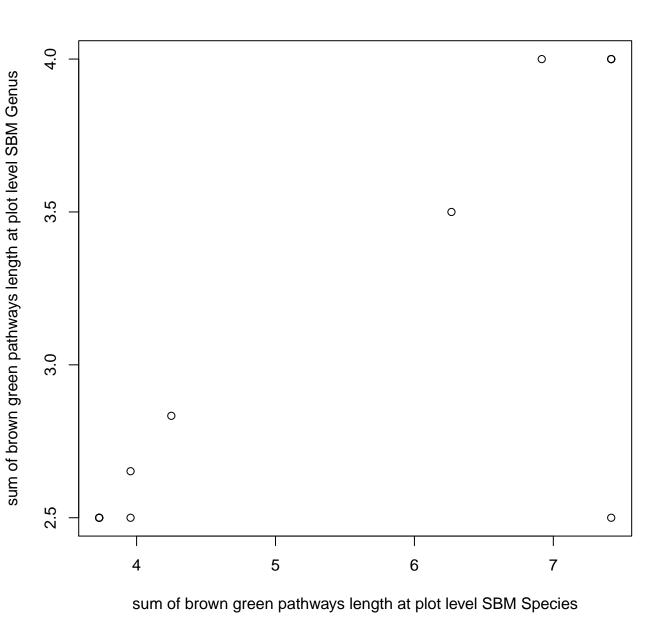
sum of brown green pathways length at point level EXPERT Species

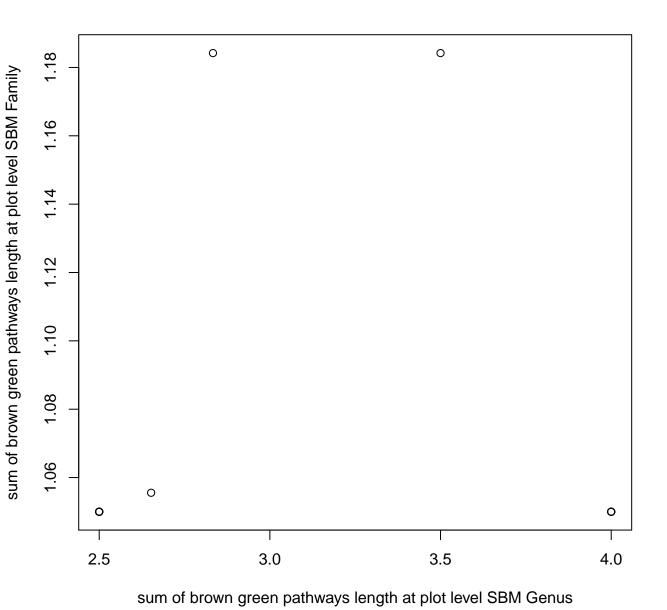


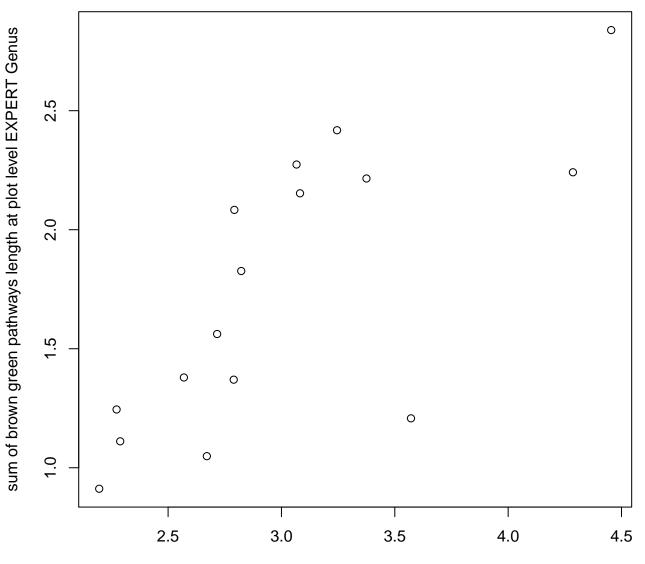




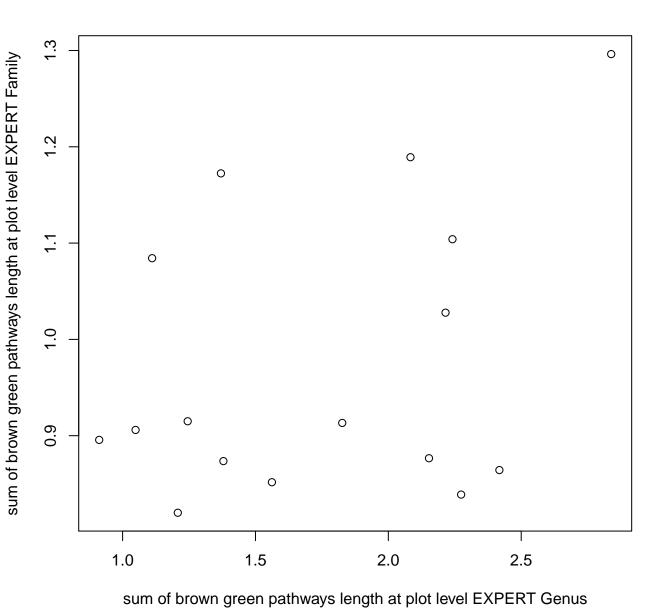
sum of brown green pathways length at plot level 0.6 -0.4 taxonomic_level \mathbb{R}^2 Genus_Family Species_Genus 0.2 -0.0 -EXPERT SBM Clustering method

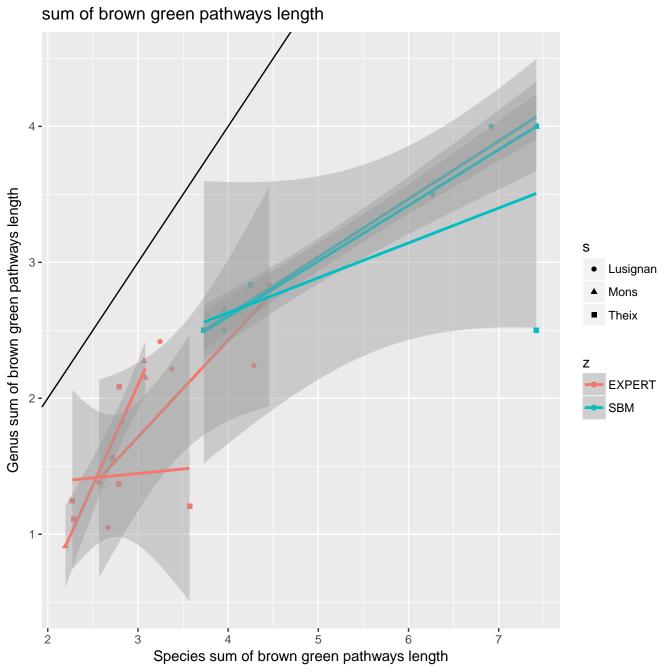


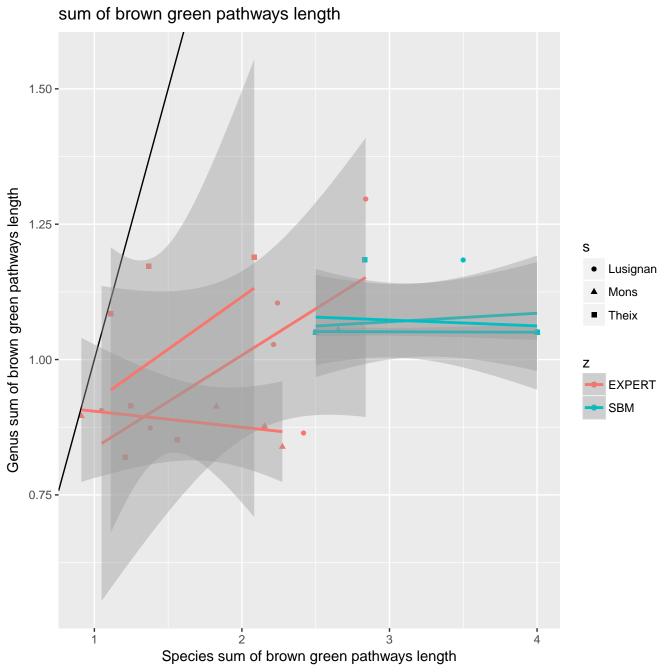




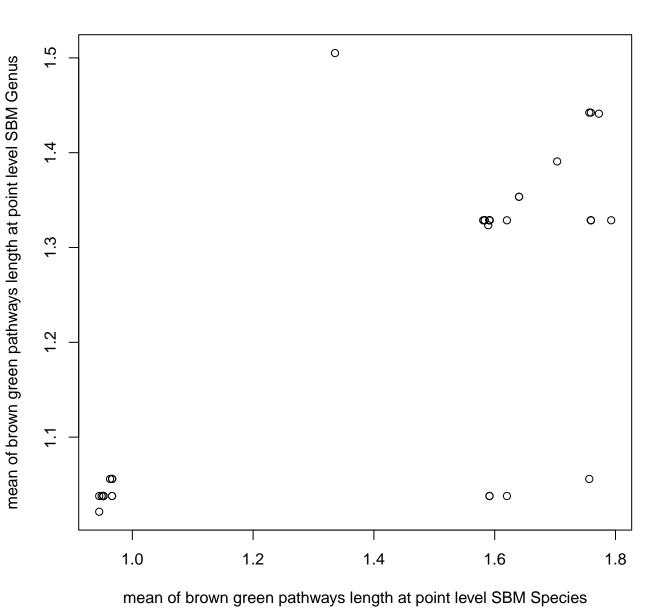
sum of brown green pathways length at plot level EXPERT Species

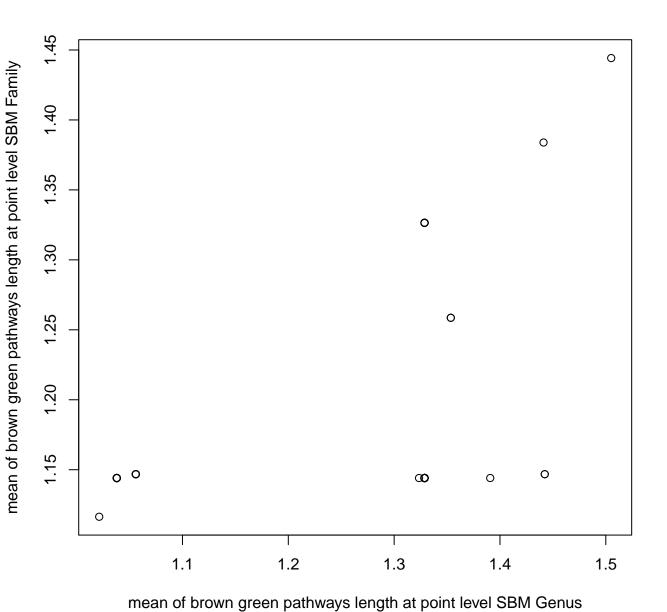


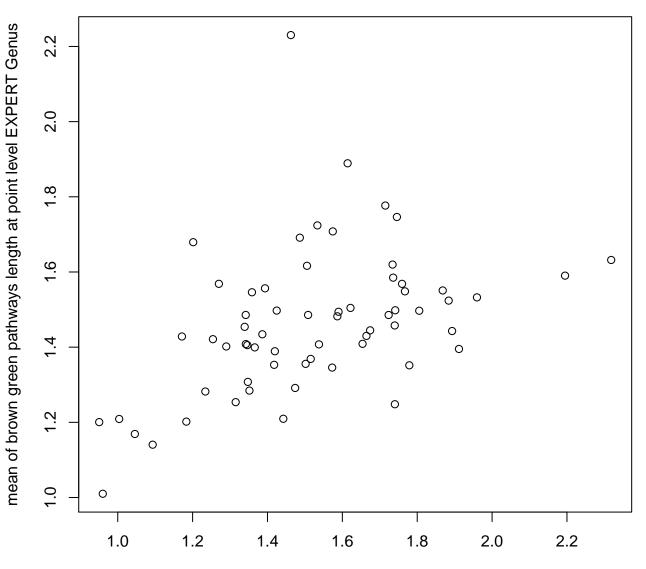




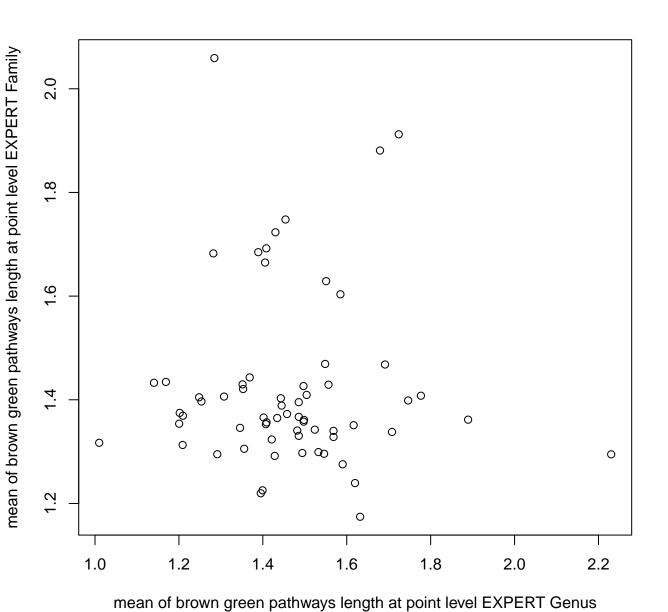
mean of brown green pathways length at point level 0.6 -0.4 taxonomic_level \mathbb{R}^2 Genus_Family Species_Genus 0.2 -0.0 -EXPERT SBM Clustering method

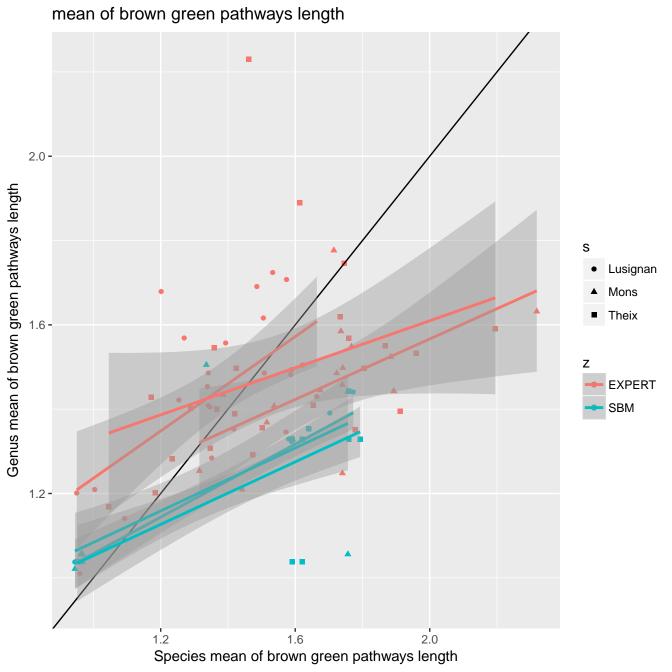


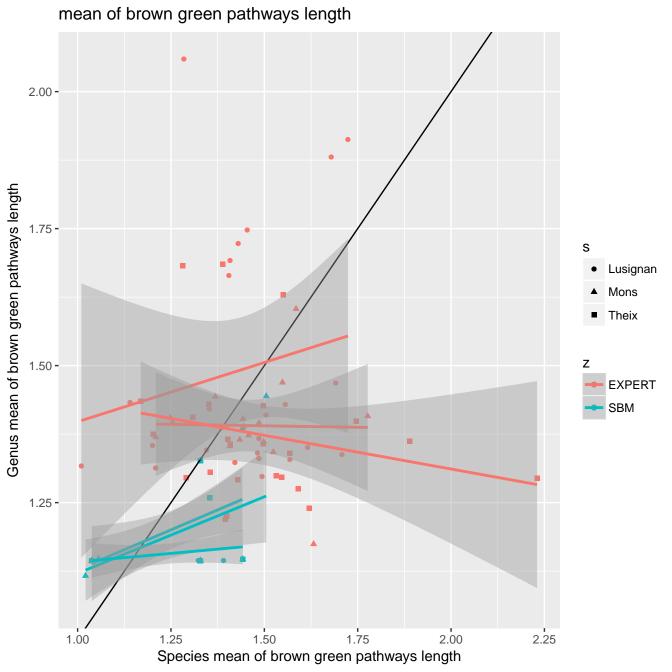




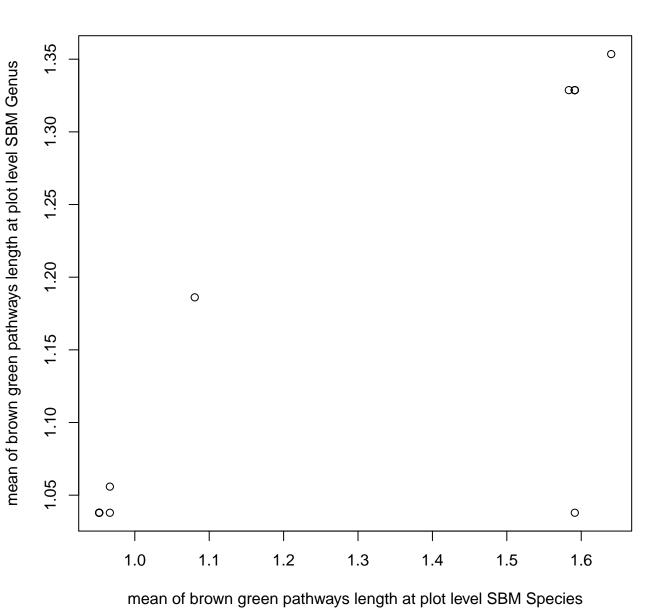
mean of brown green pathways length at point level EXPERT Species

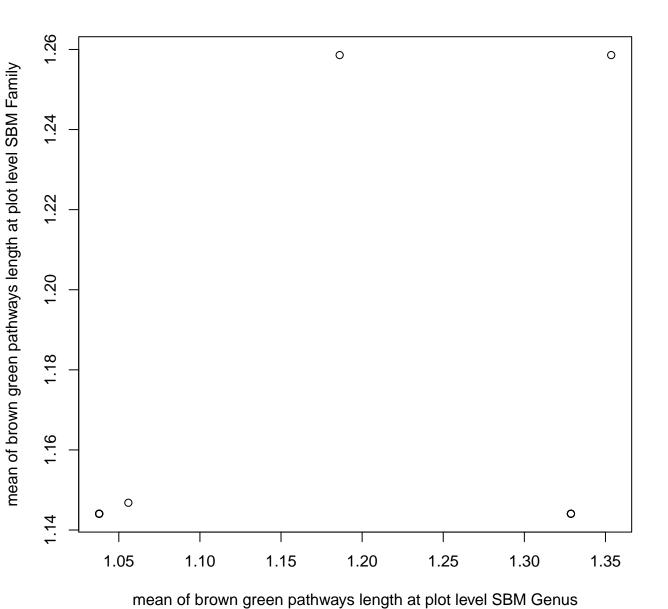


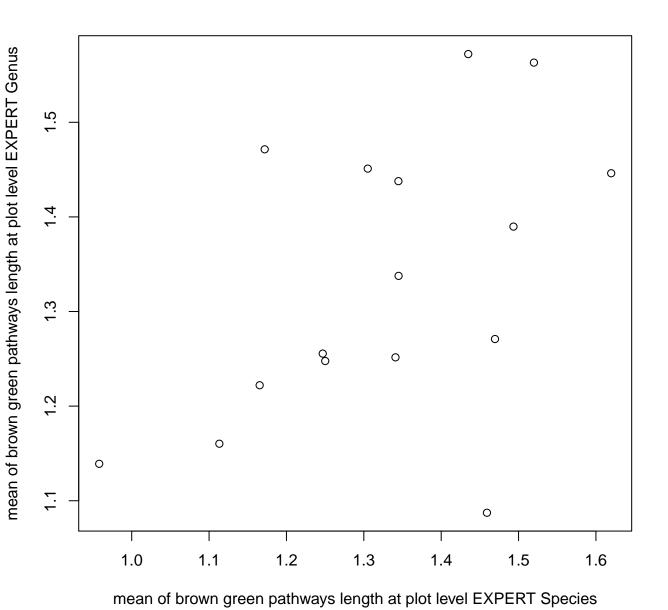


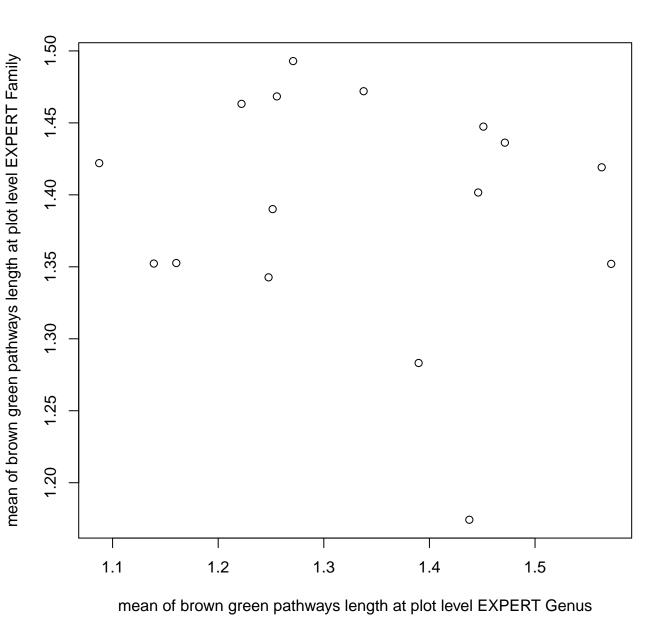


mean of brown green pathways length at plot level 0.6 taxonomic_level 0.4 -Genus_Family \mathbb{R}^2 Species_Genus 0.2 -0.0 -EXPERT SBM Clustering method

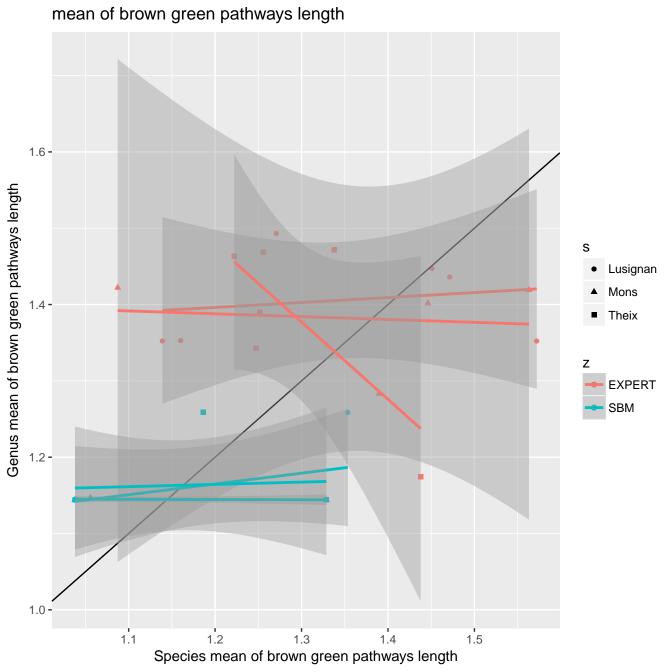




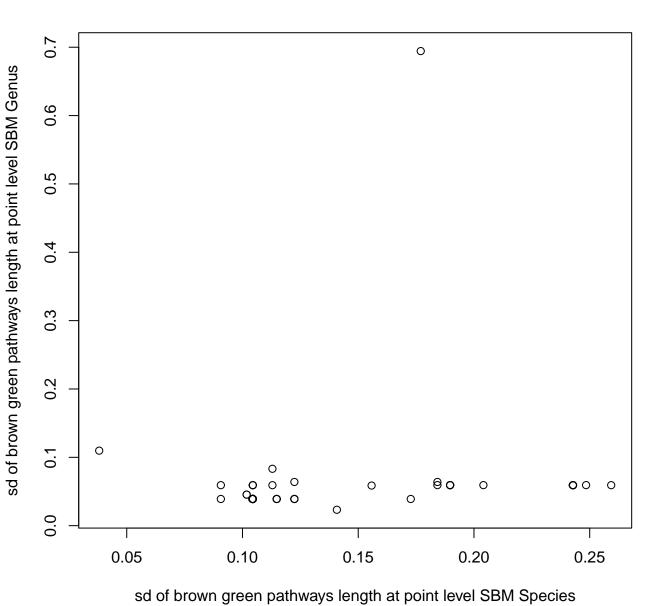


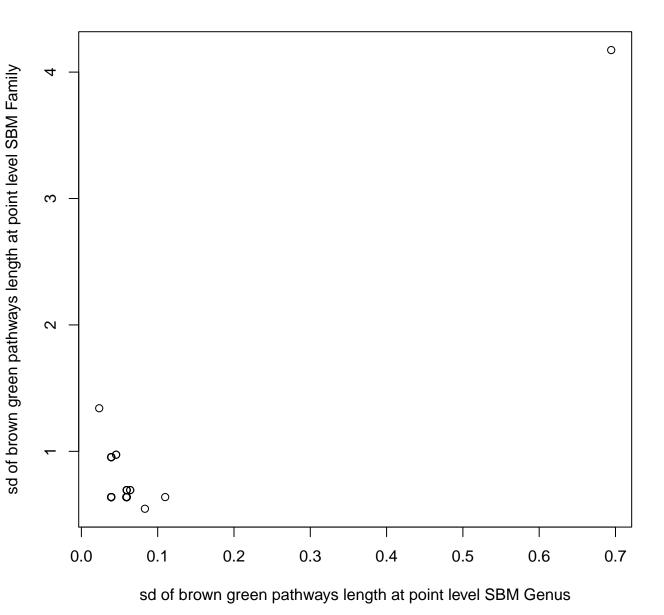


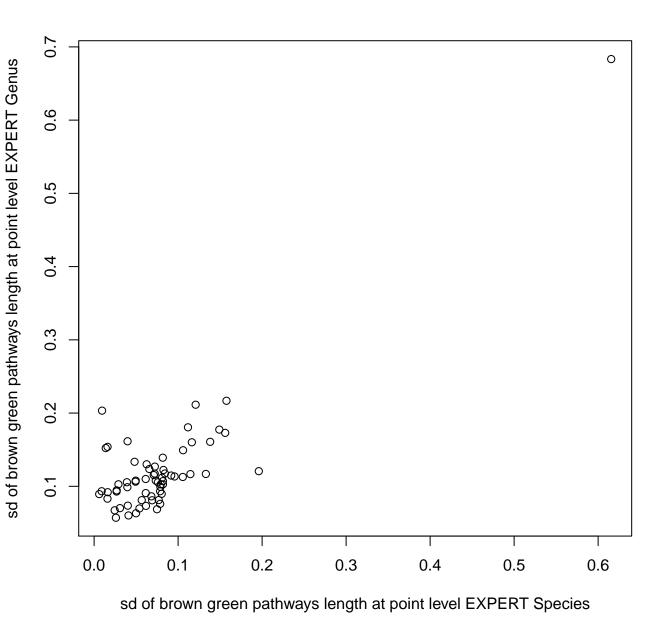
mean of brown green pathways length 2.0 -Genus mean of brown green pathways length s Lusignan Mons Theix 1.5 **-**Ζ EXPERT SBM 1.0 -1.1 1.3 1.5 Species mean of brown green pathways length

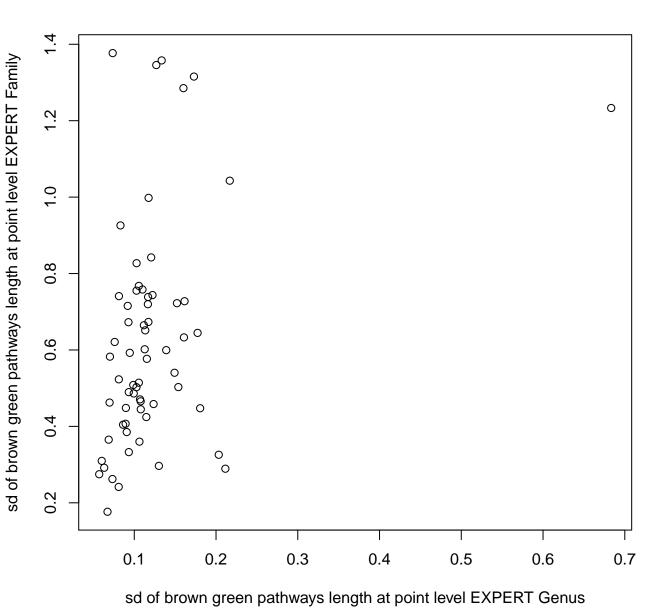


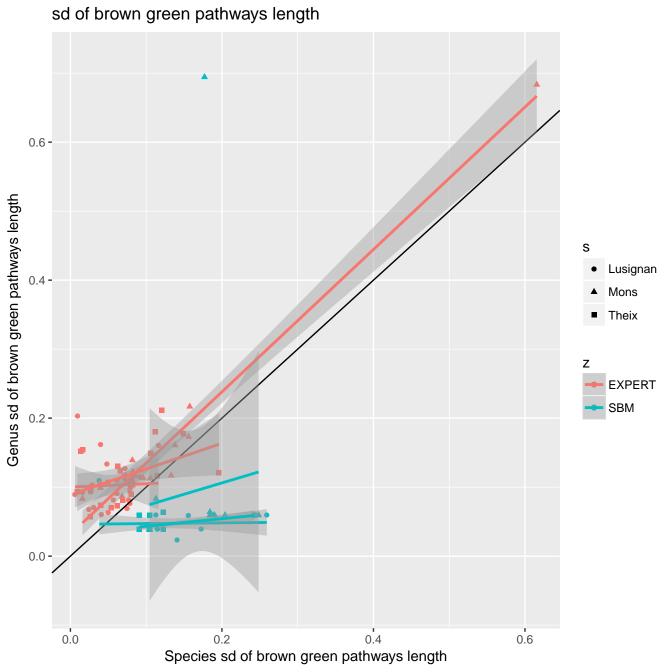
sd of brown green pathways length at point level 0.8 -0.6 taxonomic_level Genus_Family 0.4 -Species_Genus 0.2 -0.0 -EXPERT SBM Clustering method

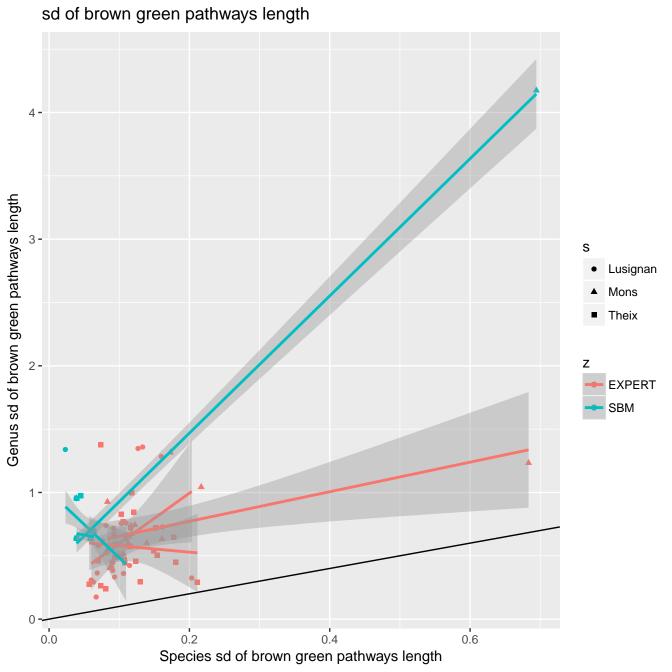




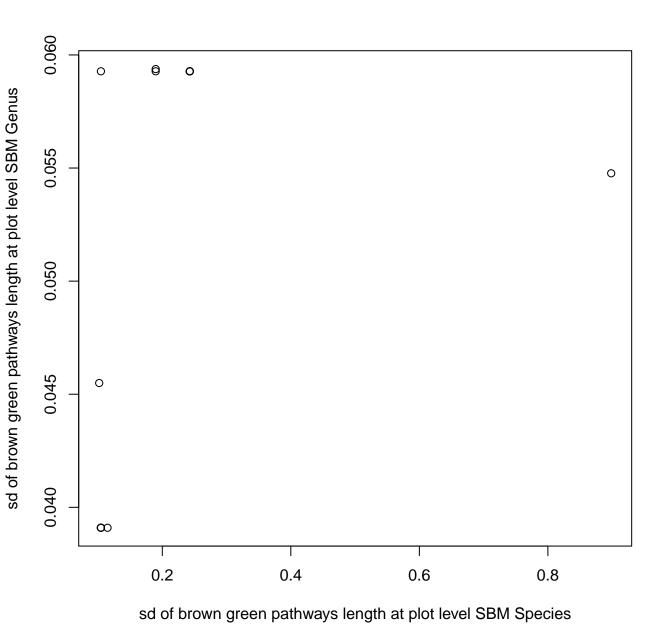


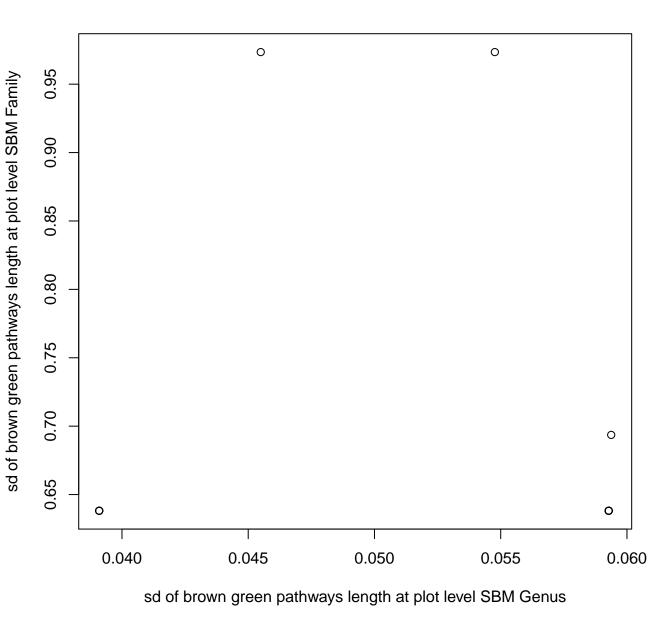


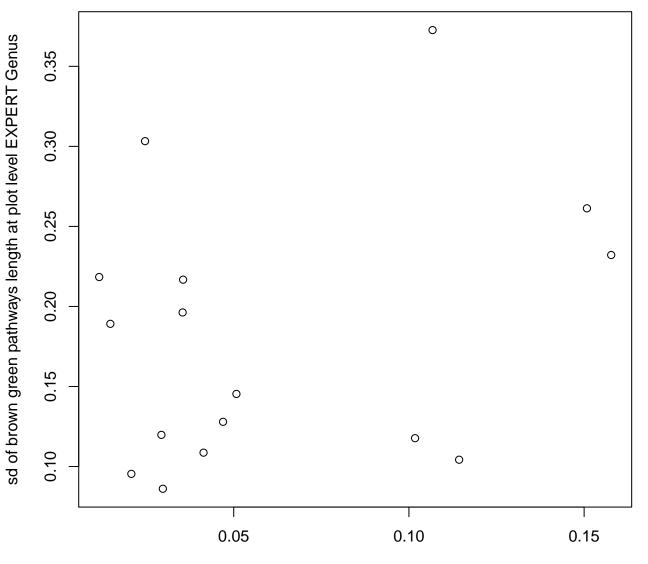




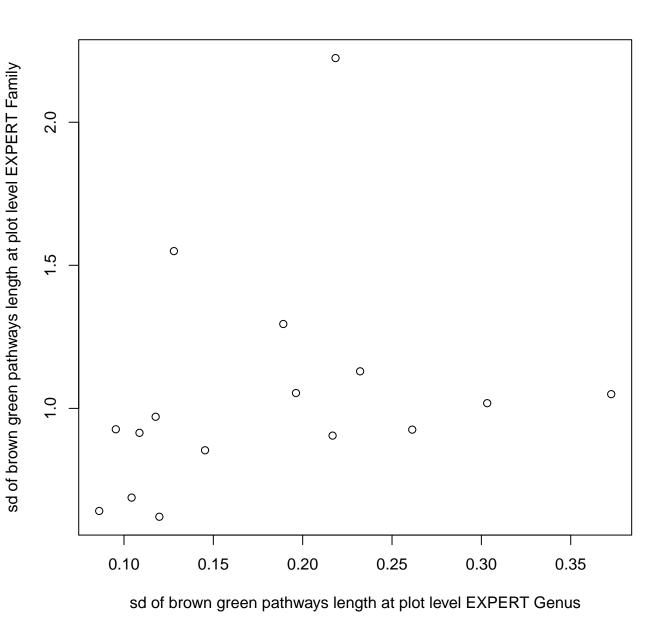
sd of brown green pathways length at plot level 0.12 -0.08 taxonomic_level \mathbb{R}^2 Genus_Family Species_Genus 0.04 -0.00 -EXPERT SBM Clustering method

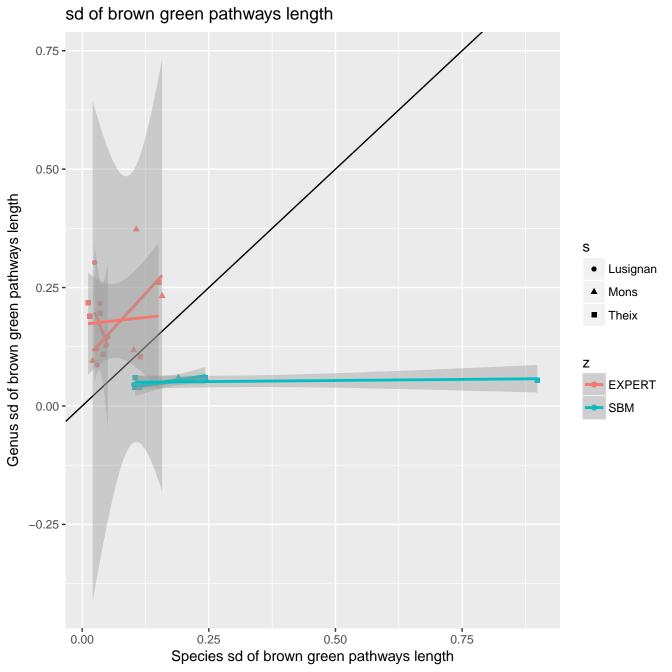


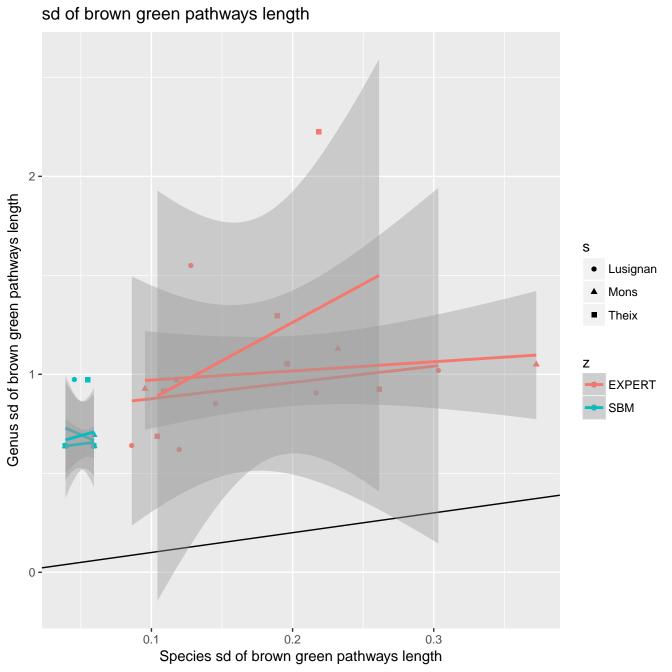


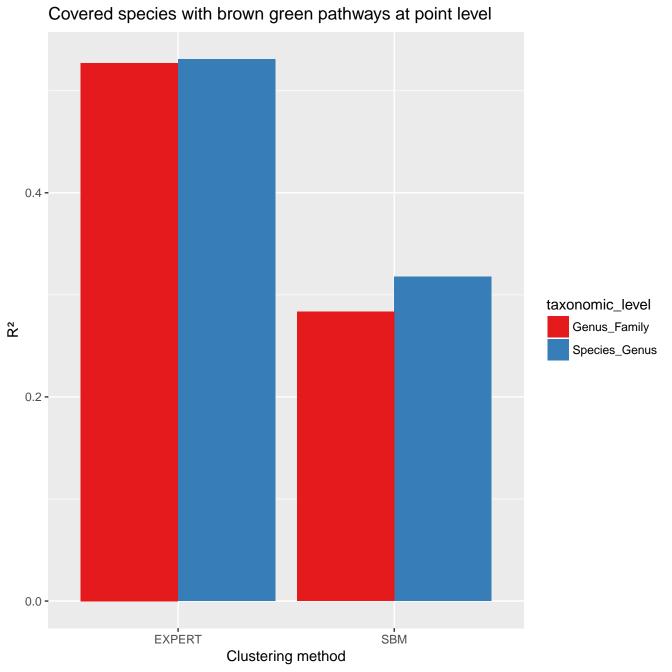


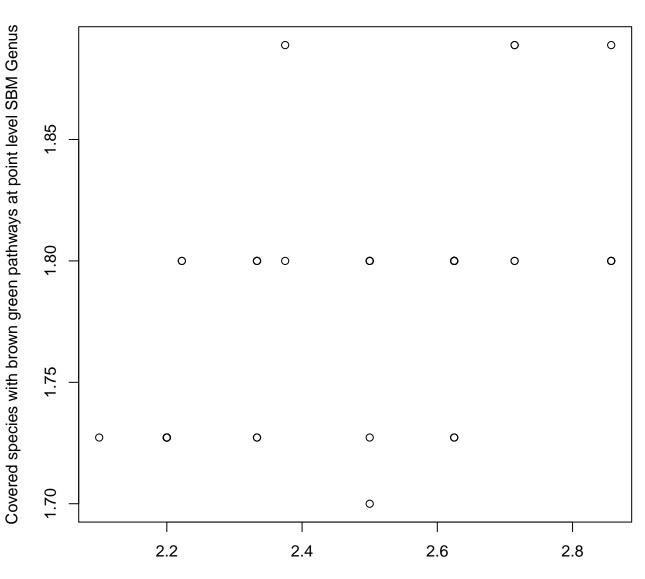
sd of brown green pathways length at plot level EXPERT Species



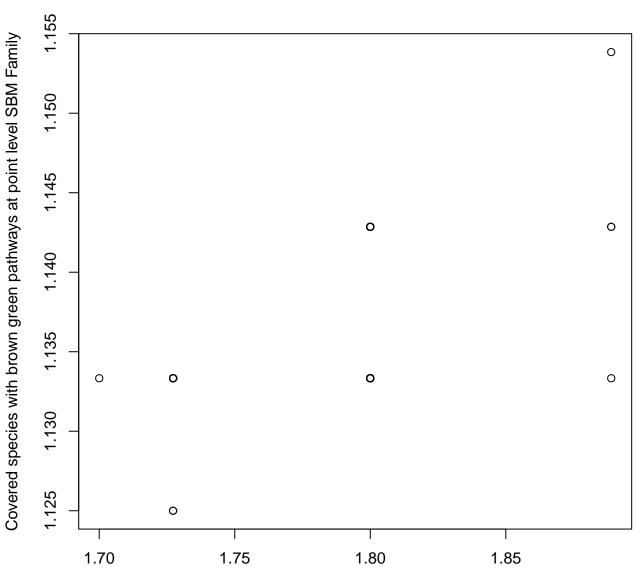




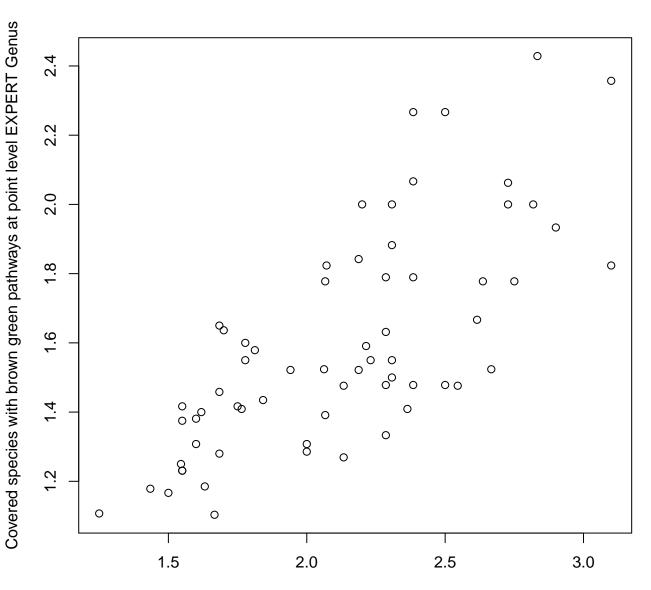




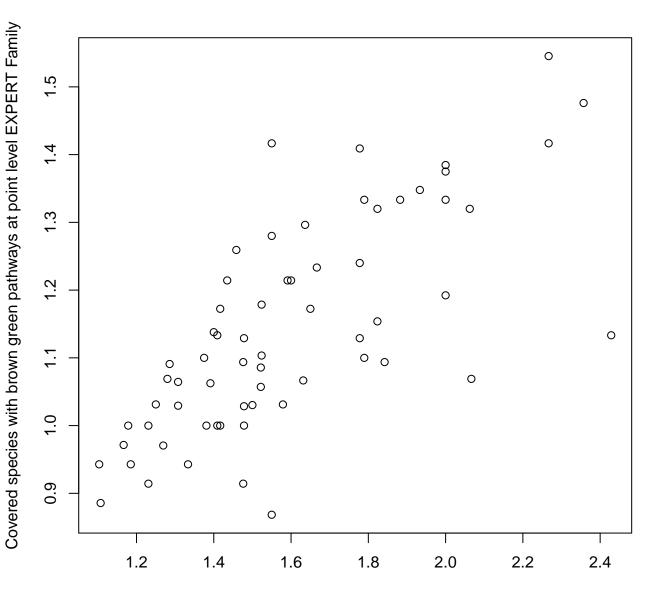
Covered species with brown green pathways at point level SBM Species



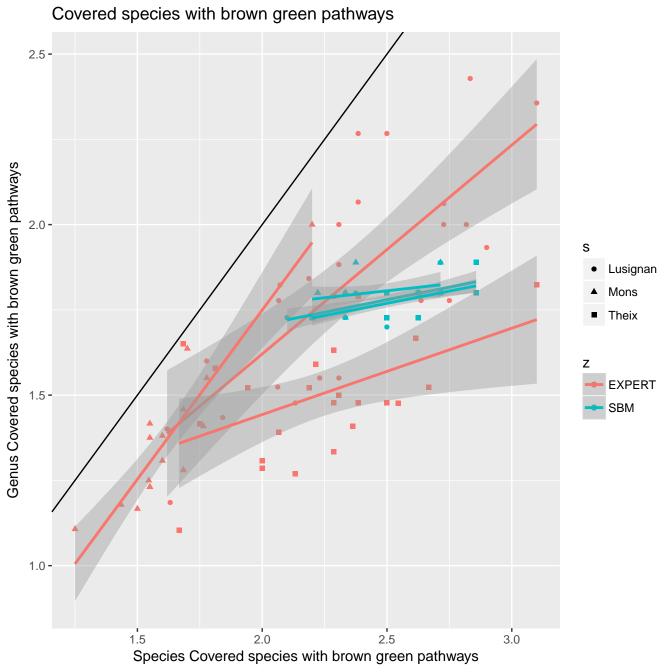
Covered species with brown green pathways at point level SBM Genus

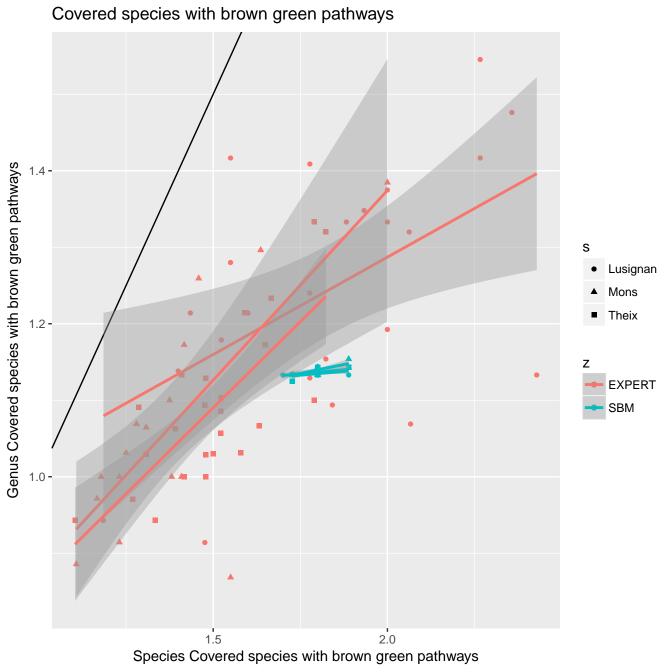


Covered species with brown green pathways at point level EXPERT Species

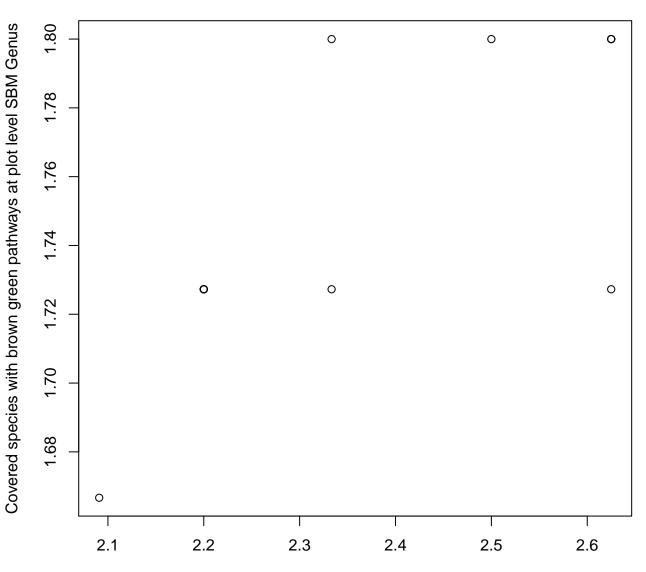


Covered species with brown green pathways at point level EXPERT Genus

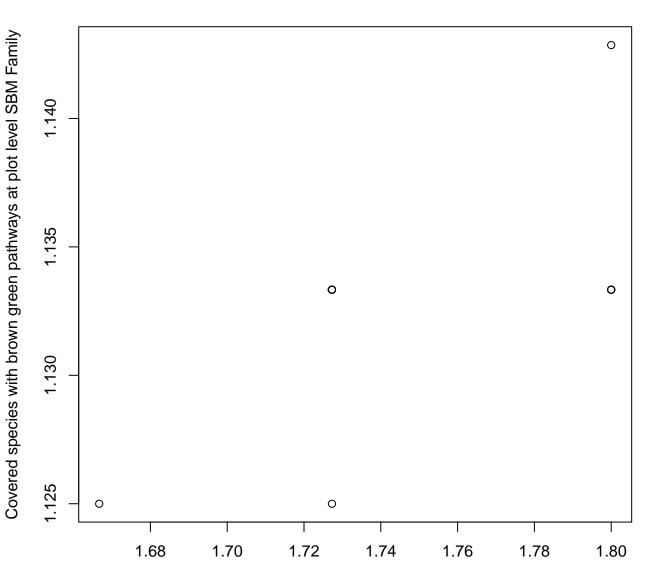




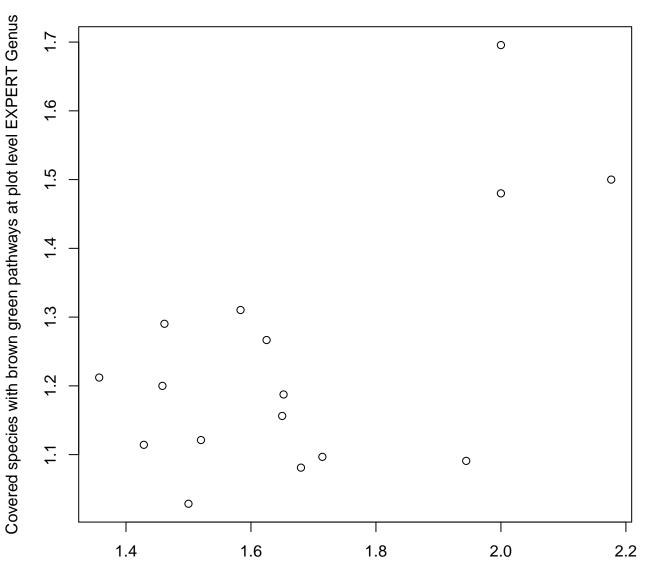
Covered species with brown green pathways at plot level 0.6 -0.4 taxonomic_level \mathbb{R}^2 Genus_Family Species_Genus 0.2 -0.0 -EXPERT SBM Clustering method



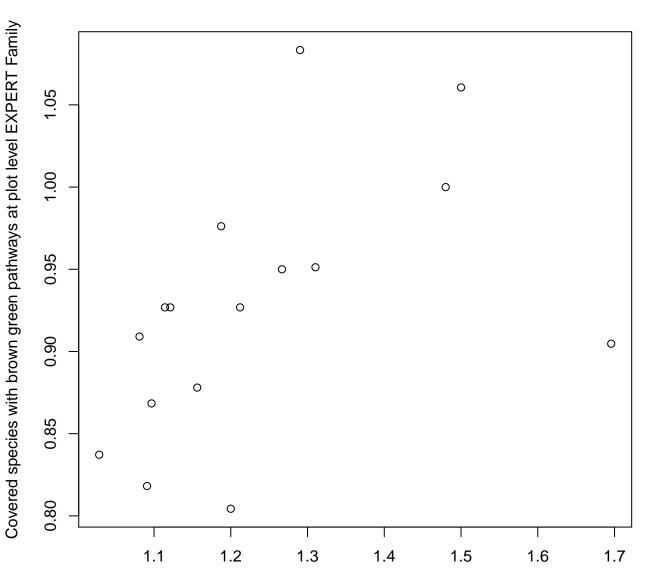
Covered species with brown green pathways at plot level SBM Species



Covered species with brown green pathways at plot level SBM Genus



Covered species with brown green pathways at plot level EXPERT Species



Covered species with brown green pathways at plot level EXPERT Genus

