

Maximum-likelihood tree of *Ssty* family members and spindlin family homologs in rat; genes shaded by chromosomal location: dark purple, NPY; light purple, NPY pseudogenes; orange, NPX; grey, autosomal pseudogenes. Heavy lines indicate branches leading to copies with intact ORFs; numbers, branches with greater than 50% bootstrap support. Scale bar; branch length in substitutions per site.

