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Cover image: The figure on the cover (from Chen et al, pages 232–243) shows the taxonomic tree of all differentially enriched species in the comparison of gut microbiota between systemic lupus erythematosus patients and healthy controls. Each circular sector filled by a different color represents a single phylum. Bar plots outside the central circle depict the group in which a specific species was enriched and how big the linear discriminant analysis score was for that species. Species that were observed to be increased in systemic lupus erythematosus patients and decreased after treatment are shown as large red dots.