

	510814-13 ♂	<i>Clethrionomys rutilus</i>	123-33-19-14-19 gms, testes 2 mm
	510814-14 ♀	<i>Microtus oeconomus</i>	111-27-18.5-11-13", ut. normal
SKULL	510814-15 ♂	<i>Spermophilus undulatus kennicottii</i>	361-102-59-22 gms, testes 13 mm
	510814-16 ♂	"	405-125-60-2 lbs.
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	510815-1 ♀	<i>Clethrionomys rutilus</i>	125-33-20-15-23 gms, ut. normal
	510815-2 ♀	"	128-34-20-14-23", " "
	510815-3 ♀	"	132-34-20-14-21", " "
	510815-4 ♀	"	126-33-20-14-17", " "
	510815-5 ♂	"	125-34-20-14-22", testes 3 mm
	510815-6 ♂	"	128-34-20-14-18", " 2 "
	510815-7 ♂	"	127-34-20-14-17", " 2 "
	510815-8 ♀	"	120-32-19.5-14-17", ut. normal
	510815-9 ♂	"	123-33-19-14-18", testes 2 mm
	510815-10 ♂	"	133-36-20-14-20", " 2 "
	510815-11 ♂	"	138-37-20-14-28", " 6 "
	510815-12 ♀	"	126-34-19-14-21", ut. normal
	510815-13 ♂	"	130-35-20-14-21", testes 3 mm
	510815-14 ♂	"	118-33-20-14-16", " 3.5 "
	510815-15 ♂	"	118-33-20-14-16", " 3.5 "
	510815-16 ♂	"	110-30-19.5-13-13", " 3.5 "
	510815-17 ♀	<i>Microtus murus muriei</i>	153-34-19-13-35", ut. normal
	510815-18 ♂	"	121-27-21.5-12-22", testes 2 mm
	510815-19 ♀	"	115-27-19-12-19 gms, ut. normal
	510815-20 ♀	"	125-28-19-12-21", " "
m.i.o.	510815-20a ♀	" <i>oeconomus</i>	128-31-19-11-18", " "
	510815-21 ♂	" <i>murus muriei</i>	125-27-21-13-24", testes 2.5 mm
	510815-22 ♀	"	122-26-19-12-20", ut. normal
	510815-23 ♂	"	115-23-19-12-20", testes 2.5 mm
m.i.o.	510815-24 ♂	"	106-24-18-10-12", " 2.5 "
"	510815-25 ♀	"	108-25-18-10-13", ut. normal
"	510815-26 ♀	"	106-24-18-10-13", " "
SKEL.	510815-27 ♂	<i>Passerculus sandwichensis</i>	L. 156, 18 gms wt.
SKULL	510815-28 ♂	<i>Spermophilus u. kennicottii</i>	392-115-64-15-558 gms, testes 6.5
	510815-29 ♂	<i>Lemmus t. subarcticus</i>	132-19-16.5-10-40", " 6 mm
	510815-30 ♂	<i>Clethrionomys rutilus</i>	120-29-20-14-20", testes 1.5 mm
	510815-31 ♂	"	118-27-20-14-20", " 1.5 "
	510815-32	<i>Sorex arcticus tundrensis</i>	98-33-13-5 gms
	510815-33 ♀	<i>Microtus murus muriei</i>	128-27-20-13-21 gms, ut. normal
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	510816-1 ♀	<i>Clethrionomys rutilus</i>	123-34-21-14-23 gms., ut. normal