

**nucleosome assembly, GO:0006334**

nucleosome, GO:0000786

chromatin remodeling, GO:0006338

nucleosomal DNA binding, GO:0031492

chromatin silencing at rDNA, GO:0000183

chromatin silencing, GO:0006342

male gonad development, GO:0008584

protein heterotetramerization, GO:0051290

chromatin DNA binding, GO:0031490

telomere organization, GO:0032200

telomere capping, GO:0016233

positive regulation of binding, GO:0051099

peptide alpha-N-acetyltransferase activity, GO:0004596